

IN THE
Supreme Court of the United States

GATEWAY CITY CHURCH, ET AL.,

Plaintiffs-Applicants,

v.

GAVIN NEWSOM, ET AL.,

Defendants-Respondents.

APPENDIX OF RESPONDENTS' EXHIBITS

OFFICE OF THE COUNTY COUNSEL
COUNTY OF SANTA CLARA

JAMES R. WILLIAMS*

County Counsel

GRETA S. HANSEN

DOUGLAS M. PRESS

TONY LOPRESTI

MELISSA KINIYALOCTS

HANNAH KIESCHNICK

70 West Hedding Street

East Wing, Ninth Floor

San José, CA 95110-1770

Telephone: (408) 299-5900

James.williams@cco.sccgov.org

**Counsel of Record*

February 24, 2021

*Attorneys for Defendants-Respondents
County of Santa Clara and Sara H.
Cody, MD*

TABLE OF CONTENTS

- Exhibit 1:** County of Santa Clara Public Health Dep't, *Mandatory Directive for Gatherings* (July 14, 2020; last rev'd Feb. 12, 2021),
<https://www.sccgov.org/sites/covid19/Documents/Mandatory-Directives-Gatherings.pdf>.
- Exhibit 2:** Declaration of Dr. Sara H. Cody in Support of Defendants County of Santa Clara and Dr. Sara H. Cody's Opposition to Plaintiffs' Motion for a Preliminary Injunction, *Gateway City Church v. Newsom*, No. 20-cv-08241-EJD, ECF No. 53-3 (N.D. Cal. Dec. 23, 2020)
- Exhibit 3:** Declaration of Dr. Marc Lipsitch in Support of Defendants County of Santa Clara and Dr. Sara H. Cody's Opposition to Plaintiffs' Motion for a Preliminary Injunction, *Gateway City Church v. Newsom*, No. 20-cv-08241-EJD, ECF No. 53-4 (N.D. Cal. Dec. 23, 2020)
- Exhibit 4:** Federal Aviation Administration, Information for Airport Sponsors Considering COVID-19 Restrictions or Accommodations (May 29, 2020), *Gateway City Church v. Newsom*, No. 20-cv-08241-EJD, ECF No. 53-1 (N.D. Cal. Dec. 23, 2020)
- Exhibit 5:** County of Santa Clara Public Health Dep't, *Mandatory Directive on Travel* (Nov. 28, 2020; last rev'd Jan. 25, 2021),
<https://www.sccgov.org/sites/covid19/Documents/Mandatory-Directives-Travel.pdf>.

Exhibit 1



Santa Clara County
**PUBLIC
HEALTH**

MANDATORY DIRECTIVE:



Gatherings

Issued July 14, 2020

sccgov.org/coronavirus

Revised and Effective: February 12, 2021

County of Santa Clara
Public Health Department

Health Officer
976 Lenzen Avenue, 2nd Floor
San José, CA 95126
408.792.5040



MANDATORY DIRECTIVE FOR GATHERINGS

Please confirm that your gathering is allowed under the State Order. Where there is a difference between the local County Order and the State Order, the more restrictive order must be followed.

Information on the State’s Order and State guidance is available at covid19.ca.gov.

Issued: July 14, 2020
Revised and Effective: February 12, 2021
Effective Upon Release

Every person, business, and entity in Santa Clara County must follow *both* the County and the State Public Health Officer Orders. Below is information on: (1) the State’s general gathering rules, and (2) the current rules for *indoor* gatherings in Santa Clara County.

1. General State Gatherings Rules:

As of January 25, 2021, the State’s Regional Stay at Home Order is ***no longer in effect*** in Santa Clara County.

On August 28, 2020, the State issued a Statewide Public Health Officer Order (“State Order,” available [here](#) and the Blueprint for a Safer Economy (“Blueprint,” available [here](#)). The State Order and Blueprint establish statewide restrictions applicable to each “tier” to which counties are assigned.

The State Health Officer has generally prohibited gatherings of all kinds statewide, with limited exceptions for worship services, cultural ceremonies like weddings and funerals, protest or political activities, and any gathering that is explicitly allowed by a State COVID-19 Industry Guidance document (<https://covid19.ca.gov/industry-guidance/>) or by the State’s “Stay home Q&A” page (<https://covid19.ca.gov/stay-home-except-for-essential-needs/>). The State also allows private gatherings consisting of no more than three households pursuant to the State’s rules, but such gatherings must be outdoors while the County is in the Purple Tier.

All gatherings are subject to the mandatory requirements in this Directive and any other applicable County Health Officer Directive, the County Health Officer's Revised Risk Reduction Order issued on October 5, 2020, the applicable restrictions under the State Order and Blueprint, the State's COVID-19 Industry Guidance documents, and any applicable health and safety regulations.

Note: While wedding ceremonies may occur outdoors subject to the mandatory requirements of this Directive, the State has clarified that “[w]edding receptions/parties/celebrations are NOT permitted at this time” under State Public Health Officer orders.

2. Current Rules for Indoor Gatherings in Santa Clara County:

Because indoor gatherings continue to pose a severe risk of COVID-19 transmission, **all indoor gatherings are currently prohibited.**

While COVID-19 is still circulating in our community, the Health Officer strongly discourages any gathering together with people from other households. Indoor gatherings are particularly risky because COVID-19 transmission occurs more easily indoors than outdoors, and COVID-19 continues to circulate widely. The most recent scientific evidence underscores the risk of transmission indoors, and **indoor gatherings are always strongly discouraged, even when allowed.** But gatherings are not prohibited by this Directive as long as everyone attending the gathering strictly complies with all the requirements set forth below to reduce risk and keep everyone who attends as safe as possible.

A “gathering” is an event, assembly, meeting, or convening that brings together multiple people from separate households in a single space, indoors or outdoors, at the same time and in a coordinated fashion—like a wedding, banquet, conference, religious service, festival, fair, party, performance, competition, movie theater operation, fitness class, barbecue, protest, or picnic. Although the County allows all types of gatherings to occur in compliance with this Directive, at this time the State generally allows gatherings only for purposes of worship services, cultural ceremonies like funerals and weddings (but not wedding receptions, which are prohibited), and protest or political activities. The State also allows gatherings for purposes identified in the State's [Industry Guidance](#) or any other State guidance document. The State also allows small gatherings of any type with no more than three households. These gatherings must follow the County's rules (contained in this Directive) and the [State's rules](#), including the State requirement that such gatherings must be outdoors when counties are in the Purple Tier. Because the stricter of the requirements applies, the only types of gatherings allowed in the County are those allowed by the State.

This Directive does not regulate whether a facility is open or closed. For example, facilities that are typically used for gatherings—such as places of worship, meeting halls, and event spaces—may remain open for purposes that do not involve gatherings, even when gatherings are prohibited indoors.

This Directive explains the local requirements for gatherings in Santa Clara County. **This Directive is *mandatory*, and failure to follow it is a violation of the Health Officer’s Order issued October 5, 2020 (“Order”).**

The Order Issued October 5, 2020

The Order imposes several restrictions on all businesses and activities to ensure that the County stays as safe as possible. All persons and businesses (including nonprofits, educational entities, and any other business entity, regardless of its corporate structure) that organize or host gatherings—such as religious institutions, wedding venues, wedding planners/coordinators, convention centers, and conference/meeting room rental facilities—must comply with the following requirements, and must ensure that participants comply with all applicable requirements:

- **Social Distancing Protocol:** All businesses and governmental entities that have not already done so must fill out and submit an updated Social Distancing Protocol under the October 5, 2020 Health Officer Order. **Social Distancing Protocols submitted prior to October 11, 2020 are no longer valid.** The Revised Social Distancing Protocol must be filled out using an updated template, which is available [here](#). The Protocol is submitted under penalty of perjury, meaning that everything written on the form must be truthful and accurate to the best of the signer’s knowledge, and submitting false information is a crime. The Protocol must be distributed to all workers, and it must be accessible to all officials who are enforcing the Order.
- **Signage:** All businesses and governmental entities must print (1) an updated COVID-19 PREPARED Sign and (2) a Social Distancing Protocol Visitor Information Sheet, and both must be posted prominently at all facility entrances. These are available for printing after submission of the Revised Social Distancing Protocol online. The Revised Social Distancing Protocol specifies additional signage requirements.
- **Face Coverings:** Everyone must wear face coverings at all times specified in the California Department of Public Health’s mandatory [Guidance for the Use of Face Coverings](#) (“Face Covering Guidance”) and in any specific directives issued by the County Health Officer. **Unless otherwise stated in this Directive, face coverings must be worn at all times when attending a gathering.**
- **Capacity Limitation:** All businesses must comply with the capacity limitations established in the [Mandatory Directive on Capacity Limitations](#).

Mandatory Requirements for All Gatherings

In general, the more people a person interacts with at a gathering, the closer the physical interaction is, the more enclosed the gathering space is, and the longer the interaction lasts, the higher the risk that a person with an unknown SARS-CoV-2 infection (the infection that causes COVID-19) might spread it to others. If not everyone follows the rules to safely gather, the risk of spreading SARS-CoV-2 is even higher. Based on those principles, the Health Officer's directives for *all* gatherings are:

1. If Gathering, the Health Officer Strongly Urges You to Gather Outdoors

- a. Gatherings that occur outdoors are significantly safer than indoor gatherings. To qualify as an outdoor gathering, the gathering must be held entirely outdoors, except that attendees may go inside to use restrooms as long as the restrooms are frequently disinfected.
- b. The maximum number of people allowed at an outdoor gathering is specified in the [Mandatory Directive on Capacity Limitations](#). This includes everyone present, such as hosts, workers, and guests. The space must be large enough so that everyone at a gathering can maintain at least 6-foot social distance from anyone (other than people from their own household).
 - i. Example 1: A small church hosts a funeral ceremony in its churchyard. The churchyard is only big enough to allow 25 people to easily maintain 6-foot social distancing between households at all times. **No more than 25 people may be present at the funeral ceremony.**
 - ii. Example 2: A couple holds their wedding ceremony outdoors at a historic hotel. The outdoor ceremony space is big enough for 1,000 people to maintain 6-foot distancing. **Even so, no more than the maximum number of people allowed by the Mandatory Directive on Capacity Limitations may be present at the wedding ceremony.**
- c. A gathering is considered an outdoor gathering only if it is held at a facility that allows the free flow of outdoor air through the entire space, as specified in the California Department of Public Health's mandatory guidance on [Use of Temporary Structures for Outdoor Business Operations](#).
- d. Fences and screens that do not impede airflow are not considered walls or sides for purposes of determining whether an area is outdoors. Partitions around or within the facility may be used and do not qualify as sides so long as they are no more than 3 feet in height as measured from the floor.

2. *The Health Officer Strongly Discourages Indoor Gatherings, Even When They Are Allowed*
 - a. Indoor gatherings may not be allowed depending on the County’s current tier under the State’s Blueprint for a Safer Economy and other local factors. See the “Current Rules for Indoor Gatherings in Santa Clara County” box at the top of this Directive for information on current rules.
 - b. When indoor gatherings *are* allowed, the maximum number of people allowed at an indoor gathering is specified in the [Mandatory Directive on Capacity Limitations](#). This includes everyone present, such as hosts, workers, and guests.
3. *Don’t Attend Gatherings If You Feel Sick or You Are in a High-Risk Group*
 - a. If you feel sick or have any COVID-19-like symptoms (fever, cough, shortness of breath, chills, night sweats, sore throat, nausea, vomiting, diarrhea, tiredness, muscle or body aches, headaches, confusion, or loss of sense of taste/smell), **you must stay home and may not attend any gatherings.**
 - b. As explained on the [People Who Need Extra Precautions](#) page, people at higher risk of severe illness or death from COVID-19 are strongly urged not to attend any gatherings.
4. *All Gatherings Must Have an Identified and Designated Host Who Is Responsible for Ensuring Compliance with All Requirements*
 - a. A specific person or business (including nonprofits, religious organizations, educational entities, or any other business entity) must be the designated host for a gathering and ensure compliance with all requirements in the Order and this Directive. **The host is responsible and subject to enforcement for any failure by participants to comply with the Order and this Directive.**
 - b. The host also must maintain a list with names and contact information of all participants at the gathering. If a participant tests positive for COVID-19, the host is legally required to assist the County Public Health Department in any case investigation and contact tracing associated with the gathering. Public Health will ask for the list of attendees *only* if an attendee tests positive for COVID-19. The County Public Health Department will keep this information confidential and use it only for case investigation and contract tracing purposes. Hosts must maintain these records for at least 21 days. The host must provide the list to any County Enforcement Officer immediately upon request.

5. Practice Physical Distancing and Hand Hygiene at Gatherings

- a. At all gatherings, **everyone must stay at least 6 feet away from other people (except people in their own household) at all times.**
- b. Seating arrangements must provide at least 6 feet of distance (in all directions—front-to-back and side-to-side) between different households. This can be done by spacing chairs apart, or for fixed seating like benches or pews, by marking off rows and indicating seating areas with tape. Seating and tables must be sanitized after each use.
- c. Everyone at a gathering should frequently wash their hands with soap and water, or use hand sanitizer if soap and water are not available. The host must make handwashing facilities or hand sanitizer available for participants to use.

6. Rules for Face Coverings, Singing, Chanting, Shouting, and Playing Wind Instruments

Current scientific evidence shows that COVID-19 spreads primarily through respiratory droplets and fine aerosols that are released from the body when people breathe, sing, shout, or otherwise expel air from their lungs. Face coverings prevent many of these droplets and aerosols from escaping into the air, and wearing a face covering has been shown to significantly decrease the risk of COVID-19 transmission. Conversely, singing, chanting, shouting, and playing wind instruments have all been shown to significantly *increase* the risk of COVID-19 transmission, because these activities all release increased amounts of respiratory droplets and fine aerosols into the air. To reduce the risk of spreading COVID-19, the following rules apply to gatherings:

- a. For all *indoor* gatherings (when indoor gatherings are allowed):
 - i. **Everyone, including performers/presenters, must wear a face covering at all times** (except for very young children, people for whom face coverings are medically inadvisable, or for communication by or with people who are hearing impaired).
 1. Food and drink may not be served at indoor gatherings—including at movie theaters—even when indoor gatherings are allowed, except as necessary to carry out a religious ceremony.
 2. Face coverings may be removed to meet urgent medical needs (for example, to use an asthma inhaler, consume items needed to manage diabetes, take medication, or if feeling light-headed).
 - ii. Singing, chanting, shouting, and playing wind instruments are **strictly prohibited**.

- b. For all *outdoor* gatherings:
- i. **Except as described below or in other directives issued by the County Health Officer, everyone must wear a face covering at all times** (except for very young children, people for whom face coverings are medically inadvisable, or for communication by or with people who are hearing impaired).
 1. Attendees may remove their face coverings to eat or drink but must put their face covering back on as soon as they are finished eating or drinking.
 2. Attendees may remove their face coverings to meet urgent medical needs (for example, to use an asthma inhaler, consume items needed to manage diabetes, take medication, or if feeling light-headed).
 - ii. If an outdoor gathering involves a performance/presentation, performers/presenters may remove their face coverings while they are performing/presenting, but they must replace their face coverings after they finish.
 1. No more than 12 performers/presenters are permitted in the performance/presentation area at a time.
 2. Until their face covering is back on, any performer/presenter who removes their face covering to speak must maintain at least **12 feet** of social distance from everyone not in their household.
 3. Until their face covering is back on, any performer/presenter who removes their face covering to sing, chant, shout, or play a wind instrument must maintain at least **12 feet** of social distance from all other performers/presenters who are not in their household and at least **25 feet** from all attendees who are not performing/presenting.
 4. Any performer/presenter playing a wind instrument must cover the opening of the instrument (e.g., with cloth) to reduce the spread of respiratory droplets from the instrument.

5. Performers/presenters who are singing or chanting are strongly encouraged to do so at a quiet volume (at or below the volume of a normal speaking voice).
- iii. All attendees who are not performing/presenting **must wear a face covering at all times** while singing, chanting, or shouting. Because these activities pose a very high risk of COVID-19 transmission, face coverings are particularly essential to reduce the spread of respiratory droplets and fine aerosols. People who cannot wear a face covering for medical or other reasons are strongly discouraged from singing, chanting, or shouting.
 1. Attendees who are singing, chanting, or shouting are strongly encouraged to maintain increased social distancing greater than 6 feet to further reduce risk.
 2. Attendees who are singing or chanting are strongly encouraged to do so at a quiet volume (at or below the volume of a normal speaking voice).

7. *Stagger Attendance at Gatherings*

- a. For gatherings that have the potential to draw larger groups, like community meetings or religious services, consider offering multiple sessions, requiring reservations that cap attendance at each session, staggering arrivals and departures, and encouraging or requiring that the same group stays together (for example, Group A attends the Sunday morning worship service every week, and Group B attends the separate Tuesday evening worship service every week).
- b. There is no limit on the number of gatherings that may be held at different times on a single day—for example, a mosque may hold prayer services five times a day—as long as (i) each gathering follows all the rules, and (ii) restrooms, chairs and tables, and any other high-touch surfaces are properly sanitized between groups.
- c. A venue may host *multiple outdoor gatherings* at the same time (for example, multiple small barbecues in a large outdoor space like a 20-acre ranch)—as long as:
 - i. Each gathering follows all the rules in the Order and in this Directive. Each gathering must, for instance, have its own designated host who must maintain a list of participant names and contact information.

- ii. Each gathering has its own area marked by prominent signage, barriers, or ropes, and there is a buffer zone of at least 100 feet between the boundaries of any two separate gatherings.
 - iii. The participants at a gathering, including hosts, workers, and guests, do not mix between or among different gatherings and stay strictly in their own area.
 - iv. There are sufficient restroom facilities, or a system of using the restroom facilities, such that participants from different gatherings do not have contact with one another when they use the restroom.
- d. When indoor gatherings are allowed, a venue may host multiple indoor gatherings at the same time (for example, multiple gatherings in separate rooms within a building)—as long as:
- i. Each gathering follows all the rules in the Order and in this Directive. Each gathering must, for instance, have its own designated host who must maintain a list of participant names and contact information.
 - ii. Each gathering is fully separated by solid, floor-to-ceiling walls or partitions from any other gathering.
 - iii. Where possible, the HVAC system for each space with a gathering should ventilate to the outdoors, rather than into a space with another gathering.
 - iv. The participants at a gathering, including hosts, workers, and guests, do not mix between or among different gatherings and do not enter into a common space with participants from any other gathering.
 - v. There are sufficient restroom facilities, or a system of using the restroom facilities, such that participants from different gatherings do not have contact with one another when they use the restroom.

//

//

8. *Livestreaming, Broadcasting, and Recording with No Audience Present*

Businesses may livestream, broadcast, or record performances, services, and classes at indoor facilities without live audiences or members of the public present. All such livestreamed, broadcasted, or recorded events at indoor facilities **must** comply with the following rules:

- a. When livestreaming under this provision, only personnel may be present at the facility. Audiences or other members of the public are strictly prohibited. The number of personnel inside the facility must be limited to the minimum necessary to conduct the event (and may never exceed 12 people or the maximum number of people allowed under the facility's current capacity limitation as dictated by the [Mandatory Directive on Capacity Limitations](#), whichever is fewer).
- b. All personnel, including performers/presenters in the performance area, must maintain at least 6 feet of physical distance from everyone outside their household at all times.
- c. People performing or presenting during a livestreamed event may remove their face coverings during the performance or presentation if everyone maintains at least 12 feet of physical distance from everyone outside their household at all times. Even so, the Health Officer strongly urges people performing or presenting to wear a face covering whenever possible. All others on-site must wear a face covering in compliance with State and County Health Officer requirements.
- d. When livestreaming under this provision with no audience present, singing, chanting, or playing wind instruments indoors is strongly discouraged but may occur so long as everyone maintains at least 12 feet of physical distance from everyone outside their household.

For clarity, the above rules for livestreaming, broadcasting, and recording do not apply if any member of the public is present for the event. Businesses may livestream, broadcast, or otherwise record an event at which members of the public are present (so long as current State and County Health Officer orders allow members of the public to be present for such an event), but there are no special rules that would apply to the livestream, broadcast, or recording. Instead, these events must comply with all rules currently governing the business's general operations. Note that these rules may be stricter those listed above.

9. Maximize Ventilation for Indoor Gatherings (When Indoor Gatherings Are Allowed)

- a. Open doors and windows to maximize circulation of outdoor air whenever environmental conditions and building requirements allow. Consider modifications to the facility to increase outdoor air exchange, such as replacing non-opening windows with openable screened windows. Contact your local Building Department for more information on permit requirements.

- b. Indoor facilities with central air handling/HVAC systems must ensure that HVAC systems are serviced and functioning properly and, to the extent feasible and appropriate to the facility:
 - i. Evaluate possibilities for and implement upgrades to the system to ensure that air filters are functioning at the highest efficiency compatible with the currently installed filter rack and air handling system (ideally MERV-13 or greater).
 - ii. Increase the percentage of outdoor air through the HVAC system, readjusting or overriding recirculation (“economizer”) dampers.
 - iii. Disable demand-control ventilation controls that reduce air supply based on temperature or occupancy.
 - iv. Implement the additional measures set forth in the County’s Guidance for Ventilation and Air Filtration Systems.
- c. Indoor facilities that do not have central air handling/HVAC systems or that do not operate or control the system must take the following measures, to the extent feasible and appropriate to the facility:
 - i. Set any ceiling fans to draw air upwards away from participants.
 - ii. If using portable fans, position them near open doors/windows and use them to draw or blow inside air to the outside of the facility. Position fans to minimize blowing air between occupants, which may spread aerosols.
 - iii. Consider installing portable air filters appropriate to the space.
 - iv. Implement additional applicable measures set forth in the County’s Guidance for Ventilation and Air Filtration Systems.
- d. Upon request by a County Enforcement Officer or County Public Health Department Staff, the facility may be required to perform a comprehensive evaluation of the facility’s ventilation and air filtration system by an appropriately licensed professional, and produce documentation regarding this evaluation to the County.

Stay Informed

For answers to frequently asked questions about this industry and other topics, please see the [FAQs page](#). **Please note that this Directive may be updated.** For up-to-date information on the Health Officer Order, please visit the County Public Health Department's website at www.sccgov.org/coronavirus.

Exhibit 2

1 JAMES R. WILLIAMS, County Counsel (S.B. #271253)
HANNAH KIESCHNICK, Deputy County Counsel (S.B. #319011)
2 KARAN SINGH DHADIALLA, Deputy County Counsel (S.B. #296313)
OFFICE OF THE COUNTY COUNSEL
3 70 West Hedding Street, East Wing, Ninth Floor
San José, California 95110-1770
4 Telephone: (408) 299-5900
Facsimile: (408) 292-7240

5 Attorneys for Defendants
6 COUNTY OF SANTA CLARA and
DR. SARA H. CODY

8 UNITED STATES DISTRICT COURT
9 NORTHERN DISTRICT OF CALIFORNIA
(San José Division)

10
11 GATEWAY CITY CHURCH, et al.,

12 Plaintiffs,

13 v.

14 GAVIN NEWSOM, et al.,

15 Defendants.

No. 20-CV-08241-EJD

**DECLARATION OF THE HEALTH
OFFICER FOR THE COUNTY OF SANTA
CLARA SARA H. CODY, M.D., IN
SUPPORT OF DEFENDANTS COUNTY
OF SANTA CLARA AND DR. SARA H
CODY'S OPPOSITION TO PLAINTIFFS'
MOTION FOR A PRELIMINARY
INJUNCTION**

Date: January 15, 2020
Time: 10:00 a.m.
Courtroom: 4
Judge: Hon. Edward J. Davila

16
17
18
19 I, SARA H. CODY, M.D., declare as follows:

20 1. I have personal knowledge of the facts set forth in this declaration. If called as a
21 witness, I could and would testify competently to the matters set forth herein.

22 **Background and Experience**

23 2. I graduated from Stanford University, where I received a degree in Human Biology. I
24 received my Doctor of Medicine from Yale University School of Medicine. Following an Internship
25 and Residency in Internal Medicine at Stanford University Hospital, I completed a two-year
26 fellowship in Epidemiology and Public Health, as an Epidemic Intelligence Service (EIS) Officer
27 with the Centers for Disease Control and Prevention (CDC).

28 3. I am currently the Director of the County of Santa Clara's Public Health Department,

1 as well as the Health Officer for the County and each of the 15 cities located within Santa Clara
2 County. I have held the Health Officer position since 2013 and the Director of Public Health
3 Department position since 2015. In these roles, I provide leadership on public health issues for all of
4 Santa Clara County and oversee approximately 400 Public Health Department employees who
5 provide a wide array of services to safeguard and promote the health of the community.

6 4. Before becoming the Health Officer for the County and each of its cities, I was
7 employed for 15 years as a Deputy Health Officer/Communicable Disease Controller at the County's
8 Public Health Department, where I oversaw surveillance and investigation of individual cases of
9 communicable diseases, investigated disease outbreaks, participated in planning for public health
10 emergencies, and responded to Severe Acute Respiratory Syndrome (SARS), influenza A virus
11 subtype H1N1 (also known as "swine flu" or H1N1), and other public health emergencies.

12 5. The mission of the Public Health Department is to promote and protect the health of
13 Santa Clara County's population of approximately 1.9 million people. None of Santa Clara
14 County's 15 cities has a health department. All 15 cities, and all Santa Clara County residents, rely
15 on the Public Health Department to perform essential public health functions. The work of the
16 Public Health Department is focused on three main areas: (1) infectious disease and emergency
17 response; (2) maternal, child, and family health; and (3) healthy communities. The Public Health
18 Department's work is guided by core public health principles of equity, collaboration and inclusion,
19 and harm prevention. This work—in particular, infectious disease control and emergency
20 response—is critical to the health of the entire community countywide.

21 **The Novel Coronavirus and COVID-19**

22 6. The current pandemic—from a novel coronavirus that was first identified in
23 December 2019—has spread to most countries in the world, including the United States.
24 Worldwide, as of December 23, 2020, authorities had confirmed at least 78,320,614 cases, and
25 1,723,502 deaths. The United States reported its first case on January 21, 2020. The disease has
26 since spread rapidly throughout the country. As of December 23, 2020, public health authorities had
27 confirmed at least 18,281,597 cases in the United States, and 323,682 deaths. Starting November 4,
28 2020, the United States has recorded more than 100,000 new cases each day with many days

1 surpassing the previous single-day record and each day's total eclipsing the summer's single-day
2 record of 77,300 new cases in mid-July. And on December 16, 2020, there were 245,000 new cases
3 and over 3,600 deaths reported. Experts consider this epidemic to be the worst public health
4 epidemic since the influenza outbreak of 1918, and recent case numbers, hospitalizations, and deaths
5 confirm the epidemic is worsening.

6 7. Coronaviruses are a large family of viruses that can cause illness ranging from the
7 common cold to more severe diseases such as Middle East Respiratory Syndrome (MERS-CoV) and
8 Severe Acute Respiratory Syndrome (SARS-CoV). A novel coronavirus is a new coronavirus that
9 has not been previously identified in humans. SARS-CoV-2 refers to the novel coronavirus
10 currently spreading throughout the world, and Coronavirus Disease-19 (COVID-19) is the disease
11 caused by SARS-CoV-2. The World Health Organization has deemed COVID-19 a pandemic. One
12 vaccine for SARS-CoV-2 received an Emergency Use Authorization on December 12, and a second
13 vaccine received Emergency Use Authorization on December 18, 2020. A small number of vaccine
14 doses have been distributed and vaccination of people in the highest risk categories, including
15 healthcare personnel at acute care facilities and residents and staff of skilled nursing and other long
16 term care facilities, began last week. There is no specific cure for COVID-19.

17 8. In mild cases, COVID-19 may cause fever, fatigue, and cough. In severe cases, it may
18 cause shortness of breath and pneumonia, multi-organ system failure, and death. In some cases,
19 COVID-19 can cause neurological symptoms such as inability to taste or smell, tingling or
20 numbness in the hands and feet, confusion, seizures, and stroke. Evidence has shown that COVID-
21 19 can also cause long-term chronic health conditions, such as cardiovascular, neurologic, renal, and
22 respiratory damage and loss of limbs from blood clotting.¹ These conditions may be experienced not
23 only by those who become seriously ill from COVID-19 but also by those who experience only mild
24 symptoms.² Public health and healthcare officials are still learning about the long-term

26 ¹ Del Rio, C., et al., *Long-term Health Consequences of COVID-19*, JAMA, Vol. 324, No. 17, pp.
27 1723–24, Oct. 5, 2020, <https://doi.org/10.1001/jama.2020.19719>.

28 ² *Id.*; Tenford, Mark W., et al., *Symptom Duration and Risk Factors for Delayed Return to Usual Health Among Outpatients with COVID-19 in a Multistate Health Care Systems Network–United*

1 complications associated with COVID-19, which was first identified in late 2019.

2 9. The SARS-CoV-2 virus is highly contagious, spreading easily between individuals.
3 While the exact mechanisms by which SARS-CoV-2 spreads are still being studied, there is
4 consensus among epidemiologists that SARS-CoV-2 is spread primarily via an airborne route. The
5 novel coronavirus is spread primarily from person to person through respiratory droplets and
6 aerosols (that is, smaller particles that travel farther than respiratory droplets) produced when an
7 infected person coughs, sneezes, sings, shouts, or talks. SARS-CoV-2 may spread to a much lesser
8 extent by touching objects or surfaces that have been contaminated with the novel coronavirus and
9 then touching one's mouth, nose, or eyes.

10 10. People who have symptomatic SARS-CoV-2 infection are contagious and can spread
11 the infection to others. People with SARS-CoV-2 infection can also spread the virus to others before
12 any symptoms develop. In addition, there is broad consensus among epidemiologists that some
13 people with SARS-CoV-2 infection never develop symptoms (i.e. asymptomatic infection) and still
14 spread the virus. An estimated 40% of people with SARS-CoV-2 infection are asymptomatic.³

15 11. There is consensus among epidemiologists that social distancing helps reduce the risk
16 of transmission.⁴ This is in part because maintaining at least six feet of physical distance between
17 persons substantially decreases the likelihood that sufficient respiratory droplets carrying SARS-
18 CoV-2 will enter a person's mouth, nose, eyes, or lungs to cause a person to contract a SARS-CoV-2
19 infection. However, six feet of social distance is likely not adequate to prevent spread of SARS-
20 CoV-2 via aerosols, which can travel greater distances and linger in the air for much longer. The
21 risk of transmission is greater in indoor environments with crowding and poor ventilation. This risk

22
23
24 *States, March–June 2020*, CDC Morbidity & Mortality Wkly. Rep., Vol. 69, No. 31, pp. 993–98,
July 31, 2020, <http://dx.doi.org/10.15585/mmwr.mm6930e1>.

25 ³ Oran, Daniel P., et al., *Prevalence of Asymptomatic SARS-CoV-2 Infection: A Narrative Review*,
26 *Annals of Internal Med.*, Vol. 173, Issue 5, pp. 362–67, Sept. 1, 2020, <https://doi.org/10.7326/M20-3012>.

27 ⁴ Alagoz, Oguzhan, et al., *Effect of Timing of and Adherence to Social Distancing Measures on*
28 *COVID-19 Burden in the United States*, *Ann. of Internal Med.*, Oct. 27, 2020,
<https://doi.org/10.7326/M20-4096>.

1 increases when people gather in indoor environments for long periods of time. As discussed below,
2 the risk increases further when people are shouting, singing, or engaging in heavy breathing because
3 these activities produce more droplets and aerosols than normal speaking. And the likelihood of
4 transmission further increases where there is widespread community transmission as this increases
5 the probability that a person at the gathering will be infectious.

6 12. There is also consensus among epidemiologists that wearing a face covering over the
7 nose and mouth reduces the number of SARS-CoV-2 infections.⁵ The CDC reports that
8 “[e]xperimental and epidemiological data support community masking to reduce the spread of
9 SARS-CoV-2” and recommends “[a]dopting universal masking policies.”⁶ The use of face
10 coverings alone, however, is not sufficient to eliminate the risk of transmission. Face coverings are
11 one prevention strategy and must be paired with other strategies such as social distancing, increasing
12 ventilation, etc. For instance, as discussed below, face coverings are not perfect at preventing
13 transmission from a person singing. Face coverings are less likely to prevent transmission if they are
14 loosely fit, as droplets or aerosols are more likely to escape.

15 13. The finding that some people with SARS-CoV-2 infection are asymptomatic
16 underscores the importance of following social distancing, always using a face covering, and
17 limiting contact with people outside of one’s household to the greatest extent possible. Because
18 people can be infected and not have symptoms, they may be unaware they are infected and
19 contagious, and therefore less likely to take steps or precautions to limit the spread of infection, like
20 staying at home. The relatively high percentage of infections that are asymptomatic means that

23 ⁵ Brooks, John T., et al., *Universal Masking to Prevent SARS-CoV-2 Transmission—The Time Is*
24 *Now*, JAMA, Vol. 324, No. 7, pp. 635–37, July 14, 2020, <https://doi.org/10.1001/jama.2020.13107>;
25 Wang, Xiaowen, et al., *Association Between Universal Masking in a Health Care System and SARS-*
CoV-2 Positivity Among Health Care Workers, JAMA, Vol. 324, No. 7, pp. 703–04, July 14, 2020,
<https://doi.org/10.1001/jama.2020.12897>.

26 ⁶ *Scientific Brief: Community Use of Cloth Masks to Control the Spread of SARS-CoV-2*, CDC (Nov.
27 10, 2020), <https://www.cdc.gov/coronavirus/2019-ncov/more/masking-science-sars-cov2.html>; Lyu,
28 Wei, et al., *Community Use of Face Masks and COVID-19: Evidence from a Natural Experiment of*
State Mandates in the US, HealthAffairs, Vol. 39, No. 8, June 16, 2020,
<https://doi.org/10.1377/hlthaff.2020.00818>.

1 SARS-CoV-2 can spread silently, fueling community transmission, increasing community
2 prevalence and risk for the most vulnerable members of a population.

3 14. Because none of these measures on their own is sufficient to eliminate the risk of
4 transmission, the CDC promotes a multipronged application of evidence-based strategies that
5 includes restrictions on “nonessential indoor spaces that pose the highest risk for transmission” in
6 addition to universal mask policies and social distancing requirements.”⁷ The restriction of indoor
7 gatherings is particularly important because the location, type, size, and duration of activity all
8 impact the risk of transmission, and indoor gatherings where people from different households
9 interact and remain in close proximity for extended periods of time create a high risk of
10 transmission.

11 **March 16, 2020 Shelter-in-Place Order**

12 15. Santa Clara County recorded the first known death from COVID-19 in the entire
13 United States (February 7, 2020) in a person who had not traveled or had contact with a known case.
14 The County also announced two of the earliest cases of travel-associated COVID-19 in the United
15 States on January 31 and February 2.

16 16. To slow community-wide virus transmission as much as possible in order to protect
17 the most vulnerable populations, prevent deaths, and prevent the health care system from being
18 overwhelmed, and after consideration of epidemiologic trends in countries with active COVID-19
19 epidemics and recognition of the potential for exponential growth in the setting of a non-immune
20 population, on March 16, 2020, I issued a shelter-in-place order directing all individuals living in the
21 County to shelter at their place of residence. Six other Bay Area jurisdictions issued similar shelter-
22 in-place orders on the same day.

23 17. The goal of sheltering in place was to reduce the number of infections, to slow the
24 spread of infection, to ensure that communities had enough hospital capacity to care for people who
25

26
27 ⁷ Honein MA, et al., *Summary of Guidance for Public Health Strategies to Address High Levels of*
28 *Community Transmission of SARS-CoV-2 and Related Deaths*, December 2020. MMWR Morb
Mortal Wkly Rep 2020;69:1860-1867. DOI: <http://dx.doi.org/10.15585/mmwr.mm6949e2>.

1 developed severe illness, and ultimately, to save lives.

2 18. The March 16 Order prohibited all individuals from leaving their place of residence,
3 except for the limited purposes of performing listed essential activities. The order prohibited all
4 travel, except essential travel, such as to carry out essential activities. All businesses with a facility
5 in the County, except certain essential businesses, were required to cease all activities at their
6 facilities except certain minimum basic operations. The order also prohibited all public and private
7 gatherings of any number, except with members of individuals' own households.

8 19. The order was based on evidence of the exponential growth of COVID-19 within the
9 County and then-current scientific evidence and best practices regarding the most effective
10 approaches to slow the transmission of the virus. It was understood then, as it is now, that a proven
11 way to decrease the probability of SARS-CoV-2 being spread from person to person is to limit
12 interactions among people to the greatest extent practicable. At the time, allowing exceptions other
13 than essential activities, businesses, and travel—even with social distancing and face coverings—
14 would have resulted in more interactions among people and more opportunity for infection to spread
15 from person to person, ultimately resulting in more infections and more illness and deaths. Limiting
16 the number, duration, and proximity of contacts between people was critical in reducing the
17 probability that the virus would spread from one person to another. Ensuring that every person
18 sheltered in place to the maximum extent feasible, therefore, was expected to reduce the rate of
19 transmission of the SARS-CoV-2 infection (i.e., “flatten the curve”) more quickly than if we had not
20 sheltered in place. This is especially true in a population—like the County’s—with no immunity to
21 the novel coronavirus.

22 **March 31, 2020 Order to Continue Sheltering in Place**

23 20. On March 31, 2020, I issued an updated shelter-in-place order that superseded the
24 March 16, 2020 Order, to reinforce social distancing requirements, and to further reduce the total
25 volume of person-to-person contact occurring in the County. The March 31 Order extended the
26 shelter-in-place requirements through May 3, 2020. Six other Bay Area jurisdictions extended their
27 shelter-in-place orders on the same day.

28 21. At that time, the public health emergency had substantially worsened since the March

1 16 Order, with a significant escalation in the number of positive cases, hospitalizations, and deaths,
2 and a corresponding increasing strain on health care resources. At the same time, evidence
3 suggested that the restrictions on mobility and social distancing requirements imposed by the prior
4 orders were slowing the rate of increase in new cases, consistent with models of infectious disease
5 transmission in a non-immune population.

6 22. The March 31 Order was more restrictive than the March 16 Order in a number of
7 ways because it included, amongst other more restrictive provisions: (1) mandatory social distancing
8 requirements; (2) additional restrictions on essential business; (3) a prohibition on the use of
9 playgrounds, dog parks, public picnic areas, and similar recreational areas; and (4) a prohibition on
10 the use of shared public recreational facilities such as golf courses, tennis and basketball courts,
11 pools, and rock walls. As a condition to operate, all open essential businesses were required to
12 prepare a protocol to implement various social distancing measures, such as limiting the number of
13 persons in the facility at a time and regularly disinfecting high-touch surfaces (“Social Distancing
14 Protocol”).

15 23. The shelter-in-place orders slowed the spread of the virus. When the March 16 Order
16 was put in place, the number of new cases was growing rapidly. That changed with the March 16
17 and March 31 shelter-in-place orders.

18 24. One of the most important indicators of the rate of growth of the SARS-CoV-2 virus
19 is what is called the “doubling time,” *i.e.*, how many days it takes for the number of cases to double.
20 A shorter doubling time means that the infection is spreading rapidly; a longer doubling time means
21 that the infection is spreading more slowly. On March 16, when the first shelter-in-place order was
22 issued, the case count was doubling approximately every five days—indicating rapid, exponential
23 growth. By the beginning of May 2020, however, the doubling time had slowed to around three and
24 a half months. The shelter-in-place orders had been successful in significantly slowing the spread of
25 the virus—that is, they had flattened the curve.

26 25. Model projections prepared by the County’s Public Health Department in
27 collaboration with infectious disease modeling experts at Stanford University’s School of Medicine
28 estimated that over the six-week period from March 16 through April 25, 2020, the County and State

1 shelter-in-place orders prevented approximately 80 percent of the infections that otherwise would
2 have occurred.

3 26. Flattening the growth curve of confirmed cases not only prevented illnesses and
4 hospitalizations and saved lives, it bought the County time to significantly increase hospital capacity
5 and healthcare resources, to improve clinical management and treatment, to better understand routes
6 of transmission, to provide resources for vulnerable populations at high risk of infection, to increase
7 testing capacity across the County, and to take other critical measures to further slow the rate of
8 spread and prevent the healthcare system from becoming overwhelmed.

9 **April 29, 2020 Extended and Revised Shelter-in-Place Order**

10 27. To continue these trends, on April 29, 2020, I issued a revised shelter-in-place order
11 that superseded the March 31 Order. The April 29 Order went into effect at 11:59 p.m. on May 3,
12 2020 and extended most shelter-in-place restrictions through May 31, 2020. Reflecting the regional
13 progress made under shelter-in-place orders, six other Bay Area jurisdictions also issued similar
14 orders extending most shelter-in-place restrictions.

15 **May 22, 2020 Order to Continue Sheltering in Place**

16 28. To further continue these trends, and based on the available epidemiological
17 evidence, on May 18, 2020, I issued another shelter-in-place order that superseded the April 29
18 Order. The order went into effect at 12:01 a.m. on May 22, 2020 and extended most shelter-in-place
19 restrictions.

20 29. By this time, we had achieved progress in slowing the spread of the SARS-CoV-2
21 virus in Santa Clara County and the neighboring counties. Even so, because widespread SARS-
22 CoV-2 diagnostic testing was still not available across the County or the Bay Area region, many
23 infections still went undetected and contributed to silent spread of infection. Thus, continuation of
24 the prior shelter-in-place order remained necessary to suppress the rate of community spread, to
25 preserve critical and limited healthcare capacity in the County, to protect vulnerable populations, and
26 to prevent death. However, in light of progress in slowing the spread of infection, the May 22 Order
27 allowed a limited number of businesses and activities to resume operations, subject to specified
28 conditions and safety precautions to reduce any associated risk of SARS-CoV-2 transmission.

1 30. I did not lift all the prior restrictions because accepted principles of infectious disease
2 epidemiology, as demonstrated by the experiences of other regions in the U.S. and countries
3 combatting the pandemic, counseled in favor of a cautious and incremental approach towards
4 relaxing public health measures, particularly since the virus continued to circulate in the community;
5 pre-symptomatic and asymptomatic transmission of the virus presented a significant risk of silent
6 spread; and researchers, clinicians, and public health officials were still learning about the virus and
7 the range of health outcomes of the disease it causes.

8 **June 5, 2020 Update to Shelter-in-Place Order**

9 31. On June 1, 2020, I announced amendments to the May 22 Order that went into effect
10 at 12:01 a.m. on June 5, 2020.

11 32. When I announced the June 5 Order, the County and the greater Bay Area had
12 continued to make substantial progress in slowing the spread of SARS-CoV-2. Because of that
13 progress, and based on our developing understanding of the virus, the June 5 Order allowed certain
14 additional businesses and activities to resume, subject to restrictions to reduce transmission risk.
15 The order also kept key restrictions in place, requiring people to stay in their homes except when
16 engaging in certain essential or allowed activities. Appendix C-2 to the order details the additional
17 activities allowed and the accompanying restrictions on those activities under the update. As with
18 the previous orders, the June 5 Order required all businesses to complete and implement a Social
19 Distancing Protocol as a condition to operate.

20 33. The June 5 Order also permitted the resumption of certain outdoor activities,
21 including non-contact recreational and athletic activities, dog parks, automobile gatherings, and
22 some types of gatherings—again all subject to restrictions. For example, outdoor ceremonies and
23 outdoor religious gatherings were allowed, subject to restrictions to mitigate the risk of transmission,
24 including social distancing and face covering requirements to be enforced by a designated “host.”
25 The June 5 Order limited these outdoor ceremonies and gatherings to 25 attendees. The order also
26 prohibited singing and shouting at ceremonies and gatherings due to the significantly increased risk
27 of SARS-CoV-2 transmission from these specific activities, described below.

28 ///

1 **Developing Evidence of Risk of Transmission Indoors and Through Singing and Shouting**

2 34. Since the issuance of the June 5 Order, the evidence has grown that the risk of
3 transmission is higher with indoor activities than outdoor activities.⁸ Research has confirmed that
4 the virus that causes COVID-19 is primarily airborne, and spreads from person to person through
5 respiratory droplets and aerosols released into the air when singing, shouting, talking, coughing, or
6 sneezing. It is more likely that one will inhale respiratory droplets and aerosols from an infected
7 person in an indoor setting because aerosols disperse less easily indoors and can remain in the air for
8 a longer period of time. When outdoors, more frequent air movement and greater air volume
9 disperse respiratory droplets and aerosols, making SARS-CoV-2 transmission less likely.

10 35. A study of COVID-19 outbreaks in China reported that all of the identified outbreaks
11 of three or more cases occurred in an indoor environment, confirming that sharing indoor space is a
12 major infection risk.⁹ Other studies have suggested that normal speaking causes airborne virus
13 transmission in confined environments,¹⁰ and that closed environments contribute to transmission of
14 COVID-19 and promote superspreading events.¹¹

15 36. The CDC advises that activities are safer when they are “held in outdoor spaces.”¹²
16 The CDC’s guidance to the public regarding whether and how to engage in public activities
17 identifies “indoor space” as a factor that can increase the risk of SARS-CoV-2 spread.¹³

18 37. Limiting the size of gatherings is another important public health intervention
19

20
21 ⁸ Leclerc, Quentin J., et al., *What settings have been linked to SARS-CoV-2 transmission clusters?*,
Wellcome Open Res., Vol. 5, No. 83, June 5, 2020,
22 <https://doi.org/10.12688/wellcomeopenres.15889.2>.

23 ⁹ Qian, Hua, et al., *Indoor transmission of SARS-CoV-2*, Indoor Air, Accepted Author Manuscript,
Oct. 31, 2020, <https://doi.org/10.1111/ina.12766>.

24 ¹⁰ Stadnytskyi, Valentyn, et al., *The airborne lifetime of small speech droplets and their potential*
importance in SARS-CoV-2 transmission, Proc. of the Nat’l Acad. of Sci., Vol. 117, No. 22, pp.
25 11875–77, June 2, 2020, <https://doi.org/10.1073/pnas.2006874117>.

26 ¹¹ Nishiura, Hiroshi, et al., *Closed environments facilitate secondary transmission of coronavirus*
disease 2019 (COVID-19), Apr. 16, 2020, <https://doi.org/10.1101/2020.02.28.20029272>.

27 ¹² *Deciding to Go Out*, CDC, [https://www.cdc.gov/coronavirus/2019-ncov/daily-life-](https://www.cdc.gov/coronavirus/2019-ncov/daily-life-coping/deciding-to-go-out.html)
[coping/deciding-to-go-out.html](https://www.cdc.gov/coronavirus/2019-ncov/daily-life-coping/deciding-to-go-out.html) (last updated October 28, 2020).

28 ¹³ *Id.*

1 because the risk of transmission increases with the size of the gathering. The larger the gathering,
2 the higher the likelihood that an infected person will be present. In addition, the larger the gathering,
3 the greater the number of people at risk of becoming infected at the gathering, who may in turn
4 infect others in the community at large. Limiting the size of gatherings in Santa Clara County is
5 particularly important now, because, as of the date of this declaration, the prevalence of SARS-CoV-
6 2 virus is higher than it has ever been, and the rate of new cases per 100,000 population per day is
7 rising faster than it ever has.

8 38. There is also evidence of risks associated with singing and shouting, which informed
9 the prohibition on singing and shouting in the June 5 order (and in subsequent orders). Research has
10 shown that singing produces more and smaller droplets, as well as aerosols that can travel a longer
11 distance, increasing the risk of infection even with social distancing.¹⁴ Research has also indicated
12 that wearing a mask significantly reduces but does not completely eliminate the increased risk from
13 singing.¹⁵ While loud singing is particularly problematic, shouting and loud talking produce more
14 and smaller droplets as well as aerosols, increasing the risk of transmission.¹⁶

15 39. In June 2020, the available evidence regarding indoor singing included a study of a
16 SARS-CoV-2 superspreading event—that is, an event where one infected person infects a larger
17 number of other people—published by the CDC in May 2020.¹⁷ As reported in the study, 61 people
18 attended a March 10, 2020 choir practice at which one person was known to have COVID-19
19 symptoms. Following that choir practice, 53 cases of SARS-CoV-2 infection were subsequently
20 identified. Three people were hospitalized, and two ultimately died. The study suggested that the
21 act of singing might have contributed to disease transmission through emission of aerosols, which is
22 affected by loudness of vocalization; and it concluded that “[t]he potential for superspreader events
23

24 ¹⁴ Alsvéd, M., et al., *Exhaled respiratory particles during singing and talking*, *Aerosol Sci. & Tech.*,
25 Vol. 54, No. 11, pp. 1245–48, Sept. 17, 2020, <https://doi.org/10.1080/02786826.2020.1812502>.

26 ¹⁵ *Id.*

27 ¹⁶ *Id.*

28 ¹⁷ Hamner, Lea, et al., *High SARS-CoV-2 Attack Rate Following Exposure at a Choir Practice—
Skagit County, Washington, March 2020*, *CDC Morbidity & Mortality Wkly. Rep.*, Vol. 69, No. 19,
pp. 606–10, May 15, 2020, [http://dx.doi.org/10.15585/mmwr.mm6919e6external icon](http://dx.doi.org/10.15585/mmwr.mm6919e6external%20icon).

1 underscores the importance of physical distancing, including avoiding gathering in large groups, to
2 control spread of COVID-19.”

3 40. Since the issuance of the June 5 Order, additional published reports have also
4 addressed the role of singing in disease transmission. A report published by the CDC examined
5 multiple SARS-CoV-2 infections at an overnight camp in Georgia.¹⁸ Five hundred ninety-seven
6 Georgia residents attended the camp, and of the 344 attendees for whom test results were available,
7 260 were positive for SARS-CoV-2. According to the report, indoor cohabitation, singing, and
8 cheering likely contributed to this outbreak: “Relatively large cohorts sleeping in the same cabin and
9 engaging in regular singing and cheering likely contributed to transmission.”

10 41. Public reports have detailed the role of “superspreader” events, including indoor
11 gatherings, in community spread of the virus.¹⁹ For example:

- 12 • In South Korea, as of March 25, at least 5,080 confirmed cases of COVID-19 were
13 traced back to a cluster of cases at a church in Daegu, arising from one 61-year-old
14 person who attended a religious service;²⁰
- 15 • In California, 71 cases of COVID-19 were linked to a church in Sacramento;²¹

16 ///

17
18
19 ¹⁸ Szablewski, et al. (August 2020) *SARS-CoV-2 Transmission and Infection Among Attendees of an
20 Overnight Camp — Georgia, June 2020*,
<https://www.cdc.gov/mmwr/volumes/69/wr/mm6931e1.htm>.

21 ¹⁹ Because SARS-CoV-2 infections in humans were first identified in December 2019, research into
22 the virus, its mechanisms of transmission, and the short, medium, and long-term consequences of
23 infection in humans remains ongoing. The same public health departments that are responding to an
24 unprecedented pandemic are also the ones trying to carve out time to publish outbreak
25 investigations, but the publication process takes time. While press reports are certainly not given the
26 same weight as peer-reviewed research, in the context of this novel pandemic and because public
27 action to stem the pandemic cannot always wait for the peer-review process, the experience of other
28 public health officials and the circumstances that they confront, do inform my thinking regarding
how to protect all people living and working in Santa Clara County.

²⁰ Shin, et al., *How a South Korean church helped fuel the spread of the coronavirus*, Washington
Post, March 25, 2020, available at
<https://www.washingtonpost.com/graphics/2020/world/coronavirus-south-korea-church/>.

²¹ Chabria, et al., *Pentecostal church in Sacramento linked to dozens of coronavirus cases*, Los
Angeles Times, April 2, 2020, available at <https://www.latimes.com/california/story/2020-04-02/pentecostal-church-in-sacramento-linked-to-dozens-of-coronavirus-cases>.

- 1 • In Kentucky, a church revival was linked to at least 28 cases and two deaths;²²
- 2 • In Texas, about 50 people contracted the virus after a pastor told congregants they
- 3 could once again hug one another;²³
- 4 • In West Virginia, there were at least 51 confirmed cases and three deaths tied to the
- 5 resumption of mask-optional services at a church in late May,²⁴ and as of October, 18
- 6 outbreaks in 13 counties have been traced to church services in the State;²⁵
- 7 • In Pennsylvania, a dozen congregants tested positive after a church resumed in-person
- 8 services;²⁶
- 9 • In Ohio, one man with SARS-CoV-2 infection attending a single church service led to
- 10 91 other people becoming infected, included 53 people who had been at the same
- 11 service;²⁷
- 12 • In North Carolina, at least 187 cases and 8 deaths have been linked to a church
- 13 revival;²⁸

14 ///

16 ²² Loosmore, et al., *Kentucky county 'hit really, really hard' by church revival that spread deadly*

17 *COVID-19*, Louisville Courier Journal, April 2, 2020, available at [https://www.courier-](https://www.courier-journal.com/story/news/2020/04/01/coronavirus-kentucky-churchrevival-leads-28-cases-2-deaths/5108111002/)

18 [journal.com/story/news/2020/04/01/coronavirus-kentucky-churchrevival-leads-28-cases-2-](https://www.courier-journal.com/story/news/2020/04/01/coronavirus-kentucky-churchrevival-leads-28-cases-2-deaths/5108111002/)

19 [deaths/5108111002/](https://www.courier-journal.com/story/news/2020/04/01/coronavirus-kentucky-churchrevival-leads-28-cases-2-deaths/5108111002/).

20 ²³ Conger, et al., *Churches Were Eager to Reopen. Now They Are a Major Source of Coronavirus*

21 *Cases*, New York Times, July 8, 2020, available at

22 <https://www.nytimes.com/2020/07/08/us/coronavirus-churches-outbreaks.html>.

23 ²⁴ *Id.*

24 ²⁵ Nakia McNabb, *At least 18 West Virginia COVID-19 Outbreaks Linked to Church Services,*

25 *Governor Says*, CNN, October 19, 2020, available at [https://www.cnn.com/2020/10/19/us/west-](https://www.cnn.com/2020/10/19/us/west-virginia-covid-churches-trnd/index.html)

26 [virginia-covid-churches-trnd/index.html](https://www.cnn.com/2020/10/19/us/west-virginia-covid-churches-trnd/index.html).

27 ²⁶ Whelan, et al., *After Philly's First Church-Linked COVID-19 Outbreak, Pastors Urge Prayers for*

28 *the Sick*, Inquirer, August 21, 2020, available at [https://www.inquirer.com/news/tacony-church-](https://www.inquirer.com/news/tacony-church-covid-outbreak-philadelphia-20200821.html)

29 [covid-outbreak-philadelphia-20200821.html](https://www.inquirer.com/news/tacony-church-covid-outbreak-philadelphia-20200821.html).

30 ²⁷ Peter Grieve, *Defiant Pastor Dismisses COVID-19 as Milder than the Flu Despite Superspreader*

31 *Event that Led to 91 Cases*, August 18, 2020, available at

32 [https://spectrumnews1.com/oh/columbus/news/2020/08/18/after-91-cases--church-pastor-says-virus-](https://spectrumnews1.com/oh/columbus/news/2020/08/18/after-91-cases--church-pastor-says-virus-is-milder-than-flu)

33 [is-milder-than-flu](https://spectrumnews1.com/oh/columbus/news/2020/08/18/after-91-cases--church-pastor-says-virus-is-milder-than-flu).

34 ²⁸ Alison Kuznitz, *More COVID-19 Deaths Linked to Super-Spreader Event at Charlotte Church,*

35 *Charlotte Observer*, November 4, 2020, available at

36 <https://www.charlotteobserver.com/news/coronavirus/article246965397.html>.

- 1 • In Maine, at least 49 cases and three hospitalizations have been linked to a church
2 fellowship event and in-person services in which attendees did not regularly wear
3 masks;²⁹ and after 62 people attended an indoor church wedding, more than 180
4 people have been infected, including at a long-term healthcare facility and a jail, and
5 8 people have died, none of whom attended the wedding.³⁰
- 6 • In Michigan, 187 infections were connected to an indoor bar and restaurant with a
7 live DJ and an open dance floor.³¹ Of the total cases traced back to the restaurant,
8 144 were people who had been to the venue and 43 were family members, friends,
9 and other contacts who had not.
- 10 • In La Cross, Wisconsin, there was a substantial spike in SARS-CoV-2 cases (more
11 than 2,000) with the return to in-person instruction at the community's three
12 universities.³² Researchers were able to use SARS-CoV-2 genomic sequencing to
13 trace COVID-19 clusters at two skilled nursing facilities—and two patient deaths—
14 back to student gatherings and parties.
- 15 • In Boston, Massachusetts, a single person infected with a strain of the virus that
16 contained a particular mutation (which allowed for tracking of related infections),
17 attended a 200-person conference in February 2020, leading to a local superspreading
18 event and ultimately downstream spread across the United States and Europe, likely
19

20 ²⁹ Dustin Wlodkowski, *COVID-19 Outbreak Centered on Maine Church Grows Again*, NBC
21 Boston, October 22, 2020, available at <https://www.nbcboston.com/news/local/covid-19-outbreak-centered-on-maine-church-grows-again/2216323/>.

22 ³⁰ Barbara A. Walsh, *The Uninvited Tragedy: A wedding – then six funerals*, The Maine Monitor,
23 September 17, 2020, available at <https://www.themainemonitor.org/the-uninvited-tragedy-a-wedding-then-six-funerals/>; Roz Plater, *How a Small Wedding in Maine Became a Deadly COVID-19 Superspreader*, Healthline, September 20, 2020, available at <https://www.healthline.com/health-news/how-a-small-wedding-in-maine-became-a-deadly-covid-19-superspreader>.

24 ³¹ Ariana E. Cha, *'Superspreading' events, triggered by people who may not even know they are infected, propel coronavirus pandemic*, The Washington Post, (July 18, 2020),
25 <https://www.washingtonpost.com/health/2020/07/18/coronavirus-superspreading-events-drive-pandemic/>.

26 ³² Richmond, Craig S., et al., *SARS-CoV-2 sequencing reveals rapid transmission from college student clusters resulting in morbidity and deaths in vulnerable populations*, Oct. 14, 2020,
27 <https://doi.org/10.1101/2020.10.12.20210294>.
28

1 causing hundreds of thousands of cases.³³

- 2 • In Massachusetts, more than 200 cases (more than 80% of them asymptomatic) have
- 3 been traced to events held at a single church on or around one specific Sunday.³⁴
- 4 • In Ohio, a wedding with 83 attendees resulted in 32 of the guests becoming infected,
- 5 including guests that had worn face coverings except when eating.³⁵
- 6 • In Washington, an indoor wedding has been linked to one death and is suspected of
- 7 leading to 23 other deaths of persons in long-term care facilities who did not
- 8 attending the wedding, as well as to numerous infections.³⁶
- 9 • In Texas, a funeral attended by more than 100 people, at which most guests wore face
- 10 coverings but did not socially distance, has been linked to more than 40 infections in
- 11 guests ranging from age 3 to age 90.³⁷
- 12 • In California, at least 64 cases have been linked to outbreaks stemming from a church
- 13 that held indoor services at three of its locations.³⁸
- 14 • In North Carolina, at least 75 people tested positive for COVID-19 following and

17
18 ³³ Jacob E. Lemieux, et. al., *Phylogenetic analysis of SARS-CoV-2 in Boston highlights the impact of*
19 *superspreading events* Science, Dec. 10, 2020, available at
20 <https://science.sciencemag.org/content/early/2020/12/09/science.abe3261>.

21 ³⁴ Kaitlin McKinley Becker, *More Than 200 COVID-19 Cases Linked to Fitchburg Church*, NBC,
22 Nov. 7, 2020, available at <https://www.nbc.com/news/local/more-than-200-covid-19-cases-linked-to-fitchburg-church/2225433/?amp>.

23 ³⁵ Jataria McGee, *Cincinnati couple shares story after wedding becomes a 'super spreader' event*,
24 WLWT5, Nov. 18, 2020, available at <https://www.wlwt.com/article/local-couple-shares-story-after-wedding-becomes-a-super-spreader-event/34694908>.

25 ³⁶ Kristin M. Kraemer, *24 COVID deaths possibly tied to Eastern WA superspreader wedding*, Tri-
26 City Herald, Dec. 11, 2020, available at <https://www.tri-cityherald.com/news/coronavirus/article247689390.html>.

27 ³⁷ Alexandria Hein, *Texas funeral became coronavirus superspreader event after 42 were sickened,*
28 *family claims*, Fox News, Nov. 27, 2020, available at <https://www.foxnews.com/health/texas-funeral-became-coronavirus-super-spreader-event-after-42-were-sickened-family-claims>.

³⁸ Brenda Gregorio-Nieto et al., *Health Officials Alert Public Over 2 Additional Outbreaks at*
Awaken Church Locations, December 3, 2020, available at
<https://www.nbcsandiego.com/news/local/health-officials-alert-public-over-2-additional-outbreaks-at-awaken-church-locations/2460223/>.

1 linked to a holiday-themed indoor church event.³⁹

2 **July 2, 2020 Risk Reduction Order and Transition to Harm Reduction Model**

3 42. On July 2, 2020, I issued a risk reduction order that superseded the May 22 Order as
4 amended on June 5. The July 2 Order went into effect at 12:01 a.m. on July 13, 2020.

5 43. By this time, we had significantly increased the County’s capacity to detect cases
6 through widespread testing and to contain disease spread through both broad and focused
7 interventions; expanded our case investigation and contact tracing program and workforce; and
8 increased hospital resources and capacity to treat infected patients. Scientific knowledge about the
9 relative risks of various activities and the primary modes of transmission had also grown
10 significantly (though in many areas it was and continues to be relatively nascent). In addition,
11 residents in the County and greater Bay Area were suffering from “pandemic fatigue.” To ensure
12 more sustained compliance, and in light of our better understanding of key routes of transmission,
13 we transitioned from the initial shelter-in-place model to a less restrictive, longer-term “harm
14 reduction” model—a well-established public health strategy aimed at reducing the risk associated
15 with certain behaviors.

16 44. In light of these circumstances, the July 2 Order allowed most activity, travel, and
17 business operations to resume, but subject to significant conditions and limitations to reduce the risk
18 of SARS-CoV-2 transmission, prevent serious illness and death, and ensure that healthcare resources
19 and capacity remained sufficient to meet the needs of the population.

20 45. The July 2 Order required facilities that pose a high risk of transmission to remain
21 closed to the public, including any indoor facility used for activities where face coverings must be
22 removed, such as indoor dining, bars, and swimming pools.

23 46. The July 2 Order permitted indoor and outdoor gatherings, but because gatherings of
24 any size outside of a single household carry significant risk of exposure to SARS-CoV-2, those
25

26 _____
27 ³⁹ Simone Jasper, *75 COVID Cases Linked to Church Event in North Carolina, Officials Say*,
28 Charlotte Observer, December 17, 2020, available at:
<https://www.charlotteobserver.com/news/state/north-carolina/article247921435.html>.

1 gatherings were subject to restrictions, including facial covering requirements and attendance limits.
2 Indoor gatherings were limited to a maximum of 20 people, or one person per 200 square feet of
3 indoor space, whichever was fewer. Outdoor gatherings were limited to areas large enough to allow
4 for social distancing of all attendees, up to a maximum of 60 people.

5 **July 8, 2020 Mandatory Directive for Gatherings**

6 47. On July 8, 2020, prior to the effective date of the July 2 Order, I issued a Mandatory
7 Directive for Gatherings. The July 8 Directive prohibited indoor gatherings, regardless of size, but
8 permitted outdoor gatherings of up to 60 people, subject to social distancing, use of face coverings,
9 and other restrictions. Under the directive, singing and shouting were not allowed at gatherings,
10 because of the evidence that these activities increase the risk of COVID-19 transmission.

11 48. The directive—including the prohibition on indoor gatherings—was part of a targeted
12 response to case counts that had abruptly started to rise again in the County, neighboring counties,
13 and the region overall. It was also based on my judgment that restrictions on gatherings of people
14 from different households continued to be necessary to reduce community transmission of the virus.
15 Indoor gatherings present a relatively high risk of transmission and infection, particularly gatherings
16 that are large, conducted in a space with poor ventilation, longer in duration, and include activities
17 like loud talking and singing.

18 **July 13, 2020 State Public Health Order**

19 49. Based on the trend of the number of new cases and hospitalizations per day, on July
20 13, 2020, the State Health Officer ordered the closure of indoor operations for particular sectors
21 across the state, including indoor dining, wineries and tasting rooms, family entertainment centers,
22 movie theaters, zoos, museums, and cardrooms. In addition, the State Health Officer ordered the
23 closure of indoor operations for additional sectors in certain counties, including Santa Clara County,
24 effective July 15, 2020. These additional closures included indoor gyms and fitness centers, worship
25 services, protest activities, personal care services, and malls.

26 **July 14, 2020 Mandatory Directives**

27 50. On July 14, 2020, I issued a number of directives in order to tailor the State's general
28 framework to local conditions. Among other directives, I issued an updated Mandatory Directive for

1 Gatherings clarifying that, in addition to the County’s restrictions, the State also prohibited indoor
 2 worship and indoor protest activities. Outdoor worship services and outdoor protests in Santa Clara
 3 County remained subject to the mandatory requirements in the directive.⁴⁰

4 **The County Begins Civil Enforcement of Public Health Orders**

5 51. On August 11, 2020, the County’s elected Board of Supervisors unanimously adopted
 6 an ordinance authorizing civil administrative fines for violations of the State and County Health
 7 Officers’ orders related to the COVID-19 pandemic, including the July 2 Order, the Mandatory
 8 Directive for Gatherings, the July 13 State Public Health Order and subsequent orders, and the
 9 mandatory provisions of the State’s industry-specific guidance. My goal has always been to inform
 10 the public regarding the COVID-19 pandemic to achieve voluntary compliance and safeguard their
 11 health, but I understand that enforcement may nonetheless prove necessary. The long-term viability
 12 of the orders and directives I have issued depends upon robust compliance in the County.

13 **August 28, 2020 State Public Health Order**

14 52. On August 28, 2020, the State Public Health Officer issued a superseding order,
 15 effective August 31, 2020, referred to as the “Blueprint for a Safer Economy.” That order
 16 established a procedure for assigning counties to one of four tiers based on average case rates and
 17 positivity rates of SARS-CoV-2, which in turn determined what activities would be allowed in the
 18 county. Santa Clara County was initially assigned to the most restrictive Tier 1 (Purple), classified
 19 as “widespread” transmission of the novel coronavirus. In Tier 1 counties, indoor church services
 20 are prohibited, but outdoor services are permitted.

21 53. On September 8, 2020, the California Department of Public Health announced that
 22 Santa Clara County had moved to Tier 2 (Red), classified as “substantial” risk of community disease
 23 transmission. Tier 2 counties can allow indoor gatherings and dining of up to 100 people or 25% of
 24 capacity (whichever is less) as well as indoor personal care services with modifications; however,
 25

26
 27 ⁴⁰ On July 20, 2020, a revised Mandatory Directive for Gatherings included in the introductory box
 28 at the top of the document clarifying information regarding the State’s exceptions to its prohibition
 of gatherings.

1 the State's August 28 Order permitted counties to issue and enforce more restrictive measures: "[a]
2 local health jurisdiction may continue to implement or maintain more restrictive public health
3 measures if the jurisdiction's Local Health Officer determines that health conditions in that
4 jurisdiction warrant such measures."

5 54. Given ongoing community transmission and the continuing risk facing county
6 residents, as well as the research establishing the significantly elevated risk associated with indoor
7 gatherings, I decided to maintain the prohibition on indoor gatherings after the County moved to Tier
8 2. Both the State and the County orders continued to prohibit singing and to require face coverings
9 and social distancing of at least six feet at all gatherings. I maintained the prohibition on indoor
10 gatherings because evidence shows that SARS-CoV-2 is spread primarily from person to person
11 through respiratory droplets and aerosols, and, as explained above, that the risk of transmission is
12 higher with indoor activities than outdoor activities.

13 55. The evidence also suggests that indoor gatherings may pose a higher risk of
14 transmission than other kinds of activities that remain subject to different restrictions, including
15 because they are of a sustained duration. For example, when people from different households are
16 together in a grocery or retail store, they typically arrive and depart at different times, and they are
17 together for a shorter duration of time as compared to attendees at a coordinated gathering where
18 attendees linger. Further, grocery and retail shoppers may be less likely to be in close proximity to
19 other shoppers, as opposed to attendees at a gathering who have social connections to one another
20 and therefore may be inclined to have extended conversations in close proximity or physically hug
21 or touch one another. These shoppers are also less likely to engage in higher-risk activities like
22 singing or chanting, as opposed to attendees at an indoor gathering like a church congregation or
23 community sponsored music concert. Thus, the risk of transmission is generally lower in a setting
24 with brief contact between individuals, particularly ones who do not personally know one another
25 and are less likely to speak to or closely interact with one another, as compared to a setting such as a
26 gathering that promotes sustained contact.

27 56. Although I decided not to remove the prohibition on indoor gatherings after the
28 County moved to Tier 2 on September 8, 2020, I revised the Mandatory Directive for Gatherings on

1 September 5, 2020, to relax restrictions on outdoor gatherings consistent with developing research
2 regarding transmission risks and the experiences in other jurisdictions easing certain restrictions.⁴¹
3 Specifically, I removed the prohibition on singing, chanting, and shouting at outdoor gatherings,
4 provided that people wore face coverings and maintained social distancing at all times when engaged
5 in those activities. The directive also permitted playing wind instruments at outdoor gatherings,
6 provided that people placed a cloth covering over the mouth of the instrument. Additionally, I
7 revised the directive to allow the use of canopies, awnings, umbrellas, tents, and other structures for
8 outdoor gatherings provided that three sides or 75% of the structure were open to the air. Finally, I
9 eased restrictions to make it easier for entities to host simultaneous but separate outdoor gatherings
10 of 60 or fewer people by allowing use of the same restroom facilities, provided that there was
11 sufficient restroom capacity or a system of use in place to avoid participants in different gatherings
12 interacting with one another while using the restrooms.

13 **October 5, 2020 Revised Risk Reduction Order**

14 57. On October 5, 2020, I issued a revised risk reduction order that would go into effect
15 and supersede the July 2 Order the day after the California Department of Public Health re-assigned
16 Santa Clara County from Tier 2 to Tier 3 (Orange), classified as “moderate” risk of community
17 disease transmission. On October 13, 2020, the State moved Santa Clara County into Tier 3. Thus,
18 the revised risk reduction order went into effect at 12:01 a.m. on October 14, 2020, and remains in
19 effect as of the date of this declaration.

20 58. To better align the County’s rules with the State’s rules, the October 5 Order allowed
21 many businesses and activities to resume to the extent allowed under the State’s rules for Tier 3
22 counties. However, I decided to maintain stricter restrictions on activities that pose a particularly
23 high risk of a superspreader event, including indoor gatherings.

24 **October 13, 2020 Revised Mandatory Directives**

25 59. On October 13, 2020, the day Santa Clara County moved into Tier 3, I revised a

26
27 ⁴¹ On September 8, 2020, I revised the Mandatory Directive for Gatherings to make clarifying
28 changes to the introductory box at the top of the document.

1 number of directives. I did so based on the trends of new cases and hospitalizations in the County,
2 growing evidence regarding the virus and how it is transmitted, and the experiences of other
3 jurisdictions that similarly eased certain restrictions.

4 60. At that time, I revised the Mandatory Directive for Gatherings to further relax
5 restrictions on outdoor gatherings and to remove the prohibition on indoor gatherings. The directive
6 limited outdoor gatherings to 200 people (up from 60 in the prior directive) and required that such
7 gatherings take place in an area large enough to allow for social distancing of all attendees.
8 Although they were strongly discouraged in the directive, I also allowed indoor gatherings to resume
9 at the level provided in the State's restrictions applicable to Tier 2 counties, which limits such
10 gatherings to 100 people or 25% of the facility's capacity, whichever is fewer. There is growing
11 evidence that reopening facilities at reduced capacity is an effective way to reduce the risk of
12 transmission indoors.⁴² Under the October 13 directive, face coverings were almost always required
13 at all times at all gatherings. Although attendees at outdoor gatherings could temporarily remove
14 their face coverings to eat or drink, food and drink could not be served at any indoor gatherings
15 except as necessary to carry out a religious ceremony given the high risk of transmission indoors.
16 For similar reasons, although permitted at outdoor gatherings subject to specific requirements,
17 singing, chanting, and playing wind instruments remained strictly prohibited at indoor gatherings
18 under this directive. However, singing was allowed indoors if it was for a livestream production,
19 presentation, or service without an audience present, and if the singer remained physically distant
20 from other persons present (who had to keep their face coverings on). As with outdoor gatherings, a
21 venue could host multiple indoor gatherings at the same time subject to specific requirements. If a
22 venue hosted multiple indoor gatherings at the same time, each gathering had to comply with the
23 requirements of the October 5 Order and directive—which included submitting and implementing a
24 Social Distancing Protocol, mandating the use of face coverings in all circumstances required by the
25 State's guidance, complying with the capacity limitations described above, maximizing ventilation,

26 _____
27 ⁴² Chang, Serina, et al., *Mobility network models of COVID-19 explain inequities and inform*
28 *reopening*, Nature, Nov. 10, 2020, <https://doi.org/10.1038/s41586-020-2923-3>.

1 and designating a host responsible for ensuring compliance with all applicable requirements—and be
2 fully separated by solid floor-to-ceiling walls or other partitions from any other gathering.

3 Participants could not mix or enter common spaces with participants from any other gatherings.

4 61. Subsequent studies published since the summer have continued to identify indoor
5 singing—especially without risk mitigation measures like face coverings and social distance—as an
6 activity with a high risk of disease transmission. For example, one study published by *Aerosol*
7 *Science and Technology* in September 2020 concluded that “singing in groups is likely to be an
8 activity at risk of transmitting infection” without appropriate measures, which include “distancing,
9 hygiene, ventilation and shielding.”⁴³ Even with certain risk mitigation measures, however, singing
10 poses an elevated risk of transmission. The study found that although wearing an ordinary surgical
11 face mask reduced the amount of aerosols and droplets a person emitted while singing, the amount
12 was still comparable to the amount emitted by a person speaking without a mask.⁴⁴ And that
13 measurement did not consider the number of particles that may have exited on the sides of masks
14 with a loose fit.⁴⁵ Another recent study published by the *International Journal of Indoor*
15 *Environment and Health* examined the same March 10, 2020 indoor choir rehearsal that was the
16 subject of the CDC study discussed above and likewise concluded that group singing indoors poses a
17 high risk of transmission.⁴⁶

18 **Orchard Community Church and The Home Church’s Compliance with the**
19 **Health Officer’s Orders and Directives**

20 62. I am aware of the lawsuit filed by Gateway City Church, The Home Church, the
21 Spectrum Church of the San Francisco Bay Area, Orchard Community Church of Campbell, and
22

23 _____
24 ⁴³ Alsved, et al. (September 2020) *Exhaled Respiratory Particles During Singing and Talking*,
<https://www.tandfonline.com/doi/full/10.1080/02786826.2020.1812502>.

25 ⁴⁴ *Id.*

26 ⁴⁵ *Id.*

27 ⁴⁶ Miller, et al. (September 2020) *Transmission of SARS-CoV-2 by inhalation of respiratory aerosol*
in the Skagit Valley Chorale superspreading event,
28 <https://onlinelibrary.wiley.com/doi/full/10.1111/ina.12751>.

1 Trinity Bible Church. I am informed that Orchard Community Church held worship services
2 indoors for six Sundays in direct violation of the County's Gatherings Directive and has held
3 worship services indoors, without masks and social distancing. I am also informed that The Home
4 Church held indoor worship services for at least three weeks where the church did not enforce social
5 distancing or require congregants to wear masks while seated next to one another.

6 63. Based on this understanding, it is my view that activities like those of the Orchard
7 Community Church and The Home Church in violation of State and County public health orders is
8 unsafe, detrimental to public health, and poses a significant risk of new infections, leading to
9 potential hospitalization and death for members of the churches as well as others in the broader
10 County community to whom they may spread the disease. It also may encourage other businesses
11 and religious institutions to similarly violate the orders and applicable directives and guidance.
12 Given the ongoing and now rapidly rising community transmission of SARS-CoV-2 in the County, it
13 is only a matter of time before large, indoor gatherings without social distancing or face coverings
14 result in disease transmission; and, unfortunately, these conditions could result in a superspreader
15 event, if they have not done so already. Adverse outcomes from a superspreader event may include
16 multiple infections, hospitalizations, chronic health problems, and even deaths. Consistent with
17 well-established public health principles, this pandemic has demonstrated that community members'
18 adherence to or disregard for public health measures, can either prevent or increase the likelihood of
19 superspreader events, here in the County and across the globe. Even one or two superspreader
20 events could set off multiple chains of transmission with far reaching and deleterious downstream
21 consequences.

22 64. I am aware that Plaintiffs in this litigation have argued there is no evidence that indoor
23 worship has led to COVID-19 in the County generally or at their churches specifically. Even if
24 Plaintiffs are not aware of any COVID-19 cases connected to their churches, that does not mean
25 there have not been any such cases or will not be any such cases. We know that people can have
26 SARS-CoV-2 infection, never develop symptoms (asymptomatic infection), and transmit the virus.
27 An estimated 40% of people with SARS-CoV-2 infection are asymptomatic. I am not aware that
28 any worship services held in the County require testing of congregants for SARS-CoV-2.

1 Moreover, we know that persons who are asymptomatic, but infected, can spread the SARS-CoV-2
2 infection to others—meaning that a person could become infected by SARS-CoV-2 at a worship
3 service held indoors, remain asymptomatic, and unknowingly transmit the infection to persons who
4 may have never set foot into any of the place of worship’s facilities. And some of those persons
5 could become ill, hospitalized, and/or die from COVID-19. Thus, even if Plaintiffs had some type of
6 data that could actually show that none of the persons who have attended their services in the last 6-7
7 months have become ill or died from SARS-CoV-2, they still cannot conclude that no person in the
8 wider community has been infected, because virus from an asymptomatic church member could
9 have been, or could be, transmitted to members of the wider community.

10 65. We need to keep the level of community transmission of SARS-CoV-2 low to protect
11 our most vulnerable residents. By way of example, we have seen a persistent and concerning
12 infection, hospitalization and mortality rate among residents of long term care facilities—both
13 skilled nursing facilities and assisted living facilities—once SARS-CoV-2 is introduced into the
14 facility. During the month of October 2020, the majority of deaths from COVID-19 occurred in
15 residents of our County’s long-term care facilities. We know from experience that once a COVID-
16 19 outbreak starts within a facility, it is exceedingly difficult to control. One pattern that we have
17 seen frequently at long-term care facilities is that a person who works at the facility becomes
18 infected in the community, remains asymptomatic, and unknowingly spreads the infection to others
19 working at or being cared for at the facility. The higher the level of transmission in the community
20 as a whole, the higher the probability that a person who works at that facility will become infected in
21 the community and introduce the virus into a long-term facility and cause an outbreak. Many of
22 these outbreaks lead to death of vulnerable residents. By suppressing community transmission to the
23 greatest extent possible, we can offer greater protection to our most vulnerable residents.

24 **Fall Surge and Revised Mandatory Directives**

25 66. The United States experienced an accelerating surge in COVID-19 cases beginning in
26 early November. As noted above, starting November 4, 2020, the United States has recorded more
27 than 100,000 new cases each day with many days surpassing the previous single-day record and each
28 day’s total eclipsing the summer’s single-day record of 77,300 new cases in mid-July.

1 67. Unfortunately, case counts, positivity rates, and hospitalizations likewise began to
2 accelerate rapidly in Santa Clara County. For example, as of October 24, 2020, there had been
3 24,014 cases confirmed in the County, the seven-day running average of new cases per day was 114,
4 and 388 County residents had died with COVID-19. In comparison, as of November 29, 2020, there
5 had been 33,732 cases confirmed in the County, the seven-day running average of new cases per day
6 was 417, and 476 County residents had died with COVID-19. In other words, the seven-day running
7 average of new cases in the County more than *tripled* during that time period. The County also saw
8 a sharp rise in hospitalizations for COVID-19 during the month of November: The number of
9 people hospitalized with COVID-19 went from 86 people on November 1, 2020; to 135 people on
10 November 14, 2020; to 272 people on November 28, 2020. And those numbers have gotten
11 dramatically worse in December: As of December 22, 2020, there had been 57,452 cases confirmed
12 in the County; the most recent seven-day running average of new cases per day was 1,183; 607
13 County residents had died with COVID-19; and 621 people were hospitalized with COVID-19. The
14 seven-day running average of new cases in the County thus almost tripled again from November to
15 December.

16 68. In response to the rapid rise in case rates and increasing positivity rate and
17 hospitalizations in Santa Clara County, the Bay Area, and California, I determined that it was
18 necessary to impose renewed restrictions on certain high-risk activities in order to blunt the rapid
19 rise in case rates and prevent the expected additional hospitalizations and deaths that generally
20 follow. Other Bay Area officials made similar determinations. As one example, effective
21 November 17, 2020, the Mandatory Directive for Dining, Bars, Wineries, and Smoking Lounges
22 prohibits indoor dining and indoor tasting activities at wineries based on the consensus among public
23 health experts that indoor activities, especially without face coverings, pose a higher risk of
24 transmission than outdoor activities.

25 69. On November 16, 2020, the State announced that Santa Clara County would move from
26 Tier 3 (Orange) directly into the most restrictive Tier 1 (Purple), effective November 17, 2020.
27 Impacts of this reassignment included the closure of indoor dining, which had already been planned
28 in the County, as well as the closure of all indoor activities associated with gyms, museums, zoos,

1 and aquariums. Indoor gatherings, including at places of worship and movie theaters, were also
2 prohibited. In addition, shopping malls and retail establishments were required to reduce their
3 indoor capacity to 25%.

4 70. In addition, effective November 17, 2020, the new Mandatory Directive on Capacity
5 Limitations established capacities for various sectors and activities.⁴⁷ I issued this directive based on
6 the growing evidence that reducing capacity is an effective way to reduce the risk of transmission
7 indoors.⁴⁸ I also revised the Mandatory Directive on Gatherings to specify that the State's
8 determination that the County has been assigned to the Purple Tier resulted in the prohibition of all
9 indoor gatherings, effective November 16, 2020.

10 **November 19, 2020 State Public Health Order**

11 71. On November 19, 2020, the State Public Health Officer issued a temporary order
12 referred to as the "Limited Stay at Home Order." This order, which was issued to address the
13 unprecedented fall surge of COVID-19 cases described above, took effect on November 21, 2020,
14 and is scheduled to remain in effect until December 21, 2020, unless extended or revised by the State
15 Public Health Officer.

16 72. To reduce opportunities for transmission of COVID-19 in the community, the State's
17 Limited Stay at Home Order placed additional restrictions on counties in Tier 1 (Purple) of the
18 State's Blueprint for a Safer Economy. In those affected counties, the State's Limited Stay at Home
19 Order prohibited all activities involving interaction or gathering—either indoors or outdoors—with
20 members of other households between the hours of 10:00 p.m. and 5:00 a.m., except (a) for certain
21 activities associated with the operation, maintenance, or use of critical infrastructure, as specified on
22 the State's Essential Critical Infrastructure Workers list, or (b) for those activities required by law.

23 73. The stated intent of the State's Limited Stay at Home Order was to decrease the amount
24 of time individuals would be mixing in the community with people outside their own households, as
25

26 ⁴⁷ Before this directive, businesses and community members had to refer to multiple directives, as
27 well as State guidance, to determine applicable capacity limitations (if any) for particular sectors or
28 activities.

⁴⁸ Chang, *supra*.

1 such interactions present opportunities for disease transmission. The Limited Stay at Home Order
2 placed restrictions from 10:00 p.m. to 5:00 a.m. because the State Public Health Officer determined
3 that activities and gatherings that occur during that timeframe are often non-essential, social
4 activities during which reduced inhibitions would more likely lead to disregard of COVID-19
5 preventative measures, such as maintaining physical distance or using face coverings. Thus, under
6 this Limited Stay at Home Order, individuals in Tier 1 (Purple) counties were prohibited from
7 engaging in non-essential activities—like dining outdoors at a restaurant or going to an outdoor
8 movie—between the hours of 10:00 p.m. and 5:00 a.m.

9 **November 30, 2020 Revised Mandatory Directives and Travel Directive**

10 74. By late November, the number of Santa Clara County residents contracting COVID-
11 19 and the number of patients hospitalized with COVID-19 had continued to rise significantly. On
12 November 28, 2020, there were 760 new cases of COVID-19 and 239 COVID-related
13 hospitalizations, 71 of whom were in the ICU. Based on infectious disease models by colleagues at
14 UCSF and Stanford, the rapidly rising numbers of patients needing hospitalization for COVID was
15 projected to exceed the available staffed hospital beds by the second or third week of December if
16 actions to dramatically decrease levels of community transmission were not taken immediately.

17 75. To reduce the likelihood of a surge in hospitalizations that would exceed the capacity
18 of hospitals in the County, I made several changes to the Mandatory Directives, including requiring
19 retail stores, limited service facilities, and most other facilities open to the public to be limited to
20 10% capacity indoors⁴⁹; requiring facilities open to the public to establish a “metering system” to
21 ensure applicable capacity limits are not exceeded; and closing cardrooms; among other changes. I
22 also reduced the maximum number of people who could attend an outdoor gathering from 200
23 people to 100 people.

24 76. The risk of COVID-19 transmission in Santa Clara County increases as people travel
25 in and out of the county because those travelers interact with members of other communities,

26 _____
27 ⁴⁹ Grocery stores, drug stores, and pharmacies were allowed to operate at 25% capacity indoors to
28 ensure adequate access to food and medicine.

1 including communities where the prevalence of COVID-19 may be higher than in Santa Clara
 2 County. Therefore, in a further attempt to stem the significant increases in COVID-19 cases and
 3 related hospitalizations in Santa Clara County, I issued, on November 28, 2020, a Mandatory
 4 Directive on Travel discouraging all travel, especially for non-essential purposes. This Mandatory
 5 Directive on Travel requires most travelers who nevertheless travel into Santa Clara County from
 6 more than 150 miles away to quarantine, meaning they are prohibited from having contact with
 7 individuals from outside of their household or their immediate traveling party. Quarantine after
 8 travel will ensure that long-distance travelers do not introduce new COVID-19 cases (and,
 9 potentially, subsequent chains of transmission) into the Santa Clara County community from their
 10 location of origin. And requiring long-distance travelers to quarantine will lower the risk that
 11 asymptomatic or pre-symptomatic COVID-19-positive travelers will unwittingly transmit the disease
 12 to others in the county.

13 77. These revised and new Mandatory Directives took effect on Monday, November 30,
 14 at 12:01 a.m. and will remain in effect until at least December 21, 2020 at 5:00 a.m. unless they are
 15 extended.⁵⁰

16 **December 3, 2020 State Regional Stay At Home Order**

17 78. In response to the continuing fall surge of COVID-19 cases in California, which had
 18 led to a dramatic increase in hospitalizations throughout the state, the State issued a Regional Stay at
 19 Home Order on December 3, 2020, in an attempt to keep hospitals across the state from being
 20 overwhelmed.

21 79. Under the Regional Stay at Home Order, the State monitors the hospital intensive
 22 care unit (“ICU”) capacity in each of five geographic regions established by the State. The State
 23 placed Santa Clara County in the “Bay Area” region with Alameda, Contra Costa, Marin, Monterey,
 24 Napa, San Francisco, San Mateo, Santa Clara, Santa Cruz, Solano, Sonoma Counties. If a region’s
 25 hospital ICU capacity falls below 15%, it triggers the Regional Stay at Home Order’s restrictions on
 26

27 ⁵⁰ On December 2, 2020, I issued an amended Mandatory Directive on Travel that clarified the
 28 exemptions from the mandatory quarantine requirement for some travelers.

1 the types of activities permitted in the region. Once a region falls below the 15% trigger, residents
2 are prohibited from gathering with members of other households—indoors or outdoors—at any time
3 of day, with limited outdoor exceptions for worship services, political activities, and cultural
4 ceremonies. Residents in the region must stay at home or at their place of residence at all times,
5 except for (a) certain activities associated with the operation, maintenance, or use of critical
6 infrastructure, as specified on the State’s Essential Critical Infrastructure Workers list, (b) activities
7 required by law, or (c) activities specifically permitted in the Regional Stay at Home Order.

8 80. When a region falls below the 15% hospital ICU capacity trigger, the Regional Stay
9 at Home Order prohibits most activities that would otherwise be permitted in the region, even under
10 Tier 1 (Purple), which is the most restrictive tier in the State’s Blueprint for a Safer Economy. For
11 example, the following activities would be prohibited in a region that has fallen under the 15%
12 trigger, even though such activities would otherwise be permitted even under Tier 1: dining outdoors
13 at a restaurant; receiving personal care services, such as a haircut, a manicure, or a tattoo; going to
14 an outdoor zoo or museum; and gaming at an outdoor cardroom. In my opinion, restricting these
15 types of activities is important to slow the rate of disease transmission and to keep the region’s
16 healthcare system from becoming overwhelmed in this moment when the region’s healthcare system
17 is under significant strain.

18 81. In addition to prohibiting certain activities in regions that have hit the 15% hospital
19 ICU capacity trigger, the Regional Stay at Home Order also imposes further restrictions on those
20 businesses that are allowed to continue operating in those regions. For example, the Regional Stay
21 at Home Order imposes stricter capacity limitations on retailers. Retailers, other than stand-alone
22 grocery stores, are required to reduce the number of people inside their indoor facility to 20% of
23 normal capacity, down from the 25% capacity limit in Tier 1 (Purple). Stand-alone grocery stores
24 must reduce their capacity to 35% of normal, down from the 50% capacity limit in Tier 1 (Purple).⁵¹

25
26
27 ⁵¹ The Regional Stay at Home Order issued on December 3, 2020, originally limited all retailers,
28 including stand-alone grocers, to a 20% capacity. However, the State amended the capacity
limitation on stand-alone grocery store, increasing it to 35%, in its December 6, 2020, Supplement to
Regional Stay at Home Order.

1 In my opinion, reducing the capacity of indoor retail spaces is an important tool to lower the
 2 opportunities for disease transmission, because reducing the number of people in an indoor space
 3 reduces the volume of respiratory droplets and aerosols being released into that space, and provides
 4 more room for individuals to stay physically distanced from one another. The use of occupancy or
 5 capacity limits is recommended by the CDC,⁵² and its efficacy is supported by current models of
 6 mobility and disease transmission.⁵³

7 **December 4, 2020 Mandatory Directive Implementing**
 8 **State's Regional Stay at Home Order**

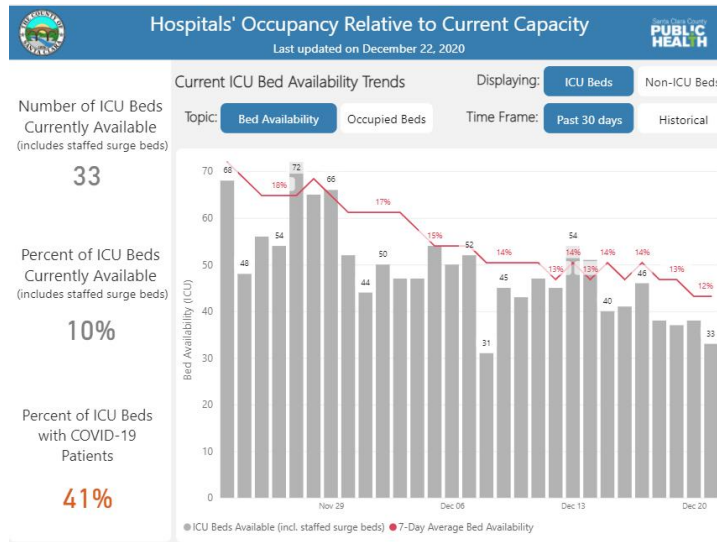
9 82. On December 4, 2020, I, along with the Health Officers for the Counties of Alameda,
 10 Contra Costa, Marin, and San Francisco, as well as the City of Berkeley, jointly announced that our
 11 respective jurisdictions would implement the State's Regional Stay at Home Order early, rather than
 12 waiting for our Bay Area region to fall below the 15% hospital ICU capacity trigger, which at that
 13 time was expected to occur in mid- to late-December.

14 83. I made this decision for Santa Clara County based on the dramatically rising case
 15 rates and hospitalization rates in the county. While the Bay Area Region's hospital ICU capacity was
 16 still above the 15% trigger at the time of my decision, Santa Clara County's hospital ICU capacity
 17 had already fallen below 15%. Furthermore, because of the significant lag between when people
 18 become infected with COVID-19 and when they may need to be hospitalized, I was concerned that
 19 Santa Clara County's healthcare system was at imminent risk of being overwhelmed. Thus, I
 20 determined that immediate further action was necessary to slow the spread of COVID-19 in the
 21 county. Since the December 4, 2020 announcement, ICU capacity has continued to fall, as reflected
 22 in the data published on the County's COVID-19 Hospitalizations Dashboard as of December 23,
 23
 24

25 _____
 26 ⁵² Honein MA, et al., *Summary of Guidance for Public Health Strategies to Address High Levels of*
 27 *Community Transmission of SARS-CoV-2 and Related Deaths*, December 2020. MMWR Morb
 28 *Mortal Wkly Rep* 2020;69:1860-1867. DOI: <http://dx.doi.org/10.15585/mmwr.mm6949e2>.

⁵³ Chang S., et al., *Mobility network models of COVID-19 explain inequities and inform reopening*.
 Nature (2020). <https://doi.org/10.1038/s41586-020-2923-3>

1 2020:



11 84. The County’s Mandatory Directive Implementing State’s Regional Stay at Home
 12 Order took effect on December 6, 2020, and will remain in place until January 4, 2021, unless
 13 otherwise rescinded, modified, or extended.

14 85. As part of my decision to implement the State’s Regional Stay at Home Order, I also
 15 amended the County’s Mandatory Directive on Capacity Limitations. The December 4, 2020,
 16 amendments to the Mandatory Directive on Capacity Limitations require all retailers to reduce the
 17 number of people inside their indoor facility to 20% of normal capacity, aligning the County’s
 18 capacity limitations on retailers with those of the State and the surrounding jurisdictions. Like the
 19 State’s Regional Stay at Home Order, the County’s revised Mandatory Directive on Capacity also
 20 prohibited or continued to prohibit restaurants, card rooms, personal care businesses, and non-
 21 essential limited services, like pet groomers, from operating indoors or outdoors.

22 86. On December 4, 2020, I also amended the County’s Mandatory Directives on
 23 Gathering to clarify that under the Regional Stay at Home Order’s, both indoor and outdoor
 24 gatherings with members of other households are prohibited, except for outdoor worship services,
 25 political events, car-based gatherings, and cultural ceremonies of up to 100 people.

26 * * *

27 87. As the surging case rates demonstrate, there is still ongoing and accelerating
 28 community transmission of SARS-CoV-2 in the County, and a recent national seroprevalence study

1 estimates that the vast majority of California residents—more than 96%—have not been exposed to
2 the virus and still have no immunity.⁵⁴ As such, our community and the greater Bay Area are still
3 facing the risk of uncontrolled new infections, illnesses, hospitalizations, and deaths. In light of
4 these ongoing risks, accepted public health principles counsel a cautious and incremental approach.
5 Keeping a risk reduction order in place remains critical to our ability to protect the community and
6 control the ongoing surge of COVID-19 cases.

7 88. I understand that a minority of public commentators have called for a more hands-off
8 approach than the one I have taken. Some of these commentators urge a “herd immunity” approach,
9 relying on the assumption that uncontrolled spread of the virus will create widespread immunity
10 against later infection and eventually bring the pandemic to an end. I agree with the vast majority of
11 infectious disease epidemiologists and public health professionals who reject the “herd immunity”
12 strategy. Achieving such immunity would take a significant amount of time—some estimate as long
13 as 18 to 24 months⁵⁵—and would almost certainly result in a staggering number of infections and
14 deaths.

15 89. Santa Clara County, the Bay Area, the State of California, the United States, and the
16 rest of the world have not seen a public health threat like this in more than 100 years. The SARS-
17 CoV-2 virus has infected persons of all ages and all health conditions. But this public health
18 emergency will not last forever. Progress has been made on therapeutic treatments for COVID-19,
19 including antivirals, monoclonal antibodies, and convalescent plasma therapy.⁵⁶ In addition, two
20 vaccines for SARS-CoV-2 have recently received Emergency Use Authorization, initial allocations
21 have already arrived in Santa Clara County, and vaccination of healthcare personnel and other

23
24 ⁵⁴ Shuchi, Anand, et al., *Prevalence of SARS-CoV-2 antibodies in a large nationwide sample of patients on dialysis in the USA: a cross-sectional study*, *The Lancet*, Vol. 396, Issue 10259, pp. 1335–44, Oct. 24, 2020, [https://doi.org/10.1016/S0140-6736\(20\)32009-2](https://doi.org/10.1016/S0140-6736(20)32009-2).

25
26 ⁵⁵ Moore, Kristine A., et al., *COVID-19: The CIDRAP Viewpoint*, Ctr. For Infectious Disease Res. & Pol. (Apr. 30, 2020), https://www.cidrap.umn.edu/sites/default/files/public/downloads/cidrap-covid19-viewpoint-part1_0.pdf.

27
28 ⁵⁶ Shawn Radcliffe, *Here’s Exactly Where We Are with Vaccines and Treatments for COVID-19*, Healthline (Dec. 8, 2020), <https://www.healthline.com/health-news/heres-exactly-where-were-at-with-vaccines-and-treatments-for-covid-19>.

1 priority groups has already begun. Numerous potential other vaccines for SARS-CoV-2 are in
2 clinical trials.⁵⁷

3 90. My staff and I continue to monitor a variety of public health indicators, including the
4 trend of the number of new cases and hospitalizations per day; the positivity rate; the number of
5 deaths; the location and character of outbreaks; especially those occurring in long-term care
6 facilities; and the capacity of the healthcare system in the County and the region, to provide care
7 during the current surge in cases and hospitalizations. We also continue to monitor our capacity to
8 effectively respond and protect the public, including our capacity to efficiently and accurately test
9 persons for SARS-CoV-2, especially in high-risk populations and settings; to conduct effective case
10 investigation and contact tracing; to support persons who are isolating or quarantining; and to
11 prevent and control outbreaks in long-term care and other settings with a concentrated population of
12 vulnerable persons. We also continue to review the developing research regarding SARS-CoV-2
13 and the disease it causes. Finally, we continue to collaborate with and monitor other jurisdictions
14 implementing various public health measures to address the pandemic.

15 91. The graph below demonstrates that throughout the pandemic, our shelter-in-place and
16 risk reduction orders, as well as the related mandatory directives, have been responsive both to the
17 state of the pandemic in the County and to the needs of residents to take part in the activities that are
18 important to their lives. As case counts have risen and fallen, my staff and I have adjusted the
19 County guidelines based on established public health principles. In particular, when the growth in
20 new cases slowed, we have been able to ease restrictions on daily activities and gatherings.

21 ///

22 ///

23 ///

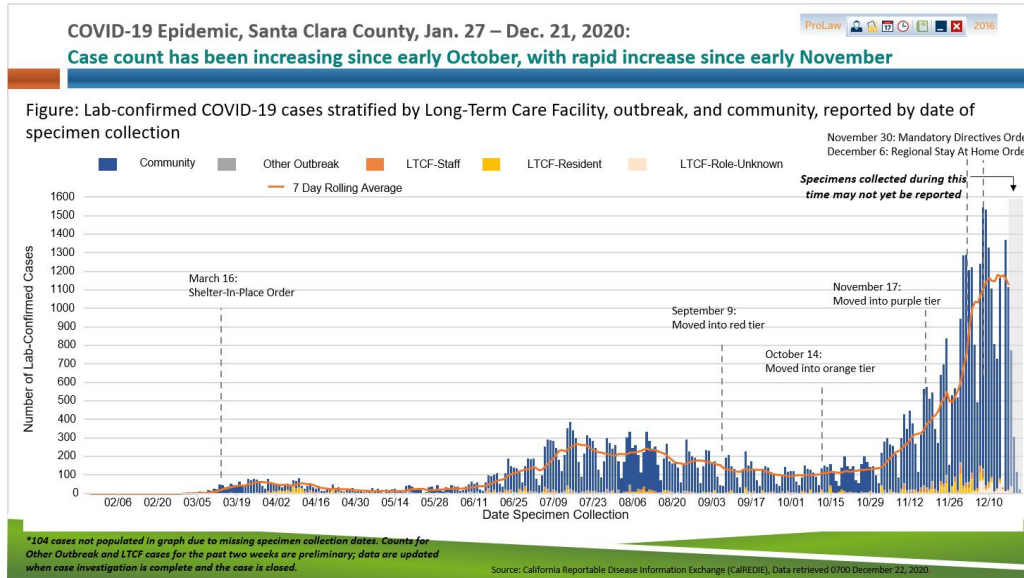
24 ///

25 ///

26 ///

27

28 _____
⁵⁷ *Id.*



92. The graph also reflects that when we have loosened restrictions, case counts have increased. For example, in response to a flattened curve in April and May 2020, on June 5, 2020, we updated our shelter-in-place order to allow additional commercial and other activity. This relaxation was followed by an increase in cases. As cases are again on the rise, now is not the time to ease restrictions on community transmission or lift the Risk Reduction Order and Mandatory Directives.

* * *

93. I—along with many health officials from all over the world—have concluded that the best course of action for managing the pandemic and preventing a surge in deaths from COVID-19 while scientists continue to develop treatments and while recently approved vaccines become available to a greater portion of the population, is to prevent the infection from spreading using the tools we currently have available. These tools include staying at home, maintaining social distancing when outside the home, conducting business and activities outdoors whenever possible, wearing a face covering when interacting with others outside of one’s household, and limiting the number and duration of contact with others to the greatest degree possible. We do not take lightly asking people to take these steps. We understand that people are suffering from real impacts on all aspects of their lives and livelihoods. We each have a part in slowing the spread of this virus. Right now, we all urgently need to reduce the number of people with whom each of us comes in contact and the intensity of those contacts. These actions will save lives. The better we all do today, the

1 sooner this pandemic will end.

2 I declare under penalty of perjury under the laws of the State of California that the foregoing
3 is true and correct. Executed in San José, California on December 23, 2020.

4

5

/s/ Sara H. Cody
SARA H. CODY, M.D.

6

7

8

9

2329525

10

11

12

13

14

15

16

17

18

19

20

21

22

23

24

25

26

27

28

Exhibit 3

1 JAMES R. WILLIAMS, County Counsel (S.B. #271253)
HANNAH KIESCHNICK, Deputy County Counsel (S.B. #319011)
2 KARAN SINGH DHADIALLA, Deputy County Counsel (S.B. #296313)
OFFICE OF THE COUNTY COUNSEL
3 70 West Hedding Street, East Wing, Ninth Floor
San José, California 95110-1770
4 Telephone: (408) 299-5900
Facsimile: (408) 292-7240

5 Attorneys for Defendants
6 COUNTY OF SANTA CLARA, SARA H. CODY

7
8 UNITED STATES DISTRICT COURT
NORTHERN DISTRICT OF CALIFORNIA
9 (San José Division)

10
11 GATEWAY CITY CHURCH, et al.,

12 Plaintiffs,

13 v.

14 GAVIN NEWSOM, et al.,

15 Defendants.

No. 20-CV-08241-EJD

DECLARATION OF DR. MARC LIPSITCH IN SUPPORT OF DEFENDANTS COUNTY OF SANTA CLARA AND DR. SARA H CODY'S OPPOSITION TO PLAINTIFFS' MOTION FOR A PRELIMINARY INJUNCTION

Date: January 15, 2020
Time: 10:00 a.m.
Courtroom: 4
Judge: Hon. Edward J. Davila

16
17
18 I, DR. MARC LIPSITCH, declare as follows:

19 1. I am a resident of Jamaica Plain, Massachusetts. I have personal knowledge of the
20 matters set forth below and could testify competently to them if called to do so.

21 **Professional Background**

22 2. I am the founding Director of the Center for Communicable Disease Dynamics
23 (CCDD) at the Harvard School of Public Health, where I am also a Professor in the Department of
24 Epidemiology and the Department of Immunology and Infectious Diseases. I have a B.A. in
25 Philosophy from Yale University and a DPhil (the Oxford equivalent of a PhD) in Zoology from the
26 University of Oxford, which I attended as a Rhodes Scholar. After receiving my DPhil, I completed
27 a postdoctoral research fellowship in Biology at Emory University working on the population
28 biology of infectious diseases. A copy of my C.V. is attached to this declaration as Exhibit A. I

1 have been elected as a Fellow of the American Academy of Microbiology and a member of the
2 National Academy of Medicine for the United States.

3 3. CCDD is a research center committed to advancing our understanding of infectious
4 disease and training the next generation of scientists. It was founded as a Center of Excellence in the
5 Models of Infectious Disease Agent Study funded by the National Institute of General Medical
6 Sciences of the U.S. National Institutes of Health (NIH). CCDD has been at the leading edge of
7 epidemiology, pioneering new approaches and methodologies for investigating both recurring and
8 emerging problems. The goal of our work at CCDD is to understand why and how infectious
9 disease persists and changes and use that knowledge to lessen its burden on people.

10 4. CCDD is closely monitoring the progress of COVID-19. CCDD faculty—including
11 me—are conducting research on the novel coronavirus SARS-CoV-2 and COVID-19, the disease it
12 causes. CCDD faculty have published over 50 peer-reviewed articles about SARS-CoV-2 and
13 COVID-19. I am the lead or a contributing author on more than 20 of those articles. CCDD faculty
14 regularly host and contribute to online events about COVID-19; appear in national and international
15 media, including print and broadcast news; participate in scientific conferences, consortia,
16 discussions, debates, and podcasts; and advise local, state, and federal officials and leaders of
17 countries around the world.

18 5. My own work on COVID-19 has included epidemiology, mathematical modeling,
19 and exploration of ethical issues related to vaccine trials and school reopenings. My research has
20 helped identify countries with undetected cases before they were reported; modeled the effects of
21 various social distancing and quarantine strategies; and contributed to some of the earliest estimates
22 of case-fatality rates. My research has also addressed new methodologies on how to study immunity
23 to COVID-19; and I am a co-lead on a large collaborative effort, led by experts at the University of
24 Chicago and involving multiple European universities, to establish best practices for estimating the
25 contagiousness of the virus. My work has also addressed the ethical aspects of COVID-19 vaccine
26 trial design, including the first published proposal for human challenge studies, which received
27 support from the World Health Organization (WHO) and the NIH, and which was implemented in
28 the U.K.

1 6. I have advised the WHO, the International Monetary Fund, the Prime Minister of
2 Israel, and senior government officials in the U.S., Canada, India, Germany, Austria, and
3 Luxembourg on COVID-19, as well as the U.S. National Governors' Association and numerous state
4 and local health officials. I am a member of the Massachusetts Governor's Medical Advisory
5 Committee and the Massachusetts COVID-19 Vaccine Working Group. I am also an ad hoc expert
6 to the COVID-19 Vaccine Working Group, which is part of the WHO Strategic Advisory Group of
7 Experts. Health departments on several continents use software that I helped develop to update their
8 estimates of trends in COVID-19 cases.

9 7. Over the course of the COVID-19 pandemic, I have been asked to provide and have
10 provided interviews and analysis to national and international media outlets, including CNN, BBC,
11 the *Guardian* and the *Wall Street Journal*; and I have published articles explaining aspects of the
12 COVID-19 pandemic in national and international media outlets, including the *New York Times* and
13 *Washington Post*. My public science communication efforts also include a Twitter account with an
14 active following. Earlier this year, physicist Jonathan Oppenheim reported that I was the second-
15 most-followed expert by other experts on the COVID-19 pandemic; and I was named by Forbes as
16 one of the "most essential people on Twitter to follow during the COVID-19 outbreak."

17 8. More generally, my research has focused on biological and mathematical approaches
18 to infectious disease questions—mainly understanding how our immune systems and medical
19 interventions such as antibiotics and vaccines exert natural selection on pathogens, and how the
20 resulting changes in pathogen populations affect human disease. My more recent work has focused
21 on antimicrobial resistance, epidemiological methods, mathematical modeling of infectious disease
22 transmission, pathogen population genomics, immunoepidemiology of *Streptococcus pneumoniae*,
23 transmission-dynamic simulations, and ethical questions surrounding vaccine trials for infectious
24 disease.

25 9. My work has addressed a number of issues relevant to modern pandemic responses.
26 My research provided modern evidence of the moderate contagiousness of the 1918 "Spanish flu."
27 During the first SARS outbreak in 2003, I led a team that provided one of the first estimates of the
28 virus' reproduction number. During the 2009 H1N1 pandemic, my research produced the first

1 reliable estimate of H1N1 flu severity. During the yellow fever outbreak in Angola and Democratic
2 Republic of Congo in 2016, my modeling work helped support fractional dosing vaccination
3 strategies, which helped extend vaccine availability in a shortage situation. I have written
4 extensively on data-driven decision making in public health.

5 10. I have worked extensively with governments and intergovernmental bodies like WHO
6 to address public health issues including pandemic response and preparedness. For example, in 2003
7 and 2004, I served on the Defense Science Board Task Force on the SARS Quarantine for the U.S.
8 Department of Defense. In 2009, I was a member of the H1N1 Working Group of the U.S.
9 President's Council of Advisors on Science and Technology; and in 2009 and 2010, I was a member
10 of the Team B Advisory Body to the CDC on the Novel H1N1 Influenza. From 2017 through today,
11 I have been a member of the Biological Agents Containment Working Group of the Board of
12 Scientific Counselors to the Office of Public Health Preparedness and Response at the CDC.

13 11. I have also worked extensively on the design and analysis of vaccine trials during
14 public health emergencies. In 2015, I served on a scientific advisory board for a major Ebola
15 vaccine trial, and as I mentioned above, I am currently advising Massachusetts and WHO on
16 COVID-19 vaccine issues.

17 12. I have published more than 330 peer-reviewed articles and a large number of other
18 publications, including book chapters, non-peer-reviewed journal articles, and popular articles in the
19 national press. I have also contributed to a number of reports, including the President's Council of
20 Advisors on Science and Technology (PCAST) H1N1 Working Group's 2010 Report to the
21 President on US Preparations for 2009-H1N1 Influenza; three reports from the Center for Infectious
22 Disease Research and Policy (CIDRAP) regarding the development of a vaccine for the Ebola virus;
23 and most recently an April 2020 CIDRAP report on COVID-19.¹ CIDRAP is based out of the
24 University of Minnesota and is a global leader in addressing public health preparedness and
25

26
27 ¹ Moore, K., et al., *COVID-19: The CIDRAP Viewpoint*, April 30, 2020, available at
28 https://www.cidrap.umn.edu/sites/default/files/public/downloads/cidrap-covid19-viewpoint-part1_0.pdf.

1 emerging infectious disease response.

2 13. I continue to teach and mentor undergraduate and graduate students at Harvard, as
3 well as supervising graduate work for doctoral candidates.

4 **Opinions Regarding Dr. Bhattacharya's Declaration**

5 14. The defendants in this case contacted me about responding to the opinions expressed
6 by Dr. Jayanta Bhattacharya in his declaration submitted by the plaintiffs. I agreed to provide a
7 declaration setting forth some of my professional opinions on the issues raised in that declaration. In
8 reaching those opinions, I have relied on my knowledge, training, experience, and the kinds of data
9 regularly relied on by experts in my field. I am working pro bono and not being compensated for my
10 time.

11 15. I have read the declaration of Dr. Jayanta Bhattacharya submitted by the plaintiffs in
12 this lawsuit. My opinions regarding that declaration are based on the available science regarding the
13 novel coronavirus SARS-CoV-2 and the disease it causes, COVID-19, as well as my training and
14 experience in infectious disease response.

15 16. As Dr. Bhattacharya discusses in his declaration, he has recommended an approach to
16 COVID-19 that is commonly referred to as "herd immunity with focused protection." This approach
17 was laid out in the so-called "Great Barrington Declaration," a document published in October at a
18 ceremony at a libertarian think tank by three scientists, including Dr. Bhattacharya. In this approach,
19 the virus would be allowed to spread among young, healthy people with little attempt to slow it
20 down, while officials would try to keep older, more vulnerable Americans from contracting it. This
21 strategy diverges sharply from the views of most infectious-disease epidemiologists and has been
22 rejected by the National Institute of Allergy and Infectious Diseases Director Dr. Anthony Fauci,²
23 WHO Director-General Tedros Adhanom Ghebreyesus,³ and the more than 6,900 scientists,

24 _____
25 ² Mandavilli A., et al., *A Viral Theory Cited by Health Officials Draws Fire From Scientists*, New
26 York Times, Oct. 19, 2020, available at <https://www.nytimes.com/2020/10/19/health/coronavirus-great-barrington.html> (accessed Nov. 16, 2020).

27 ³ *WHO chief says herd immunity approach to pandemic 'unethical'*, The Guardian, Oct. 12, 2020,
28 available at <https://www.theguardian.com/world/2020/oct/12/who-chief-says-herd-immunity-approach-to-pandemic-unethical> (accessed Nov. 16, 2020).

1 researchers, and healthcare professionals who have signed a formal response called the John Snow
 2 Memorandum.⁴ Without a vaccine, this strategy also risks the deaths of a million or more
 3 Americans. With the FDA's recent emergency-use approval of Pfizer's and Moderna's vaccines,
 4 and given the likelihood of FDA approval of additional effective vaccines, the strategy risks
 5 significant avoidable illness and death.

6 **COVID-19 Is Not Harmless for Younger Populations**

7 17. The assumption underlying the herd immunity approach—that COVID-19 is harmless
 8 to most people and risky only for defined groups—is false.

9 18. The impact of a pandemic on health depends not only on the infection-fatality rate,
 10 but also on other measures of severity such as the risk of hospitalization or ICU admission among
 11 those infected. Crucially, it also depends on the number of people who become infected, because a
 12 small risk of death, ICU or hospitalization multiplied by a large number of people infected can result
 13 in large numbers of deaths and high burdens on health care resources. Indeed, the extraordinarily
 14 high peak demand for intensive care in Wuhan, China⁵ and in Northern Italy⁶ were two of the
 15 earliest warnings that uncontrolled SARS-CoV-2 spread could result in horrific burdens on the
 16 health care system. The intense stress on even very high-quality health systems is being felt across
 17 Europe and in many parts of the U.S. as of December 2020, with overloaded intensive care units in
 18 multiple locations due to COVID-19 surges. The U.S. has hit its highest number to date of
 19 hospitalizations, with more than 115,000 COVID-19 patients in hospital as of December 21, 2020:⁷

20 ///

21 ///

22 _____
 23 ⁴ Available at <https://www.johnsnowmemo.com/>.

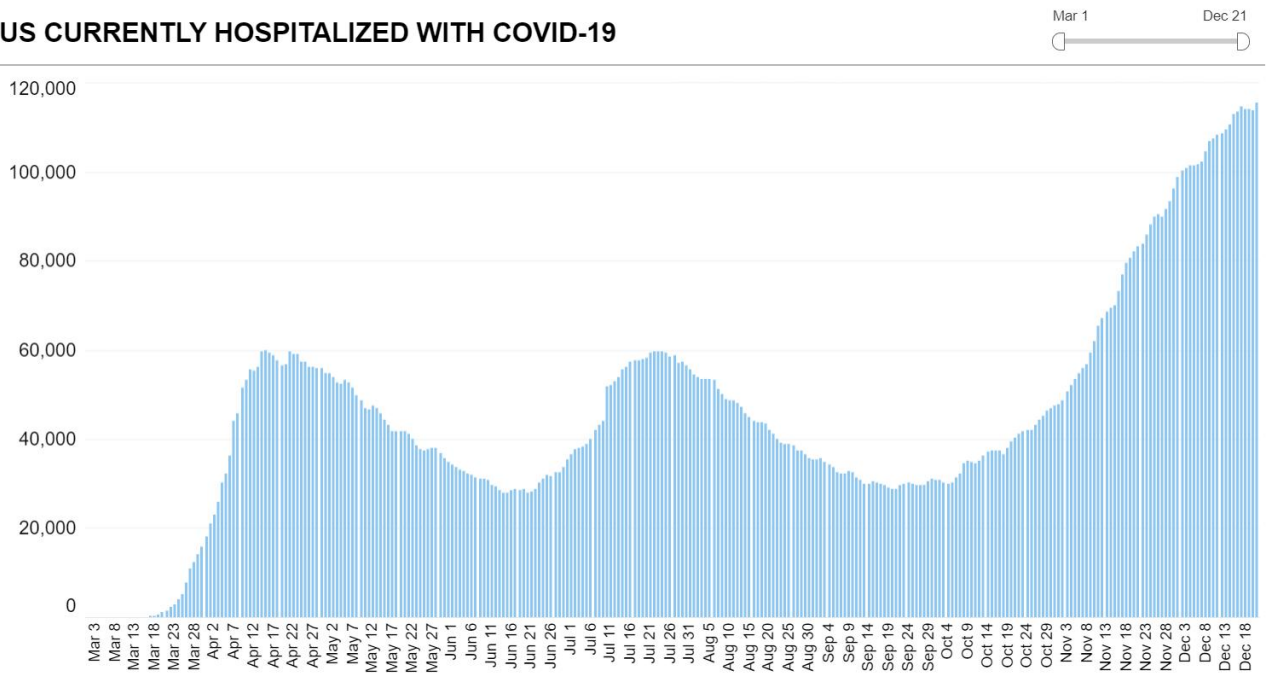
24 ⁵ Li, R., et al., *Beds for Patients With COVID-19 Based on Comparisons With Wuhan and*
 25 *Guangzhou, China*, May 6, 2020, JAMA Netw. Open. 2020;3(5): e208297, available at
<https://jamanetwork.com/journals/jamanetworkopen/fullarticle/2765575>.

26 ⁶ Grasselli, G., et al., *Critical Care Utilization for the COVID-19 Outbreak in Lombardy, Italy:*
 27 *Early Experience and Forecast During an Emergency Response*, March 13, 2020, JAMA.
 2020;323(16):1545-1546, available at <https://jamanetwork.com/journals/jama/fullarticle/2763188>.

28 ⁷ The COVID Tracking Project, The Atlantic, available at <https://covidtracking.com/data/charts/us-currently-hospitalized> (accessed Nov. 23, 2020).

1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
22
23
24
25
26
27
28

US CURRENTLY HOSPITALIZED WITH COVID-19



Note: Florida began reporting this figure on July 10.

19. In this context, academic debates about the risk of severe outcomes per individual, while relevant, are better understood in the context of the total burden created: individual risk times number of individuals infected.

20. COVID-19 is unquestionably worse for someone who is male, older, sicker, or lacks access to health care. Younger, healthier demographics do better than older demographics.⁸ These facts do not mean, however, that COVID-19 is harmless for younger cohorts. To date, more than 52,000 Americans under 65 have died from the disease⁹—more than four times as many as typically die in that age group from seasonal flu in an entire year¹⁰—and we have only had about eleven

⁸ CDC, *COVID-19 Hospitalization and Death by Age*, <https://www.cdc.gov/coronavirus/2019-ncov/covid-data/investigations-discovery/hospitalization-death-by-age.html> (accessed Dec. 12, 2020).

⁹ CDC, *Weekly Updates by Select Demographic and Geographic Characteristics*, available at https://www.cdc.gov/nchs/nvss/vsrr/covid_weekly/index.htm (accessed Dec. 12, 2020)

¹⁰ Quandelacy, T., et al., *Age- and Sex-related Risk Factors for Influenza-associated Mortality in the United States Between 1997–2007*, *Am. J. Epidemiol.* 2014 Jan. 15; 179(2): 156–167, doi: 10.1093/aje/kwt235, available at <https://www.ncbi.nlm.nih.gov/pmc/articles/PMC3873104/>.

1 months of intense COVID-19 activity, so that number will continue to grow.

2 21. Dr. Bhattacharya’s declaration, by focusing only on those at lowest risk, significantly
3 underestimates the SARS-CoV-2 infection-fatality rate. The best estimate to date of the overall
4 infection-fatality rate for SARS-CoV-2 infection is by Dr. Gideon Meyerowitz-Katz and colleagues,
5 and is approximately 0.7%.¹¹ Importantly, while the risk is age specific, and the infection-fatality
6 rate increases sharply with age, there is no cutoff at age 70; rather, the risk of dying if infected with
7 this virus “increases progressively to 0.4% at age 55, 1.4% at age 65, 4.6% at age 75, and 15% at age
8 85.”¹² It is misleading to call the infection-fatality rate below age 70 “vanishingly small” given
9 these risk estimates and given that over 83% of the U.S. population or nearly 274 million people are
10 under 65 (2019 census estimate). Dr. Bhattacharya relies on a meta-analysis by Dr. John P.A.
11 Ioannidis that estimates a lower infection-fatality rate (Para. 18), but Dr. Meyerowitz-Katz’s meta-
12 analysis is distinguished by more rigorous criteria for including studies than that by Dr. Ioannidis.
13 While the former excludes studies expected to be heavily biased by a nonrepresentative sample, the
14 latter does no such quality checks. On this basis, I judge the conclusions of Dr. Meyerowitz-Katz’s
15 meta-analysis more reliable.

16 22. If COVID-19 were judged on the criteria established for evaluating the severity of an
17 influenza pandemic, it would land at the top—the most severe end—of the scale. The CDC in 2017
18 defined influenza pandemics along a scale of transmissibility (from 1-5) and clinical severity (from
19 1-7). Based on the reproduction number and approximately half of infections being symptomatic,
20 COVID-19 would exceed the specifications for the highest transmissibility category ($R_0 > 1.8$) (for
21
22
23

24 ¹¹ Meyerowitz-Katz, G., *A systematic review and meta-analysis of published research data on*
25 *COVID-19 infection fatality rates*, International Journal of Infectious Diseases, Vol. 101, P138-148,
26 December 01, 2020, available at [https://www.ijidonline.com/article/S1201-9712\(20\)32180-](https://www.ijidonline.com/article/S1201-9712(20)32180-9/fulltext)
27 [9/fulltext](https://www.ijidonline.com/article/S1201-9712(20)32180-9/fulltext).

28 ¹² Levin, A., *Assessing the Age Specificity of Infection Fatality Rates for COVID-19: Systematic*
Review, Meta-Analysis, and Public Policy Implications medRxiv 2020.07.23.20160895; doi:
<https://doi.org/10.1101/2020.07.23.20160895>, available at
<https://www.medrxiv.org/content/10.1101/2020.07.23.20160895v7>.

1 COVID-19 R0 is thought to be at least 2¹³ and up to 6 in some places¹⁴). It would also likely exceed
 2 the specification for the highest clinical severity category, which is a case-fatality ratio of 1% or
 3 more.¹⁵ (Given the underascertainment of infections relative to cases, this criterion would be
 4 satisfied by an observed infection-fatality ratio of well under 1%, consistent with even the
 5 downwardly biased estimates of Dr. Ioannidis.) In short, the COVID-19 pandemic is at the upper
 6 end, and arguably at the very top, of the severity scale for influenza pandemics. It was for those
 7 pandemics that community mitigation strategies based on nonpharmaceutical interventions have
 8 been planned at the federal¹⁶ and state¹⁷ levels. A decade or more of pandemic planning envisioned
 9 exactly the kinds of measures being challenged by the defendants in response to a pandemic of a
 10 similar viral infection, even with lower severity than COVID-19.

11 23. In every pandemic, decisions about control must be made before comprehensive
 12 evidence is available on the characteristics of the infection in affected populations.¹⁸ Evidence-
 13 gathering and mitigation efforts must proceed in parallel.¹⁹ Waiting for definitive evidence on
 14 severity, transmissibility, and other characteristics of the pathogen and the population before
 15

16
 17 ¹³ CDC, *COVID-19 Pandemic Planning Scenarios*, available at
<https://www.cdc.gov/coronavirus/2019-ncov/hcp/planning-scenarios.html>.

18 ¹⁴ Sanche, S., et al., *High Contagiousness and Rapid Spread of Severe Acute Respiratory Syndrome*
Coronavirus 2, *Emerg Infect Dis.* 2020;26(7):1470-1477, available at
 19 https://wwwnc.cdc.gov/eid/article/26/7/20-0282_article.

20 ¹⁵ Qualls, N., et al., *Community Mitigation Guidelines to Prevent Pandemic Influenza — United*
States, 2017, *Recommendations and Reports / April 21, 2017 / 66(1);1–34* (Table 6), available at
 21 <https://www.cdc.gov/mmwr/volumes/66/rr/rr6601a1.htm>.

22 ¹⁶ Qualls (2017), *supra*.

23 ¹⁷ California Governor's Office of Emergency Services, *Statewide Concept of Operations for*
Pandemic Influenza, available at
 24 <https://www.caloes.ca.gov/PlanningPreparednessSite/Documents/StatewideConOpsforPandemicInfluenza%202009.pdf>.

25 ¹⁸ Lipsitch, M., et al., *Managing and Reducing Uncertainty in an Emerging Influenza Pandemic*, *N.*
Engl. J. Med. 2009; 361:112-115, doi: 10.1056/NEJMp0904380, available at
 26 <https://www.nejm.org/doi/full/10.1056/nejmp0904380>.

27 ¹⁹ Lipsitch, M., et al., for the 2009 H1N1 Surveillance Group. *Improving the Evidence Base for*
Decision Making During a Pandemic: The Example of 2009 Influenza A/H1N1, 2011, *Biosecurity*
 28 *and Bioterrorism: Biodefense Strategy, Practice, and Science* Vol. 9, No. 2, available at
<https://www.liebertpub.com/doi/full/10.1089/bsp.2011.0007>.

1 adopting control measures is not a viable option, because exponential or near-exponential spread of
2 infection in new pandemics in highly susceptible populations can rapidly transform a small public
3 threat into a large one, and the impact of control measures is often delayed. Thus, it is rational to
4 take action to avert possible negative outcomes before there is certainty about the likelihood and
5 timing of these outcomes. There is room for legitimate disagreement about the strength of evidence
6 and the justification for particular control measures. Yet in the face of a growing pandemic with
7 clear ability to cause severe illnesses, to kill, and to cause health care disruption, it would be
8 irresponsible public health policy to await definitive evidence before taking control measures that
9 are expected to blunt the impact of the pandemic. This was the very clear situation in March 2020 in
10 the United States, as we watched the impact of the pandemic in other countries that had been struck
11 earlier.²⁰ Indeed, there is a compelling argument in my view that many state authorities were too
12 slow, not too fast, to impose restrictions to slow the spread of SARS-CoV-2 during the early months
13 of 2020. On this view, California is a model, while other states deserve criticism for slower
14 reactions.²¹

15 24. The actions taken by California's public health authorities appear to have reduced
16 transmission. Not only do seroprevalence estimates indicate that the virus has been less widespread
17 in California than other regions of the U.S.; but a recent analysis showed that states with stricter
18 containment measures to reduce the spread of the virus—like California—had fewer new cases and
19 hospitalizations per capita than states that imposed few restrictions, which developed some of the
20 worst outbreaks:²²

23 ²⁰ Lipsitch, M., *We know enough now to act decisively against Covid-19. Social distancing is a good*
24 *place to start*, March 18, 2020, STAT, available at <https://www.statnews.com/2020/03/18/we-know-enough-now-to-act-decisively-against-covid-19/>.

25 ²¹ Sexton, J., et al., *Two Coasts. One Virus. How New York Suffered Nearly 10 Times the Number of*
26 *Deaths as California*, May 16, 2020, ProPublica, available at
<https://www.propublica.org/article/two-coasts-one-virus-how-new-york-suffered-nearly-10-times-the-number-of-deaths-as-california>.

27 ²² Leatherby L., et al., *States That Imposed Few restrictions Now Have the Worst Outbreaks*, New
28 *York Times*, Nov. 18, 2020, available at <https://www.nytimes.com/interactive/2020/11/18/us/covid-state-restrictions.html?action=click&module=RelatedLinks&pgtype=Article>.



25. The size of the vulnerable population in the United States is large. Not only are there significant numbers of American over 65—according to the U.S. Census Bureau, about 16.5% of the population was 65 or over in 2019²³—but the CDC estimates that nearly 50 percent of Americans live with underlying conditions that predispose them to serious outcomes from COVID-19.

26. Letting the virus spread unchecked in younger populations—which include Americans with underlying conditions—will result in more serious illness and deaths, in addition to increasing the risk of transmission to older populations.

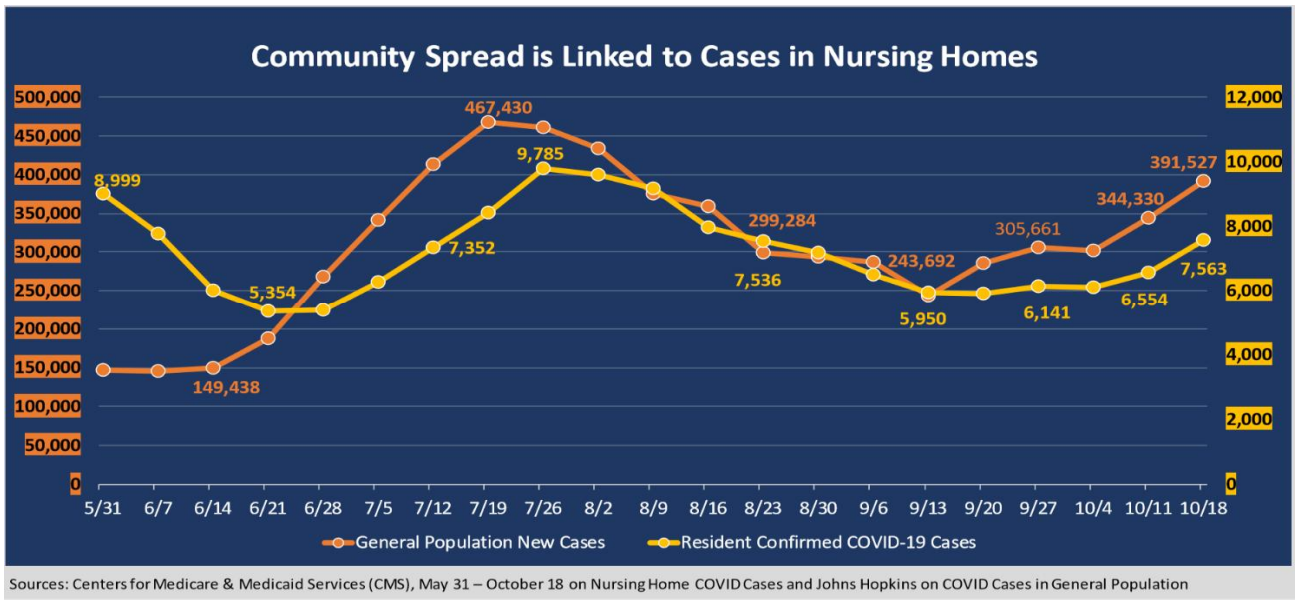
There Is No Proven Means to Protect the Vulnerable Without Restraining Transmission in The General Population

27. No one has yet devised an effective approach to protecting vulnerable populations when there is widespread community transmission. Vulnerable individuals—including older Americans and those with pre-existing conditions—live and work with, and receive care from, members of the larger community. Many of the vulnerable live in a multigenerational home, are

²³ U.S. Census Bureau, <https://www.census.gov/data/tables/time-series/demo/peppest/2010s-national-detail.html> (accessed Nov. 9, 2020).

1 cared for by others in nursing and long-term care facilities, and/or are essential workers with
 2 comorbidities; and these individuals cannot be completely isolated from the larger community.
 3 Nonetheless, scientists, clinicians, and policy makers have been working hard to protect these
 4 groups, with little or modest success, for most of the year, while also attempting to minimize the
 5 threat that community transmission poses to them and to all of us. This “belt-and-suspenders”
 6 approach is the current consensus approach among infectious-disease epidemiologists.

7 28. Reducing or eliminating community transmission is critical to protecting vulnerable
 8 populations, including those in long-term care facilities. The American Health Care Association
 9 (AHCA) and the National Center for Assisted Living (NCAL) recently released a report stating that
 10 COVID-19 cases in U.S. nursing homes have risen with the community spread of COVID-19 since
 11 mid-September.²⁴ That report explicitly links cases in nursing homes to community spread of the
 12 virus:



23 29. Without reducing community transmission, strategies focused on protecting
 24 vulnerable populations are unlikely to succeed. Sweden, the best-known exemplar of the “age-

25
 26
 27 ²⁴ AHCA and NCAL, *Report: COVID-19 Cases In U.S. Nursing Homes*, updated Nov. 2, 2020,
 28 available at <https://www.ahcanal.org/News-and-Communications/Fact-Sheets/FactSheets/Report-Nursing-Homes-Cases-Nov2-2020.pdf> (accessed Nov. 10, 2020).

1 targeted” approach, was unable to protect people in nursing homes. Ostensibly the goal in Sweden
 2 was to protect the elderly and other high-risk groups while slowing viral spread enough to avoid
 3 hospitals being overwhelmed; although it has been widely reported that the goal was in fact to
 4 develop herd immunity.²⁵ The strategy failed to meet its goal of protecting the elderly: the virus ran
 5 rampant in nursing homes, and Stockholm’s nursing homes lost 7% of their residents. Sweden’s
 6 policies are now falling back in line with its European neighbors.²⁶ Vulnerable individuals living in
 7 multigenerational households present a distinct challenge to “focused protection” in the absence of
 8 community control, particularly given that transmission of SARS-CoV-2 in households is
 9 common,²⁷ and households are the single greatest known source of transmission in many locales.²⁸

10 30. The elderly, and nursing home residents in particular, are only a fraction of the truly
 11 vulnerable population. As noted above, over 52,000 deaths have occurred in those under 65 in the
 12 US, about 18% of the total death toll. Some comorbidities that predispose to severe outcomes, such
 13 as diabetes and certain cancers, may be invisible to those who are charged with protecting the
 14 vulnerable. Nonwhite race/ethnicity,²⁹ low socioeconomic status,³⁰ and other variables are also
 15 associated with high vulnerability to severe outcomes, making the logistics of “shielding the
 16

17
 18 ²⁵ Vogel, G., ‘It’s been so, so surreal.’ *Critics of Sweden’s lax pandemic policies face fierce*
 19 *backlash*, Science, Oct. 6, 2020, available at [https://www.sciencemag.org/news/2020/10/it-s-been-](https://www.sciencemag.org/news/2020/10/it-s-been-so-so-surreal-critics-sweden-s-lax-pandemic-policies-face-fierce-backlash)
 20 [so-so-surreal-critics-sweden-s-lax-pandemic-policies-face-fierce-backlash](https://www.sciencemag.org/news/2020/10/it-s-been-so-so-surreal-critics-sweden-s-lax-pandemic-policies-face-fierce-backlash); Bjorklund, K., *The*
 21 *Swedish COVID-19 Response Is a Disaster. It Shouldn’t Be a Model for the Rest of the World*, Time,
 22 Oct. 14, 2020.

23 ²⁶ Vogel 2020, *supra*.

24 ²⁷ Grijalva, C., *Transmission of SARS-COV-2 Infections in Households — Tennessee and Wisconsin,*
 25 *April–September 2020*, Nov. 6, 2020, MMWR, 69(44);1631–1634.

26 ²⁸ Bebinger, M., *Mass. Takes Close Look At Cluster Origins To Stop Coronavirus Spread*, WBUR,
 27 Oct. 20, 2020, <https://www.wbur.org/commonhealth/2020/10/30/massachusetts-covid-cluster-data>;
 28 *Infektionsumfeld von COVID-19-Ausbrüchen in Deutschland*, Epidemiologisches Bulletin, 38 2020,
 29 Sept. 17, 2020,
 30 [https://www.rki.de/DE/Content/Infekt/EpidBull/Archiv/2020/Ausgaben/38_20.pdf;jsessionid=B8D1](https://www.rki.de/DE/Content/Infekt/EpidBull/Archiv/2020/Ausgaben/38_20.pdf;jsessionid=B8D1D66F6ECEC4B21EFDF40B78C8FB74.internet071?_blob=publicationFile)
 31 [D66F6ECEC4B21EFDF40B78C8FB74.internet071?_blob=publicationFile](https://www.rki.de/DE/Content/Infekt/EpidBull/Archiv/2020/Ausgaben/38_20.pdf;jsessionid=B8D1D66F6ECEC4B21EFDF40B78C8FB74.internet071?_blob=publicationFile).

32 ²⁹ APM Research Lab, *The color of coronavirus: COVID-19 deaths by race and ethnicity in the U.S.*,
 33 Oct. 15, 2020, available at <https://www.apmresearchlab.org/covid/deaths-by-race>.

34 ³⁰ Finch, W., et al., *Poverty and Covid-19: Rates of Incidence and Deaths in the United States*
 35 *During the First 10 Weeks of the Pandemic*, Front. Sociol., June 15, 2020, available at
 36 <https://www.frontiersin.org/articles/10.3389/fsoc.2020.00047/full>.

1 vulnerable” even more challenging.

2 31. Until we have a proven means to protect those most at risk and put those safeguards
3 in place, it would be reckless to remove the protections against unmitigated community transmission
4 and plunge ahead in pursuit of herd immunity via massive infection rates. Reducing community
5 transmission remains one of the best ways to protect vulnerable populations.

6 **Any Herd Immunity May Be Short-Lived and Partial**

7 32. In the modern era, herd immunity is best achieved by vaccination—that is, when
8 enough people acquire immunity to an infection through a shot in the arm to protect the whole
9 community. That is our public health goal every flu season; and it is the reason we vaccinate infants
10 against many childhood diseases.

11 33. The “herd immunity with focused protection” approach that Dr. Bhattacharya
12 champions is to allow the spread of COVID-19 in younger populations in order to create immunity
13 against later infection—the theory being that previously infected individuals will carry COVID-19
14 antibodies that will prevent reinfection. If carrying COVID-19 antibodies confers immunity, then
15 half³¹ or more³² of the population must be seropositive—*i.e.*, COVID-19 antibody carriers—before
16 we can control the virus without special measures, such as face coverings, social distancing,
17 surveillance, and contact tracing.

18 34. It is possible that letting the virus spread uncontrolled in the younger population will
19 build up some level of herd immunity and reduce further spread—for some period of time and with
20 the significant cost of serious illness and death discussed above. However, the process of building
21 up herd immunity could take a significant amount of time. The length of the pandemic could be 18
22 to 24 months or more, as herd immunity gradually develops in the human population.³³ The

24 _____
25 ³¹ Britton, T., et al., *A mathematical model reveals the influence of population heterogeneity on herd
immunity to SARS-CoV-2*, Science 14 Aug 2020; 846-849.

26 ³² Sanche S, et al.. *High Contagiousness and Rapid Spread of Severe Acute Respiratory Syndrome
Coronavirus 2*. Emerg Infect Dis. 2020;26(7):1470-1477.
27 <https://dx.doi.org/10.3201/eid2607.200282>.

28 ³³ Moore 2020, *supra*.

1 serosurveillance data available to date suggests that a relatively small fraction of the population has
 2 been infected and infection rates likely vary substantially by geographic area. In late September
 3 2020, CDC Director Robert Redfield told Congress that over 90 percent of the U.S. population
 4 remains susceptible to this coronavirus,³⁴ citing published data.³⁵ Another recent seroprevalence
 5 study estimated that that the percentage of people exposed to the virus ranged from 1% to 23%
 6 depending on jurisdiction, and that overall less than 10% of people had detectable SARS-CoV-2
 7 antibodies.³⁶ Given the transmissibility of SARS-CoV-2, half to two-thirds of the population may
 8 need to be immune to reach a critical threshold of herd immunity to halt the pandemic.³⁷

9 35. Unfortunately, however, coronavirus immunity is notoriously short-lived and partial.
 10 Other coronaviruses are called “seasonal” because, like the flu, they circulate every year. Based on
 11 seasonal coronaviruses, we can anticipate that even if immunity declines after exposure, there may
 12 still be some protection against disease severity and reduced contagiousness, but this remains to be
 13 assessed for SARS-CoV-2.³⁸ As a result, widespread infection in the general population is unlikely
 14 to eliminate the disease but will more likely result in a persistent problem until an effective vaccine
 15 is available and widely adopted.

16 36. The quality of the seroprevalence studies conducted to date has varied widely, as have
 17 their results. The widely varying results of early seroprevalence studies emphasized the very local
 18 nature of the pandemic. For example, in a study that received extensive criticism for its sampling

19 ///

20 ///

22 ³⁴ C-SPAN, *CDC Director Redfield Says 90% U.S. Population Susceptible to Coronavirus Infection*,
 23 *Sept. 23, 2020*, <https://www.c-span.org/video/?c4909117/cdc-director-redfield-90-us-population-susceptible-coronavirus-infection>.

24 ³⁵ Anand, S., et al., *Prevalence of SARS-CoV-2 antibodies in a large nationwide sample of patients on dialysis in the USA: a cross-sectional study*, *The Lancet*, Vol. 396, Issue 10259, pp. 1335-1344,
 25 October 24, 2020, DOI: [https://doi.org/10.1016/S0140-6736\(20\)32009-2](https://doi.org/10.1016/S0140-6736(20)32009-2).

26 ³⁶ Bajema K. et al., *Estimated SARS-CoV-2 Seroprevalence in the US as of September 2020*, *JAMA Intern Med.* (published online Nov. 24, 2020), doi:10.1001/jamainternmed.2020.7976.

27 ³⁷ Britton 2020, *supra*; Sanche 2020; *supra*.

28 ³⁸ Moore 2020, *supra*.

1 methods, statistics, and biased interpretation of the data obtained,³⁹ Dr. Bhattacharya and his co-
 2 authors found that 1.5 percent of Santa Clara County’s population sampled tested positive for
 3 antibodies in the spring of 2020.⁴⁰ Other locales—in studies reflecting different kinds of
 4 imperfections in sampling—have shown much more widespread evidence of past infection,
 5 including 21 percent of those tested in New York City⁴¹ and nearly a third in Chelsea, Massachusetts
 6 in April 2020.⁴²

7 37. More recent national studies have continued to show regional variation in
 8 seroprevalence. The CDC conducted a commercial laboratory seroprevalence survey using blood
 9 samples collected from 10 U.S. sites from March to July 2020. The surveys estimated
 10 seroprevalence of 0.7% (San Francisco Bay Area) to as high as 23.2% (New York City Metro
 11 Area).⁴³ Another seroprevalence study of dialysis patients estimated that during the first wave of the
 12 COVID-19 pandemic, fewer than 10% of the U.S. adult population formed antibodies against SARS-
 13 CoV-2, with large regional variances.⁴⁴ In California, the study estimated seroprevalence of 3.8%.⁴⁵
 14 The low seroprevalence estimates in California and the Bay Area suggest both that those regions
 15 have been more successful in limiting community transmission and that any measure of herd
 16 immunity is a distant prospect.

17 ///

18 _____
 19 ³⁹ Vogel, G., *Antibody surveys suggesting vast undercount of coronavirus infections may be*
 20 *unreliable*. Science, Apr. 21, 2020, [https://www.sciencemag.org/news/2020/04/antibody-surveys-](https://www.sciencemag.org/news/2020/04/antibody-surveys-suggesting-vast-undercount-coronavirus-infections-may-be-unreliable)
 21 [suggesting-vast-undercount-coronavirus-infections-may-be-unreliable](https://www.sciencemag.org/news/2020/04/antibody-surveys-suggesting-vast-undercount-coronavirus-infections-may-be-unreliable).

22 ⁴⁰ Bendavid, E., et al., *COVID-19 Antibody Seroprevalence in Santa Clara County, California*,
 medRxiv, doi: <https://doi.org/10.1101/2020.04.14.20062463>.

23 ⁴¹ Goodman, J., et al., *1 in 5 New Yorkers May Have Had Covid-19, Antibody Tests Suggest*,
 24 <https://www.nytimes.com/2020/04/23/nyregion/coronavirus-antibodies-test-ny.html>.

25 ⁴² Saltzman, J., *Nearly a third of 200 blood samples taken in Chelsea show exposure*, Boston Globe,
 Apr. 17, 2020, [https://www.bostonglobe.com/2020/04/17/business/nearly-third-200-blood-samples-](https://www.bostonglobe.com/2020/04/17/business/nearly-third-200-blood-samples-taken-chelsea-show-exposure-coronavirus/)
 26 [taken-chelsea-show-exposure-coronavirus/](https://www.bostonglobe.com/2020/04/17/business/nearly-third-200-blood-samples-taken-chelsea-show-exposure-coronavirus/).

27 ⁴³ CDC, *Commercial Laboratory Seroprevalence Surveys*, [https://www.cdc.gov/coronavirus/2019-](https://www.cdc.gov/coronavirus/2019-ncov/cases-updates/commercial-lab-surveys.html#surveymap)
 28 [ncov/cases-updates/commercial-lab-surveys.html#surveymap](https://www.cdc.gov/coronavirus/2019-ncov/cases-updates/commercial-lab-surveys.html#surveymap)

⁴⁴ Anand 2020, *supra*.

⁴⁵ *Id.*

Risks of Indoor Gatherings

1
2 38. SARS-CoV-2 spreads through contact (via larger droplets and aerosols), and longer-
3 range transmission via aerosols, especially in conditions where ventilation is poor. This makes
4 large, indoor gatherings particularly high-risk activities for transmission of the virus, especially
5 where mitigation measures like the use of face coverings and social distancing are not being
6 observed. I understand that the plaintiffs in this lawsuit want to hold church services indoors, and
7 that some of them have done so in the past few months without requiring face coverings and social
8 distancing, and while permitting singing. These circumstances present a relatively high risk of
9 transmission. I disagree with Dr. Bhattacharya's assertion that that permitting high risk activities
10 like large, indoor gatherings is consistent with good public health practice at this point in the
11 COVID-19 pandemic.

12 39. A number of factors make this type of conduct particularly high risk. First, longer
13 duration contacts increase the risk of transmission. For direct interpersonal interactions—that is,
14 close contact without social distancing—the risk of transmission is proportional to the duration of
15 the interaction. Longer duration contacts create a higher risk of transmission. Epidemiologists often
16 distinguish between contacts below fifteen minutes (lower risk) and at or beyond fifteen minutes
17 (higher risk)—hence the “fifteen-minute rule.” For viruses like SARS-CoV-2 that are spread through
18 respiratory droplets and aerosols, microbial risk assessment experts use estimates of breathing rates
19 and duration of exposure to develop control strategies to reduce transmission.⁴⁶ Models of COVID-
20 19 transmission using cell phone mobility data identify locations associated with longer duration
21 indoor contacts—including full-service restaurants and religious organizations—as producing the
22 largest predicted increases in infections when reopened.⁴⁷ In contrast, shorter duration and more
23 transitory interpersonal interactions—such as those one would expect in grocery stores, retail stores,
24 and while transiting at an airport—would be expected to create a lower risk of transmission. Dr.

25
26
27 ⁴⁶ Hass C, *Action Levels for SARS-CoV-2 in Air: Preliminary Approach*, Preprint (Aug. 14, 2020).

28 ⁴⁷ Chang S, et al., *Mobility network models of COVID-19 explain inequities and inform reopening*.
Nature (2020), doi: <https://doi.org/10.1038/s41586-020-2923-3>.

1 Bhattacharya does not address the relative risks of transmission associated with different activities.

2 40. The aerosol component of COVID-19 transmission would also be expected to
3 increase this risk of transmission over time. Simply stated, the longer people are in an enclosed
4 space, the more viral particles will build up and be available to infect others. The amount of virus
5 per liter of air will depend on a number of factors, including the size of the space, the number of
6 people in that space, and the frequency of air changes. For these reasons, more crowded gatherings,
7 and poor ventilation would both be expected to increase the risk of transmission in an indoor space.

8 41. Second, larger gatherings in indoor spaces increase the risk of transmission both for
9 the reasons explained above, and because the larger the gathering, the more likely that infectious
10 individuals will in fact be present, depending on the prevalence of the disease in the community.

11 42. Third, singing is a well-described risk factor for transmitting respiratory disease.⁴⁸
12 Dr. Bhattacharya previously opined in another lawsuit, on behalf of another church plaintiff, that
13 churchgoers can “safely” hold indoor services that include “singing and chanting.” Dr. Bhattacharya
14 appears to have retreated from this prior opinion, which was not supported by the research on
15 singing. It is understood that singing, chanting, shouting, and similar vocalizations can cause the
16 release of a larger number of virus-bearing respiratory droplets and aerosols and may also increase
17 the distance that droplets or aerosolized particles can travel compared to speaking at a normal
18 volume. There are documented COVID-19 outbreaks where singing is presumed to have been a
19 factor, including one involving a choir practice in Skagit County, Washington.⁴⁹ Because COVID-
20 19 is spread via respiratory droplets and aerosols, singing indoors is not a safe activity at multi-
21 household gatherings, particularly where there is widespread community transmission of the disease
22 and the use of social distancing and face coverings are not required.

23 ///

24
25
26 ⁴⁸ Buonanno G., et al., *Quantitative assessment of the risk of airborne transmission of SARS-CoV-2*
27 *infection: Prospective and retrospective applications*, Environ Int. (2020) Sep. 6;145:106112, doi:
28 10.1016/j.envint.2020.106112.

⁴⁹ Hamner, L., *High SARS-CoV-2 Attack Rate Following Exposure at a Choir Practice - Skagit*
County, Washington, March 2020. Morbidity and Mortality Weekly Report 69.

1 43. Instead, Dr. Bhattacharya now argues that the psychological benefits of communal
2 singing should be considered against the costs of gathering indoors (Para. 47), and that permitting in-
3 person worship is consistent with good public health practice (Para. 49). Again, Dr. Bhattacharya
4 does not appear to consider the research on singing or the COVID-19 outbreaks that have been
5 traced back to gatherings where attendees sang together. He also appears to ignore the CDC
6 guidance for people hosting holiday gatherings: “avoid singing or shouting, especially indoors” and
7 “[k]eep music levels down so people don’t have to shout or speak loudly to be heard.”⁵⁰

8 44. I understand that some of the plaintiffs have been permitting singing at their services.
9 This would increase the risk of transmission, especially where mitigation measures including face
10 coverings and social distancing are not enforced consistently. Public health measures to limit or
11 prohibit indoor singing, chanting, shouting, and similar vocalizations, would decrease the risk of
12 transmission of COVID-19, which is principally spread by respiratory droplets and aerosols.

13 45. While face coverings and social distancing would mitigate the risk of COVID-19
14 transmission from singing indoors, it would not eliminate that risk. Wearing a surgical mask while
15 singing can reduce the amount of measured exhaled aerosol particles and droplets to levels
16 comparable with normal talking,⁵¹ but that still presents a risk of transmission indoors.

17 46. Dr. Bhattacharya states in his declaration that he has reviewed the CDC’s May 23,
18 2020 guidance titled “Considerations for Communities of Faith.” The current version of that
19 guidance was updated on October 29, 2020.⁵² The CDC offers this guidance to faith communities
20 “in the course of preparing to reconvene for in-person gatherings while still working to prevent the
21 spread of COVID-19.” As this prefatory statement indicates, the CDC guidance begins with the
22 understanding that many faith communities have not been gathering during the pandemic and
23

24 _____
25 ⁵⁰ CDC, *Holiday Celebrations and Small Gatherings*, <https://www.cdc.gov/coronavirus/2019-ncov/daily-life-coping/holidays.html> (accessed Dec. 21, 2020).

26 ⁵¹ Alsvéd, M., et al., *Exhaled respiratory particles during singing and talking*, *Aerosol Sci. & Tech.*,
27 Vol. 54, No. 11, pp. 1245–48, Sept. 17, 2020, <https://doi.org/10.1080/02786826.2020.1812502>.

28 ⁵² CDC, *Considerations for Communities of Faith*, updated Oct. 29, 2020, available at
<https://www.cdc.gov/coronavirus/2019-ncov/community/faith-based.html> (visited Dec. 12, 2020).

1 provides instructions on how to begin to gather again safely. The first point in the guidance under
2 the heading “Scaling Up Operations,” advises faith communities to “[e]stablish and maintain
3 communication with local and State authorities to determine current mitigation levels in your
4 community.” In response to the current surge in COVID-19 cases and the pressure on hospital
5 resources and ICU capacity in California and Santa Clara County in particular, it is my
6 understanding that the current mitigation measures prohibit indoor gatherings, including church
7 services. I do not understand the CDC’s guidance to instruct faith communities to ignore or act
8 contrary to the current local mitigation strategy.

9 47. The CDC recently published a *Summary of Guidance for Public Health Strategies to*
10 *Address High Levels of Community Transmission of SARS-CoV-2 and Related Deaths.*⁵³ That
11 guidance lists recommended public health strategies as well as recommendations for community-
12 level implementation. The guidance recommends the “universal use of face masks” as a public
13 health strategy, and for that strategy recommends the following community-level implementation:
14 “Issue policies or directives mandating universal use of face masks in indoor (nonhousehold)
15 settings.” The CDC guidance also recommends “[a]void[ing] nonessential indoor spaces and
16 crowded outdoor settings” as a public health strategy, and for that strategy recommends the
17 following community-level implementation: “Promoting flexible worksites (e.g., telework); apply
18 limits to occupancy of indoor spaces and to the size of social gatherings.” In my view, these are
19 critical national and local strategies to mitigate the risk of COVID-19 transmission.

20 48. Any region experiencing moderate, high, or increasing levels of community
21 transmission should do everything possible to lower transmission. The path to low transmission in
22 other countries has included adherence to stringent community control measures—including closure
23 of nonessential indoor work and recreational spaces.⁵⁴ Such measures along with universal mask
24

25
26 ⁵³ Honein M., et al., *Summary of Guidance for Public Health Strategies to Address High Levels of*
27 *Community Transmission of SARS-CoV-2 and Related Deaths*, December 2020. MMWR Morb
28 Mortal Wkly Rep 2020;69:1860-1867, doi: <http://dx.doi.org/10.15585/mmwr.mm6949e2>.

⁵⁴ Couzin-Frankel J, Vogel G, Weiland M., *School openings across globe suggest ways to keep coronavirus at bay, despite outbreaks*, American Association for the Advancement of Science.

1 wearing (with specific exceptions⁵⁵) are essential to bring case numbers down to safe levels for
2 communities to reopen.⁵⁶

3 49. The United States is in the midst of a large surge in cases, recording over 200,000
4 new cases and more than 3,000 deaths per day. These levels of transmission threaten to overwhelm
5 hospitals and ICU capacity in many areas. As the CDC reported last week, all age groups have
6 reached their highest weekly hospitalization rate since the start of the pandemic, with those rates
7 expected to increase as additional data are reported.⁵⁷ And ICU capacity is dwindling in many areas,
8 including Santa Clara County, where ICU beds have been filling up and available capacity has
9 dropped from around 25% to under 15% in the past month.⁵⁸ Given these circumstances, it would be
10 reckless to lift restrictions on community transmission, for example, by permitting the large, indoor
11 gatherings that may include singing that plaintiffs want to hold.

12 50. “Flatten the curve” was a good idea when the world first heard the concept in March,
13 and it is a particularly good one right now. A flatter curve, with more infections delayed, will help
14 the health-care system better cope with the cases it does have. Whereas an overwhelmed health-care
15 system will mean there is little reserve to care for the seriously ill, including all the other diseases
16 hospitals were created to treat. Finally, the first vaccine is here, and more vaccines appear to be on
17 their way. These vaccines appear to be effective enough to protect us, if we can stay uninfected long
18 enough to get our shots.

19
20
21 July 7, 2020, available at <https://www.sciencemag.org/news/2020/07/school-openings-across-globe-suggest-ways-keep-coronavirus-bay-despite-outbreaks>.

22 ⁵⁵ CDC, *Coronavirus Disease 2019 (COVID-19), Frequently Asked Questions*, available at
23 https://www.cdc.gov/coronavirus/2019-ncov/faq.html?CDC_AA_refVal=https%3A%2F%2Fwww.cdc.gov%2Fcoronavirus%2F2019-ncov%2Fneed-extra-precautions%2Fpeople-with-seasonal-allergies-faqs.html#People-with-Seasonal-Allergies.

24
25 ⁵⁶ Levinson, M., *Reopening Primary Schools during the Pandemic* (2020) N. Engl. J. Med. 2020; 383:981-985, DOI: 10.1056/NEJMms2024920.

26 ⁵⁷ CDC, COVIDView, *Key Updates for Week 49, ending December 5, 2020*, available at
27 <https://www.cdc.gov/coronavirus/2019-ncov/covid-data/covidview/index.html>.

28 ⁵⁸ Santa Clara County Public Health Department, COVID-19 Hospitalizations Dashboard, available at <https://www.sccgov.org/sites/covid19/Pages/dashboard-hospitals.aspx>.

1
2
3
4
5
6
7
8
9
10
11
12
13
14
15
16
17
18
19
20
21
22
23
24
25
26
27
28

I declare under penalty of perjury under the laws of the United States that the foregoing is true and correct. Executed at Jamaica Plain, Massachusetts on December 22, 2020.

/s/ Marc Lipsitch
MARC LIPSITCH

2329409

Exhibit A

CURRICULUM VITAE

DATE: October 9, 2020

NAME: Marc Lipsitch

ADDRESS: Department of Epidemiology
Harvard T.H. Chan School of Public Health
677 Huntington Avenue
Boston, MA 02115

DATE & PLACE OF BIRTH: November 15, 1969, New Haven, CT, USA

EDUCATION:

<i>Date</i>	<i>Discipline</i>	<i>Degree</i>	<i>Institution</i>
1991	Philosophy	B.A. <i>summa cum laude</i>	Yale University
1995	Zoology	D.Phil.	University of Oxford

POSTDOCTORAL TRAINING:

1995-1999	Biology	Postdoc with Dr. Bruce Levin	Emory University
-----------	---------	------------------------------	------------------

ACADEMIC APPOINTMENTS:

1997-1999	Visiting Scientist, Respiratory Diseases Immunology Section, Centers for Disease Control and Prevention
1999-2004	Assistant Professor, Department of Epidemiology, Harvard School of Public Health
2004-2006	Associate Professor, Department of Epidemiology and Department of Immunology and Infectious Diseases, Harvard School of Public Health
2006-present	Professor, Department of Epidemiology and Department of Immunology and Infectious Diseases, Harvard School of Public Health
2009-present	Director, Center for Communicable Disease Dynamics, Harvard School of Public Health
2009-present	Associate Member, Broad Institute, Cambridge, MA
2012-2018	External Faculty Member, Santa Fe Institute, Santa Fe, NM

HONORS AND DISTINCTIONS:

1991 Phi Beta Kappa, Yale College
 1992-1995 Rhodes Scholar, University of Oxford, England
 2002 Ellison Medical Foundation New Scholar in Global Infectious Disease
 2002 PhRMA Foundation Research Starter Award in Health Outcomes
 2002 ICAAC Young Investigator Award, American Academy of Microbiology
 2006 Mentoring Award, Harvard School of Public Health
 2009 Thompson Science Hall of Fame, Westminster Schools, Atlanta, GA
 2011 Kenneth Rothman Award, Best Paper in *Epidemiology* in 2010
 2012 Junior Faculty Mentoring Award, Harvard School of Public Health
 2013 Reviewer of the Year in *Epidemiology* in 2012
 2014 Member, winning team (PI Shaman), CDC Predict the Influenza Season Challenge
 2015 Elected Fellow, American Academy of Microbiology
 2016 Robert Austrian Lecturer, International Symposium on Pneumococci and Pneumococcal Diseases
 2018 2017 Article of the Year, American Journal of Epidemiology
 2019 23rd Annual Robert M. Fekety, Jr., MD Lecturer, University of Michigan
 2020 Elected Member, National Academy of Medicine

PROFESSIONAL SERVICE:

1999 Temporary Advisor, WHO. Priorities for Pneumococcal and Hib Vaccine Development and Introduction. Geneva, Switzerland
 2000, 2002 National Institutes of Health, National Center for Research Resources, Special Emphasis Panel, Centers of Biomedical Research Excellence
 2001-2003 Consultant and invited speaker for three public meetings, FDA Center for Veterinary Medicine. Topic: regulation of antimicrobial drugs in veterinary medicine
 2002 Member, WHO Pneumococcal Vaccine Trials Nasopharyngeal Carriage Study Group
 2003 Member, WHO Working Group on SARS Epidemiology and Modeling
 2003-2004 US Department of Defense, Defense Science Board Task Force on SARS Quarantine
 2005 Consultant, Ministry of Foreign Affairs, Canada – Pandemic Influenza
 2005 Consultant, Congressional Budget Office – Pandemic Influenza
 2006, 2007 National Institutes of Health Study Section on Genetic Variation and Evolution
 2007, 2008, 2012 National Institutes of Health Special Emphasis Panels, various
 2007-2011 Report reviewer, National Research Council, NEIDL Risk Assessments
 2008 Food and Drug Administration, Antiviral Advisory Committee, guest member
 2008-2009 Member, World Economic Forum Global Agenda Council on Pandemics
 2009 Consultant, Mexico Ministry of Health, Pneumococcal Conjugate Vaccine
 2009 US President's Council of Advisors on Science and Technology (PCAST)-H1N1 Working Group
 2009 Massachusetts Department of Public Health H1N1 Advisory Group
 2009-2010 Team B Advisory Body to CDC on Novel H1N1 Influenza

Lipsitch, Marc

- 2010-2013 Member, Informal Advisory Group on 2009 Pandemic Influenza Mortality, WHO
- 2010- Member, Pneumococcal Serotype Replacement Technical Advisory Group, WHO
- 2011- Member, Scientific Advisory Board, Pneumococcal Global Sequencing Project (Gates Foundation, Keith Klugman, PI)
- 2014-2016 Member, CIDRAP/Wellcome Trust Team B on Ebola Vaccines
2015 Member, Scientific Advisory Group, Norwegian Institute of Public Health/WHO/MSF Ebola Virus Vaccine Trial
- 2015 Member, Scientific Review Committee, Wellcome Trust Sanger Institute 5-Year Review (4-day evaluation visit)
- 2017- Member, Biological Agents Containment Working Group, Board of Scientific Counselors, Office of Public Health Preparedness and Response, CDC
- 2018- Member, Advisory Board, Vaccines and Immunotherapies, CARB-X
2018- Member, Technical Advisory Group, Pneumococcal Serotype Replacement and Distribution Project (PSERENADE), International Vaccine Action Center
- 2019- Co-chair, WHO Working Group on Vaccines and Antimicrobial Resistance (VAC-AMR)
- 2019 Member, Steering Committee, Scorecard on Progress on Recommendations of the Review of Antimicrobial Resistance, Chatham House
- 2020 Member, Massachusetts Governor's Medical Advisory Committee
2020 Member, Massachusetts COVID-19 Vaccine Working Group
2020 Ad hoc expert, WHO Strategic Advisory Group of Experts, COVID-19 Vaccine Working Group

EDITORIAL BOARDS:

- 2002-2012 Associate Editor, *American Journal of Epidemiology*
- 2004-2008 Associate Faculty Editor, *Emerging Themes in Epidemiology*
- 2009-2010 Member, Faculty of 1000 Biology
- 2006-2016 Editorial Board, *PLoS Medicine*
- 2008-2011 Associate Editor, *Epidemics*
- 2008-2011 Editorial Board, *Emerging Health Threats*
- 2009-2010 Board of Editorial Advisors, *Journal of Infectious Diseases*
- 2009-present Editorial Board, *Epidemiology*
- 2015-present Board of Reviewing Editors, *eLife*

PROFESSIONAL SOCIETIES:

Society for Epidemiologic Research
 American Society for Microbiology
 National Center for Science Education
 Union of Concerned Scientists

PUBLIC HEALTH ORGANIZATIONS:

Founder, Cambridge Working Group, 2014
Founder, Society for Safe Science, 2014

SCIENTIFIC COMMITTEES AND CONFERENCE ORGANIZING:

Scientific Committee: 4th International Symposium on Pneumococci and Pneumococcal Diseases, Helsinki, Finland, May 2004
Scientific Committee: 5th International Symposium on Pneumococci and Pneumococcal Diseases, Alice Springs, Australia, May 2006
Scientific Committee: 6th International Symposium on Pneumococci and Pneumococcal Diseases, Reykjavik, Iceland, June 2008
Scientific Committee: 7th International Symposium on Pneumococci and Pneumococcal Diseases, Tel Aviv, Israel, March 2010
Scientific Committee: 9th International Symposium on Pneumococci and Pneumococcal Diseases, Hyderabad, India, March 2014
Conference Chair: First Annual Center for Communicable Disease Dynamics Symposium: Surveillance for Decision Making in Emerging Diseases: Lessons from the 2009 H1N1 Pandemic Influenza. Boston, June 2010
Organizing Committee: Epistemology of Modeling and Simulation. University of Pittsburgh, April 2011.
Conference Chair: Second Annual Center for Communicable Disease Dynamics Symposium: Antimicrobial Resistance: Biology, Population Dynamics and Policy Options. Boston, October 2011
Conference Chair: Epidemics³, Boston, November 2011
Scientific Committee: Epidemics⁴, Amsterdam, November 2013
Conference organizer: Workshop on Modeling and Simulation for Infectious Disease Trial Design, Seattle, August 2016 (with Betz Halloran)
Workshop organizer: Ethical Design of Vaccine Trials in Emerging Infections, ETHOX, Oxford, UK, July 2017 (with Rebecca Kahn, Nir Eyal, Annette Rid)
Scientific Committee: 12th International Symposium on Pneumococci and Pneumococcal Diseases, Toronto, 2020
Advisory group on COVID-19, Science Philanthropy Alliance, August 2020

GRANT REVIEWER SINCE 2003:

Research Fund for the Control of Infectious Disease (RFCID), Hong Kong Semi-Autonomous Region, China
UK Medical Research Council
Wellcome Trust (UK)
Department of Veterans Affairs, USA
Alliance for the Prudent Use of Antibiotics
Swiss National Science Foundation
RIVM (National Institute of Health and Environment), Netherlands
Innovational Research Incentives Scheme, Royal Netherlands Academy of Sciences
National Institutes of Health, ad hoc member, GVE study section (4x), IRAP study section (1x), several telephone special review groups
Health Research Council of New Zealand

AXA Foundation Fellowships
 Canadian Institutes of Health Research
 Royal Society of New Zealand
 French National Research Agency (ANR)
 Royal Society (UK)
 NIH Special Emphasis Panels (2013/01 ZRG1 IDM-A (02) S; 2014/08 ZRG1 RPHB-W
 (53) R - RFA-RM-13-009: NIH Director's Early Independence Awards
 Review)
 Dutch Research Council NOW (2011, 2017)
 NIH Special Emphasis Panel (2014/10 ZRG1 IDM-S (02) M)
 Chair of NIH Infectious Diseases and Microbiology Integrated Review Group, ZRG1 IDM
 S02 10/2014 council
 NIH CRFS Study section, ad hoc member, 2015
 UK Medical Research Council (2016)

INVITED TALKS (SINCE 2015):

01/2015 Otto Wolff Lecture, Institute of Child Health, University College London
 01/2015 London School of Hygiene and Tropical Medicine, invited lecture
 01/2015 Centre for the Study of Existential Risk, Cambridge University, UK, invited
 lecture
 02/2015 Public Health England, Colindale, London UK, invited lecture
 03/2015 London School of Hygiene and Tropical Medicine, Health Protection
 Research Unit Annual Conference, Invited Lecture
 04/2015 University of Bristol Department of Social Medicine
 04/2015 University of Pittsburgh Marcella L. Finegold Memorial Public Debate
 Series
 05/2015 Applied Bioinformatics and Public Health Conference, Wellcome Trust
 Sanger Institute, Keynote Lecture
 06/2015 Médecins sans Frontières Science Day, Panel Discussion, Paris, France
 06/2015 Eijkman Lecture, UMC Utrecht, Netherlands
 07/2015 ETH Zurich Latsis Symposium, plenary talk
 07/2015 Jenner Lecture, Jenner Vaccine Institute, Oxford University, UK
 09/2015 [popular presentation] HubWeek *Four Global Health Threats, Four Global
 Health Opportunities*, Harvard University
 01/2016 National Science Advisory Board on Biosecurity
 02/2016 WHO Technical Expert Consultation: Alternate Dosing Schedules of
 Pneumococcal Conjugate Vaccines, Geneva (by videolink)
 02/2016 PATH Scientific Advisory Board, PATHwSP Vaccine Trial, Geneva (by
 videolink)
 05/2016 Department of Microbiology and Immunology, Emory University, Atlanta
 05/2016 Causal Inference in the Presence of Interference, Department of
 Biostatistics, Harvard T.H. Chan School of Public Health, Boston, MA
 06/2016 Robert Austrian Award Lecture, International Symposium on
 Pneumococci and Pneumococcal Diseases-10
 07/2016 Glaxo SmithKline Vaccines, Rockville, MD
 07/2016 White House Pandemic Prediction and Forecasting Science and
 Technology Working Group (PPFST WG)

Lipsitch, Marc

09/2016 Keynote Address, Project Prometheus Workshop on Multi-Strain Modeling, RIVM (National Institute of Public Health and the Environment), Bilthoven, Netherlands

01/2017 Postdocs in Complexity Conference, Santa Fe Institute, NM (not delivered due to travel delays)

02/2017 Department of Epidemiology of Microbial Diseases, Yale School of Public Health, New Haven, CT

03/2017 Department of Mathematics, University of Utah

03/2017 The Value of Vaccines in the Avoidance of Antimicrobial Resistance, Chatham House, London

03/2017 WHO Workshop on Vaccines and Antimicrobial Resistance, London, UK

04/2017 National Math Festival, Washington, DC (two talks)

05/2017 Memorial Symposium for Ellis McKenzie, Fogarty International Center, NIH, Bethesda, MD

06/2017 EA Global, Society for Effective Altruism, Cambridge, MA

06/2017 Panelist, Surveillance workshop, Simons Foundation, New York, NY

09/2017 Emerging Leaders in Biosecurity meeting, Johns Hopkins Center for Health Security (held Cambridge, MA)

12/2017 Risk in Complex Systems: Models, Applications, Perceptions, and Policy Implications, Centre de Recherches Mathematiques, University of Montreal

3/2018 Department of Ecology and Evolutionary Biology, Princeton University

3/2018 17th Annual Symposium, Institute for Systems Biology, Seattle

4/2018 Gates Vaccine Impact Modeling Consortium, Keynote Address, annual meeting in Cambridge, MA

5/2018 "Antibiotic resistance: Evolutionary concepts versus clinical realities," Wenner-Gren Center, Stockholm

6/2018 London School of Hygiene and Tropical Medicine

6/2018 Institut Pasteur, Paris

6/2018 Big Data Institute, University of Oxford

8/2018 PRISM (Policy relevant infectious disease simulation and mathematical modelling) Annual Meeting, Palm Cove, Australia

8/2018 David Danks Seminar, Murdoch Children's Research Institute, Melbourne, Australia

11/2018 Berkman-Klein Center, Harvard Law School, Data and Health Seminar

6/2019 Chan-Zuckerberg Biohub, San Francisco

6/2019 Stanford Medical School, Stanford, CA

6/2019 Proctor Foundation, UC San Francisco, CA

7/2019 Wellcome Meeting on the Global Burden of Disease from Antimicrobial Resistance, London

10/2019 23rd Annual Robert M. Fekety, MD Lecture, Department of Medicine, Division of Infectious Diseases, University of Michigan

10/2019 Microbiology and Infectious Diseases Seminar, University of Geneva, Switzerland

3/2020 National Academy of Medicine/American Public Health Association First Webinar on COVID-19, Washington DC (via Zoom)

3/2020 Harvard Kennedy School Growth Lab, Cambridge, MA (via Zoom)

3/2020 White House Modeling Consortium, Washington DC (via Zoom)

4/2020 European Central Bank (via Zoom)

4/2020 *USA Today* Editorial Board (via Zoom)

4/2020 *New York Times* Editorial Board (via Zoom)

Lipsitch, Marc

4/2020 Harvard Medical School Department of Medicine Grand Rounds (one of many short talks, via Zoom)

5/2020 Massachusetts Coalition for Pathogen Research (one of many short talks, via Zoom)

5/2020 Covid symposium, National Institute of Statistical Science/American Statistical Association

5/2020 Private briefing, Deputy Prime Minister Chrystia Freeman, Ottawa, Canada

5/2020 Briefing, New Democrat Coalition, by Zoom

5/2020 Isaac Newton Institute, Cambridge University, UK, by Zoom

5/2020 Futureproofing Public Health, University of Stellenbosch, by Zoom

5/2020 Private briefing, Rahul Gandhi, Leader, Congress Party, India

5/2020 Harvard Club of Boston, by Zoom

5/2020 Tsinghua University / AAAS Symposium, Beijing, by Zoom

5/2020 Institute for Genome Sciences, University of Maryland, by Zoom

5/2020 National Academy of Sciences, Section 43, by Zoom

5/2020 Medical Grand Rounds, Boston Children's Hospital, by Zoom

5/2020 Vaccine Research Center, Beth Israel Deaconess Medical Center, by Zoom

5/2020 COVID-19 and the Role of Modeling, American Statistical Association and the National Institute of Statistical Sciences (via Zoom)

6/2020 International Monetary Fund, by Zoom

6/2020 Webinarium: Role of a Medical University in a Pandemic, Karolinska Institutet, Stockholm, by Zoom

7/2020 Bipartisan Commission on Biosecurity, by Zoom

7/2020 National Bureau of Economic Research, by Zoom

7/2020 SAGE Working Group on COVID-19 Vaccines, by Zoom (panelist)

7/2020 Congressional Briefing on Human Challenge Trials, organized by 1DaySooner and Rep. Bill Foster, by Zoom

7/2020 International Symposium on Novel Ideas in Science and Ethics of Vaccines against COVID-19 Pandemic, India Council of Medical Research, by Zoom

7/2020 Private and Public Science, Advisory, and Consumer Food Policy Group (PAPSAC), Harvard Kennedy School, by Zoom

8/2020 National Academies (NASEM) Committee on Equitable Allocation of Vaccine for the Novel Coronavirus (panel), by Zoom

8/2020 Mathematical Sciences Research Institute, University of California, Berkeley, by Zoom

8/2020 Coronavirus Conversations, Science and Society, Duke University (panel), by Zoom

8/2020 Giving Pledge Meeting – Q&A with Scott Dowell, Bill and Melinda Gates Foundation, by Zoom

8/2020 Janelia Farm, Howard Hughes Medical Institute, via Zoom

9/2020 COVID-19, Public Health Ethics, and Policy for Pandemics, Harvard Medical School Center for Bioethics, Cambridge MA (via Zoom)

9/2020 From testing to distribution: the importance of, and challenges to, estimating the protective effects of vaccines, National Institute of Statistical Sciences, Research Triangle, NC (via Zoom)

9/2020 Epidemiology of COVID-19: Implications for Control, American Physical Society, College Park, MD (via Zoom)

10/2020 COVID-19 and Vaccines: Clinical Trials, Immunity and Immunization,
American Lung Association, Chicago, IL (via Zoom)
10/2020 Board of Directors, Blue Cross-Blue Shield of Massachusetts

RESEARCH SUPPORT:**Past Funding**

1997-1999 NIH postdoctoral fellowship 1 F32 GM019182 Population Genetics of Bacterial Infection and Treatment. Role: PI

1997-1999 SmithKlineBeecham unrestricted educational grant. Effects of Antiviral Usage on Resistance in Herpes Simplex Virus, Type 1. Bruce R. Levin, PI. Role: Co-PI

2001-2005 NIH research grant R01 AI051929. Drug Resistance in Tuberculosis: Genetics and Dynamics. Eric Rubin, PI. Role: Co-PI

2001-2006 NIH research grant R01 AI48935. Vaccination and the Evolutionary Dynamics of Pneumococci. Role: PI

2001-2011 NIH research grant R01 AI048935 Mechanisms of Capsular Diversity in *Streptococcus pneumoniae*. Role: PI

2002 PhRMA Foundation Research Starter Grant. Planning and Assessing Antimicrobial Cycling and Other Interventions to Control Resistance in Hospitals. Role: PI

2002-2006 Ellison Foundation New Scientist in Global Infectious Diseases Award. Antibiotic Resistance in *Streptococcus pneumoniae*: Transmission Dynamics and Consequences for Public Health. Role: PI

2003-2006 NIH research grant 5 R21 AI055825. Epidemiologic Methods: Resistant Nosocomial Infections. Role: PI

2004-2013 NIH/NIAID R01 AI058736 (Freedberg). Optimizing HIV care in less developed countries. Role: Consortium Co-Investigator

2006 Taplin Foundation Equipment Grant, Harvard School of Public Health.

2006 NIAID/TIGR Pathogen Functional Genomics Resource Center grant of access to free microarrays. Effects of Host Immunity on Pneumococcal Gene Expression. Role: PI

2006-2009 NIH cooperative agreement U01 GM076497. Methodological Approaches to Planning and Analysis of New Infectious Diseases. Role: PI. Replaced by U54 GM088558.

Lipsitch, Marc

- 2006-2016 NIH/NIAID R01 AI066304 (Finkelstein). Conjugate Vaccine Impact of Pneumococcal Carriage, Disease, and Population (SPARC2). Role: Consortium PI
- 2010-2014 NIH/NIMH R01 MH087328 (Seage). Modeling the Impact of HIV Prevention Interventions (CEPAC Dynamic). Role: Co-PI
- 2011-2016 NIH/NIGMS R01 GM100467 (Shaman). Influenza Outbreak Prediction: Applying Data Assimilation Methodologies to Make Skillful Forecasts of an Inherently Chaotic, Nonlinear System. Role: Consortium PI
- 2012-2018 NIH/NIAID R01 AI048935-15 NCE. Mechanisms and Population Genomics of Pneumococcal Antigenic Diversity. Role: PI
- 2013-2019 NIH/NIGMS R01 AI106786-05 NCE (PI: Hanage). Ecological and Genetic Contributions to the Spread of Resistance in Pnumococcus. Role: Co-PI
- 2014-2018 Pfizer Inc. (No number). Modeling serotype replacement with Prevnar13 using an agent-based model (Phase 2). Role: PI
- 2014-2018 NIH/NIGMS R01 GM116525-03 (Seage). Calibration and Simulation of the Botswana Combination Prevention Project. Role: Co-PI
- 2014-2018 Program for Appropriate Technology in Health. 1773-00460733-COL. Modeling pneumococcal population dynamics under serotype-nonspecific vaccination. Role: PI
- 2015-2018 NIH/NIAID R21 AI112991-02 NCE (Huttenhower). Staphylococcus Aureus Carriage and the Nasal Microbiome. Role: Co-PI
- 2015-2020 NIH/NIGMS R01 GM113233 (Wargo). The impacts of host vaccination and selective breeding for disease resistance on pathogen transmission and ecology in freshwater aquaculture. Role: Consortium PI
- 2017-2019 Pfizer Inc. CP147216 (Lipsitch/Lewnard). Quantifying pneumococcal conjugate vaccine impact against otitis media. Role: Co-PI
- 2017-2020 CDC 1 U01 CK000538-01 (Samore). Modeling and simulation to support antibiotic stewardship and epidemiological decision-making in healthcare settings. Role: Consortium PI

Current Funding

- 2014-2021 NIH/NIGMS U54 GM088558-10 NCE. MIDAS Center for Communicable Disease Dynamics. Role: PI
- 2018-2022 NIH/NIAID 5 R01 AI128344-01 (Hanage). Deep sequencing of pathogens to precisely define transmission networks using rare variants. Role: Co-PI

Lipsitch, Marc

- 2018-2021 National Institute for Health Research. PR-OD-1017-20006 (PI: Cooper / University of Oxford). Leveraging Pathogen Sequence Data and Adaptive Designs to Improve Vaccine Trials in Emerging Epidemics in LMICs. Role: Consortium PI
- 2018-2020 Pfizer Inc. A34479 (Lewnard). Changes in antimicrobial prescribing for otitis media in the era of pneumococcal conjugate vaccination. Role: Co-PI
- 2019-2024 CDC U01 IP001121-02 (PI: Rosenfeld / Carnegie Mellon University). Delphi Influenza Forecasting Center of Excellence. Role: Consortium PI
- 2020-2022 Wellcome Trust. 219759/Z/19/Z. Vaccine-avertable antimicrobial prescribing from influenza and RSV: a mixed-methods observational study. Role: PI
- 2020-2022 Wellcome Trust. 219812/Z/19/Z (Grad). Reducing antibiotic prescribing through a prioritized vaccination strategy. Role: Co-PI
- 2020-2023 Open Philanthropy Project / Silicon Valley Community Foundation. 2020-211809. Research and Policy Activities on Biosafety and Biosecurity. Role: PI
- 2020-2022 NIH/NCI U01 CA261277-01. Causal, Statistical and Mathematical Modeling with Serologic Data
- 2020-2025 CDC U01 CK000585-01 (Samore). Modeling and Simulation to support Epidemiological decision-making in Healthcare settings. Role: Consortium PI
- 2020-2022 Morris-Singer Foundation. Morris-Singer Fund for the Center for Communicable Disease Dynamics. Role: PI

TEACHING EXPERIENCE:**Full Courses**

Biostatistics 516: Inferential Methods for Infectious Diseases. Co-developer and co-instructor of course.

Taught 2011 (Spring 2)

Epidemiology 260: Mathematical Models of Infectious Diseases: Developed course, sole instructor.

Taught 2001, 2002, 2003, 2005, 2007, 2009, 2011, 2014, 2016, 2018, 2020 always d period (Spring 2).

Interdepartmental 267-268: Seminar in Infectious Disease Epidemiology: Developed and taught course jointly with Dr. Megan Murray in 2000; sole instructor in 2001 and 2002.

Taught 2000-2001 as full year; taught 2001 and 2002 as fall only (ID267ab).

Interdepartmental 298: Inference in Infectious Disease Epidemiology. Developed course, sole instructor.

Taught 2005, 2007 winter session.

Epidemiology 502: Biology and Epidemiology of Antibiotic Resistance. Co-developed and co-taught course with Dr. Gili Regev-Yochay.

Taught 2008, 2010, 2012, 2014, 2016, 2018, 2020 winter session.

Participation

Epidemiology 201: Introduction to Epidemiology. One week each year (2 2-hr. lectures).

Taught 2000-2007; coinstructor 2005-7.

Epidemiology 200: Principles of Epidemiology. One lecture per year.

Taught 2000-2005; coinstructor 2005-7.

Epidemiology 289: two lectures on Infectious Disease Epidemiology, 2010

Interdepartmental 229: Epidemiology of Infectious Diseases of Importance in Developing Countries (and predecessors). One lecture per year.

Taught 2000-2006.

DBS205: Biological Sciences Seminars. One presentation of research per year.

Taught 2002, 2005, 2015.

RDS281 (now 285): Decision Analysis Methods in Public Health and Medicine. One lecture on infectious diseases and dynamic modeling.

Taught 2002-2007, 2010.

IMI225 Design and Development of a vaccine. One lecture per year.

Taught 2007.

IMI227 Genetics of Infectious Disease. One lecture per year.

Taught 2007-8.

Epidemiology 205: Practice of Epidemiology. Supervised one student.

Participated 2000.

Epidemiology 294: Screening. Two lectures on introductory Infectious Disease Epidemiology

2011, (Spring 2)

ID517 Public Health Response to Mass Emergencies. One lecture.

Taught 2008.

Life Sciences 120 (Harvard College): Global Health Threats.

One to three lectures, 2011, 2012, 2016

Probabilistic Risk Analysis (HSPH Continuing Education). One lecture per year.

Taught 2002-2004.

Health Science & Technology Microbiology (HMS). One lecture per year.
Taught 2002, 2003, 2005, 2007, 2008, 2009.

Modern Medical Microbe Hunters (HMS). One lecture per year.
Taught 2000, 2001, 2002, 2004.

Epidemiology 513: Issues in the Reporting of Clinical Trials. *Guest lecture 2011.*

Epidemiology 203. Four, 2-hour lectures per year on infectious disease epidemiology
2012, 2014, 2016, 2017, 2018, 2019

ID250 Ethical Basis of Public Health Practice. One lecture per year
Guest lecture 2015

GHP539 Control of Infectious Diseases in Low/Mid Income Countries: Social, Political
and Economic dimensions
Guest lecture 2017

MPH100
Guest lecture 2019, 2020 (Spring and Fall)

Harvard College Gen Ed 1098 Natural Disasters
Guest lecture 2020

Short Courses Outside Harvard

Harvard-Karolinska Summer School on Modern Methods in Biostatistics and
Epidemiology, Treviso, Italy. Infectious Disease Epidemiology. Developed a 1-week
intensive introductory course with exercises and was sole instructor. *Taught 2005.*

Hong Kong Centre for Health Protection Short Course in Mathematical Modelling of
Infectious Diseases. Participated in course development and taught three lectures.
Taught 2006.

Infection and Immunity in Children, Department of Paediatrics, Oxford University, UK.
Delivered 1 lecture by videolink. *Taught June 2010.*

Winter Forum – Pandemic 2011. Duke Global Health Institute and Office for
Undergraduate Education, Durham, NC. Delivered one lecture by videolink. *Taught
January 2011.*

Practical Short Course in Infectious Disease Modeling. National Center for
Immunization and Respiratory Diseases, US Centers for Disease Control & Prevention
(CDC). Course director (collaboratively with Hong Kong University and Imperial College
London) and instructor. *Taught June 2011.*

Erasmus Summer Program, Erasmus University, Rotterdam, Netherlands. Master Class
taught by videolink. *Taught August 2011.*

Lipsitch, Marc

Short Course in Infectious Disease Modeling: Hong Kong University and HSPH CCDD:
Kuala Lumpur 2014, Bangkok 2012

Course organizer and faculty: Short Course in Infectious Disease Modeling: HSPH
CCDD, Imperial College London, Hong Kong University
Centers for Disease Control and Prevention 2011, 2014

Faculty Guest Lecturer: ICARe (International course on Antibiotic Resistance), Pasteur
Institute, at Fondation Merieux, Les Pensieres, France.
2018, 2019

Online Modules

Herd Immunity in: Vaccines 101, Harvard School of Public Health online course
Recorded Summer 2014

Heterogeneity in: Epidemics, University of Hong Kong HKUx EdX course
Recorded Summer 2014

Guest Teaching Outside Harvard

Georgetown University Department of Microbiology and Immunology: guest lecture in
Science Diplomacy and World Health, Tomoko Steen instructor (2014, 2016)

Princeton University Woodrow Wilson School: guest lecture in Topics in Development:
Global Challenges of Infection, Burden and Control, Adel Mahmoud and Bryan Grenfell
instructors (2016, 2018)

Supervision

Research Scientist supervisor:

2005-2008	Krzysztof Trzcinski, D.V.M., Ph.D., Research Scientist
2008-2009	Krzysztof Trzcinski, D.V.M., Ph.D., Senior Research Scientist (now Assistant Professor, University of Utrecht, Netherlands)
2008-2010	Edward Goldstein, Ph.D., Research Scientist
2010-2018	Edward Goldstein, Ph.D., Senior Research Scientist
2019-present	Rene Niehus, Ph.D, Research Associate

Postdoctoral supervisor:

2001-2005	Krzysztof Trzcinski, D.V.M., Ph.D. Assistant Professor, University of Utrecht, Netherlands.
2001-2002	Noman Siddiqi, Ph.D. (Co-supervisor with Eric Rubin). Currently BL3 Manager, Harvard School of Public Health
2001-2002	Susan Huang, M.D., M.P.H. (Secondary advisor) (currently Professor of Infectious Disease, University of California, Irvine).

Lipsitch, Marc

2002-2003	Ben Cooper, Ph.D. Professor, Nuffield Department of Medicine, Mahidol-Oxford Tropical Research Unit, Bangkok, Thailand
2003-2005	Michael Palmer, Ph.D. Currently working in the IT Industry
2004-2008	Gili Regev-Yochay, M.D. Currently Assistant Professor, Tel Aviv University and Head of Infectious Diseases Epidemiology Unit, Gertner Institute, Tel Aviv, Israel.
2006-2008	Debby Bogaert, M.D., Ph.D. Professor of Pediatric Infectious Diseases, University of Edinburgh
2008	Edward Goldstein, Ph.D. Senior Research Scientist, HSPH
2010	Daniel Weinberger, Ph.D. Assistant Professor, Yale School of Public Health
2009-2011	Joel Miller, Ph.D. Currently Senior Research Scientist, Institute for Disease Modeling, Seattle
2010-2013	Sarah Cobey, Ph.D. Associate Professor of Ecology and Evolution, University of Chicago
2010-2014	Yuan Li, Ph.D. Epidemiologist, CDC
2010-2014	Yonatan Grad, M.D., Ph.D. Assistant Professor, Harvard TH Chan School of Public Health
2011-2013	Nicholas Croucher, Ph.D. (co-advisor with W. Hanage). Senior Lecturer and Henry Dale Fellow, Imperial College
2013-2016	Colin Worby, Ph.D. (co-advisor with W. Hanage) Staff Scientist, Broad Institute.
2013-2019	Hsiao-Han Chang, Ph.D. (co-advisor with C. Buckee). Starting 2019: Assistant Professor, National Tsing Hua University, Taiwan
2014-2016	Nadia Abuelezam, Sc.D. (secondary advisor with George Seage). Currently Assistant Professor, Boston College School of Nursing.
2014	Ben Althouse, Ph.D. (external faculty advisor for his Santa Fe Institute Postdoc)
2015-2017	Kate Langwig, Ph.D. Assistant Professor of Ecology, Virginia Tech
2015-2018	Taj Azarian, Ph.D. Assistant Professor of Medicine, Burnett School of Biomedical Sciences, Department of Molecular Microbiology, University of Central Florida College of Medicine
2015-2018	Brian Arnold, Ph.D. Bioinformatics Scientist, Faculty of Arts and Sciences, Harvard University
2015-2018	Maria Georgieva, Ph.D. Postdoctoral Fellow, Department of Physiology, University of Lausanne
2016-present	Samantha Palace, Ph.D. (co-supervisor with Y. Grad)
2017-2018	Lucy Li, Ph.D. Bioinformatics Scientist I, Chan Zuckerberg Biohub
2017-2018	Joseph Lewnard, Ph.D. Assistant Professor, Department of Epidemiology, UC Berkeley
2017-2019	Ayesha Mahmud, Ph.D. (Co-advisor with C. Buckee). Assistant Professor, Department of Demography, UC Berkeley
2018-present	Pamela Martinez, Ph.D. (Co-advisor with C. Buckee)
2019-present	Xueting Qiu, Ph.D. (Co-advisor with W. Hanage)
2019-present	Lerato Magosi, D.Phil.
2020	Lee Kennedy-Shaffer, Ph.D., Assistant Professor, Vassar College
2020-present	Rebecca Kahn, Ph.D.

Doctoral student supervisor:

Lipsitch, Marc

2000-2001	Ivo Foppa (Epi). Currently Epidemiologist, Influenza Division, CDC
2000-2005	Hailay Teklehaimenot (Epi). Currently at Ministry of Health, Addis Ababa, Ethiopia
2000-2001	Robert Suruki (Epi): completed doctoral studies with another advisor; currently at GlaxoSmithKline.
2001-2006	Christina Mills (Epi). Currently Attending Physician, Boston Children's Hospital Boston
2002-2005	Alethea McCormick (Epi). Currently Research Associate, Harvard School of Public Health
2002-2011	Sibel Ascioğlu (Epi). Currently at Glaxo SmithKline
2003-2008	Virginia Pitzer (Epi). Currently Assistant Professor, Yale School of Public Health
2004-2012	Jessica Hartman Jacobs (Epi)
2006-2012	Justin O'Hagan (Epi). Currently Head of Outcomes Research, Dengue, Merck.
2006-2010	Daniel Weinberger (BPH). Currently Assistant Professor, Yale School of Public Health
2007-2008	Chris Ford (BPH). Currently Postdoc, Broad Institute.
2007-2008	Karell Pelle (BPH)
2014-2018	Matthew Hitchings (Epi). Currently postdoctoral fellow, Emerging Pathogens Institute, University of Florida
2016-present	Christine Tedijanto (Population Health Sciences/Epi)
2016-present	Emma Accorsi (Population Health Sciences/Epi)
2018-2020	Rebecca Kahn (Population Health Sciences/Epi)
2019-present	Keya Joshi (Population Health Sciences/Epi)

Master's student supervisor:

2000-2002	Alison Han (MPH)
2000-2008	Catherine Laine (Epi MSc). Currently Founder and Deputy Director, Appropriate Infrastructure Development Group.
2001-2002	Benjamin Ip, MD (MPH)
2001-2002	Rajneesh Hazarika, MD (MPH)
2002-2004	Hoa Nguyen, MD (MS)
2003-2005	Dereje Dengela (MS)
2004-2005	Heather Green (MS)
2004-2005	Wei-yen Lim, MD (MPH)
2004-2005	Phil James, MD (MPH)
2005-2006	Jeffrey Cloud, MD (MPH)
2005-2006	Yen-Tsung Huang, MD (MPH)
2005-2006	Minghua Chen, MD (MPH)
2006-2007	Chou-Cheng Lai, MD (MS)
2006-2007	Jennifer Shuford, MD (MPH)
2006-2007	Chih-Hao Chen, MD (MS)
2006-2007	Mark Brady (MPH)
2007-2008	Indrajit Hazarika, MD (MPH)
2007-2008	Hyun Joon Shin, MD (MPH)
2007-2008	Amit Vora (MPH)
2005-2006	Christie Jeon (MS)

Lipsitch, Marc

2004-2007	Gili Regev-Yochay, MD (MS)
2009-2011	Weixiong Ke
2010-2012	Karen Aanensen
2011-2013	Talia Quandelacy
2011-2013	Patrick Mitchell
2013-2014	Fausto Bustos
2016-2017	Say Tat Ooi, MD (MPH)
2016-2018	Michael Martin (MS)
2016-2018	Inga Holmdahl (MS)
2016-2018	Rebecca Kahn (MS)
2017-2019	Sarah Lapidus (MS)
2018-2020	Nancy Li (MS)
2019-2020	Eva Rumppler (MS)
2019-present	Rafia Bosan (MS)

Undergraduate supervisor:

Summer 2001	Eneida Villanueva (Summer Minority Intern)
Fall 2001	Jonathan Burton-MacLeod (Bio 91r supervised reading, FAS)
2016-2018	Alan Yang (Supervised research)
Summer 2018	Tara E. Gallagher, Dartmouth College (Summer intern)

Thesis committees:

2000-2001	Megan Murray (Dr. P.H., completed 2000-1)
2004-2007	Eben Kenah (Epidemiology)
2004-2005	Y. Claire Wang (Health Policy and Management)
2005	Seema Thakore Meloni (Ph.D., Biological Sciences in Public Health)
2005-2007	Mary Farrow (Ph.D., Biological Sciences in Public Health)
2007-2010	Kevin Chan (Population & International Health)
2008-2011	Amy Bei (Ph.D., Biological Sciences in Public Health)
2010-2013	Chris Ford (Ph.D., Biological Sciences in Public Health)
2010-2013	Regina Joice (Ph.D., Biological Sciences in Public Health)
2011	Rachel Daniels (Ph.D., Biological Sciences in Public Health)
2011-2017	Freeman Suber MD (Ph.D., Biological Sciences in Public Health)
2011-2012	Tami Lieberman (Ph.D., Systems Biology)
2011-2013	Wei Wu (Epidemiology)
2013	Opponent, PhD of Rolf Ypma, University of Utrecht
2014	Clare Louise Kinnear, PhD, The University of Melbourne (external examiner)
2015-2017	Corey Peak (SD, Epidemiology)
2015-2016	Patrick Mitchell (SD, Epidemiology)
2016	Hattie Chung (PhD, Systems Biology, Harvard GSAS), examination committee
2016	Nicole Espy (PhD, BPH, defense committee)
2015-2017	Quizhi Chang (SD, Epidemiology)
2017-2019	Eric Mooring (SD, Epidemiology)
2017-present	Rebecca Mandt (PhD, BPH)

Lipsitch, Marc

2017-2020 Sarah McGough (PhD, Population Health Sciences)
 2018-2020 Lee Kennedy-Shaffer (Biostatistics)

Oral exam committees:

2001 Yemane Yihdego (IID)
 2001 Chris Mores (IID)
 2002 Pride Chigwedere, MD (IID)
 2004 Beth Ann Griffin (Biostatistics)
 2004 Eben Kenah (Epidemiology)
 2004 Laura Forsberg (Biostatistics)
 2007 Kevin Chan (Population & International Health)
 2008 Hsien-Ho Lin (Epidemiology)
 2009 Regina Joice (BPH)
 2009 Celene Chang (BPH)
 2009 Kathleen Wirth (Epi)
 2011 Freeman Suber (BPH)
 2011 Wei Wu (Epi)
 2012 Nicanor Rodriguez, DVM (IID)
 2016 Corey Peak (Epi)
 2016 Qiuzhi Chang (Epi)

Laboratory rotations supervised:

2000 Chun Chao (MS student, Immunology & Infectious Diseases)
 2004 Adam MacNeil (PhD, Biological Sciences in Public Health)
 2006 Daniel Weinberger (PhD, Biological Sciences in Public Health)
 2006 Amy Bei (PhD, Biological Sciences in Public Health)
 2008 Chris Ford (PhD, Biological Sciences in Public Health)
 2009 Richa Gawande (PhD, Biological Sciences in Public Health)
 2010 Sri Kalyanamaran (PhD, Biological Sciences in Public Health)
 2011 Wen Xie (PhD, Biological Sciences in Public Health)
 2016 Rebecca Mandt (PhD, Biological Sciences in Public Health)

SCHOOL AND DEPARTMENTAL SERVICE

Interdisciplinary Program in the Epidemiology of Infectious Disease

- Steering committee, 2000-present
- Seminar Coordinator, 2000-present
- Associate Director, 2004-present

Biological Sciences in Public Health Program

- Admissions interviewer, 2001-present
- Curriculum Committee, 2006-2012

Department of Epidemiology

- Co-leader, department retreat, 2001

- Admissions committee, 2001-2014

HSPH Epidemiology and Biostatistics Planning Committee: member, 2003-2004

HSPH Allston Planning Committee: member, 2003-2004

HSPH Information Technology Advisory Committee: member, 2004-2005

HSPH Committee on Educational Policy: member, 2005-2008

HSPH Standing Committee on Appointments, Reappointments, and Promotions: member, 2008-present, vice-chair 2010-2012, chair 2012-2013

University Pandemic Response Planning Committee: member, 2005-present

Bioinformatics Junior Faculty Search Committee: member, 2007-2008

Epidemiology Methods Junior Faculty Search Committee: member, 2008

Epidemiology Infectious Diseases Junior Faculty Search Committee: chair, 2008

HSPH Committee on the Concerns of Women Faculty: member, 2010-2012

HMS Subcommittee on Admissions for the MD/PhD: member, 2010-2012, 2014-present

Harvard University Office of Scholarly Communication Advisory Committee, member, 2013-present

Harvard T.H. Chan School of Public Health Dean Search Advisory Committee, member, 2015

Harvard T.H. Chan School of Public Health Faculty Judge, Postdoctoral Association Travel Awards, member, 2016

Epidemiology Junior Faculty Search Committee, 2017-2018

BIBLIOGRAPHY

Peer-Reviewed Articles

1. Petrie M, Lipsitch M. Avian polygyny is most likely in populations with high variability in heritable male fitness. *Proc R Soc Lond B*. 1994 Jun 22;256(1347):275-80. doi: [10.1098/rspb.1994.0081](https://doi.org/10.1098/rspb.1994.0081).
2. Lipsitch M, Nowak MA. The evolution of virulence in sexually transmitted HIV/AIDS. *J Theor Biol*. 1995 Jun 21;174(4):427-40. doi: [10.1006/jtbi.1995.0109](https://doi.org/10.1006/jtbi.1995.0109). PMID: 7666673.
3. Lipsitch M, Nowak MA, Ebert D, May RM. The population dynamics of vertically and horizontally transmitted parasites. *Proc Biol Sci*. 1995 Jun 22;260(1359):321-7. doi: [10.1098/rspb.1995.0099](https://doi.org/10.1098/rspb.1995.0099). PMID: 7630898.
4. Lipsitch M, Herre EA, Nowak MA. Host Population Structure and the Evolution of Virulence: A "Law Of Diminishing Returns". *Evolution*. 1995 Aug;49(4):743-748. doi: [10.1111/j.1558-5646.1995.tb02310.x](https://doi.org/10.1111/j.1558-5646.1995.tb02310.x). PMID: 28565133.
5. Mangin KL, Lipsitch M, Ebert D. Virulence and transmission modes of two microsporidia in *Daphnia magna*. *Parasitology*. 1995;111(2):133-142. doi: [10.1017/S0031182000064878](https://doi.org/10.1017/S0031182000064878).
6. Lipsitch M, Siller S, Nowak MA. The Evolution of Virulence in Pathogens With Vertical and Horizontal Transmission. *Evolution*. 1996 Oct;50(5):1729-1741. doi: [10.1111/j.1558-5646.1996.tb03560.x](https://doi.org/10.1111/j.1558-5646.1996.tb03560.x). PMID: 28565576.
7. Lipsitch M, Moxon ER. Virulence and transmissibility of pathogens: what is the relationship? *Trends Microbiol*. 1997 Jan;5(1):31-7. doi: [10.1016/S0966-842X\(97\)81772-6](https://doi.org/10.1016/S0966-842X(97)81772-6). PMID: 9025233.
8. Antia R, Lipsitch M. Mathematical models of parasite responses to host immune defences. *Parasitology*. 1997;115 Suppl:S155-67. doi: [10.1017/s003118209700200x](https://doi.org/10.1017/s003118209700200x). PMID: 9571700.
9. Levin BR, Lipsitch M, Perrot V, Schrag S, Antia R, Simonsen L, Walker NM, Stewart FM. The population genetics of antibiotic resistance. *Clin Infect Dis*. 1997 Jan;24 Suppl 1:S9-16. doi: [10.1093/clinids/24.supplement_1.s9](https://doi.org/10.1093/clinids/24.supplement_1.s9). PMID: 8994776.
10. Lipsitch M, Levin BR. The within-host population dynamics of antibacterial chemotherapy: conditions for the evolution of resistance. *Ciba Found Symp*. 1997;207:112-27; discussion 127-30. doi: [10.1002/9780470515358.ch8](https://doi.org/10.1002/9780470515358.ch8). PMID: 9189638.
11. Lipsitch M, Levin BR. The population dynamics of antimicrobial chemotherapy. *Antimicrob Agents Chemother*. 1997 Feb;41(2):363-73. doi: [10.1128/AAC.41.2.363](https://doi.org/10.1128/AAC.41.2.363). PMID: 9021193; PMCID: PMC163715.
12. Lipsitch M. Vaccination against colonizing bacteria with multiple serotypes. *Proc Natl Acad Sci U S A*. 1997 Jun 10;94(12):6571-6. doi: [10.1073/pnas.94.12.6571](https://doi.org/10.1073/pnas.94.12.6571). PMID: 9177259; PMCID: PMC21091.
13. Lipsitch M. Evolution in health and disease. *Trends Microbiol*. 1997 Aug;5(8):303-5. doi: [10.1016/S0966-842X\(97\)01087-1](https://doi.org/10.1016/S0966-842X(97)01087-1). PMID: 9263405.
14. Datta A, Hendrix M, Lipsitch M, Jinks-Robertson S. Dual roles for DNA sequence identity and the mismatch repair system in the regulation of mitotic crossing-over in yeast. *Proc Natl Acad Sci U S A*. 1997 Sep 2;94(18):9757-62. doi: [10.1073/pnas.94.18.9757](https://doi.org/10.1073/pnas.94.18.9757). PMID: 9275197; PMCID: PMC23263.
15. Bonhoeffer S, Lipsitch M, Levin BR. Evaluating treatment protocols to prevent antibiotic resistance. *Proc Natl Acad Sci U S A*. 1997 Oct 28;94(22):12106-11. doi: [10.1073/pnas.94.22.12106](https://doi.org/10.1073/pnas.94.22.12106). PMID: 9342370; PMCID: PMC23718.

16. Lipsitch M. Transmission Rates and HIV Virulence: Comments to Massad. *Evolution*. 1997 Feb;51(1):319-320. doi: [10.1111/j.1558-5646.1997.tb02416.x](https://doi.org/10.1111/j.1558-5646.1997.tb02416.x). PMID: 28568797.
17. Levin BR, Antia R, Berliner E, Bloland P, Bonhoeffer S, Cohen M, DeRouin T, Fields PI, Jafari H, Jernigan D, Lipsitch M, McGowan JE Jr, Mead P, Nowak M, Porco T, Sykora P, Simonsen L, Spitznagel J, Tauxe R, Tenover F. Resistance to antimicrobial chemotherapy: a prescription for research and action. *Am J Med Sci*. 1998 Feb;315(2):87-94. doi: [10.1097/00000441-199802000-00004](https://doi.org/10.1097/00000441-199802000-00004). PMID: 9472907.
18. Lipsitch M, Levin BR. Population dynamics of tuberculosis treatment: mathematical models of the roles of non-compliance and bacterial heterogeneity in the evolution of drug resistance. *Int J Tuberc Lung Dis*. 1998 Mar;2(3):187-99. PMID: [9526190](https://pubmed.ncbi.nlm.nih.gov/9526190/).
19. Stewart FM, Antia R, Levin BR, Lipsitch M, Mittler JE. The population genetics of antibiotic resistance. II: Analytic theory for sustained populations of bacteria in a community of hosts. *Theor Popul Biol*. 1998 Apr;53(2):152-65. doi: [10.1006/tpbi.1997.1352](https://doi.org/10.1006/tpbi.1997.1352). PMID: 9615474.
20. Levin BR, Lipsitch M, Bonhoeffer S. Population biology, evolution, and infectious disease: convergence and synthesis. *Science*. 1999 Feb 5;283(5403):806-9. doi: [10.1126/science.283.5403.806](https://doi.org/10.1126/science.283.5403.806). PMID: 9933155.
21. Lipsitch M. Bacterial vaccines and serotype replacement: lessons from *Haemophilus influenzae* and prospects for *Streptococcus pneumoniae*. *Emerg Infect Dis*. 1999 May-Jun;5(3):336-45. doi: [10.3201/eid0503.990304](https://doi.org/10.3201/eid0503.990304). Erratum in: *Emerg Infect Dis* 1999 Sep-Oct;5(5):734. PMID: 10341170; PMCID: PMC2640786.
22. Lipsitch M, Bergstrom CT, Levin BR. The epidemiology of antibiotic resistance in hospitals: paradoxes and prescriptions. *Proc Natl Acad Sci U S A*. 2000 Feb 15;97(4):1938-43. doi: [10.1073/pnas.97.4.1938](https://doi.org/10.1073/pnas.97.4.1938). PMID: 10677558; PMCID: PMC26540.
23. Lipsitch M, Dykes JK, Johnson SE, Ades EW, King J, Briles DE, Carlone GM. Competition among *Streptococcus pneumoniae* for intranasal colonization in a mouse model. *Vaccine*. 2000 Jun 15;18(25):2895-901. doi: [10.1016/s0264-410x\(00\)00046-3](https://doi.org/10.1016/s0264-410x(00)00046-3). Erratum in: *Vaccine* 2000 Oct 15;19(4-5):598. PMID: 10812233.
24. Bergstrom CT, Lipsitch M, Levin BR. Natural selection, infectious transfer and the existence conditions for bacterial plasmids. *Genetics*. 2000 Aug;155(4):1505-19. PMID: [10924453](https://pubmed.ncbi.nlm.nih.gov/10924453/); PMCID: PMC1461221.
25. Negri MC, Lipsitch M, Blázquez J, Levin BR, Baquero F. Concentration-dependent selection of small phenotypic differences in TEM beta-lactamase-mediated antibiotic resistance. *Antimicrob Agents Chemother*. 2000 Sep;44(9):2485-91. doi: [10.1128/aac.44.9.2485-2491.2000](https://doi.org/10.1128/aac.44.9.2485-2491.2000). PMID: 10952599; PMCID: PMC90089.
26. Lipsitch M, Bacon TH, Leary JJ, Antia R, Levin BR. Effects of antiviral usage on transmission dynamics of herpes simplex virus type 1 and on antiviral resistance: predictions of mathematical models. *Antimicrob Agents Chemother*. 2000 Oct;44(10):2824-35. doi: [10.1128/aac.44.10.2824-2835.2000](https://doi.org/10.1128/aac.44.10.2824-2835.2000). PMID: 10991866; PMCID: PMC90157.
27. Ebert D, Lipsitch M, Mangin KL. The Effect of Parasites on Host Population Density and Extinction: Experimental Epidemiology with *Daphnia* and Six Microparasites. *Am Nat*. 2000 Nov;156(5):459-477. doi: [10.1086/303404](https://doi.org/10.1086/303404). PMID: 29587512.

28. Lipsitch M. Measuring and interpreting associations between antibiotic use and penicillin resistance in *Streptococcus pneumoniae*. *Clin Infect Dis*. 2001 Apr 1;32(7):1044-54. doi: [10.1086/319604](https://doi.org/10.1086/319604). Epub 2001 Mar 23. PMID: 11264033.
29. Lipsitch M. Microbiology. Bacterial population genetics and disease. *Science*. 2001 Apr 6;292(5514):59-60. doi: [10.1126/science.1060498](https://doi.org/10.1126/science.1060498). PMID: 11294216.
30. Francis KP, Yu J, Bellinger-Kawahara C, Joh D, Hawkinson MJ, Xiao G, Purchio TF, Caparon MG, Lipsitch M, Contag PR. Visualizing pneumococcal infections in the lungs of live mice using bioluminescent *Streptococcus pneumoniae* transformed with a novel gram-positive lux transposon. *Infect Immun*. 2001 May;69(5):3350-8. doi: [10.1128/IAI.69.5.3350-3358.2001](https://doi.org/10.1128/IAI.69.5.3350-3358.2001). PMID: 11292758; PMCID: PMC98294.
31. Lipsitch M. Interpreting results from trials of pneumococcal conjugate vaccines: a statistical test for detecting vaccine-induced increases in carriage of nonvaccine serotypes. *Am J Epidemiol*. 2001 Jul 1;154(1):85-92. doi: [10.1093/aje/154.1.85](https://doi.org/10.1093/aje/154.1.85). PMID: 11427408.
32. Malley R, Lipsitch M, Stack A, Saladino R, Fleisher G, Pelton S, Thompson C, Briles D, Anderson P. Intranasal immunization with killed unencapsulated whole cells prevents colonization and invasive disease by capsulated pneumococci. *Infect Immun*. 2001 Aug;69(8):4870-3. doi: [10.1128/IAI.69.8.4870-4873.2001](https://doi.org/10.1128/IAI.69.8.4870-4873.2001). PMID: 11447162; PMCID: PMC98576.
33. Lipsitch M. The rise and fall of antimicrobial resistance. *Trends Microbiol*. 2001 Sep;9(9):438-44. doi: [10.1016/s0966-842x\(01\)02130-8](https://doi.org/10.1016/s0966-842x(01)02130-8). PMID: 11553456.
34. Bonten MJ, Austin DJ, Lipsitch M. Understanding the spread of antibiotic resistant pathogens in hospitals: mathematical models as tools for control. *Clin Infect Dis*. 2001 Nov 15;33(10):1739-46. doi: [10.1086/323761](https://doi.org/10.1086/323761). Epub 2001 Oct 10. PMID: 11595995.
35. Lipsitch M, Samore MH. Antimicrobial use and antimicrobial resistance: a population perspective. *Emerg Infect Dis*. 2002 Apr;8(4):347-54. doi: [10.3201/eid0804.010312](https://doi.org/10.3201/eid0804.010312). Erratum in: *Emerg Infect Dis* 2002 May;8(5):540. PMID: 11971765; PMCID: PMC2730242.
36. Lipsitch M, Singer RS, Levin BR. Antibiotics in agriculture: when is it time to close the barn door? *Proc Natl Acad Sci U S A*. 2002 Apr 30;99(9):5752-4. doi: [10.1073/pnas.092142499](https://doi.org/10.1073/pnas.092142499). PMID: 11983874; PMCID: PMC122845.
37. Harris AD, Samore MH, Lipsitch M, Kaye KS, Perencevich E, Carmeli Y. Control-group selection importance in studies of antimicrobial resistance: examples applied to *Pseudomonas aeruginosa*, Enterococci, and *Escherichia coli*. *Clin Infect Dis*. 2002 Jun 15;34(12):1558-63. doi: [10.1086/340533](https://doi.org/10.1086/340533). Epub 2002 May 23. PMID: 12032889.
38. Lipsitch M, Davis G, Corey L. Potential benefits of a serodiagnostic test for herpes simplex virus type 1 (HSV-1) to prevent neonatal HSV-1 infection. *Sex Transm Dis*. 2002 Jul;29(7):399-405. doi: [10.1097/00007435-200207000-00007](https://doi.org/10.1097/00007435-200207000-00007). PMID: 12170129.
39. Lipsitch M, Sousa AO. Historical intensity of natural selection for resistance to tuberculosis. *Genetics*. 2002 Aug;161(4):1599-607. PMID: [12196403](https://pubmed.ncbi.nlm.nih.gov/12196403/); PMCID: PMC1462208.
40. Fisman DN, Lipsitch M, Hook EW 3rd, Goldie SJ. Projection of the future dimensions and costs of the genital herpes simplex type 2 epidemic in the United States. *Sex Transm Dis*. 2002 Oct;29(10):608-22. doi: [10.1097/00007435-200210000-00008](https://doi.org/10.1097/00007435-200210000-00008). PMID: 12370529.
41. Singer RS, Finch R, Wegener HC, Bywater R, Walters J, Lipsitch M. Antibiotic resistance--the interplay between antibiotic use in animals and human beings.

- Lancet Infect Dis. 2003 Jan;3(1):47-51. doi: [10.1016/s1473-3099\(03\)00490-0](https://doi.org/10.1016/s1473-3099(03)00490-0). PMID: 12505035.
42. Lipsitch M, Bergstrom CT, Antia R. Effect of human leukocyte antigen heterozygosity on infectious disease outcome: the need for allele-specific measures. *BMC Med Genet.* 2003 Jan 24;4:2. doi: [10.1186/1471-2350-4-2](https://doi.org/10.1186/1471-2350-4-2). Epub 2003 Jan 24. PMID: 12542841; PMCID: PMC149356.
 43. Malley R, Henneke P, Morse SC, Cieslewicz MJ, Lipsitch M, Thompson CM, Kurt-Jones E, Paton JC, Wessels MR, Golenbock DT. Recognition of pneumolysin by Toll-like receptor 4 confers resistance to pneumococcal infection. *Proc Natl Acad Sci U S A.* 2003 Feb 18;100(4):1966-71. doi: [10.1073/pnas.0435928100](https://doi.org/10.1073/pnas.0435928100). Epub 2003 Feb 4. PMID: 12569171; PMCID: PMC149942.
 44. O'Brien KL, Nohynek H; World Health Organization Pneumococcal Vaccine Trials Carraige Working Group. Report from a WHO working group: standard method for detecting upper respiratory carriage of *Streptococcus pneumoniae*. *Pediatr Infect Dis J.* 2003 Feb;22(2):133-40. doi: [10.1097/01.inf.0000048676.93549.d1](https://doi.org/10.1097/01.inf.0000048676.93549.d1). PMID: 12586977.
 45. Lipsitch M, Murray MB. Multiple equilibria: tuberculosis transmission require unrealistic assumptions. *Theor Popul Biol.* 2003 Mar;63(2):169-70. doi: [10.1016/s0040-5809\(02\)00037-0](https://doi.org/10.1016/s0040-5809(02)00037-0). PMID: 12615499.
 46. McCormick AW, Whitney CG, Farley MM, Lynfield R, Harrison LH, Bennett NM, Schaffner W, Reingold A, Hadler J, Cieslak P, Samore MH, Lipsitch M. Geographic diversity and temporal trends of antimicrobial resistance in *Streptococcus pneumoniae* in the United States. *Nat Med.* 2003 Apr;9(4):424-30. doi: [10.1038/nm839](https://doi.org/10.1038/nm839). Epub 2003 Mar 10. PMID: 12627227.
 47. Lipsitch M, Cohen T, Cooper B, Robins JM, Ma S, James L, Gopalakrishna G, Chew SK, Tan CC, Samore MH, Fisman D, Murray M. Transmission dynamics and control of severe acute respiratory syndrome. *Science.* 2003 Jun 20;300(5627):1966-70. doi: [10.1126/science.1086616](https://doi.org/10.1126/science.1086616). Epub 2003 May 23. PMID: 12766207; PMCID: PMC2760158.
 48. Trzcinski K, Thompson CM, Lipsitch M. Construction of otherwise isogenic serotype 6B, 7F, 14, and 19F capsular variants of *Streptococcus pneumoniae* strain TIGR4. *Appl Environ Microbiol.* 2003 Dec;69(12):7364-70. doi: [10.1128/aem.69.12.7364-7370.2003](https://doi.org/10.1128/aem.69.12.7364-7370.2003). PMID: 14660386; PMCID: PMC309976.
 49. Cooper B, Lipsitch M. The analysis of hospital infection data using hidden Markov models. *Biostatistics.* 2004 Apr;5(2):223-37. doi: [10.1093/biostatistics/5.2.223](https://doi.org/10.1093/biostatistics/5.2.223). PMID: 15054027.
 50. Perencevich EN, Fisman DN, Lipsitch M, Harris AD, Morris JG Jr, Smith DL. Projected benefits of active surveillance for vancomycin-resistant enterococci in intensive care units. *Clin Infect Dis.* 2004 Apr 15;38(8):1108-15. doi: [10.1086/382886](https://doi.org/10.1086/382886). Epub 2004 Apr 5. PMID: 15095215.
 51. Trzciński K, Thompson CM, Lipsitch M. Single-step capsular transformation and acquisition of penicillin resistance in *Streptococcus pneumoniae*. *J Bacteriol.* 2004 Jun;186(11):3447-52. doi: [10.1128/JB.186.11.3447-3452.2004](https://doi.org/10.1128/JB.186.11.3447-3452.2004). PMID: 15150231; PMCID: PMC415782.
 52. Laine C, Mwangi T, Thompson CM, Obiero J, Lipsitch M, Scott JA. Age-specific immunoglobulin G (IgG) and IgA to pneumococcal protein antigens in a population in coastal Kenya. *Infect Immun.* 2004 Jun;72(6):3331-5. doi: [10.1128/IAI.72.6.3331-3335.2004](https://doi.org/10.1128/IAI.72.6.3331-3335.2004). PMID: 15155637; PMCID: PMC415695.
 53. Teklehaimanot HD, Schwatz J, Teklehaimanot A, Lipsitch M. Alert threshold algorithms and malaria epidemic detection. *Emerg Infect Dis.* 2004

- Jul;10(7):1220-6. doi: [10.3201/eid1007.030722](https://doi.org/10.3201/eid1007.030722). PMID: 15324541; PMCID: PMC3323320.
54. Bergstrom CT, Lo M, Lipsitch M. Ecological theory suggests that antimicrobial cycling will not reduce antimicrobial resistance in hospitals. *Proc Natl Acad Sci U S A*. 2004 Sep 7;101(36):13285-90. doi: [10.1073/pnas.0402298101](https://doi.org/10.1073/pnas.0402298101). Epub 2004 Aug 12. PMID: 15308772; PMCID: PMC516561.
 55. Lipsitch M, Bergstrom CT. Invited commentary: real-time tracking of control measures for emerging infections. *Am J Epidemiol*. 2004 Sep 15;160(6):517-9; discussion 520. doi: [10.1093/aje/kwh256](https://doi.org/10.1093/aje/kwh256). PMID: 15353410.
 56. Teklehaimanot HD, Lipsitch M, Teklehaimanot A, Schwartz J. Weather-based prediction of *Plasmodium falciparum* malaria in epidemic-prone regions of Ethiopia I. Patterns of lagged weather effects reflect biological mechanisms. *Malar J*. 2004 Nov 12;3:41. doi: [10.1186/1475-2875-3-41](https://doi.org/10.1186/1475-2875-3-41). PMID: 15541174; PMCID: PMC535540.
 57. Teklehaimanot HD, Schwartz J, Teklehaimanot A, Lipsitch M. Weather-based prediction of *Plasmodium falciparum* malaria in epidemic-prone regions of Ethiopia II. Weather-based prediction systems perform comparably to early detection systems in identifying times for interventions. *Malar J*. 2004 Nov 19;3:44. doi: [10.1186/1475-2875-3-44](https://doi.org/10.1186/1475-2875-3-44). PMID: 15555061; PMCID: PMC535541.
 58. Mills CE, Robins JM, Lipsitch M. Transmissibility of 1918 pandemic influenza. *Nature*. 2004 Dec 16;432(7019):904-6. doi: [10.1038/nature03063](https://doi.org/10.1038/nature03063). PMID: 15602562; PMCID: PMC7095078.
 59. Lipsitch M, Whitney CG, Zell E, Kaijalainen T, Dagan R, Malley R. Are anticapsular antibodies the primary mechanism of protection against invasive pneumococcal disease? *PLoS Med*. 2005 Jan;2(1):e15. doi: [10.1371/journal.pmed.0020015](https://doi.org/10.1371/journal.pmed.0020015). Epub 2005 Jan 25. PMID: 15696204; PMCID: PMC545206.
 60. Lipsitch M. Ethics of rationing the flu vaccine. *Science*. 2005 Jan 7;307(5706):41. doi: [10.1126/science.307.5706.41b](https://doi.org/10.1126/science.307.5706.41b). PMID: 15637252.
 61. Malley R, Trzciński K, Srivastava A, Thompson CM, Anderson PW, Lipsitch M. CD4+ T cells mediate antibody-independent acquired immunity to pneumococcal colonization. *Proc Natl Acad Sci U S A*. 2005 Mar 29;102(13):4848-53. doi: [10.1073/pnas.0501254102](https://doi.org/10.1073/pnas.0501254102). Epub 2005 Mar 21. PMID: 15781870; PMCID: PMC555733.
 62. Trzciński K, MacNeil A, Klugman KP, Lipsitch M. Capsule homology does not increase the frequency of transformation of linked penicillin binding proteins PBP 1a and PBP 2x in *Streptococcus pneumoniae*. *Antimicrob Agents Chemother*. 2005 Apr;49(4):1591-2. doi: [10.1128/AAC.49.4.1591-1592.2005](https://doi.org/10.1128/AAC.49.4.1591-1592.2005). PMID: 15793147; PMCID: PMC1068637.
 63. Lipsitch M. Pandemic flu: we are not prepared. *MedGenMed*. 2005 Apr 15;7(2):56. PMID: [16369434](https://pubmed.ncbi.nlm.nih.gov/16369434/); PMCID: PMC1681602.
 64. Huang SS, Finkelstein JA, Lipsitch M. Modeling community- and individual-level effects of child-care center attendance on pneumococcal carriage. *Clin Infect Dis*. 2005 May 1;40(9):1215-22. doi: [10.1086/428580](https://doi.org/10.1086/428580). Epub 2005 Mar 23. PMID: 15825020.
 65. Dagan R, Givon-Lavi N, Fraser D, Lipsitch M, Siber GR, Kohberger R. Serum serotype-specific pneumococcal anticapsular immunoglobulin g concentrations after immunization with a 9-valent conjugate pneumococcal vaccine correlate with nasopharyngeal acquisition of pneumococcus. *J Infect Dis*. 2005 Aug 1;192(3):367-76. doi: [10.1086/431679](https://doi.org/10.1086/431679). Epub 2005 Jun 28. PMID: 15995949.

66. Trzcinski K, Thompson C, Malley R, Lipsitch M. Antibodies to conserved pneumococcal antigens correlate with, but are not required for, protection against pneumococcal colonization induced by prior exposure in a mouse model. *Infect Immun*. 2005 Oct;73(10):7043-6. doi: [10.1128/IAI.73.10.7043-7046.2005](https://doi.org/10.1128/IAI.73.10.7043-7046.2005). PMID: 16177389; PMCID: PMC1230924.
67. Samore MH, Lipsitch M, Alder SC, Haddadin B, Stoddard G, Williamson J, Sebastian K, Carroll K, Ergonul O, Carmeli Y, Sande MA. Mechanisms by which antibiotics promote dissemination of resistant pneumococci in human populations. *Am J Epidemiol*. 2006 Jan 15;163(2):160-70. doi: [10.1093/aje/kwj021](https://doi.org/10.1093/aje/kwj021). Epub 2005 Nov 30. PMID: 16319292.
68. Asquith B, Edwards CT, Lipsitch M, McLean AR. Inefficient cytotoxic T lymphocyte-mediated killing of HIV-1-infected cells in vivo. *PLoS Biol*. 2006 Apr;4(4):e90. doi: [10.1371/journal.pbio.0040090](https://doi.org/10.1371/journal.pbio.0040090). Epub 2006 Mar 14. PMID: 16515366; PMCID: PMC1395353.
69. Malley R, Srivastava A, Lipsitch M, Thompson CM, Watkins C, Tzianabos A, Anderson PW. Antibody-independent, interleukin-17A-mediated, cross-serotype immunity to pneumococci in mice immunized intranasally with the cell wall polysaccharide. *Infect Immun*. 2006 Apr;74(4):2187-95. doi: [10.1128/IAI.74.4.2187-2195.2006](https://doi.org/10.1128/IAI.74.4.2187-2195.2006). PMID: 16552049; PMCID: PMC1418935.
70. Trzcinski K, Thompson CM, Gilbey AM, Dowson CG, Lipsitch M. Incremental increase in fitness cost with increased beta -lactam resistance in pneumococci evaluated by competition in an infant rat nasal colonization model. *J Infect Dis*. 2006 May 1;193(9):1296-303. doi: [10.1086/501367](https://doi.org/10.1086/501367). Epub 2006 Mar 17. PMID: 16586368.
71. Palmer ME, Lipsitch M. The influence of hitchhiking and deleterious mutation upon asexual mutation rates. *Genetics*. 2006 May;173(1):461-72. doi: [10.1534/genetics.105.049445](https://doi.org/10.1534/genetics.105.049445). Epub 2006 Feb 19. PMID: 16489233; PMCID: PMC1461451.
72. Cohen T, Lipsitch M, Walensky RP, Murray M. Beneficial and perverse effects of isoniazid preventive therapy for latent tuberculosis infection in HIV-tuberculosis coinfecting populations. *Proc Natl Acad Sci U S A*. 2006 May 2;103(18):7042-7. doi: [10.1073/pnas.0600349103](https://doi.org/10.1073/pnas.0600349103). Epub 2006 Apr 21. PMID: 16632605; PMCID: PMC1459015.
73. Lipsitch M, Robins JM, Mills CE, Bergstrom CT. Multiple outbreaks and flu containment plans. *Science*. 2006 May 12;312(5775):845. doi: [10.1126/science.312.5775.845b](https://doi.org/10.1126/science.312.5775.845b). PMID: 16690840.
74. Mills CE, Robins JM, Bergstrom CT, Lipsitch M. Pandemic influenza: risk of multiple introductions and the need to prepare for them. *PLoS Med*. 2006 Jun;3(6):e135. doi: [10.1371/journal.pmed.0030135](https://doi.org/10.1371/journal.pmed.0030135). PMID: 17214503; PMCID: PMC1370924.
75. Wang YC, Lipsitch M. Upgrading antibiotic use within a class: tradeoff between resistance and treatment success. *Proc Natl Acad Sci U S A*. 2006 Jun 20;103(25):9655-60. doi: [10.1073/pnas.0600636103](https://doi.org/10.1073/pnas.0600636103). Epub 2006 Jun 13. PMID: 16772381; PMCID: PMC1480462.
76. Regev-Yochay G, Trzcinski K, Thompson CM, Malley R, Lipsitch M. Interference between *Streptococcus pneumoniae* and *Staphylococcus aureus*: In vitro hydrogen peroxide-mediated killing by *Streptococcus pneumoniae*. *J Bacteriol*. 2006 Jul;188(13):4996-5001. doi: [10.1128/JB.00317-06](https://doi.org/10.1128/JB.00317-06). PMID: 16788209; PMCID: PMC1482988.

77. Lipsitch M, Cohen T, Murray M, Levin BR. Antiviral resistance and the control of pandemic influenza. *PLoS Med.* 2007 Jan;4(1):e15. doi: [10.1371/journal.pmed.0040015](https://doi.org/10.1371/journal.pmed.0040015). PMID: 17253900; PMCID: PMC1779817.
78. Hanage WP, Huang SS, Lipsitch M, Bishop CJ, Godoy D, Pelton SI, Goldstein R, Huot H, Finkelstein JA. Diversity and antibiotic resistance among nonvaccine serotypes of *Streptococcus pneumoniae* carriage isolates in the post-heptavalent conjugate vaccine era. *J Infect Dis.* 2007 Feb 1;195(3):347-52. doi: [10.1086/510249](https://doi.org/10.1086/510249). Epub 2006 Dec 27. PMID: 17205472.
79. Wallinga J, Lipsitch M. How generation intervals shape the relationship between growth rates and reproductive numbers. *Proc Biol Sci.* 2007 Feb 22;274(1609):599-604. doi: [10.1098/rspb.2006.3754](https://doi.org/10.1098/rspb.2006.3754). PMID: 17476782; PMCID: PMC1766383.
80. Högberg L, Geli P, Ringberg H, Melander E, Lipsitch M, Ekdahl K. Age- and serogroup-related differences in observed durations of nasopharyngeal carriage of penicillin-resistant pneumococci. *J Clin Microbiol.* 2007 Mar;45(3):948-52. doi: [10.1128/JCM.01913-06](https://doi.org/10.1128/JCM.01913-06). Epub 2007 Jan 3. PMID: 17202280; PMCID: PMC1829115.
81. McCormick AW, Walensky RP, Lipsitch M, Losina E, Hsu H, Weinstein MC, Paltiel AD, Freedberg KA, Seage GR 3rd. The effect of antiretroviral therapy on secondary transmission of HIV among men who have sex with men. *Clin Infect Dis.* 2007 Apr 15;44(8):1115-22. doi: [10.1086/512816](https://doi.org/10.1086/512816). Epub 2007 Mar 9. PMID: 17366461; PMCID: PMC2365722.
82. Hatchett RJ, Mecher CE, Lipsitch M. Public health interventions and epidemic intensity during the 1918 influenza pandemic. *Proc Natl Acad Sci U S A.* 2007 May 1;104(18):7582-7. doi: [10.1073/pnas.0610941104](https://doi.org/10.1073/pnas.0610941104). Epub 2007 Apr 6. PMID: 17416679; PMCID: PMC1849867.
83. Basset A, Trzcinski K, Hermos C, O'Brien KL, Reid R, Santosham M, McAdam AJ, Lipsitch M, Malley R. Association of the pneumococcal pilus with certain capsular serotypes but not with increased virulence. *J Clin Microbiol.* 2007 Jun;45(6):1684-9. doi: [10.1128/JCM.00265-07](https://doi.org/10.1128/JCM.00265-07). Epub 2007 Mar 28. PMID: 17392439; PMCID: PMC1933072.
84. Pitzer VE, Olsen SJ, Bergstrom CT, Dowell SF, Lipsitch M. Little evidence for genetic susceptibility to influenza A (H5N1) from family clustering data. *Emerg Infect Dis.* 2007 Jul;13(7):1074-6. doi: [10.3201/eid1307.061538](https://doi.org/10.3201/eid1307.061538). PMID: 18214184; PMCID: PMC2878232.
85. Pitzer VE, Leung GM, Lipsitch M. Estimating variability in the transmission of severe acute respiratory syndrome to household contacts in Hong Kong, China. *Am J Epidemiol.* 2007 Aug 1;166(3):355-63. doi: [10.1093/aje/kwm082](https://doi.org/10.1093/aje/kwm082). Epub 2007 May 10. PMID: 17493952; PMCID: PMC7110150.
86. Regev-Yochay G, Trzcinski K, Thompson CM, Lipsitch M, Malley R. SpxB is a suicide gene of *Streptococcus pneumoniae* and confers a selective advantage in an in vivo competitive colonization model. *J Bacteriol.* 2007 Sep;189(18):6532-9. doi: [10.1128/JB.00813-07](https://doi.org/10.1128/JB.00813-07). Epub 2007 Jul 13. PMID: 17631628; PMCID: PMC2045178.
87. Malley R, Lipsitch M, Bogaert D, Thompson CM, Hermans P, Watkins AC, Sethi S, Murphy TF. Serum antipneumococcal antibodies and pneumococcal colonization in adults with chronic obstructive pulmonary disease. *J Infect Dis.* 2007 Sep 15;196(6):928-35. doi: [10.1086/520937](https://doi.org/10.1086/520937). Epub 2007 Aug 7. PMID: 17703425.
88. Lipsitch M, O'Neill K, Cordy D, Bugalter B, Trzcinski K, Thompson CM, Goldstein R, Pelton S, Huot H, Bouchet V, Reid R, Santosham M, O'Brien KL. Strain

- characteristics of *Streptococcus pneumoniae* carriage and invasive disease isolates during a cluster-randomized clinical trial of the 7-valent pneumococcal conjugate vaccine. *J Infect Dis.* 2007 Oct 15;196(8):1221-7. doi: [10.1086/521831](https://doi.org/10.1086/521831). Epub 2007 Sep 17. PMID: 17955441; PMCID: PMC3350793.
89. Lipsitch M, O'Hagan JJ. Patterns of antigenic diversity and the mechanisms that maintain them. *J R Soc Interface.* 2007 Oct 22;4(16):787-802. doi: [10.1098/rsif.2007.0229](https://doi.org/10.1098/rsif.2007.0229). PMID: 17426010; PMCID: PMC2394542.
 90. Basset A, Thompson CM, Hollingshead SK, Briles DE, Ades EW, Lipsitch M, Malley R. Antibody-independent, CD4+ T-cell-dependent protection against pneumococcal colonization elicited by intranasal immunization with purified pneumococcal proteins. *Infect Immun.* 2007 Nov;75(11):5460-4. doi: [10.1128/IAI.00773-07](https://doi.org/10.1128/IAI.00773-07). Epub 2007 Aug 13. PMID: 17698570; PMCID: PMC2168310.
 91. Regev-Yochay G, Malley R, Rubinstein E, Raz M, Dagan R, Lipsitch M. In vitro bactericidal activity of *Streptococcus pneumoniae* and bactericidal susceptibility of *Staphylococcus aureus* strains isolated from cocolonized versus noncocolonized children. *J Clin Microbiol.* 2008 Feb;46(2):747-9. doi: [10.1128/JCM.01781-07](https://doi.org/10.1128/JCM.01781-07). Epub 2007 Nov 26. PMID: 18039795; PMCID: PMC2238136.
 92. Dagan R, Barkai G, Givon-Lavi N, Sharf AZ, Vardy D, Cohen T, Lipsitch M, Greenberg D. Seasonality of antibiotic-resistant streptococcus pneumoniae that causes acute otitis media: a clue for an antibiotic-restriction policy? *J Infect Dis.* 2008 Apr 15;197(8):1094-102. doi: [10.1086/528995](https://doi.org/10.1086/528995). PMID: 18419528; PMCID: PMC2652754.
 93. Walensky RP, Wood R, Weinstein MC, Martinson NA, Losina E, Fofana MO, Goldie SJ, Divi N, Yazdanpanah Y, Wang B, Paltiel AD, Freedberg KA; CEPAC-International Investigators. Scaling up antiretroviral therapy in South Africa: the impact of speed on survival. *J Infect Dis.* 2008 May 1;197(9):1324-32. doi: [10.1086/587184](https://doi.org/10.1086/587184). PMID: 18422445; PMCID: PMC2423492.
 94. Kenah E, Lipsitch M, Robins JM. Generation interval contraction and epidemic data analysis. *Math Biosci.* 2008 May;213(1):71-9. doi: [10.1016/j.mbs.2008.02.007](https://doi.org/10.1016/j.mbs.2008.02.007). Epub 2008 Feb 29. PMID: 18394654; PMCID: PMC2365921.
 95. Weinberger DM, Dagan R, Givon-Lavi N, Regev-Yochay G, Malley R, Lipsitch M. Epidemiologic evidence for serotype-specific acquired immunity to pneumococcal carriage. *J Infect Dis.* 2008 Jun 1;197(11):1511-8. doi: [10.1086/587941](https://doi.org/10.1086/587941). PMID: 18471062.
 96. Trzciński K, Thompson CM, Srivastava A, Basset A, Malley R, Lipsitch M. Protection against nasopharyngeal colonization by *Streptococcus pneumoniae* is mediated by antigen-specific CD4+ T cells. *Infect Immun.* 2008 Jun;76(6):2678-84. doi: [10.1128/IAI.00141-08](https://doi.org/10.1128/IAI.00141-08). Epub 2008 Apr 7. PMID: 18391006; PMCID: PMC2423086.
 97. Cohen T, Lipsitch M. Too little of a good thing: a paradox of moderate infection control. *Epidemiology.* 2008 Jul;19(4):588-9. doi: [10.1097/EDE.0b013e31817734ba](https://doi.org/10.1097/EDE.0b013e31817734ba). PMID: 18552592; PMCID: PMC2652751.
 98. Regev-Yochay G, Bogaert D, Malley R, Hermans PW, Veenhoven RH, Sanders EA, Lipsitch M, Rubinstein E. Does pneumococcal conjugate vaccine influence *Staphylococcus aureus* carriage in children? *Clin Infect Dis.* 2008 Jul 15;47(2):289-91; author reply 291-2. doi: [10.1086/589573](https://doi.org/10.1086/589573). PMID: 18564933.

99. McDonnell Norms Group. Antibiotic overuse: the influence of social norms. *J Am Coll Surg*. 2008 Aug;207(2):265-75. doi: [10.1016/j.jamcollsurg.2008.02.035](https://doi.org/10.1016/j.jamcollsurg.2008.02.035). Epub 2008 May 12. PMID: 18656057.
100. Lu YJ, Gross J, Bogaert D, Finn A, Bagraade L, Zhang Q, Kolls JK, Srivastava A, Lundgren A, Forte S, Thompson CM, Harney KF, Anderson PW, Lipsitch M, Malley R. Interleukin-17A mediates acquired immunity to pneumococcal colonization. *PLoS Pathog*. 2008 Sep 19;4(9):e1000159. doi: [10.1371/journal.ppat.1000159](https://doi.org/10.1371/journal.ppat.1000159). PMID: 18802458; PMCID: PMC2528945.
101. Klugman KP, Astley CM, Lipsitch M. Time from illness onset to death, 1918 influenza and pneumococcal pneumonia. *Emerg Infect Dis*. 2009 Feb;15(2):346-7. doi: [10.3201/eid1502.081208](https://doi.org/10.3201/eid1502.081208). PMID: 19193293; PMCID: PMC2657896.
102. Lipsitch M, Viboud C. Influenza seasonality: lifting the fog. *Proc Natl Acad Sci U S A*. 2009 Mar 10;106(10):3645-6. doi: [10.1073/pnas.0900933106](https://doi.org/10.1073/pnas.0900933106). PMID: 19276125; PMCID: PMC2656132.
103. Regev-Yochay G, Lipsitch M, Basset A, Rubinstein E, Dagan R, Raz M, Malley R. The pneumococcal pilus predicts the absence of *Staphylococcus aureus* co-colonization in pneumococcal carriers. *Clin Infect Dis*. 2009 Mar 15;48(6):760-3. doi: [10.1086/597040](https://doi.org/10.1086/597040). PMID: 19207082; PMCID: PMC2674784.
104. Bogaert D, Weinberger D, Thompson C, Lipsitch M, Malley R. Impaired innate and adaptive immunity to *Streptococcus pneumoniae* and its effect on colonization in an infant mouse model. *Infect Immun*. 2009 Apr;77(4):1613-22. doi: [10.1128/IAI.00871-08](https://doi.org/10.1128/IAI.00871-08). Epub 2009 Jan 21. PMID: 19168741; PMCID: PMC2663178.
105. Wu JT, Leung GM, Lipsitch M, Cooper BS, Riley S. Hedging against antiviral resistance during the next influenza pandemic using small stockpiles of an alternative chemotherapy. *PLoS Med*. 2009 May 19;6(5):e1000085. doi: [10.1371/journal.pmed.1000085](https://doi.org/10.1371/journal.pmed.1000085). Epub 2009 May 19. PMID: 19440354; PMCID: PMC2680070.
106. Weinberger DM, Trzciński K, Lu YJ, Bogaert D, Brandes A, Galagan J, Anderson PW, Malley R, Lipsitch M. Pneumococcal capsular polysaccharide structure predicts serotype prevalence. *PLoS Pathog*. 2009 Jun;5(6):e1000476. doi: [10.1371/journal.ppat.1000476](https://doi.org/10.1371/journal.ppat.1000476). Epub 2009 Jun 12. PMID: 19521509; PMCID: PMC2689349.
107. Huang SS, Hinrichsen VL, Stevenson AE, Rifas-Shiman SL, Kleinman K, Pelton SI, Lipsitch M, Hanage WP, Lee GM, Finkelstein JA. Continued impact of pneumococcal conjugate vaccine on carriage in young children. *Pediatrics*. 2009 Jul;124(1):e1-11. doi: [10.1542/peds.2008-3099](https://doi.org/10.1542/peds.2008-3099). PMID: 19564254; PMCID: PMC2782668.
108. Lipsitch M, Riley S, Cauchemez S, Ghani AC, Ferguson NM. Managing and reducing uncertainty in an emerging influenza pandemic. *N Engl J Med*. 2009 Jul 9;361(2):112-5. doi: [10.1056/NEJMp0904380](https://doi.org/10.1056/NEJMp0904380). Epub 2009 May 27. PMID: 19474417; PMCID: PMC3066026.
109. Lee GM, Huang SS, Rifas-Shiman SL, Hinrichsen VL, Pelton SI, Kleinman K, Hanage WP, Lipsitch M, McAdam AJ, Finkelstein JA. Epidemiology and risk factors for *Staphylococcus aureus* colonization in children in the post-PCV7 era. *BMC Infect Dis*. 2009 Jul 11;9:110. doi: [10.1186/1471-2334-9-110](https://doi.org/10.1186/1471-2334-9-110). PMID: 19594890; PMCID: PMC2716346.
110. McCaw JM, Wood JG, McBryde ES, Nolan TM, Wu JT, Lipsitch M, McVernon J. Understanding Australia's influenza pandemic policy on the strategic use of the antiviral drug stockpile. *Med J Aust*. 2009 Aug 3;191(3):136-7. doi: [10.1186/1471-2334-9-110](https://doi.org/10.1186/1471-2334-9-110). PMID: 19645639; PMCID: PMC3073016.

111. Walensky RP, Wolf LL, Wood R, Fofana MO, Freedberg KA, Martinson NA, Paltiel AD, Anglaret X, Weinstein MC, Losina E; CEPAC (Cost-Effectiveness of Preventing AIDS Complications)-International Investigators. When to start antiretroviral therapy in resource-limited settings. *Ann Intern Med.* 2009 Aug 4;151(3):157-66. doi: [10.7326/0003-4819-151-3-200908040-00138](https://doi.org/10.7326/0003-4819-151-3-200908040-00138). Epub 2009 Jul 20. PMID: 19620143; PMCID: PMC3092478.
112. Goldstein E, Paur K, Fraser C, Kenah E, Wallinga J, Lipsitch M. Reproductive numbers, epidemic spread and control in a community of households. *Math Biosci.* 2009 Sep;221(1):11-25. doi: [10.1016/j.mbs.2009.06.002](https://doi.org/10.1016/j.mbs.2009.06.002). Epub 2009 Jun 25. PMID: 19559715; PMCID: PMC2731010.
113. Lipsitch M, Lajous M, O'Hagan JJ, Cohen T, Miller JC, Goldstein E, Danon L, Wallinga J, Riley S, Dowell SF, Reed C, McCarron M. Use of cumulative incidence of novel influenza A/H1N1 in foreign travelers to estimate lower bounds on cumulative incidence in Mexico. *PLoS One.* 2009 Sep 9;4(9):e6895. doi: [10.1371/journal.pone.0006895](https://doi.org/10.1371/journal.pone.0006895). PMID: 19742302; PMCID: PMC2731883.
114. Pitzer VE, Lipsitch M. Exploring the relationship between incidence and the average age of infection during seasonal epidemics. *J Theor Biol.* 2009 Sep 21;260(2):175-85. doi: [10.1016/j.jtbi.2009.06.008](https://doi.org/10.1016/j.jtbi.2009.06.008). Epub 2009 Jun 13. PMID: 19527734; PMCID: PMC2745250.
115. Losina E, Touré H, Uhler LM, Anglaret X, Paltiel AD, Balestre E, Walensky RP, Messou E, Weinstein MC, Dabis F, Freedberg KA; ART-LINC Collaboration of International Epidemiological Databases to Evaluate AIDS (IeDEA); CEPAC International investigators. Cost-effectiveness of preventing loss to follow-up in HIV treatment programs: a Côte d'Ivoire appraisal. *PLoS Med.* 2009 Oct;6(10):e1000173. doi: [10.1371/journal.pmed.1000173](https://doi.org/10.1371/journal.pmed.1000173). Epub 2009 Oct 27. PMID: 19859538; PMCID: PMC2762030.
116. Lipsitch M, Hayden FG, Cowling BJ, Leung GM. How to maintain surveillance for novel influenza A H1N1 when there are too many cases to count. *Lancet.* 2009 Oct 3;374(9696):1209-11. doi: [10.1016/S0140-6736\(09\)61377-5](https://doi.org/10.1016/S0140-6736(09)61377-5). Epub 2009 Aug 11. PMID: 19679345.
117. White LF, Wallinga J, Finelli L, Reed C, Riley S, Lipsitch M, Pagano M. Estimation of the reproductive number and the serial interval in early phase of the 2009 influenza A/H1N1 pandemic in the USA. *Influenza Other Respir Viruses.* 2009 Nov;3(6):267-76. doi: [10.1111/j.1750-2659.2009.00106.x](https://doi.org/10.1111/j.1750-2659.2009.00106.x). PMID: 19903209; PMCID: PMC2782458.
118. Presanis AM, De Angelis D; New York City Swine Flu Investigation Team, Hagy A, Reed C, Riley S, Cooper BS, Finelli L, Biedrzycki P, Lipsitch M. The severity of pandemic H1N1 influenza in the United States, from April to July 2009: a Bayesian analysis. *PLoS Med.* 2009 Dec;6(12):e1000207. doi: [10.1371/journal.pmed.1000207](https://doi.org/10.1371/journal.pmed.1000207). Epub 2009 Dec 8. PMID: 19997612; PMCID: PMC2784967.
119. Reed C, Angulo FJ, Swerdlow DL, Lipsitch M, Meltzer MI, Jernigan D, Finelli L. Estimates of the prevalence of pandemic (H1N1) 2009, United States, April-July 2009. *Emerg Infect Dis.* 2009 Dec;15(12):2004-7. doi: [10.3201/eid1512.091413](https://doi.org/10.3201/eid1512.091413). PMID: 19961687; PMCID: PMC3375879.
120. Lipsitch M, Colijn C, Cohen T, Hanage WP, Fraser C. No coexistence for free: neutral null models for multistrain pathogens. *Epidemics.* 2009 Mar;1(1):2-13. doi: [10.1016/j.epidem.2008.07.001](https://doi.org/10.1016/j.epidem.2008.07.001). Epub 2008 Nov 4. PMID: 21352747; PMCID: PMC3099423.
121. Goldstein E, Dushoff J, Ma J, Plotkin JB, Earn DJ, Lipsitch M. Reconstructing influenza incidence by deconvolution of daily mortality time series. *Proc Natl*

- Acad Sci U S A. 2009 Dec 22;106(51):21825-9. doi: [10.1073/pnas.0902958106](https://doi.org/10.1073/pnas.0902958106). Epub 2009 Dec 18. PMID: 20080801; PMCID: PMC2796142.
122. Bogaert D, Thompson CM, Trzcinski K, Malley R, Lipsitch M. The role of complement in innate and adaptive immunity to pneumococcal colonization and sepsis in a murine model. *Vaccine*. 2010 Jan 8;28(3):681-5. doi: [10.1016/j.vaccine.2009.10.085](https://doi.org/10.1016/j.vaccine.2009.10.085). Epub 2009 Nov 3. PMID: 19892042; PMCID: PMC2810519.
 123. Wallinga J, van Boven M, Lipsitch M. Optimizing infectious disease interventions during an emerging epidemic. *Proc Natl Acad Sci U S A*. 2010 Jan 12;107(2):923-8. doi: [10.1073/pnas.0908491107](https://doi.org/10.1073/pnas.0908491107). Epub 2009 Dec 28. PMID: 20080777; PMCID: PMC2818907.
 124. Shaman J, Pitzer VE, Viboud C, Grenfell BT, Lipsitch M. Absolute humidity and the seasonal onset of influenza in the continental United States. *PLoS Biol*. 2010 Feb 23;8(2):e1000316. doi: [10.1371/journal.pbio.1000316](https://doi.org/10.1371/journal.pbio.1000316). Erratum in: *PLoS Biol*. 2010;8(3). doi: [10.1371/annotation/35686514-b7a9-4f65-9663-7baefc0d63c0](https://doi.org/10.1371/annotation/35686514-b7a9-4f65-9663-7baefc0d63c0). PMID: 20186267; PMCID: PMC2826374.
 125. Goldstein E, Miller JC, O'Hagan JJ, Lipsitch M. Pre-dispensing of antivirals to high-risk individuals in an influenza pandemic. *Influenza Other Respir Viruses*. 2010 Mar;4(2):101-12. doi: [10.1111/j.1750-2659.2009.00128.x](https://doi.org/10.1111/j.1750-2659.2009.00128.x). PMID: 20167050; PMCID: PMC3075926.
 126. Lipsitch M, Tchetgen Tchetgen E, Cohen T. Negative controls: a tool for detecting confounding and bias in observational studies. *Epidemiology*. 2010 May;21(3):383-8. doi: [10.1097/EDE.0b013e3181d61eeb](https://doi.org/10.1097/EDE.0b013e3181d61eeb). Erratum in: *Epidemiology*. 2010 Jul;21(4):589. PMID: 20335814; PMCID: PMC3053408.
 127. Miller JC, Danon L, O'Hagan JJ, Goldstein E, Lajous M, Lipsitch M. Student behavior during a school closure caused by pandemic influenza A/H1N1. *PLoS One*. 2010 May 5;5(5):e10425. doi: [10.1371/journal.pone.0010425](https://doi.org/10.1371/journal.pone.0010425). PMID: 20463960; PMCID: PMC2864742.
 128. Goldstein E, Apolloni A, Lewis B, Miller JC, Macauley M, Eubank S, Lipsitch M, Wallinga J. Distribution of vaccine/antivirals and the 'least spread line' in a stratified population. *J R Soc Interface*. 2010 May 6;7(46):755-64. doi: [10.1098/rsif.2009.0393](https://doi.org/10.1098/rsif.2009.0393). Epub 2009 Oct 14. PMID: 19828505; PMCID: PMC2874227.
 129. Lau LL, Cowling BJ, Fang VJ, Chan KH, Lau EH, Lipsitch M, Cheng CK, Houck PM, Uyeki TM, Peiris JS, Leung GM. Viral shedding and clinical illness in naturally acquired influenza virus infections. *J Infect Dis*. 2010 May 15;201(10):1509-16. doi: [10.1086/652241](https://doi.org/10.1086/652241). PMID: 20377412; PMCID: PMC3060408.
 130. Van Kerkhove MD, Asikainen T, Becker NG, Bjorge S, Desenclos JC, dos Santos T, Fraser C, Leung GM, Lipsitch M, Longini IM Jr, McBryde ES, Roth CE, Shay DK, Smith DJ, Wallinga J, White PJ, Ferguson NM, Riley S; WHO Informal Network for Mathematical Modelling for Pandemic Influenza H1N1 2009 (Working Group on Data Needs). Studies needed to address public health challenges of the 2009 H1N1 influenza pandemic: insights from modeling. *PLoS Med*. 2010 Jun 1;7(6):e1000275. doi: [10.1371/journal.pmed.1000275](https://doi.org/10.1371/journal.pmed.1000275). PMID: 20532237; PMCID: PMC2879409.
 131. Colijn C, Cohen T, Fraser C, Hanage W, Goldstein E, Givon-Lavi N, Dagan R, Lipsitch M. What is the mechanism for persistent coexistence of drug-susceptible and drug-resistant strains of *Streptococcus pneumoniae*? *J R Soc Interface*. 2010 Jun 6;7(47):905-19. doi: [10.1098/rsif.2009.0400](https://doi.org/10.1098/rsif.2009.0400). Epub 2009 Nov 25. PMID: 19940002; PMCID: PMC2871802.

132. Regev-Yochay G, Hanage WP, Trzcinski K, Rifas-Shiman SL, Lee G, Bessolo A, Huang SS, Pelton SI, McAdam AJ, Finkelstein JA, Lipsitch M, Malley R. Re-emergence of the type 1 pilus among *Streptococcus pneumoniae* isolates in Massachusetts, USA. *Vaccine*. 2010 Jul 5;28(30):4842-6. doi: [10.1016/j.vaccine.2010.04.042](https://doi.org/10.1016/j.vaccine.2010.04.042). Epub 2010 Apr 29. PMID: 20434550; PMCID: PMC2897942.
133. Goldstein E, Cowling BJ, O'Hagan JJ, Danon L, Fang VJ, Hagy A, Miller JC, Reshef D, Robins J, Biedrzycki P, Lipsitch M. Oseltamivir for treatment and prevention of pandemic influenza A/H1N1 virus infection in households, Milwaukee, 2009. *BMC Infect Dis*. 2010 Jul 20;10:211. doi: [10.1186/1471-2334-10-211](https://doi.org/10.1186/1471-2334-10-211). PMID: 20642862; PMCID: PMC2919545.
134. Walensky RP, Paltiel AD, Losina E, Morris BL, Scott CA, Rhode ER, Seage GR, Freedberg KA; CEPAC Investigators. Test and treat DC: forecasting the impact of a comprehensive HIV strategy in Washington DC. *Clin Infect Dis*. 2010 Aug 15;51(4):392-400. doi: [10.1086/655130](https://doi.org/10.1086/655130). Erratum in: *Clin Infect Dis*. 2011 Sep;53(5):502. PMID: 20617921; PMCID: PMC2906630.
135. Lajous M, Danon L, López-Ridaura R, Astley CM, Miller JC, Dowell SF, O'Hagan JJ, Goldstein E, Lipsitch M. Mobile messaging as surveillance tool during pandemic (H1N1) 2009, Mexico. *Emerg Infect Dis*. 2010 Sep;16(9):1488-9. doi: [10.3201/eid1609.100671](https://doi.org/10.3201/eid1609.100671). PMID: 20735942; PMCID: PMC3294993.
136. Rydzak CE, Cotich KL, Sax PE, Hsu HE, Wang B, Losina E, Freedberg KA, Weinstein MC, Goldie SJ; CEPAC Investigators. Assessing the performance of a computer-based policy model of HIV and AIDS. *PLoS One*. 2010 Sep 9;5(9):e12647. doi: [10.1371/journal.pone.0012647](https://doi.org/10.1371/journal.pone.0012647). PMID: 20844741; PMCID: PMC2936574.
137. Uhler LM, Kumarasamy N, Mayer KH, Saxena A, Losina E, Muniyandi M, Stoler AW, Lu Z, Walensky RP, Flanigan TP, Bender MA, Freedberg KA, Swaminathan S; CEPAC International investigators. Cost-effectiveness of HIV testing referral strategies among tuberculosis patients in India. *PLoS One*. 2010 Sep 16;5(9):e12747. doi: [10.1371/journal.pone.0012747](https://doi.org/10.1371/journal.pone.0012747). PMID: 20862279; PMCID: PMC2940842.
138. Weinberger DM, Harboe ZB, Sanders EA, Ndiritu M, Klugman KP, Rückinger S, Dagan R, Adegbola R, Cutts F, Johnson HL, O'Brien KL, Scott JA, Lipsitch M. Association of serotype with risk of death due to pneumococcal pneumonia: a meta-analysis. *Clin Infect Dis*. 2010 Sep 15;51(6):692-9. doi: [10.1086/655828](https://doi.org/10.1086/655828). PMID: 20715907; PMCID: PMC2927802.
139. Huang SS, Avery TR, Song Y, Elkins KR, Nguyen CC, Nutter SK, Nafday AA, Condon CJ, Chang MT, Chrest D, Boos J, Bobashev G, Wheaton W, Frank SA, Platt R, Lipsitch M, Bush RM, Eubank S, Burke DS, Lee BY. Quantifying interhospital patient sharing as a mechanism for infectious disease spread. *Infect Control Hosp Epidemiol*. 2010 Nov;31(11):1160-9. doi: [10.1086/656747](https://doi.org/10.1086/656747). PMID: 20874503; PMCID: PMC3064463.
140. Goldhaber-Fiebert JD, Lipsitch M, Mahal A, Zaslavsky AM, Salomon JA. Quantifying child mortality reductions related to measles vaccination. *PLoS One*. 2010 Nov 4;5(11):e13842. doi: [10.1371/journal.pone.0013842](https://doi.org/10.1371/journal.pone.0013842). PMID: 21079809; PMCID: PMC2973966.
141. Shaman J, Goldstein E, Lipsitch M. Absolute humidity and pandemic versus epidemic influenza. *Am J Epidemiol*. 2011 Jan 15;173(2):127-35. doi: [10.1093/aje/kwq347](https://doi.org/10.1093/aje/kwq347). Epub 2010 Nov 16. PMID: 21081646; PMCID: PMC3011950.

142. Yildirim I, Hanage WP, Lipsitch M, Shea KM, Stevenson A, Finkelstein J, Huang SS, Lee GM, Kleinman K, Pelton SI. Serotype specific invasive capacity and persistent reduction in invasive pneumococcal disease. *Vaccine*. 2010 Dec 16;29(2):283-8. doi: [10.1016/j.vaccine.2010.10.032](https://doi.org/10.1016/j.vaccine.2010.10.032). Epub 2010 Oct 26. PMID: 21029807; PMCID: PMC3139683.
143. Weinberger DM, Harboe ZB, Flasche S, Scott JA, Lipsitch M. Prediction of serotypes causing invasive pneumococcal disease in unvaccinated and vaccinated populations. *Epidemiology*. 2011 Mar;22(2):199-207. doi: [10.1097/EDE.0b013e3182087634](https://doi.org/10.1097/EDE.0b013e3182087634). PMID: 21646962; PMCID: PMC3142570.
144. Hanage WP, Bishop CJ, Huang SS, Stevenson AE, Pelton SI, Lipsitch M, Finkelstein JA. Carried pneumococci in Massachusetts children: the contribution of clonal expansion and serotype switching. *Pediatr Infect Dis J*. 2011 Apr;30(4):302-8. doi: [10.1097/INF.0b013e318201a154](https://doi.org/10.1097/INF.0b013e318201a154). PMID: 21085049; PMCID: PMC3175614.
145. Ciaranello AL, Lockman S, Freedberg KA, Hughes M, Chu J, Currier J, Wood R, Holmes CB, Pillay S, Conradie F, McIntyre J, Losina E, Walensky RP; CEPAC-International and OCTANE Investigators. First-line antiretroviral therapy after single-dose nevirapine exposure in South Africa: a cost-effectiveness analysis of the OCTANE trial. *AIDS*. 2011 Feb 20;25(4):479-92. doi: [10.1097/QAD.0b013e3283428cbe](https://doi.org/10.1097/QAD.0b013e3283428cbe). PMID: 21293199; PMCID: PMC3068908.
146. Hsu HE, Ryzak CE, Cotich KL, Wang B, Sax PE, Losina E, Freedberg KA, Goldie SJ, Lu Z, Walensky RP; CEPAC Investigators. Quantifying the risks and benefits of efavirenz use in HIV-infected women of childbearing age in the USA. *HIV Med*. 2011 Feb;12(2):97-108. doi: [10.1111/j.1468-1293.2010.00856.x](https://doi.org/10.1111/j.1468-1293.2010.00856.x). PMID: 20561082; PMCID: PMC3010302.
147. Ford CB, Lin PL, Chase MR, Shah RR, Iartchouk O, Galagan J, Mohaideen N, Iøerger TR, Sacchettini JC, Lipsitch M, Flynn JL, Fortune SM. Use of whole genome sequencing to estimate the mutation rate of *Mycobacterium tuberculosis* during latent infection. *Nat Genet*. 2011 May;43(5):482-6. doi: [10.1038/ng.811](https://doi.org/10.1038/ng.811). Epub 2011 Apr 24. PMID: 21516081; PMCID: PMC3101871.
148. Lipsitch M, Finelli L, Heffernan RT, Leung GM, Redd SC; 2009 H1n1 Surveillance Group. Improving the evidence base for decision making during a pandemic: the example of 2009 influenza A/H1N1. *Biosecur Bioterror*. 2011 Jun;9(2):89-115. doi: [10.1089/bsp.2011.0007](https://doi.org/10.1089/bsp.2011.0007). PMID: 21612363; PMCID: PMC3102310.
149. Smith J, Lipsitch M, Almond JW. Vaccine production, distribution, access, and uptake. *Lancet*. 2011 Jul 30;378(9789):428-38. doi: [10.1016/S0140-6736\(11\)60478-9](https://doi.org/10.1016/S0140-6736(11)60478-9). Epub 2011 Jun 12. PMID: 21664680; PMCID: PMC3164579.
150. Goldstein E, Cobey S, Takahashi S, Miller JC, Lipsitch M. Predicting the epidemic sizes of influenza A/H1N1, A/H3N2, and B: a statistical method. *PLoS Med*. 2011 Jul;8(7):e1001051. doi: [10.1371/journal.pmed.1001051](https://doi.org/10.1371/journal.pmed.1001051). Epub 2011 Jul 5. PMID: 21750666; PMCID: PMC3130020.
151. Mostofsky E, Lipsitch M, Regev-Yochay G. Is methicillin-resistant *Staphylococcus aureus* replacing methicillin-susceptible *S. aureus*? *J Antimicrob Chemother*. 2011 Oct;66(10):2199-214. doi: [10.1093/jac/dkr278](https://doi.org/10.1093/jac/dkr278). Epub 2011 Jul 7. PMID: 21737459; PMCID: PMC3172038.
152. Shaman J, Jeon CY, Giovannucci E, Lipsitch M. Shortcomings of vitamin D-based model simulations of seasonal influenza. *PLoS One*. 2011;6(6):e20743. doi: [10.1371/journal.pone.0020743](https://doi.org/10.1371/journal.pone.0020743). Epub 2011 Jun 3. PMID: 21677774; PMCID: PMC3108988.

153. Hernán MA, Lipsitch M. Oseltamivir and risk of lower respiratory tract complications in patients with flu symptoms: a meta-analysis of eleven randomized clinical trials. *Clin Infect Dis*. 2011 Aug 1;53(3):277-9. doi: [10.1093/cid/cir400](https://doi.org/10.1093/cid/cir400). Epub 2011 Jun 15. PMID: 21677258; PMCID: PMC3137795.
154. Goldstein E, Cowling BJ, Aiello AE, Takahashi S, King G, Lu Y, Lipsitch M. Estimating incidence curves of several infections using symptom surveillance data. *PLoS One*. 2011;6(8):e23380. doi: [10.1371/journal.pone.0023380](https://doi.org/10.1371/journal.pone.0023380). Epub 2011 Aug 24. PMID: 21887246; PMCID: PMC3160845.
155. Presanis AM, Pebody RG, Paterson BJ, Tom BD, Birrell PJ, Charlett A, Lipsitch M, De Angelis D. Changes in severity of 2009 pandemic A/H1N1 influenza in England: a Bayesian evidence synthesis. *BMJ*. 2011 Sep 8;343:d5408. doi: [10.1136/bmj.d5408](https://doi.org/10.1136/bmj.d5408). PMID: 21903689; PMCID: PMC3168935.
156. Hanage WP, Bishop CJ, Lee GM, Lipsitch M, Stevenson A, Rifas-Shiman SL, Pelton SI, Huang SS, Finkelstein JA. Clonal replacement among 19A *Streptococcus pneumoniae* in Massachusetts, prior to 13 valent conjugate vaccination. *Vaccine*. 2011 Nov 8;29(48):8877-81. doi: [10.1016/j.vaccine.2011.09.075](https://doi.org/10.1016/j.vaccine.2011.09.075). Epub 2011 Sep 29. PMID: 21964059; PMCID: PMC3221484.
157. Weinberger DM, Malley R, Lipsitch M. Serotype replacement in disease after pneumococcal vaccination. *Lancet*. 2011 Dec 3;378(9807):1962-73. doi: [10.1016/S0140-6736\(10\)62225-8](https://doi.org/10.1016/S0140-6736(10)62225-8). Epub 2011 Apr 12. PMID: 21492929; PMCID: PMC3256741.
158. O'Hagan JJ, Hernán MA, Walensky RP, Lipsitch M. Apparent declining efficacy in randomized trials: examples of the Thai RV144 HIV vaccine and South African CAPRISA 004 microbicide trials. *AIDS*. 2012 Jan 14;26(2):123-6. doi: [10.1097/QAD.0b013e32834e1ce7](https://doi.org/10.1097/QAD.0b013e32834e1ce7). PMID: 22045345; PMCID: PMC3319457.
159. Shaman J, Lipsitch M. The El Niño-Southern Oscillation (ENSO)-pandemic influenza connection: coincident or causal? *Proc Natl Acad Sci U S A*. 2013 Feb 26;110 Suppl 1(Suppl 1):3689-91. doi: [10.1073/pnas.1107485109](https://doi.org/10.1073/pnas.1107485109). Epub 2012 Jan 17. PMID: 22308322; PMCID: PMC3586607.
160. Grad YH, Lipsitch M, Aiello AE. Secular trends in *Helicobacter pylori* seroprevalence in adults in the United States: evidence for sustained race/ethnic disparities. *Am J Epidemiol*. 2012 Jan 1;175(1):54-9. doi: [10.1093/aje/kwr288](https://doi.org/10.1093/aje/kwr288). Epub 2011 Nov 15. PMID: 22085628; PMCID: PMC3244610.
161. Scott JR, Millar EV, Lipsitch M, Moulton LH, Weatherholtz R, Perilla MJ, Jackson DM, Beall B, Craig MJ, Reid R, Santosham M, O'Brien KL. Impact of more than a decade of pneumococcal conjugate vaccine use on carriage and invasive potential in Native American communities. *J Infect Dis*. 2012 Jan 15;205(2):280-8. doi: [10.1093/infdis/jir730](https://doi.org/10.1093/infdis/jir730). Epub 2011 Nov 29. PMID: 22128315; PMCID: PMC3244367.
162. Abdullahi O, Karani A, Tigoi CC, Mugo D, Kungu S, Wanjiru E, Jomo J, Musyimi R, Lipsitch M, Scott JA. The prevalence and risk factors for pneumococcal colonization of the nasopharynx among children in Kilifi District, Kenya. *PLoS One*. 2012;7(2):e30787. doi: [10.1371/journal.pone.0030787](https://doi.org/10.1371/journal.pone.0030787). Epub 2012 Feb 20. PMID: 22363489; PMCID: PMC3282706.
163. Cobey S, Lipsitch M. Niche and neutral effects of acquired immunity permit coexistence of pneumococcal serotypes. *Science*. 2012 Mar 16;335(6074):1376-80. doi: [10.1126/science.1215947](https://doi.org/10.1126/science.1215947). Epub 2012 Mar 1. PMID: 22383809; PMCID: PMC3341938.
164. Scott JR, Hanage WP, Lipsitch M, Millar EV, Moulton LH, Hinds J, Reid R, Santosham M, O'Brien KL. Pneumococcal sequence type replacement among

- American Indian children: a comparison of pre- and routine-PCV7 eras. *Vaccine*. 2012 Mar 16;30(13):2376-81. doi: [10.1016/j.vaccine.2011.11.004](https://doi.org/10.1016/j.vaccine.2011.11.004). Epub 2011 Nov 15. PMID: 22094283.
165. Lipsitch M, Abdullahi O, D'Amour A, Xie W, Weinberger DM, Tchetgen Tchetgen E, Scott JA. Estimating rates of carriage acquisition and clearance and competitive ability for pneumococcal serotypes in Kenya with a Markov transition model. *Epidemiology*. 2012 Jul;23(4):510-9. doi: [10.1097/EDE.0b013e31824f2f32](https://doi.org/10.1097/EDE.0b013e31824f2f32). Erratum in: *Epidemiology*. 2013 Jan;24(1):177. PMID: 22441543; PMCID: PMC3670084.
 166. Wroe PC, Lee GM, Finkelstein JA, Pelton SI, Hanage WP, Lipsitch M, Stevenson AE, Rifas-Shiman SL, Kleinman K, Dutta-Linn MM, Hinrichsen VL, Lakoma M, Huang SS. Pneumococcal carriage and antibiotic resistance in young children before 13-valent conjugate vaccine. *Pediatr Infect Dis J*. 2012 Mar;31(3):249-54. doi: [10.1097/INF.0b013e31824214ac](https://doi.org/10.1097/INF.0b013e31824214ac). PMID: 22173142; PMCID: PMC3288953.
 167. Grad YH, Lipsitch M, Feldgarden M, Arachchi HM, Cerqueira GC, Fitzgerald M, Godfrey P, Haas BJ, Murphy CI, Russ C, Sykes S, Walker BJ, Wortman JR, Young S, Zeng Q, Abouelleil A, Bochicchio J, Chauvin S, Desmet T, Gujja S, McCowan C, Montmayeur A, Steelman S, Frimodt-Møller J, Petersen AM, Struve C, Krogfelt KA, Bingen E, Weill FX, Lander ES, Nusbaum C, Birren BW, Hung DT, Hanage WP. Genomic epidemiology of the *Escherichia coli* O104:H4 outbreaks in Europe, 2011. *Proc Natl Acad Sci U S A*. 2012 Feb 21;109(8):3065-70. doi: [10.1073/pnas.1121491109](https://doi.org/10.1073/pnas.1121491109). Epub 2012 Feb 6. Erratum in: *Proc Natl Acad Sci U S A*. 2012 Apr 3;109(14):5547. PMID: 22315421; PMCID: PMC3286951.
 168. Pitzer VE, Burgner D, Viboud C, Simonsen L, Andreasen V, Steiner CA, Lipsitch M. Modelling seasonal variations in the age and incidence of Kawasaki disease to explore possible infectious aetiologies. *Proc Biol Sci*. 2012 Jul 22;279(1739):2736-43. doi: [10.1098/rspb.2011.2464](https://doi.org/10.1098/rspb.2011.2464). Epub 2012 Mar 7. PMID: 22398170; PMCID: PMC3367771.
 169. Ke W, Huang SS, Hudson LO, Elkins KR, Nguyen CC, Spratt BG, Murphy CR, Avery TR, Lipsitch M. Patient sharing and population genetic structure of methicillin-resistant *Staphylococcus aureus*. *Proc Natl Acad Sci U S A*. 2012 Apr 24;109(17):6763-8. doi: [10.1073/pnas.1113578109](https://doi.org/10.1073/pnas.1113578109). Epub 2012 Mar 19. PMID: 22431601; PMCID: PMC3340079.
 170. Grad YH, Miller JC, Lipsitch M. Cholera modeling: challenges to quantitative analysis and predicting the impact of interventions. *Epidemiology*. 2012 Jul;23(4):523-30. doi: [10.1097/EDE.0b013e3182572581](https://doi.org/10.1097/EDE.0b013e3182572581). PMID: 22659546; PMCID: PMC3380087.
 171. Van Kerkhove MD, Riley S, Lipsitch M, Guan Y, Monto AS, Webster RG, Zambon M, Nicoll A, Peiris JS, Ferguson NM. Comment on "Seroevidence for H5N1 influenza infections in humans: meta-analysis". *Science*. 2012 Jun 22;336(6088):1506; author reply 1506. doi: [10.1126/science.1221434](https://doi.org/10.1126/science.1221434). PMID: 22723396.
 172. Lipsitch M, Plotkin JB, Simonsen L, Bloom B. Evolution, safety, and highly pathogenic influenza viruses. *Science*. 2012 Jun 22;336(6088):1529-31. doi: [10.1126/science.1223204](https://doi.org/10.1126/science.1223204). PMID: 22723411; PMCID: PMC3467308.
 173. Goldstein E, Wallinga J, Lipsitch M. Vaccine allocation in a declining epidemic. *J R Soc Interface*. 2012 Nov 7;9(76):2798-803. doi: [10.1098/rsif.2012.0404](https://doi.org/10.1098/rsif.2012.0404). Epub 2012 Jul 6. PMID: 22772378; PMCID: PMC3479926.
 174. Abdullahi O, Karani A, Tigoi CC, Mugo D, Kungu S, Wanjiru E, Jomo J, Musyimi R, Lipsitch M, Scott JA. Rates of acquisition and clearance of pneumococcal

- serotypes in the nasopharynges of children in Kilifi District, Kenya. *J Infect Dis.* 2012 Oct 1;206(7):1020-9. doi: [10.1093/infdis/jis447](https://doi.org/10.1093/infdis/jis447). Epub 2012 Jul 24. PMID: 22829650; PMCID: PMC3433858.
175. Jacobs JH, Archer BN, Baker MG, Cowling BJ, Heffernan RT, Mercer G, Uez O, Hanshaworakul W, Viboud C, Schwartz J, Tchetgen Tchetgen E, Lipsitch M. Searching for sharp drops in the incidence of pandemic A/H1N1 influenza by single year of age. *PLoS One.* 2012;7(8):e42328. doi: [10.1371/journal.pone.0042328](https://doi.org/10.1371/journal.pone.0042328). Epub 2012 Aug 2. PMID: 22876316; PMCID: PMC3410923.
 176. Goldstein E, Kirkcaldy RD, Reshef D, Berman S, Weinstock H, Sabeti P, Del Rio C, Hall G, Hook EW, Lipsitch M. Factors related to increasing prevalence of resistance to ciprofloxacin and other antimicrobial drugs in *Neisseria gonorrhoeae*, United States. *Emerg Infect Dis.* 2012 Aug;18(8):1290-7. doi: [10.3201/eid1808.111202](https://doi.org/10.3201/eid1808.111202). PMID: 22840274; PMCID: PMC3414012.
 177. Lipsitch M, Bloom BR. Rethinking biosafety in research on potential pandemic pathogens. *mBio.* 2012 Oct 9;3(5):e00360-12. doi: [10.1128/mBio.00360-12](https://doi.org/10.1128/mBio.00360-12). PMID: 23047752; PMCID: PMC3484391.
 178. Li Y, Gierahn T, Thompson CM, Trzciński K, Ford CB, Croucher N, Gouveia P, Flechtner JB, Malley R, Lipsitch M. Distinct effects on diversifying selection by two mechanisms of immunity against *Streptococcus pneumoniae*. *PLoS Pathog.* 2012;8(11):e1002989. doi: [10.1371/journal.ppat.1002989](https://doi.org/10.1371/journal.ppat.1002989). Epub 2012 Nov 8. PMID: 23144610; PMCID: PMC3493470. **Highlighted in the PLoS Pathogens Tenth Anniversary Collection of the Editors' Choice of 42 articles from the first 10 years.**
 179. Goldstein E, Viboud C, Charu V, Lipsitch M. Improving the estimation of influenza-related mortality over a seasonal baseline. *Epidemiology.* 2012 Nov;23(6):829-38. doi: [10.1097/EDE.0b013e31826c2dda](https://doi.org/10.1097/EDE.0b013e31826c2dda). PMID: 22992574; PMCID: PMC3516362.
 - A commentary on this paper was published alongside it, to which we published a rejoinder: Goldstein E, Viboud C, Charu V, Lipsitch M. The authors respond. *Epidemiology.* 2012 Nov;23(6):829-38.
 180. Hyams C, Trzcinski K, Camberlein E, Weinberger DM, Chimalapati S, Noursadeghi M, Lipsitch M, Brown JS. *Streptococcus pneumoniae* capsular serotype invasiveness correlates with the degree of factor H binding and opsonization with C3b/iC3b. *Infect Immun.* 2013 Jan;81(1):354-63. doi: [10.1128/IAI.00862-12](https://doi.org/10.1128/IAI.00862-12). Epub 2012 Nov 12. PMID: 23147038; PMCID: PMC3536142.
 181. Palmer ME, Lipsitch M, Moxon ER, Bayliss CD. Broad conditions favor the evolution of phase-variable loci. *mBio.* 2013 Jan 8;4(1):e00430-12. doi: [10.1128/mBio.00430-12](https://doi.org/10.1128/mBio.00430-12). PMID: 23300246; PMCID: PMC3546556.
 182. Cobey S, Lipsitch M. Pathogen diversity and hidden regimes of apparent competition. *Am Nat.* 2013 Jan;181(1):12-24. doi: [10.1086/668598](https://doi.org/10.1086/668598). Epub 2012 Nov 27. PMID: 23234842; PMCID: PMC3716377.
 183. Grad YH, Godfrey P, Cerquiera GC, Mariani-Kurkdjian P, Gouali M, Bingen E, Shea TP, Haas BJ, Griggs A, Young S, Zeng Q, Lipsitch M, Waldor MK, Weill FX, Wortman JR, Hanage WP. Comparative genomics of recent Shiga toxin-producing *Escherichia coli* O104:H4: short-term evolution of an emerging pathogen. *mBio.* 2013 Jan 22;4(1):e00452-12. doi: [10.1128/mBio.00452-12](https://doi.org/10.1128/mBio.00452-12). PMID: 23341549; PMCID: PMC3551546.
 184. Koep TH, Enders FT, Pierret C, Ekker SC, Krageschmidt D, Neff KL, Lipsitch M, Shaman J, Huskins WC. Predictors of indoor absolute humidity and estimated

- effects on influenza virus survival in grade schools. *BMC Infect Dis*. 2013 Feb 5;13:71. doi: [10.1186/1471-2334-13-71](https://doi.org/10.1186/1471-2334-13-71). PMID: 23383620; PMCID: PMC3568414.
185. O'Hagan JJ, Hernán MA, Walensky RP, Lipsitch M. Apparent declining efficacy in randomized trials: examples of the Thai RV144 HIV vaccine and South African CAPRISA 004 microbicide trials. *AIDS*. 2012 Jan 14;26(2):123-6. doi: [10.1097/QAD.0b013e32834e1ce7](https://doi.org/10.1097/QAD.0b013e32834e1ce7). PMID: 22045345; PMCID: PMC3319457.
 186. Grad YH, Lipsitch M, Griggs AD, Haas BJ, Shea TP, McCowan C, Montmayeur A, FitzGerald M, Wortman JR, Krogfelt KA, Bingen E, Weill FX, Tietze E, Flieger A, Lander ES, Nusbaum C, Birren BW, Hung DT, Hanage WP. Reply to Guy et al.: Support for a bottleneck in the 2011 *Escherichia coli* O104:H4 outbreak in Germany. *Proc Natl Acad Sci U S A*. 2012 Dec 26;109(52):E3629-30. doi: [10.1073/pnas.1209419110](https://doi.org/10.1073/pnas.1209419110). PMID: 23479789; PMCID: PMC3535640.
 187. Croucher NJ, Finkelstein JA, Pelton SI, Mitchell PK, Lee GM, Parkhill J, Bentley SD, Hanage WP, Lipsitch M. Population genomics of post-vaccine changes in pneumococcal epidemiology. *Nat Genet*. 2013 Jun;45(6):656-63. doi: [10.1038/ng.2625](https://doi.org/10.1038/ng.2625). Epub 2013 May 5. PMID: 23644493; PMCID: PMC3725542.
 188. Joice R, Lipsitch M. Targeting imperfect vaccines against drug-resistance determinants: a strategy for countering the rise of drug resistance. *PLoS One*. 2013 Jul 25;8(7):e68940. doi: [10.1371/journal.pone.0068940](https://doi.org/10.1371/journal.pone.0068940). PMID: 23935910; PMCID: PMC3723804.
 189. Ford CB, Shah RR, Maeda MK, Gagneux S, Murray MB, Cohen T, Johnston JC, Gardy J, Lipsitch M, Fortune SM. Mycobacterium tuberculosis mutation rate estimates from different lineages predict substantial differences in the emergence of drug-resistant tuberculosis. *Nat Genet*. 2013 Jul;45(7):784-90. doi: [10.1038/ng.2656](https://doi.org/10.1038/ng.2656). Epub 2013 Jun 9. PMID: 23749189; PMCID: PMC3777616.
 190. Link-Gelles R, Thomas A, Lynfield R, Petit S, Schaffner W, Harrison L, Farley MM, Aragon D, Nicols M, Kirley PD, Zansky S, Jorgensen J, Juni BA, Jackson D, Moore MR, Lipsitch M. Geographic and temporal trends in antimicrobial nonsusceptibility in *Streptococcus pneumoniae* in the post-vaccine era in the United States. *J Infect Dis*. 2013 Oct 15;208(8):1266-73. doi: [10.1093/infdis/jit315](https://doi.org/10.1093/infdis/jit315). Epub 2013 Jul 12. PMID: 23852588; PMCID: PMC3778966.
 191. Weinberger DM, Bruden DT, Grant LR, Lipsitch M, O'Brien KL, Pelton SI, Sanders EA, Feikin DR. Using pneumococcal carriage data to monitor postvaccination changes in invasive disease. *Am J Epidemiol*. 2013 Nov 1;178(9):1488-95. doi: [10.1093/aje/kwt156](https://doi.org/10.1093/aje/kwt156). Epub 2013 Sep 7. PMID: 24013204; PMCID: PMC3813314.
 192. Li Y, Weinberger DM, Thompson CM, Trzciński K, Lipsitch M. Surface charge of *Streptococcus pneumoniae* predicts serotype distribution. *Infect Immun*. 2013 Dec;81(12):4519-24. doi: [10.1128/IAI.00724-13](https://doi.org/10.1128/IAI.00724-13). Epub 2013 Sep 30. PMID: 24082068; PMCID: PMC3837974.
 193. Li Y, Thompson CM, Trzciński K, Lipsitch M. Within-host selection is limited by an effective population of *Streptococcus pneumoniae* during nasopharyngeal colonization. *Infect Immun*. 2013 Dec;81(12):4534-43. doi: [10.1128/IAI.00527-13](https://doi.org/10.1128/IAI.00527-13). Epub 2013 Sep 30. PMID: 24082074; PMCID: PMC3837969.
 194. Feikin DR, Kagucia EW, Loo JD, Link-Gelles R, Puhon MA, Cherian T, Levine OS, Whitney CG, O'Brien KL, Moore MR; Serotype Replacement Study Group. Serotype-specific changes in invasive pneumococcal disease after pneumococcal conjugate vaccine introduction: a pooled analysis of multiple surveillance sites. *PLoS Med*. 2013;10(9):e1001517. doi:

- [10.1371/journal.pmed.1001517](https://doi.org/10.1371/journal.pmed.1001517). Epub 2013 Sep 24. PMID: 24086113; PMCID: PMC3782411.
195. Quandelacy TM, Viboud C, Charu V, Lipsitch M, Goldstein E. Age- and sex-related risk factors for influenza-associated mortality in the United States between 1997-2007. *Am J Epidemiol*. 2014 Jan 15;179(2):156-67. doi: [10.1093/aje/kwt235](https://doi.org/10.1093/aje/kwt235). Epub 2013 Nov 4. PMID: 24190951; PMCID: PMC3873104.
 196. O'Hagan JJ, Lipsitch M, Hernán MA. Estimating the per-exposure effect of infectious disease interventions. *Epidemiology*. 2014 Jan;25(1):134-8. doi: [10.1097/EDE.0000000000000003](https://doi.org/10.1097/EDE.0000000000000003). PMID: 24240656; PMCID: PMC3898464.
 197. Shaman J, Karspeck A, Yang W, Tamerius J, Lipsitch M. Real-time influenza forecasts during the 2012-2013 season. *Nat Commun*. 2013;4:2837. doi: [10.1038/ncomms3837](https://doi.org/10.1038/ncomms3837). PMID: 24302074; PMCID: PMC3873365.
 198. Huang KE, Lipsitch M, Shaman J, Goldstein E. The US 2009 A(H1N1) influenza epidemic: quantifying the impact of school openings on the reproductive number. *Epidemiology*. 2014 Mar;25(2):203-6. doi: [10.1097/EDE.0000000000000055](https://doi.org/10.1097/EDE.0000000000000055). PMID: 24434751; PMCID: PMC3960948.
 199. Grad YH, Kirkcaldy RD, Trees D, Dordel J, Harris SR, Goldstein E, Weinstock H, Parkhill J, Hanage WP, Bentley S, Lipsitch M. Genomic epidemiology of *Neisseria gonorrhoeae* with reduced susceptibility to cefixime in the USA: a retrospective observational study. *Lancet Infect Dis*. 2014 Mar;14(3):220-6. doi: [10.1016/S1473-3099\(13\)70693-5](https://doi.org/10.1016/S1473-3099(13)70693-5). Epub 2014 Jan 22. PMID: 24462211; PMCID: PMC4030102.
 200. Patterson-Lomba O, Van Noort S, Cowling BJ, Wallinga J, Gomes MG, Lipsitch M, Goldstein E. Utilizing syndromic surveillance data for estimating levels of influenza circulation. *Am J Epidemiol*. 2014 Jun 1;179(11):1394-401. doi: [10.1093/aje/kwu061](https://doi.org/10.1093/aje/kwu061). Epub 2014 Apr 18. PMID: 24748609; PMCID: PMC4036214.
 201. Gomes MG, Lipsitch M, Wargo AR, Kurath G, Rebelo C, Medley GF, Coutinho A. A missing dimension in measures of vaccination impacts. *PLoS Pathog*. 2014 Mar 6;10(3):e1003849. doi: [10.1371/journal.ppat.1003849](https://doi.org/10.1371/journal.ppat.1003849). PMID: 24603721; PMCID: PMC3946326.
 202. Worby CJ, Lipsitch M, Hanage WP. Within-host bacterial diversity hinders accurate reconstruction of transmission networks from genomic distance data. *PLoS Comput Biol*. 2014 Mar 27;10(3):e1003549. doi: [10.1371/journal.pcbi.1003549](https://doi.org/10.1371/journal.pcbi.1003549). PMID: 24675511; PMCID: PMC3967931.
 203. Grad YH, Newman R, Zody M, Yang X, Murphy R, Qu J, Malboeuf CM, Levin JZ, Lipsitch M, DeVincenzo J. Within-host whole-genome deep sequencing and diversity analysis of human respiratory syncytial virus infection reveals dynamics of genomic diversity in the absence and presence of immune pressure. *J Virol*. 2014 Jul;88(13):7286-93. doi: [10.1128/JVI.00038-14](https://doi.org/10.1128/JVI.00038-14). Epub 2014 Apr 16. PMID: 24741088; PMCID: PMC4054443.
 204. McCormick AW, Abuelezam NN, Rhode ER, Hou T, Walensky RP, Pei PP, Becker JE, DiLorenzo MA, Losina E, Freedberg KA, Lipsitch M, Seage GR 3rd. Development, calibration and performance of an HIV transmission model incorporating natural history and behavioral patterns: application in South Africa. *PLoS One*. 2014 May 27;9(5):e98272. doi: [10.1371/journal.pone.0098272](https://doi.org/10.1371/journal.pone.0098272). PMID: 24867402; PMCID: PMC4035281.
 205. Lipsitch M, Galvani AP. Ethical alternatives to experiments with novel potential pandemic pathogens. *PLoS Med*. 2014 May 20;11(5):e1001646. doi: [10.1371/journal.pmed.1001646](https://doi.org/10.1371/journal.pmed.1001646). PMID: 24844931; PMCID: PMC4028196.

206. Li Y, Thompson CM, Lipsitch M. A modified Janus cassette (Sweet Janus) to improve allelic replacement efficiency by high-stringency negative selection in *Streptococcus pneumoniae*. *PLoS One*. 2014 Jun 24;9(6):e100510. doi: [10.1371/journal.pone.0100510](https://doi.org/10.1371/journal.pone.0100510). PMID: 24959661; PMCID: PMC4068995.
207. Ascioğlu S, Samore MH, Lipsitch M. A new approach to the analysis of antibiotic resistance data from hospitals. *Microb Drug Resist*. 2014 Dec;20(6):583-90. doi: [10.1089/mdr.2013.0173](https://doi.org/10.1089/mdr.2013.0173). PMID: 25055133.
208. Johnson SR, Grad Y, Ganakammal SR, Burroughs M, Frace M, Lipsitch M, Weil R, Trees D. In Vitro selection of *Neisseria gonorrhoeae* mutants with elevated MIC values and increased resistance to cephalosporins. *Antimicrob Agents Chemother*. 2014 Nov;58(11):6986-9. doi: [10.1128/AAC.03082-14](https://doi.org/10.1128/AAC.03082-14). Epub 2014 Sep 8. PMID: 25199775; PMCID: PMC4249396.
209. Jacobs JH, Viboud C, Tchetgen ET, Schwartz J, Steiner C, Simonsen L, Lipsitch M. The association of meningococcal disease with influenza in the United States, 1989-2009. *PLoS One*. 2014 Sep 29;9(9):e107486. doi: [10.1371/journal.pone.0107486](https://doi.org/10.1371/journal.pone.0107486). PMID: 25265409; PMCID: PMC4180274.
210. Russell CA, Kasson PM, Donis RO, Riley S, Dunbar J, Rambaut A, Asher J, Burke S, Davis CT, Garten RJ, Gnanakaran S, Hay SI, Herfst S, Lewis NS, Lloyd-Smith JO, Macken CA, Maurer-Stroh S, Neuhaus E, Parrish CR, Pepin KM, Shepard SS, Smith DL, Suarez DL, Trock SC, Widdowson MA, George DB, Lipsitch M, Bloom JD. Improving pandemic influenza risk assessment. *Elife*. 2014 Oct 16;3:e03883. doi: [10.7554/eLife.03883](https://doi.org/10.7554/eLife.03883). PMID: 25321142; PMCID: PMC4199076.
211. Worby CJ, Chang HH, Hanage WP, Lipsitch M. The distribution of pairwise genetic distances: a tool for investigating disease transmission. *Genetics*. 2014 Dec;198(4):1395-404. doi: [10.1534/genetics.114.171538](https://doi.org/10.1534/genetics.114.171538). Epub 2014 Oct 13. PMID: 25313129; PMCID: PMC4256759.
212. Grad YH, Lipsitch M. Epidemiologic data and pathogen genome sequences: a powerful synergy for public health. *Genome Biol*. 2014 Nov 18;15(11):538. doi: [10.1186/s13059-014-0538-4](https://doi.org/10.1186/s13059-014-0538-4). PMID: 25418119; PMCID: PMC4282151.
213. Abel S, Abel zur Wiesch P, Chang HH, Davis BM, Lipsitch M, Waldor MK. Sequence tag-based analysis of microbial population dynamics. *Nat Methods*. 2015 Mar;12(3):223-6, 3 p following 226. doi: [10.1038/nmeth.3253](https://doi.org/10.1038/nmeth.3253). Epub 2015 Jan 19. PMID: 25599549; PMCID: PMC4344388.
214. Lipsitch M, Inglesby TV. Reply to "Studies on influenza virus transmission between ferrets: the public health risks revisited". *mBio*. 2015 Jan 23;6(1):e00041-15. doi: [10.1128/mBio.00041-15](https://doi.org/10.1128/mBio.00041-15). PMID: 25616376; PMCID: PMC4323416.
215. Lipsitch M, Inglesby TV. Moratorium on research intended to create novel potential pandemic pathogens. *mBio*. 2014 Dec 12;5(6):e02366-14. doi: [10.1128/mBio.02366-14](https://doi.org/10.1128/mBio.02366-14). Erratum in: *MBio*. 2015;6(1). pii: e02534-14. doi: [10.1128/mBio.02534-14](https://doi.org/10.1128/mBio.02534-14). PMID: 25505122; PMCID: PMC4271556.
216. Chang Q, Stevenson AE, Croucher NJ, Lee GM, Pelton SI, Lipsitch M, Finkelstein JA, Hanage WP. Stability of the pneumococcal population structure in Massachusetts as PCV13 was introduced. *BMC Infect Dis*. 2015 Feb 18;15:68. doi: [10.1186/s12879-015-0797-z](https://doi.org/10.1186/s12879-015-0797-z). PMID: 25887323; PMCID: PMC4336693.
217. Chang HH, Cohen T, Grad YH, Hanage WP, O'Brien TF, Lipsitch M. Origin and proliferation of multiple-drug resistance in bacterial pathogens. *Microbiol Mol Biol Rev*. 2015 Mar;79(1):101-16. doi: [10.1128/MMBR.00039-14](https://doi.org/10.1128/MMBR.00039-14). PMID: 25652543; PMCID: PMC4402963.

218. Croucher NJ, Kagedan L, Thompson CM, Parkhill J, Bentley SD, Finkelstein JA, Lipsitch M, Hanage WP. Selective and genetic constraints on pneumococcal serotype switching. *PLoS Genet*. 2015 Mar 31;11(3):e1005095. doi: [10.1371/journal.pgen.1005095](https://doi.org/10.1371/journal.pgen.1005095). PMID: 25826208; PMCID: PMC4380333.
219. Yang W, Lipsitch M, Shaman J. Inference of seasonal and pandemic influenza transmission dynamics. *Proc Natl Acad Sci U S A*. 2015 Mar 3;112(9):2723-8. doi: [10.1073/pnas.1415012112](https://doi.org/10.1073/pnas.1415012112). Epub 2015 Feb 17. PMID: 25730851; PMCID: PMC4352784.
220. Lipsitch M, Eyal N, Halloran ME, Hernán MA, Longini IM, Perencevich EN, Grais RF. Ebola and beyond. *Science*. 2015 Apr 3;348(6230):46-8. doi: [10.1126/science.aaa3178](https://doi.org/10.1126/science.aaa3178). PMID: 25838371; PMCID: PMC4408019.
221. Li Y, Croucher NJ, Thompson CM, Trzciński K, Hanage WP, Lipsitch M. Identification of pneumococcal colonization determinants in the stringent response pathway facilitated by genomic diversity. *BMC Genomics*. 2015 May 9;16(1):369. doi: [10.1186/s12864-015-1573-6](https://doi.org/10.1186/s12864-015-1573-6). PMID: 25956132; PMCID: PMC4424882.
222. Chang Q, Wang W, Regev-Yochay G, Lipsitch M, Hanage WP. Antibiotics in agriculture and the risk to human health: how worried should we be? *Evol Appl*. 2015 Mar;8(3):240-7. doi: [10.1111/eva.12185](https://doi.org/10.1111/eva.12185). Epub 2014 Aug 2. PMID: 25861382; PMCID: PMC4380918.
223. Kunkel A, Colijn C, Lipsitch M, Cohen T. How could preventive therapy affect the prevalence of drug resistance? Causes and consequences. *Philos Trans R Soc Lond B Biol Sci*. 2015 Jun 5;370(1670):20140306. doi: [10.1098/rstb.2014.0306](https://doi.org/10.1098/rstb.2014.0306). PMID: 25918446; PMCID: PMC4424438.
224. Mitchell PK, Lipsitch M, Hanage WP. Carriage burden, multiple colonization and antibiotic pressure promote emergence of resistant vaccine escape pneumococci. *Philos Trans R Soc Lond B Biol Sci*. 2015 Jun 5;370(1670):20140342. doi: [10.1098/rstb.2014.0342](https://doi.org/10.1098/rstb.2014.0342). PMID: 25918447; PMCID: PMC4424439.
225. Goldstein E, Greene SK, Olson DR, Hanage WP, Lipsitch M. Estimating the hospitalization burden associated with influenza and respiratory syncytial virus in New York City, 2003-2011. *Influenza Other Respir Viruses*. 2015 Sep;9(5):225-33. doi: [10.1111/irv.12325](https://doi.org/10.1111/irv.12325). PMID: 25980600; PMCID: PMC4548992.
226. Worby CJ, Chaves SS, Wallinga J, Lipsitch M, Finelli L, Goldstein E. On the relative role of different age groups in influenza epidemics. *Epidemics*. 2015 Dec;13:10-16. doi: [10.1016/j.epidem.2015.04.003](https://doi.org/10.1016/j.epidem.2015.04.003). PMID: 26097505; PMCID: PMC4469206.
227. Lipsitch M, Donnelly CA, Fraser C, Blake IM, Cori A, Dorigatti I, Ferguson NM, Garske T, Mills HL, Riley S, Van Kerkhove MD, Hernán MA. Potential Biases in Estimating Absolute and Relative Case-Fatality Risks during Outbreaks. *PLoS Negl Trop Dis*. 2015 Jul 16;9(7):e0003846. doi: [10.1371/journal.pntd.0003846](https://doi.org/10.1371/journal.pntd.0003846). PMID: 26181387; PMCID: PMC4504518. **Awarded 2nd Prize for Publication with Impact on the Field in 2015-6 by International Society for Disease Surveillance.**
228. Evans NG, Lipsitch M, Levinson M. The ethics of biosafety considerations in gain-of-function research resulting in the creation of potential pandemic pathogens. *J Med Ethics*. 2015 Nov;41(11):901-8. doi: [10.1136/medethics-2014-102619](https://doi.org/10.1136/medethics-2014-102619). Epub 2015 Aug 28. PMID: 26320212; PMCID: PMC4623968.
229. Worby CJ, Kenyon C, Lynfield R, Lipsitch M, Goldstein E. Examining the role of different age groups, and of vaccination during the 2012 Minnesota pertussis

- outbreak. *Sci Rep.* 2015 Aug 17;5:13182. doi: [10.1038/srep13182](https://doi.org/10.1038/srep13182). PMID: 26278132; PMCID: PMC4538373.
230. Trzciński K, Li Y, Weinberger DM, Thompson CM, Cordy D, Bessolo A, Malley R, Lipsitch M. Effect of Serotype on Pneumococcal Competition in a Mouse Colonization Model. *mBio.* 2015 Sep 15;6(5):e00902-15. doi: [10.1128/mBio.00902-15](https://doi.org/10.1128/mBio.00902-15). PMID: 26374118; PMCID: PMC4600102.
231. Althouse BM, Scarpino SV, Meyers LA, Ayers JW, Bargsten M, Baumbach J, Brownstein JS, Castro L, Clapham H, Cummings DA, Del Valle S, Eubank S, Fairchild G, Finelli L, Generous N, George D, Harper DR, Hébert-Dufresne L, Johansson MA, Konty K, Lipsitch M, Milinovich G, Miller JD, Nsoesie EO, Olson DR, Paul M, Polgreen PM, Priedhorsky R, Read JM, Rodríguez-Barrquer I, Smith DJ, Stefansen C, Swerdlow DL, Thompson D, Vespignani A, Wesolowski A. Enhancing disease surveillance with novel data streams: challenges and opportunities. *EPJ Data Sci.* 2015;4(1):17. doi: [10.1140/epjds/s13688-015-0054-0](https://doi.org/10.1140/epjds/s13688-015-0054-0). Epub 2015 Oct 16. PMID: 27990325; PMCID: PMC5156315.
232. Chang Q, Lipsitch M, Hanage WP. Impact of Host Heterogeneity on the Efficacy of Interventions to Reduce *Staphylococcus aureus* Carriage. *Infect Control Hosp Epidemiol.* 2016 Feb;37(2):197-204. doi: [10.1017/ice.2015.269](https://doi.org/10.1017/ice.2015.269). Epub 2015 Nov 24. Erratum in: *Infect Control Hosp Epidemiol.* 2016 Feb;37(2):244. PMID: 26598029; PMCID: PMC4760641.
233. Grad YH, Goldstein E, Lipsitch M, White PJ. Improving Control of Antibiotic-Resistant Gonorrhea by Integrating Research Agendas Across Disciplines: Key Questions Arising From Mathematical Modeling. *J Infect Dis.* 2016 Mar 15;213(6):883-90. doi: [10.1093/infdis/jiv517](https://doi.org/10.1093/infdis/jiv517). Epub 2015 Oct 30. PMID: 26518045; PMCID: PMC4760416.
234. García-Albéniz X, Hsu J, Lipsitch M, Logan RW, Hernández-Díaz S, Hernán MA. Infective endocarditis and cancer in the elderly. *Eur J Epidemiol.* 2016 Jan;31(1):41-9. doi: [10.1007/s10654-015-0111-9](https://doi.org/10.1007/s10654-015-0111-9). Epub 2015 Dec 18. PMID: 26683995; PMCID: PMC5354127.
235. Chang HH, Dordel J, Donker T, Worby CJ, Feil EJ, Hanage WP, Bentley SD, Huang SS, Lipsitch M. Identifying the effect of patient sharing on between-hospital genetic differentiation of methicillin-resistant *Staphylococcus aureus*. *Genome Med.* 2016 Feb 13;8(1):18. doi: [10.1186/s13073-016-0274-3](https://doi.org/10.1186/s13073-016-0274-3). PMID: 26873713; PMCID: PMC4752745.
236. Lipsitch M, Siber GR. How Can Vaccines Contribute to Solving the Antimicrobial Resistance Problem? *mBio.* 2016 Jun 7;7(3):e00428-16. doi: [10.1128/mBio.00428-16](https://doi.org/10.1128/mBio.00428-16). PMID: 27273824; PMCID: PMC4959668.
237. Lipsitch M, Evans NG, Cotton-Barratt O. Underprotection of Unpredictable Statistical Lives Compared to Predictable Ones. *Risk Anal.* 2017 May;37(5):893-904. doi: [10.1111/risa.12658](https://doi.org/10.1111/risa.12658). Epub 2016 Jul 9. PMID: 27393181; PMCID: PMC5222861.
238. Lipsitch M, Jha A, Simonsen L. Observational studies and the difficult quest for causality: lessons from vaccine effectiveness and impact studies. *Int J Epidemiol.* 2016 Dec 1;45(6):2060-2074. doi: [10.1093/ije/dyw124](https://doi.org/10.1093/ije/dyw124). PMID: 27453361; PMCID: PMC5841615.
239. Abuelezam NN, McCormick AW, Fussell T, Afriyie AN, Wood R, DeGruttola V, Freedberg KA, Lipsitch M, Seage GR 3rd. Can the Heterosexual HIV Epidemic be Eliminated in South Africa Using Combination Prevention? A Modeling Analysis. *Am J Epidemiol.* 2016 Aug 1;184(3):239-48. doi: [10.1093/aje/kwv344](https://doi.org/10.1093/aje/kwv344). Epub 2016 Jul 13. PMID: 27416841; PMCID: PMC4967594.

240. García-Albéniz X, Hsu J, Lipsitch M, Bretthauer M, Logan RW, Hernández-Díaz S, Hernán MA. Colonoscopy and Risk of Infective Endocarditis in the Elderly. *J Am Coll Cardiol*. 2016 Aug 2;68(5):570-571. doi: [10.1016/j.jacc.2016.05.041](https://doi.org/10.1016/j.jacc.2016.05.041). PMID: 27470461; PMCID: PMC5287292.
241. Lipsitch M, Cowling BJ. Zika vaccine trials. *Science*. 2016 Sep 9;353(6304):1094-5. doi: [10.1126/science.aai8126](https://doi.org/10.1126/science.aai8126). PMID: 27609872.
242. Grad YH, Harris SR, Kirkcaldy RD, Green AG, Marks DS, Bentley SD, Trees D, Lipsitch M. Genomic Epidemiology of Gonococcal Resistance to Extended-Spectrum Cephalosporins, Macrolides, and Fluoroquinolones in the United States, 2000-2013. *J Infect Dis*. 2016 Nov 15;214(10):1579-1587. doi: [10.1093/infdis/jiw420](https://doi.org/10.1093/infdis/jiw420). Epub 2016 Sep 16. PMID: 27638945; PMCID: PMC5091375. **Nominee, Charles R. Shepard Science Award, CDC.**
243. Goldstein E, Pitzer VE, O'Hagan JJ, Lipsitch M. Temporally Varying Relative Risks for Infectious Diseases: Implications for Infectious Disease Control. *Epidemiology*. 2017 Jan;28(1):136-144. doi: [10.1097/EDE.0000000000000571](https://doi.org/10.1097/EDE.0000000000000571). PMID: 27748685; PMCID: PMC5131868.
244. Wu JT, Peak CM, Leung GM, Lipsitch M. Fractional dosing of yellow fever vaccine to extend supply: a modelling study. *Lancet*. 2016 Dec 10;388(10062):2904-2911. doi: [10.1016/S0140-6736\(16\)31838-4](https://doi.org/10.1016/S0140-6736(16)31838-4). Epub 2016 Nov 10. PMID: 27837923; PMCID: PMC5161610.
245. Lipsitch M, Barclay W, Raman R, Russell CJ, Belser JA, Cobey S, Kasson PM, Lloyd-Smith JO, Maurer-Stroh S, Riley S, Beauchemin CA, Bedford T, Friedrich TC, Handel A, Herfst S, Murcia PR, Roche B, Wilke CO, Russell CA. Viral factors in influenza pandemic risk assessment. *Elife*. 2016 Nov 11;5:e18491. doi: [10.7554/eLife.18491](https://doi.org/10.7554/eLife.18491). PMID: 27834632; PMCID: PMC5156527.
246. Leung K, Lipsitch M, Yuen KY, Wu JT. Monitoring the fitness of antiviral-resistant influenza strains during an epidemic: a mathematical modelling study. *Lancet Infect Dis*. 2017 Mar;17(3):339-347. doi: [10.1016/S1473-3099\(16\)30465-0](https://doi.org/10.1016/S1473-3099(16)30465-0). Epub 2016 Dec 1. PMID: 27914853; PMCID: PMC5470942.
247. Croucher NJ, Campo JJ, Le TQ, Liang X, Bentley SD, Hanage WP, Lipsitch M. Diverse evolutionary patterns of pneumococcal antigens identified by pangenome-wide immunological screening. *Proc Natl Acad Sci U S A*. 2017 Jan 17;114(3):E357-E366. doi: [10.1073/pnas.1613937114](https://doi.org/10.1073/pnas.1613937114). Epub 2017 Jan 4. PMID: 28053228; PMCID: PMC5255586.
248. Lehtinen S, Blanquart F, Croucher NJ, Turner P, Lipsitch M, Fraser C. Evolution of antibiotic resistance is linked to any genetic mechanism affecting bacterial duration of carriage. *Proc Natl Acad Sci U S A*. 2017 Jan 31;114(5):1075-1080. doi: [10.1073/pnas.1617849114](https://doi.org/10.1073/pnas.1617849114). Epub 2017 Jan 17. PMID: 28096340; PMCID: PMC5293062.
249. Azarian T, Grant LR, Georgieva M, Hammitt LL, Reid R, Bentley SD, Goldblatt D, Santosham M, Weatherholtz R, Burbidge P, Goklish N, Thompson CM, Hanage WP, O'Brien KL, Lipsitch M. Association of Pneumococcal Protein Antigen Serology With Age and Antigenic Profile of Colonizing Isolates. *J Infect Dis*. 2017 Mar 1;215(5):713-722. doi: [10.1093/infdis/jiw628](https://doi.org/10.1093/infdis/jiw628). PMID: 28035010; PMCID: PMC6005115.
250. Hitchings MD, Grais RF, Lipsitch M. Using simulation to aid trial design: Ring-vaccination trials. *PLoS Negl Trop Dis*. 2017 Mar 22;11(3):e0005470. doi: [10.1371/journal.pntd.0005470](https://doi.org/10.1371/journal.pntd.0005470). PMID: 28328984; PMCID: PMC5378415.
251. McCormick AW, Abuelezam NN, Fussell T, Seage GR 3rd, Lipsitch M. Displacement of sexual partnerships in trials of sexual behavior interventions: A model-based assessment of consequences. *Epidemics*. 2017 Sep;20:94-101.

- doi: [10.1016/j.epidem.2017.03.007](https://doi.org/10.1016/j.epidem.2017.03.007). Epub 2017 Apr 2. PMID: 28416219; PMCID: PMC5610917.
252. Eyal N, Lipsitch M. Vaccine testing for emerging infections: the case for individual randomisation. *J Med Ethics*. 2017 Sep;43(9):625-631. doi: [10.1136/medethics-2015-103220](https://doi.org/10.1136/medethics-2015-103220). Epub 2017 Apr 10. PMID: 28396558; PMCID: PMC5577361.
 253. Worby CJ, Wallinga J, Lipsitch M, Goldstein E. Population effect of influenza vaccination under co-circulation of non-vaccine variants and the case for a bivalent A/H3N2 vaccine component. *Epidemics*. 2017 Jun;19:74-82. doi: [10.1016/j.epidem.2017.02.008](https://doi.org/10.1016/j.epidem.2017.02.008). Epub 2017 Feb 21. PMID: 28262588; PMCID: PMC5533618.
 254. Cobey S, Baskerville EB, Colijn C, Hanage W, Fraser C, Lipsitch M. Host population structure and treatment frequency maintain balancing selection on drug resistance. *J R Soc Interface*. 2017 Aug;14(133):20170295. doi: [10.1098/rsif.2017.0295](https://doi.org/10.1098/rsif.2017.0295). PMID: 28835542; PMCID: PMC5582124.
 255. Lipsitch M, Eyal N. Improving vaccine trials in infectious disease emergencies. *Science*. 2017 Jul 14;357(6347):153-156. doi: [10.1126/science.aam8334](https://doi.org/10.1126/science.aam8334). PMID: 28706038; PMCID: PMC5568786.
 256. Corander J, Fraser C, Gutmann MU, Arnold B, Hanage WP, Bentley SD, Lipsitch M, Croucher NJ. Frequency-dependent selection in vaccine-associated pneumococcal population dynamics. *Nat Ecol Evol*. 2017 Dec;1(12):1950-1960. doi: [10.1038/s41559-017-0337-x](https://doi.org/10.1038/s41559-017-0337-x). Epub 2017 Oct 16. PMID: 29038424; PMCID: PMC5708525.
 257. Lee GM, Kleinman K, Pelton S, Lipsitch M, Huang SS, Lakoma M, Dutta-Linn M, Rett M, Hanage WP, Finkelstein JA. Immunization, Antibiotic Use, and Pneumococcal Colonization Over a 15-Year Period. *Pediatrics*. 2017 Nov;140(5):e20170001. doi: [10.1542/peds.2017-0001](https://doi.org/10.1542/peds.2017-0001). Epub 2017 Oct 4. PMID: 28978716; PMCID: PMC5654389.
 258. Langwig KE, Wargo AR, Jones DR, Viss JR, Rutan BJ, Egan NA, Sá-Guimarães P, Kim MS, Kurath G, Gomes MGM, Lipsitch M. Vaccine Effects on Heterogeneity in Susceptibility and Implications for Population Health Management. *mBio*. 2017 Nov 21;8(6):e00796-17. doi: [10.1128/mBio.00796-17](https://doi.org/10.1128/mBio.00796-17). PMID: 29162706; PMCID: PMC5698548.
 259. Lewnard JA, Givon-Lavi N, Weinberger DM, Lipsitch M, Dagan R. Pan-serotype Reduction in Progression of *Streptococcus pneumoniae* to Otitis Media After Rollout of Pneumococcal Conjugate Vaccines. *Clin Infect Dis*. 2017 Nov 13;65(11):1853-1861. doi: [10.1093/cid/cix673](https://doi.org/10.1093/cid/cix673). PMID: 29020218; PMCID: PMC6248775.
 260. Goldstein E, Nguyen HH, Liu P, Viboud C, Steiner CA, Worby CJ, Lipsitch M. On the Relative Role of Different Age Groups During Epidemics Associated With Respiratory Syncytial Virus. *J Infect Dis*. 2018 Jan 4;217(2):238-244. doi: [10.1093/infdis/jix575](https://doi.org/10.1093/infdis/jix575). PMID: 29112722; PMCID: PMC5853559.
 261. Masala GL, Lipsitch M, Bottomley C, Flasche S. Exploring the role of competition induced by non-vaccine serotypes for herd protection following pneumococcal vaccination. *J R Soc Interface*. 2017 Nov;14(136):20170620. doi: [10.1098/rsif.2017.0620](https://doi.org/10.1098/rsif.2017.0620). PMID: 29093131; PMCID: PMC5721164.
 262. Kahn R, Hitchings M, Bellan S, Lipsitch M. Impact of stochastically generated heterogeneity in hazard rates on individually randomized vaccine efficacy trials. *Clin Trials*. 2018 Apr;15(2):207-211. doi: [10.1177/1740774517752671](https://doi.org/10.1177/1740774517752671). Epub 2018 Jan 27. PMID: 29374974; PMCID: PMC5891371.
 263. Halloran ME, Auranen K, Baird S, Basta NE, Bellan SE, Brookmeyer R, Cooper BS, DeGruttola V, Hughes JP, Lessler J, Lofgren ET, Longini IM, Onnela JP,

- Özler B, Seage GR, Smith TA, Vespignani A, Vynnycky E, Lipsitch M. Simulations for designing and interpreting intervention trials in infectious diseases. *BMC Med.* 2017 Dec 29;15(1):223. doi: [10.1186/s12916-017-0985-3](https://doi.org/10.1186/s12916-017-0985-3). PMID: 29287587; PMCID: PMC5747936.
264. Lipsitch M, Li LM, Patterson S, Trammel J, Juergens C, Gruber WC, Scott DA, Dagan R. Serotype-specific immune responses to pneumococcal conjugate vaccine among children are significantly correlated by individual: Analysis of randomized controlled trial data. *Vaccine.* 2018 Jan 25;36(4):473-478. doi: [10.1016/j.vaccine.2017.12.015](https://doi.org/10.1016/j.vaccine.2017.12.015). Epub 2017 Dec 14. PMID: 29248266; PMCID: PMC5767551.
265. Arnold BJ, Gutmann MU, Grad YH, Sheppard SK, Corander J, Lipsitch M, Hanage WP. Weak Epistasis May Drive Adaptation in Recombining Bacteria. *Genetics.* 2018 Mar;208(3):1247-1260. doi: [10.1534/genetics.117.300662](https://doi.org/10.1534/genetics.117.300662). Epub 2018 Jan 12. PMID: 29330348; PMCID: PMC5844334.
266. Hitchings MDT, Lipsitch M, Wang R, Bellan SE. Competing Effects of Indirect Protection and Clustering on the Power of Cluster-Randomized Controlled Vaccine Trials. *Am J Epidemiol.* 2018 Aug 1;187(8):1763-1771. doi: [10.1093/aje/kwy047](https://doi.org/10.1093/aje/kwy047). PMID: 29522080; PMCID: PMC6070038.
267. Worby CJ, Lipsitch M, Hanage WP. Shared Genomic Variants: Identification of Transmission Routes Using Pathogen Deep-Sequence Data. *Am J Epidemiol.* 2017 Nov 15;186(10):1209-1216. doi: [10.1093/aje/kwx182](https://doi.org/10.1093/aje/kwx182). PMID: 29149252; PMCID: PMC5860558. **Selected as one of the 2017 Articles of the Year, *The American Journal of Epidemiology*.**
268. Johansson MA, Reich NG, Meyers LA, Lipsitch M. Preprints: An underutilized mechanism to accelerate outbreak science. *PLoS Med.* 2018 Apr 3;15(4):e1002549. doi: [10.1371/journal.pmed.1002549](https://doi.org/10.1371/journal.pmed.1002549). PMID: 29614073; PMCID: PMC5882117.
269. Azarian T, Grant LR, Arnold BJ, Hammitt LL, Reid R, Santosham M, Weatherholtz R, Goklish N, Thompson CM, Bentley SD, O'Brien KL, Hanage WP, Lipsitch M. The impact of serotype-specific vaccination on phylodynamic parameters of *Streptococcus pneumoniae* and the pneumococcal pan-genome. *PLoS Pathog.* 2018 Apr 4;14(4):e1006966. doi: [10.1371/journal.ppat.1006966](https://doi.org/10.1371/journal.ppat.1006966). PMID: 29617440; PMCID: PMC5902063.
270. Goldstein E, Worby CJ, Lipsitch M. On the Role of Different Age Groups and Pertussis Vaccines During the 2012 Outbreak in Wisconsin. *Open Forum Infect Dis.* 2018 Apr 16;5(5):ofy082. doi: [10.1093/ofid/ofy082](https://doi.org/10.1093/ofid/ofy082). PMID: 29942818; PMCID: PMC5961225.
271. Blanquart F, Lehtinen S, Lipsitch M, Fraser C. The evolution of antibiotic resistance in a structured host population. *J R Soc Interface.* 2018 Jun;15(143):20180040. doi: [10.1098/rsif.2018.0040](https://doi.org/10.1098/rsif.2018.0040). PMID: 29925579; PMCID: PMC6030642.
272. Lewnard JA, Tähtinen PA, Laine MK, Lindholm L, Jalava J, Huovinen P, Lipsitch M, Ruohola A. Impact of Antimicrobial Treatment for Acute Otitis Media on Carriage Dynamics of Penicillin-Susceptible and Penicillin-Nonsusceptible *Streptococcus pneumoniae*. *J Infect Dis.* 2018 Sep 22;218(9):1356-1366. doi: [10.1093/infdis/jiy343](https://doi.org/10.1093/infdis/jiy343). PMID: 29873739; PMCID: PMC6151080.
273. Olesen SW, Barnett ML, MacFadden DR, Lipsitch M, Grad YH. Trends in outpatient antibiotic use and prescribing practice among US older adults, 2011-15: observational study. *BMJ.* 2018 Jul 27;362:k3155. doi: [10.1136/bmj.k3155](https://doi.org/10.1136/bmj.k3155). PMID: 30054353; PMCID: PMC6062849.

274. Kahn R, Rid A, Smith PG, Eyal N, Lipsitch M. Choices in vaccine trial design in epidemics of emerging infections. *PLoS Med*. 2018 Aug 7;15(8):e1002632. doi: [10.1371/journal.pmed.1002632](https://doi.org/10.1371/journal.pmed.1002632). PMID: 30086139; PMCID: PMC6080746.
275. Eyal N, Lipsitch M, Bärnighausen T, Wikler D. Opinion: Risk to study nonparticipants: A procedural approach. *Proc Natl Acad Sci U S A*. 2018 Aug 7;115(32):8051-8053. doi: [10.1073/pnas.1810920115](https://doi.org/10.1073/pnas.1810920115). PMID: 30087210; PMCID: PMC6094093.
276. Cai FY, Fussell T, Cobey S, Lipsitch M. Use of an individual-based model of pneumococcal carriage for planning a randomized trial of a whole-cell vaccine. *PLoS Comput Biol*. 2018 Oct 1;14(10):e1006333. doi: [10.1371/journal.pcbi.1006333](https://doi.org/10.1371/journal.pcbi.1006333). PMID: 30273332; PMCID: PMC6181404.
277. Lewnard JA, Tedijanto C, Cowling BJ, Lipsitch M. Measurement of Vaccine Direct Effects Under the Test-Negative Design. *Am J Epidemiol*. 2018 Dec 1;187(12):2686-2697. doi: [10.1093/aje/kwy163](https://doi.org/10.1093/aje/kwy163). PMID: 30099505; PMCID: PMC6269249.
278. Relman DA, Lipsitch M. Microbiome as a tool and a target in the effort to address antimicrobial resistance. *Proc Natl Acad Sci U S A*. 2018 Dec 18;115(51):12902-12910. doi: [10.1073/pnas.1717163115](https://doi.org/10.1073/pnas.1717163115). PMID: 30559176; PMCID: PMC6304941.
279. Tedijanto C, Olesen SW, Grad YH, Lipsitch M. Estimating the proportion of bystander selection for antibiotic resistance among potentially pathogenic bacterial flora. *Proc Natl Acad Sci U S A*. 2018 Dec 18;115(51):E11988-E11995. doi: [10.1073/pnas.1810840115](https://doi.org/10.1073/pnas.1810840115). PMID: 30559213; PMCID: PMC6304942.
280. Sevilla JP, Bloom DE, Cadarette D, Jit M, Lipsitch M. Toward economic evaluation of the value of vaccines and other health technologies in addressing AMR. *Proc Natl Acad Sci U S A*. 2018 Dec 18;115(51):12911-12919. doi: [10.1073/pnas.1717161115](https://doi.org/10.1073/pnas.1717161115). PMID: 30559203; PMCID: PMC6305008.
281. Campo JJ, Le TQ, Pablo JV, Hung C, Teng AA, Tettelin H, Tate A, Hanage WP, Alderson MR, Liang X, Malley R, Lipsitch M, Croucher NJ. Panproteome-wide analysis of antibody responses to whole cell pneumococcal vaccination. *Elife*. 2018 Dec 28;7:e37015. doi: [10.7554/eLife.37015](https://doi.org/10.7554/eLife.37015). PMID: 30592459; PMCID: PMC6344088.
282. Olesen SW, Barnett ML, MacFadden DR, Brownstein JS, Hernández-Díaz S, Lipsitch M, Grad YH. The distribution of antibiotic use and its association with antibiotic resistance. *Elife*. 2018 Dec 18;7:e39435. doi: [10.7554/eLife.39435](https://doi.org/10.7554/eLife.39435). PMID: 30560781; PMCID: PMC6307856.
283. Georgieva M, Buckee CO, Lipsitch M. Models of immune selection for multi-locus antigenic diversity of pathogens. *Nat Rev Immunol*. 2019 Jan;19(1):55-62. doi: [10.1038/s41577-018-0092-5](https://doi.org/10.1038/s41577-018-0092-5). PMID: 30479379; PMCID: PMC6352731.
284. Abuelezam NN, McCormick AW, Surface ED, Fussell T, Freedberg KA, Lipsitch M, Seage GR. Modelling the epidemiologic impact of achieving UNAIDS fast-track 90-90-90 and 95-95-95 targets in South Africa. *Epidemiol Infect*. 2019 Jan;147:e122. doi: [10.1017/S0950268818003497](https://doi.org/10.1017/S0950268818003497). PMID: 30869008; PMCID: PMC6452860.
285. Mahmud AS, Lipsitch M, Goldstein E. On the role of different age groups during pertussis epidemics in California, 2010 and 2014. *Epidemiol Infect*. 2019 Jan;147:e184. doi: [10.1017/S0950268819000761](https://doi.org/10.1017/S0950268819000761). PMID: 31063110; PMCID: PMC6518560.
286. Olesen SW, Torrone EA, Papp JR, Kirkcaldy RD, Lipsitch M, Grad YH. Azithromycin Susceptibility Among *Neisseria gonorrhoeae* Isolates and Seasonal Macrolide Use. *J Infect Dis*. 2019 Jan 29;219(4):619-623. doi: [10.1093/infdis/jiy551](https://doi.org/10.1093/infdis/jiy551). PMID: 30239814; PMCID: PMC6350947.

287. Gallagher T, Lipsitch M. Postexposure Effects of Vaccines on Infectious Diseases. *Epidemiol Rev.* 2019 Jan 31;41(1):13-27. doi: [10.1093/epirev/mxz014](https://doi.org/10.1093/epirev/mxz014). PMID: 31680134; PMCID: PMC7159179.
288. Kahn R, Hitchings M, Wang R, Bellan SE, Lipsitch M. Analyzing Vaccine Trials in Epidemics With Mild and Asymptomatic Infection. *Am J Epidemiol.* 2019 Feb 1;188(2):467-474. doi: [10.1093/aje/kwy239](https://doi.org/10.1093/aje/kwy239). PMID: 30329134; PMCID: PMC6357804.
289. Mitchell PK, Azarian T, Croucher NJ, Callendrello A, Thompson CM, Pelton SI, Lipsitch M, Hanage WP. Population genomics of pneumococcal carriage in Massachusetts children following introduction of PCV-13. *Microb Genom.* 2019 Feb;5(2):e000252. doi: [10.1099/mgen.0.000252](https://doi.org/10.1099/mgen.0.000252). Epub 2019 Feb 19. PMID: 30777813; PMCID: PMC6421351.
290. Langwig KE, Gomes MGM, Clark MD, Kwitny M, Yamada S, Wargo AR, Lipsitch M. Limited available evidence supports theoretical predictions of reduced vaccine efficacy at higher exposure dose. *Sci Rep.* 2019 Mar 1;9(1):3203. doi: [10.1038/s41598-019-39698-x](https://doi.org/10.1038/s41598-019-39698-x). PMID: 30824732; PMCID: PMC6397254.
291. Yang A, Cai F, Lipsitch M. Herd immunity alters the conditions for performing dose schedule comparisons: an individual-based model of pneumococcal carriage. *BMC Infect Dis.* 2019 Mar 5;19(1):227. doi: [10.1186/s12879-019-3833-6](https://doi.org/10.1186/s12879-019-3833-6). PMID: 30836941; PMCID: PMC6402138.
292. Hitchings MDT, Coldiron ME, Grais RF, Lipsitch M. Analysis of a meningococcal meningitis outbreak in Niger - potential effectiveness of reactive prophylaxis. *PLoS Negl Trop Dis.* 2019 Mar 11;13(3):e0007077. doi: [10.1371/journal.pntd.0007077](https://doi.org/10.1371/journal.pntd.0007077). PMID: 30856166; PMCID: PMC6428357.
293. Ray GT, Lewis N, Klein NP, Daley MF, Lipsitch M, Fireman B. Depletion-of-susceptibles Bias in Analyses of Intra-season Waning of Influenza Vaccine Effectiveness. *Clin Infect Dis.* 2020 Mar 17;70(7):1484-1486. doi: [10.1093/cid/ciz706](https://doi.org/10.1093/cid/ciz706). PMID: 31351439; PMCID: PMC7318775.
294. Lehtinen S, Blanquart F, Lipsitch M, Fraser C; with the Maela Pneumococcal Collaboration. On the evolutionary ecology of multidrug resistance in bacteria. *PLoS Pathog.* 2019 May 13;15(5):e1007763. doi: [10.1371/journal.ppat.1007763](https://doi.org/10.1371/journal.ppat.1007763). PMID: 31083687; PMCID: PMC6532944.
295. McAdams D, Wollein Waldetoft K, Tedijanto C, Lipsitch M, Brown SP. Resistance diagnostics as a public health tool to combat antibiotic resistance: A model-based evaluation. *PLoS Biol.* 2019 May 16;17(5):e3000250. doi: [10.1371/journal.pbio.3000250](https://doi.org/10.1371/journal.pbio.3000250). PMID: 31095567; PMCID: PMC6522007.
296. MacFadden DR, Fisman DN, Hanage WP, Lipsitch M. The Relative Impact of Community and Hospital Antibiotic Use on the Selection of Extended-spectrum Beta-lactamase-producing *Escherichia coli*. *Clin Infect Dis.* 2019 Jun 18;69(1):182-188. doi: [10.1093/cid/ciy978](https://doi.org/10.1093/cid/ciy978). PMID: 30462185; PMCID: PMC6771767.
297. Goldstein E, MacFadden DR, Karaca Z, Steiner CA, Viboud C, Lipsitch M. Antimicrobial resistance prevalence, rates of hospitalization with septicemia and rates of mortality with sepsis in adults in different US states. *Int J Antimicrob Agents.* 2019 Jul;54(1):23-34. doi: [10.1016/j.ijantimicag.2019.03.004](https://doi.org/10.1016/j.ijantimicag.2019.03.004). Epub 2019 Mar 6. PMID: 30851403; PMCID: PMC6571064.
298. Goldstein E, Olesen SW, Karaca Z, Steiner CA, Viboud C, Lipsitch M. Levels of outpatient prescribing for four major antibiotic classes and rates of septicemia hospitalization in adults in different US states - a statistical analysis. *BMC Public Health.* 2019 Aug 19;19(1):1138. doi: [10.1186/s12889-019-7431-8](https://doi.org/10.1186/s12889-019-7431-8). PMID: 31426780; PMCID: PMC6701127.

299. Lee RTC, Chang HH, Russell CA, Lipsitch M, Maurer-Stroh S. Influenza A Hemagglutinin Passage Bias Sites and Host Specificity Mutations. *Cells*. 2019 Aug 22;8(9):958. doi: [10.3390/cells8090958](https://doi.org/10.3390/cells8090958). PMID: 31443542; PMCID: PMC6770435.
300. Abuelezam NN, Reshef YA, Novak D, Grad YH, Seage lii GR, Mayer K, Lipsitch M. Interaction Patterns of Men Who Have Sex With Men on a Geosocial Networking Mobile App in Seven United States Metropolitan Areas: Observational Study. *J Med Internet Res*. 2019 Sep 12;21(9):e13766. doi: [10.2196/13766](https://doi.org/10.2196/13766). PMID: 31516124; PMCID: PMC6746104.
301. Eyal N, Kimmelman J, Holtzman LG, Lipsitch M. Regulating impact on bystanders in clinical trials: An unsettled frontier. *Clin Trials*. 2019 Oct;16(5):450-454. doi: [10.1177/1740774519862783](https://doi.org/10.1177/1740774519862783). Epub 2019 Aug 1. PMID: 31368813; PMCID: PMC6742522.
302. Goldstein E, Finelli L, O'Halloran A, Liu P, Karaca Z, Steiner CA, Viboud C, Lipsitch M. Hospitalizations Associated with Respiratory Syncytial Virus and Influenza in Children, Including Children Diagnosed with Asthma. *Epidemiology*. 2019 Nov;30(6):918-926. doi: [10.1097/EDE.0000000000001092](https://doi.org/10.1097/EDE.0000000000001092). PMID: 31469696; PMCID: PMC6768705.
303. Hicks AL, Kissler SM, Lipsitch M, Grad YH. Surveillance to maintain the sensitivity of genotype-based antibiotic resistance diagnostics. *PLoS Biol*. 2019 Nov 12;17(11):e3000547. doi: [10.1371/journal.pbio.3000547](https://doi.org/10.1371/journal.pbio.3000547). PMID: 31714937; PMCID: PMC6874359.
304. Lipsitch M, Goldstein E, Ray GT, Fireman B. Depletion-of-susceptibles bias in influenza vaccine waning studies: how to ensure robust results. *Epidemiol Infect*. 2019 Nov 27;147:e306. doi: [10.1017/S0950268819001961](https://doi.org/10.1017/S0950268819001961). PMID: 31774051; PMCID: PMC7003633.
305. Knight GM, Davies NG, Colijn C, Coll F, Donker T, Gifford DR, Glover RE, Jit M, Klemm E, Lehtinen S, Lindsay JA, Lipsitch M, Llewelyn MJ, Mateus ALP, Robotham JV, Sharland M, Stekel D, Yakob L, Atkins KE. Mathematical modelling for antibiotic resistance control policy: do we know enough? *BMC Infect Dis*. 2019 Nov 29;19(1):1011. doi: [10.1186/s12879-019-4630-y](https://doi.org/10.1186/s12879-019-4630-y). PMID: 31783803; PMCID: PMC6884858.
306. Ryu S, Cowling BJ, Wu P, Olesen S, Fraser C, Sun DS, Lipsitch M, Grad YH. Case-based surveillance of antimicrobial resistance with full susceptibility profiles. *JAC Antimicrob Resist*. 2019 Dec;1(3):dlz070. doi: [10.1093/jacamr/dlz070](https://doi.org/10.1093/jacamr/dlz070). Epub 2019 Dec 10. PMID: 32280945; PMCID: PMC7134534.
307. Chua H, Feng S, Lewnard JA, Sullivan SG, Blyth CC, Lipsitch M, Cowling BJ. The Use of Test-negative Controls to Monitor Vaccine Effectiveness: A Systematic Review of Methodology. *Epidemiology*. 2020 Jan;31(1):43-64. doi: [10.1097/EDE.0000000000001116](https://doi.org/10.1097/EDE.0000000000001116). PMID: 31609860; PMCID: PMC6888869.
308. Tedijanto C, Grad YH, Lipsitch M. Potential impact of outpatient stewardship interventions on antibiotic exposures of common bacterial pathogens. *Elife*. 2020 Feb 5;9:e52307. doi: [10.7554/eLife.52307](https://doi.org/10.7554/eLife.52307). PMID: 32022685; PMCID: PMC7025820.
309. Lipsitch M, Swerdlow DL, Finelli L. Defining the Epidemiology of Covid-19 - Studies Needed. *N Engl J Med*. 2020 Mar 26;382(13):1194-1196. doi: [10.1056/NEJMp2002125](https://doi.org/10.1056/NEJMp2002125). Epub 2020 Feb 19. PMID: 32074416.
310. Kennedy-Shaffer L, de Gruttola V, Lipsitch M. Novel methods for the analysis of stepped wedge cluster randomized trials. *Stat Med*. 2020 Mar 30;39(7):815-844.

- doi: [10.1002/sim.8451](https://doi.org/10.1002/sim.8451). Epub 2019 Dec 26. PMID: 31876979; PMCID: PMC7247054.
311. Goldstein E, Lipsitch M. Temporal rise in the proportion of younger adults and older adolescents among coronavirus disease (COVID-19) cases following the introduction of physical distancing measures, Germany, March to April 2020. *Euro Surveill.* 2020 Apr;25(17):2000596. doi: [10.2807/1560-7917.ES.2020.25.17.2000596](https://doi.org/10.2807/1560-7917.ES.2020.25.17.2000596). PMID: 32372753; PMCID: PMC7201953.
 312. Wu JT, Leung K, Bushman M, Kishore N, Niehus R, de Salazar PM, Cowling BJ, Lipsitch M, Leung GM. Estimating clinical severity of COVID-19 from the transmission dynamics in Wuhan, China. *Nat Med.* 2020 Apr;26(4):506-510. doi: [10.1038/s41591-020-0822-7](https://doi.org/10.1038/s41591-020-0822-7). Epub 2020 Mar 19. Erratum in: *Nat Med.* 2020 Jul;26(7):1149-1150. PMID: 32284616; PMCID: PMC7094929.
 313. McGough SF, Johansson MA, Lipsitch M, Menzies NA. Nowcasting by Bayesian Smoothing: A flexible, generalizable model for real-time epidemic tracking. *PLoS Comput Biol.* 2020 Apr 6;16(4):e1007735. doi: [10.1371/journal.pcbi.1007735](https://doi.org/10.1371/journal.pcbi.1007735). PMID: 32251464; PMCID: PMC7162546.
 314. MacFadden DR, Coburn B, Břinda K, Corbeil A, Daneman N, Fisman D, Lee RS, Lipsitch M, McGeer A, Melano RG, Mubareka S, Hanage WP. Using Genetic Distance from Archived Samples for the Prediction of Antibiotic Resistance in *Escherichia coli*. *Antimicrob Agents Chemother.* 2020 Apr 21;64(5):e02417-19. doi: [10.1128/AAC.02417-19](https://doi.org/10.1128/AAC.02417-19). PMID: 32152083; PMCID: PMC7179619.
 315. Li R, Rivers C, Tan Q, Murray MB, Toner E, Lipsitch M. Estimated Demand for US Hospital Inpatient and Intensive Care Unit Beds for Patients With COVID-19 Based on Comparisons With Wuhan and Guangzhou, China. *JAMA Netw Open.* 2020 May 1;3(5):e208297. doi: [10.1001/jamanetworkopen.2020.8297](https://doi.org/10.1001/jamanetworkopen.2020.8297). PMID: 32374400; PMCID: PMC7203604.
 316. Eyal N, Lipsitch M, Smith PG. Human Challenge Studies to Accelerate Coronavirus Vaccine Licensure. *J Infect Dis.* 2020 May 11;221(11):1752-1756. doi: [10.1093/infdis/jiaa152](https://doi.org/10.1093/infdis/jiaa152). PMID: 32232474; PMCID: PMC7184325.
 317. Lehtinen S, Chewapreecha C, Lees J, Hanage WP, Lipsitch M, Croucher NJ, Bentley SD, Turner P, Fraser C, Mostowy RJ. Horizontal gene transfer rate is not the primary determinant of observed antibiotic resistance frequencies in *Streptococcus pneumoniae*. *Sci Adv.* 2020 May 20;6(21):eaaz6137. doi: [10.1126/sciadv.aaz6137](https://doi.org/10.1126/sciadv.aaz6137). PMID: 32671212; PMCID: PMC7314567.
 318. Kissler SM, Tedijanto C, Goldstein E, Grad YH, Lipsitch M. Projecting the transmission dynamics of SARS-CoV-2 through the postpandemic period. *Science.* 2020 May 22;368(6493):860-868. doi: [10.1126/science.abb5793](https://doi.org/10.1126/science.abb5793). Epub 2020 Apr 14. PMID: 32291278; PMCID: PMC7164482.
 319. Lipsitch M. Good science is good science: we need specialists, not sects. *Eur J Epidemiol.* 2020 Jun;35(6):519-522. doi: [10.1007/s10654-020-00651-2](https://doi.org/10.1007/s10654-020-00651-2). Epub 2020 Jun 20. PMID: 32564181; PMCID: PMC7305476.
 320. Flasche S, Lipsitch M, Ojal J, Pinsent A. Estimating the contribution of different age strata to vaccine serotype pneumococcal transmission in the pre vaccine era: a modelling study. *BMC Med.* 2020 Jun 10;18(1):129. doi: [10.1186/s12916-020-01601-1](https://doi.org/10.1186/s12916-020-01601-1). PMID: 32517683; PMCID: PMC7285529.
 321. Hicks AL, Kissler SM, Mortimer TD, Ma KC, Taiaroa G, Ashcroft M, Williamson DA, Lipsitch M, Grad YH. Targeted surveillance strategies for efficient detection of novel antibiotic resistance variants. *Elife.* 2020 Jun 30;9:e56367. doi: [10.7554/eLife.56367](https://doi.org/10.7554/eLife.56367). PMID: 32602459; PMCID: PMC7326491.
 322. De Salazar PM, Niehus R, Taylor A, Buckee CO, Lipsitch M. Identifying Locations with Possible Undetected Imported Severe Acute Respiratory

- Syndrome Coronavirus 2 Cases by Using Importation Predictions. *Emerg Infect Dis.* 2020 Jul;26(7):1465-1469. doi: [10.3201/eid2607.200250](https://doi.org/10.3201/eid2607.200250). Epub 2020 Jun 21. PMID: 32207679; PMCID: PMC7323530.
323. Niehus R, De Salazar PM, Taylor AR, Lipsitch M. Using observational data to quantify bias of traveller-derived COVID-19 prevalence estimates in Wuhan, China. *Lancet Infect Dis.* 2020 Jul;20(7):803-808. doi: [10.1016/S1473-3099\(20\)30229-2](https://doi.org/10.1016/S1473-3099(20)30229-2). Epub 2020 Apr 1. PMID: 32246905; PMCID: PMC7270516.
324. Kennedy-Shaffer L, Lipsitch M. Statistical Properties of Stepped Wedge Cluster-Randomized Trials in Infectious Disease Outbreaks. *Am J Epidemiol.* 2020 Jul 10;kwaa141. doi: [10.1093/aje/kwaa141](https://doi.org/10.1093/aje/kwaa141). Epub ahead of print. PMID: 32648891.
325. Eyal N, Lipsitch M. Opinion: It's ethical to test promising coronavirus vaccines against less-promising ones. *Proc Natl Acad Sci U S A.* 2020 Aug 11;117(32):18898-18901. doi: [10.1073/pnas.2014154117](https://doi.org/10.1073/pnas.2014154117). Epub 2020 Jul 22. PMID: 32699147; PMCID: PMC7431044.
326. Peak CM, Kahn R, Grad YH, Childs LM, Li R, Lipsitch M, Buckee CO. Individual quarantine versus active monitoring of contacts for the mitigation of COVID-19: a modelling study. *Lancet Infect Dis.* 2020 Sep;20(9):1025-1033. doi: [10.1016/S1473-3099\(20\)30361-3](https://doi.org/10.1016/S1473-3099(20)30361-3). Epub 2020 May 20. PMID: 32445710; PMCID: PMC7239635.
327. Kahn R, Kennedy-Shaffer L, Grad YH, Robins JM, Lipsitch M. Potential Biases Arising from Epidemic Dynamics in Observational Seroprotection Studies. *Am J Epidemiol.* 2020 Sep 1;kwaa188. doi: [10.1093/aje/kwaa188](https://doi.org/10.1093/aje/kwaa188). Epub ahead of print. PMID: 32870977; PMCID: PMC7499481.
328. Levinson M, Cevik M, Lipsitch M. Reopening Primary Schools during the Pandemic. *N Engl J Med.* 2020 Sep 3;383(10):981-985. doi: [10.1056/NEJMms2024920](https://doi.org/10.1056/NEJMms2024920). Epub 2020 Jul 29. PMID: 32726550.
329. Messinger CJ, Lipsitch M, Bateman BT, He M, Huybrechts KF, MacDonald S, Mogun H, Mott K, Hernández-Díaz S. Association Between Congenital Cytomegalovirus and the Prevalence at Birth of Microcephaly in the United States. *JAMA Pediatr.* 2020 Sep 14:e203009. doi: [10.1001/jamapediatrics.2020.3009](https://doi.org/10.1001/jamapediatrics.2020.3009). Epub ahead of print. PMID: 32926077; PMCID: PMC7490747.
330. Lipsitch M, Grad YH, Sette A, Crotty S. Cross-reactive memory T cells and herd immunity to SARS-CoV-2. *Nat Rev Immunol.* 2020 Oct 6. doi: [10.1038/s41577-020-00460-4](https://doi.org/10.1038/s41577-020-00460-4). Epub ahead of print. PMID: 33024281.

Articles in Press

1. Selection of Macrolide and Non-Macrolide Resistance with Mass Azithromycin Distribution: A Community-Randomized Trial. *N Engl J Med.* Forthcoming 2020.
2. Lipsitch M. COVID-19 vaccines: Understanding the details of how well they work. *Science.* Forthcoming 2020.

Data Sets

1. Croucher NJ, Finkelstein JA, Pelton SI, Parkhill J, Bentley SD, Lipsitch M, Hanage WP. Population genomic datasets describing the post-vaccine evolutionary epidemiology of *Streptococcus pneumoniae*. *Sci Data.* 2015 Oct

27;2:150058. doi: 10.1038/sdata.2015.58. eCollection 2015. PMID: 26528397; PMCID: PMC4622223.

Preprints

1. Mooring EQ, Marks M, Mitja O, Castro M, Lipsitch M, Murray MB. Programmatic goals and spatial epidemiology influence the merit of targeted versus of population-wide interventions for yaws eradication. *bioRxiv* [Preprint]. 2019 May 17. doi.org/10.1101/640326
2. Luo W, Majumder MS, Liu D, Poirier C, Mandel KD, Lipsitch M, Santillana M. The role of absolute humidity on transmission rates of the COVID-19 outbreak. *medRxiv*. [Preprint]. 2020 Feb 17. doi.org/10.1101/2020.02.12.20022467
3. Niehus R, De Salazar PM, Taylor AR, Lipsitch M. Quantifying bias of COVID-19 prevalence and severity estimates in Wuhan, China that depend on reported cases in international travelers. *medRxiv* [Preprint]. 2020 Feb 18:2020.02.13.20022707. doi: 10.1101/2020.02.13.20022707. PMID: 32511442; PMCID: PMC7239063.
4. Azarian T, Martinez PP, Arnold BJ, Grant LR, Corander J, Fraser C, Croucher NJ, Hammitt LL, Reid R, Santosham M, Weatherholtz, RC, Bentley SD, O'Brien KL, Lipsitch M, Hanage WP. Predicting evolution using frequency-dependent selection in bacterial populations. *bioRxiv* [Preprint]. 2020 Feb 25. doi.org/10.1101/420315
5. Li R, Rivers C, Tan Q, Murray MB, Toner E, Lipsitch M. The demand for inpatient and ICU beds for COVID-19 in the US: lessons from Chinese cities. *medRxiv* [Preprint]. 2020 Mar 16:2020.03.09.20033241. doi: [10.1101/2020.03.09.20033241](https://doi.org/10.1101/2020.03.09.20033241). PMID: 32511447; PMCID: PMC7239072.
6. Kissler SM, Tedijanto C, Lipsitch M, Grad Y. Social distancing strategies for curbing the COVID-19 epidemic. *medRxiv* [Preprint]. 2020 Mar 24: 2020.03.22.20041079v1. doi.org/10.1101/2020.03.22.20041079
7. Menkir TF, Chin T, Hay JA, Surface E, Martinez de Salazar P, Buckee C, Mina MJ, Khan K, Watts A, Lipsitch M, Niehus R. Estimating the number of undetected COVID-19 cases exported internationally from all of China. *medRxiv* [Preprint]. 2020 Mar 26:2020.03.23.20038331. doi: [10.1101/2020.03.23.20038331](https://doi.org/10.1101/2020.03.23.20038331). PMID: 32511613; PMCID: PMC7276040.
8. Lu FS, Nguyen AT, Link NB, Lipsitch M, Santillana M. Estimating the Early Outbreak Cumulative Incidence of COVID-19 in the United States: Three Complementary Approaches. *medRxiv* [Preprint]. 2020 Jun 18:2020.04.18.20070821. doi: [10.1101/2020.04.18.20070821](https://doi.org/10.1101/2020.04.18.20070821). PMID: 32587997; PMCID: PMC7310656.
9. Gostic KM, McGough L, Baskerville E, Abbott S, Joshi K, Tedijanto C, Kahn R, Niehus R, Hay JA, De Salazar PM, Hellewell J, Meakin S, Munday J, Bosse N, Sherratt K, Thompson RM, White LF, Huisman J, Scire J, Bonhoeffer S, Stadler T, Wallinga J, Funk S, Lipsitch M, Cobey S. Practical considerations for measuring the effective reproductive number, Rt. *medRxiv* [Preprint]. 2020 Jun 20:2020.06.18.20134858. doi: [10.1101/2020.06.18.20134858](https://doi.org/10.1101/2020.06.18.20134858). PMID: 32607522; PMCID: PMC7325187.
10. Olesen SW, Lipsitch M, Grad Y. The role of “spillover” in antibiotic resistance. *bioRxiv* [Preprint]. 2020 Jun 30. doi.org/10.1101/536714
11. Martinez de Salazar P, Gomez-Barroso D, Pampaka D, Gil JM, Penalver B, Fernandez-Escobar C, Lipsitch M, Larrauri A, Goldstein E, Hernan M.

- Lockdown measures and relative changes in the age-specific incidence of SARS-CoV-2 in Spain. medRxiv [Preprint]. 2020 Jul 2:2020.06.30.20143560. doi: [10.1101/2020.06.30.20143560](https://doi.org/10.1101/2020.06.30.20143560). PMID: 32637975; PMCID: PMC7340201.
12. Goldstein E, Lipsitch M, Cevik M. On the effect of age on the transmission of SARS-CoV-2 in households, schools and the community. medRxiv [Preprint]. 2020 Jul 24:2020.07.19.20157362. doi: [10.1101/2020.07.19.20157362](https://doi.org/10.1101/2020.07.19.20157362). PMID: 32743609; PMCID: PMC7386533.
 13. Lu FS, Nguyen AT, Link NB, Lipsitch M, Santillana M. Estimating the Early Outbreak Cumulative Incidence of COVID-19 in the United States: Three Complementary Approaches. medRxiv [Preprint]. 2020 Jun 18:2020.04.18.20070821. doi: [10.1101/2020.04.18.20070821](https://doi.org/10.1101/2020.04.18.20070821). PMID: 32587997; PMCID: PMC7310656.
 14. Bubar KM, Kissler SM, Lipsitch M, Cobey S, Grad Y, Larremore DB. Model-informed COVID-19 vaccine prioritization strategies by age and serostatus. medRxiv [Preprint]. 2020 Sept 10. doi.org/[10.1101/2020.09.08.20190629](https://doi.org/10.1101/2020.09.08.20190629)
 15. Kahn R, Wang R, Leavitt S, Hanage WP, Lipsitch M. Leveraging pathogen sequence and contact tracing data to enhance vaccine trials in emerging epidemics. medRxiv [Preprint]. 2020 Sept 16. doi.org/[10.1101/2020.09.14.20193789](https://doi.org/10.1101/2020.09.14.20193789)
 16. Hay J, Kennedy-Shaffer L, Kanjilal S, Lipsitch M, Mina M. Estimating epidemiologic dynamics from single cross-sectional viral load distributions. Harvard DASH [Preprint]. 2020. <https://nrs.harvard.edu/URN-3:HUL.INSTREPOS:37365444>
 17. de Kraker MEA and Lipsitch M. Burden of antimicrobial resistance: compared to what? Harvard DASH [Preprint]. 2020. <https://nrs.harvard.edu/URN-3:HUL.INSTREPOS:37365584>
 18. Eyal N and Lipsitch M. Testing SARS-CoV-2 vaccine efficacy through deliberate natural viral exposure. Harvard DASH [Preprint]. 2020. <https://nrs.harvard.edu/URN-3:HUL.INSTREPOS:37365585>

Other Publications

(a) Book Chapters

1. Lipsitch M, Levin BR. 1997. The within-host population dynamics of anti-bacterial chemotherapy: conditions for the evolution of resistance. Pp. 112-127 in Ciba Foundation Symposium No. 207: Antibiotic Resistance: Origins, Evolution, Selection and Spread. Chichester, UK: John Wiley & Sons.
2. Bangham C, Anderson R, Baquero F, Bax R, Hastings I, Koella J, Lipsitch M, McLean A, Smith T, Taddei F, Levin B. 1999. Evolution of infectious diseases: The impact of vaccines, drugs and social factors. Chapter 13 (pp. 152-160) in Evolution in Health and Disease, ed. S.C. Stearns. Oxford: Oxford University Press.
3. Lipsitch M, Bergstrom CT. Modeling of antibiotic resistance in the ICU. Chapter 18, pp. 231-43 in Infection Control in the ICU Environment, ed. R.A. Weinstein and M. Bonten. Kluwer Press. 2002.
4. Lipsitch M. Vaccination against Haemophilus influenzae and Streptococcus pneumoniae: a problem in virulence management. Chapter 26 in Virulence Management: The Adaptive Dynamics of Pathogen-Host Interactions, ed. U.

- Dieckmann, H. Metz, M. Sabelis & K. Sigmund. Cambridge: Cambridge University Press. 2002.
5. Lipsitch M. Antibiotic resistance: Strategies for managing resistance. Vol. 1, pp. 57-61 in *The Oxford Encyclopedia of Evolution*, ed. M. Pagel. Oxford University Press. 2002.
 6. Dagan R, Lipsitch M. Ecological Effects of Vaccines and Antibiotics. Chapter 18 in *The Pneumococcus*, ed. E. Tuomanen, T. Mitchell, D. Morrison, B. Spratt. Washington, DC: ASM Press. 2004.
 7. Lipsitch M. Infectious Disease Epidemiology. Chapter 16 in *Teaching Epidemiology*, 3rd Edition, ed. J. Olsen, G. Saracci, D. Trichopoulos. New York: Oxford University 2010.
 8. Chapter 16 in 4th Ed., ed. Olsen J, Greene N, Saracci G, Trichopoulos D. New York: Oxford University 2015.
 9. Lipsitch M and Smith D. Application of quantitative modeling to influenza virus transmission dynamics, antigenic and genetic evolution, and molecular structure. Chapter 27, pp. 434-452 in Webster R et al., eds. *Textbook of Influenza*. Oxford: Wiley-Blackwell. 2013.
 10. Lipsitch M. (2018) Why Do Exceptionally Dangerous Gain-of-Function Experiments in Influenza? In: Yamauchi Y. (eds) *Influenza Virus. Methods in Molecular Biology*, vol 1836. Humana Press, New York, NY doi: 10.1007/978-1-4939-8678-1_29. 2018
 11. Lipsitch M, Santillana M. Enhancing Situational Awareness to Prevent Infectious Disease Outbreaks from Becoming Catastrophic. *Curr Top Microbiol Immunol*. 2019 Jul 11. doi: 10.1007/82_2019_172. [Epub ahead of print] PubMed PMID: 31292726.
- (b) Non peer-reviewed journal articles and working papers and letters
12. Levin BR, Antia R, Berliner E, Bloland P, Bonhoeffer S, Cohen M, DeRouin T, Fields PI, Jafari H, Jernigan D, Lipsitch M, McGowan JE, Mead P, Nowak M, Porco T, Sykora P, Simonsen L, Spitznagel J, Tauxe R, Tenover F. 1998. Resistance to antimicrobial chemotherapy: A prescription for research and action. *American Journal of the Medical Sciences* 315: 87-94. PMID: 9472907.
 13. Lipsitch M. Evolution in health and disease (meeting report). *Trends in Microbiology* 1997; 5: 303-4. PMID: 9263405.
 14. Lipsitch M. Fifty Years of Antimicrobials: Past Perspectives and Future Trends, ed. P.A. Hunter, G.K. Darby, and N.J. Russell. (review) *Quarterly Review of Biology* 1997; 71: 570-1.
 15. Lipsitch M. Modelling the AIDS Epidemic: Planning, Policy and Prevention, ed. E.H. Kaplan and Margaret L. Brandeau. (review) *Quarterly Review of Biology* 1995; 70: 123.
 16. Lipsitch M. Microbiology: Bacterial Population Genetics and Disease. *Science* 2001; 292:59-60. PMID: 11294216.
 17. Lipsitch M, Singer RS, Levin BR. Antibiotics in Agriculture: When is it Time to Close the Barn Door? *Proceedings of the National Academy of Sciences, USA* 2002; 99:5752-4. PMID: 11983874. PMCID: PMC122845.
 18. Lipsitch M. Antibiotic Resistance – the Interplay between Antibiotic Use in Animals and Human Beings (contribution to a “Forum”). *Lancet Infectious Diseases* 2003; 3: 51. PMID: 12505035.

19. Lipsitch M, Bergstrom CT. Real-time tracking of control measures for emerging infections [commentary]. *American Journal of Epidemiology* 2004;160(6):517-9. PMID: 15353410.
20. Halloran ME, Lipsitch M. Infectious disease modeling contributions to the American Journal of Epidemiology [commentary]. *American Journal of Epidemiology* 2005; 161: 997-8.
21. Lipsitch M. Pandemic flu: We are not prepared. *Medscape General Medicine*. 2005; 7(2). <http://www.medscape.com/viewarticle/502709>. PMID: 16369434. PMCID: PMC1681602.
22. Lipsitch M. Ethics of rationing the flu vaccine. [letter]. *Science*. 2005; 307(5706):41. PMID: 15637252.
23. Lipsitch M. How Do Antimicrobial Agents Lead to Resistance in Pathogens Causing Acute Respiratory Tract Infections? *Infectious Diseases in Clinical Practice* 2006; 14 Supp 4: S6-S10.
24. Goossens H, Lipsitch M. Global Burden of Antimicrobial Resistance. *Johns Hopkins Advanced Studies in Medicine* 2006; 6(7C): S644-S651.
25. Regev-Yochay G, Bogaert D, Malley R, Hermans PW, Veenhoven RH, Sanders EA, Lipsitch M, Rubinstein E. Does pneumococcal conjugate vaccine influence *Staphylococcus aureus* carriage in children? [letter] *Clin Infect Dis*. 2008 Jul 15;47(2):289-91; author reply 291-2. PMID: 18564933.
26. Cohen T, Lipsitch M. Too little of a good thing: a paradox of moderate infection control. *Epidemiology*. 2008 Jul;19(4):588-9. PMID: 18552592. PMCID: PMC2652751.
27. Klugman KP, Astley CM, Lipsitch M. Time from illness onset to death, 1918 influenza and pneumococcal pneumonia (letter). *Emerg Infect Dis* 2009; 15(2):346-7. PMID: 19193293. PMCID: PMC2657896.
28. Lipsitch M, Viboud C. Influenza seasonality: lifting the fog. *Proc Natl Acad Sci U S A*. 2009 Mar 10;106(10):3645-6. PMID: 19276125.
29. Goldstein E, Lipsitch M. Antiviral usage for H1N1 treatment: pros, cons and an argument for broader prescribing guidelines in the United States. *PLoS Curr. Influenza* 2009 Oct 29:RRN1122.2.
30. Goldstein E, Lipsitch M. H1N1 vaccination and adults with underlying health conditions in the US. *PLoS Curr. Influenza*, 2009 November: RRN1132.
31. Holmes E, Palese P, Rambaut A, Moscona A, Viboud C, Webby RJ, Riley S, Katze M, Lipsitch M, Salzberg SL, Garcia-Sastre A, Miller M, Fouchier RA, Wolf
32. YI, Lipman DJ, Graeff A, Parrish CR, Donis R. PLoS Currents: Influenza: A moderated collection for rapid and open sharing of useful new scientific data, analyses, and ideas. *PLoS Curr. Influenza* 2010 Jan 4:RRN1142.
33. Hernán MA, Lipsitch M. Reply to cochrane neuraminidase inhibitors review team. [letter] *Clin Infect Dis*. 2011 Dec;53(12):1303-4. PMID: 22080125.
34. Lipsitch M, Hernán MA. Oseltamivir Effect on Antibiotic-Treated Lower Respiratory Tract Complications in Virologically Positive Randomized Trial Participants. [letter] *Clin Infect Dis*. 2013 Aug 9. [Epub ahead of print] PMID: 23883518. PMCID: PMC3792722.
35. Lipsitch M. Avian influenza: Ferret H7N9 flu model questioned. *Nature*. 2013 Sep 5;501(7465):33. doi: 10.1038/501033e. PMID: 24005404.
36. Leung N, Worby C, Hanage WP, Lipsitch M, Cowling BJ. Probable person to person transmission of novel avian influenza A (H7N9) virus in Eastern China, 2013: epidemiological investigation. *BMJ*. 2013 Aug 6;347:f4752. doi: 10.1136/bmj.f4752. <http://www.bmj.com/content/347/bmj.f4752?tab=responses>.

37. Fisman DN, Leung GM, Lipsitch M. Nuanced risk assessment for emerging infectious diseases. *Lancet*. 2014 Jan 18;383(9913):189-90. doi:10.1016/S0140-6736(13)62123-6. PMID: 24439726.
38. Lipsitch M. Can limited scientific value of potential pandemic pathogen experiments justify the risks? *MBio*. 2014 Oct 14;5(5):e02008-14. doi: 10.1128/mBio.02008-14. PMID: 25316701; PMCID: PMC4205796.
39. Lipsitch M, Inglesby TV. Moratorium on research intended to create novel potential pandemic pathogens. *MBio*. 2014 Dec 12;5(6). pii: e02366-14. doi: 10.1128/mBio.02366-14. PMID: 25505122; PMCID: PMC4271556.
40. Duprex WP, Fouchier RA, Imperiale MJ, Lipsitch M, Relman DA. Gain-of-function experiments: time for a real debate. *Nat Rev Microbiol*. 2015 Jan;13(1):58-64. doi: 10.1038/nrmicro3405. Epub 2014 Dec 8. PMID: 25482289.
41. Lipsitch M, Esvelt K, Inglesby T. Calls for Caution in Genome Engineering Should Be a Model for Similar Dialogue on Pandemic Pathogen Research. *Ann Intern Med*. 2015 Sep 8. doi: 10.7326/M15-1048. [Epub ahead of print] PMID: 26344802.
42. Frank GM, Adalja A, Barbour A, Casadevall A, Dormitzer PR, Duchin J, Hayden FG, Hirsch MS, Hynes NA, Lipsitch M, Pavia AT, Relman DA. IDSA and Gain-of-Function Experiments with Pathogens having Pandemic Potential1. *J Infect Dis*. 2015 Sep 27. pii: jiv474. [Epub ahead of print] PMID: 26416656.
43. Lipsitch M, Relman DA, Inglesby TV. Six policy options for conducting gain-of-function research [commentary]. *CIDRAP*. 2016 Mar 8. <http://www.cidrap.umn.edu/news-perspective/2016/03/commentary-six-policy-options-conducting-gain-function-research>
44. Lipsitch M. Comment on Gain-of-Function Research and the Relevance to Clinical Practice. *J Infect Dis*. 2016 Aug 8. <http://jid.oxfordjournals.org/content/early/2016/08/05/infdis.jiw348.full.pdf?keytype=ref&ijkey=3UezwCqIHkSHZIP>
45. Lipsitch M. Zika vaccine trials: There are new and familiar challenges in the race for timely and effective vaccines. *Science*. 2016 Sep 9;353(6304):1094-5. doi: 10.1126/science.aai8126. <http://science.sciencemag.org/content/353/6304/1094.full>
46. Lipsitch M. If a Global Catastrophic Biological Risk Materializes, at What Stage Will We Recognize It? *Health Secur*. 2017 Jul/Aug;15(4):331-4. doi: 10.1089/hs.2017.0037. Epub 2017 Jul 26. doi:10.1089/hs.2017.0037.[Epub ahead of print] PMID: 28745911.
47. MacFadden DR, Lipsitch M, Olesen SW, Grad Yonatan. Multidrug-resistant *Neisseria Gonorrhoeae*: Implications for Future Treatment Strategies. *The Lancet Infect Dis*. June 2018, Vol 18, No. 6, p.599. Letter to The Editor. [https://www.thelancet.com/journals/laninf/article/PIIS1473-3099\(18\)30274-3/fulltext](https://www.thelancet.com/journals/laninf/article/PIIS1473-3099(18)30274-3/fulltext)
48. Atkins KE, Lipsitch M. Can antibiotic resistance be reduced by vaccinating against respiratory disease? *Lancet Respir Med*. 2018 Jul 31. pii: S2213-2600(18)30328-X. doi: 10.1016/S2213-2600(18)30328-X. PubMed PMID: 30076121.
49. Lipsitch M. Challenges of vaccine effectiveness and waning studies. *Clin Infect Dis*. 2018 Sep 10. doi: 10.1093/cid/ciy773. PubMed PMID: 30204853.
50. Lewnard JA, Tedijanto C, Cowling BJ, Lipsitch M. Accounting for unobserved and differential susceptible time at risk in retrospective studies: response to Dean (2019). *Am J Epidemiol*. 2019 Jan 25. doi: 10.1093/aje/kwz018. [Epub ahead of print] PMID: 30689694.

51. Lipsitch M, Shaman J. Comment on: 'Antibiotic footprint' as a communication tool to aid reduction of antibiotic consumption, *J Antimicrob Chemother.* 2019 Jul 17. pii: dkz320. doi: 10.1093/jac/dkz320. [Epub ahead of print] PubMed PMID: 31314102.
52. Inglesby T and Lipsitch M. 2020. Proposed Changes to US Policy on Potential Pandemic Pathogen Oversight and Implementation. *mSphere* 5:e00990-19.
53. Buckee CO, Balsari S, Chan J, Crosas M, Dominici F, Gasser U, Grad YH, Grenfell B, Halloran ME, Kraemer MUG, Lipsitch M, Metcalf CJE, Meyers LA, Perkins TA, Santillana M, Scarpino SV, Viboud C, Wesolowski A, Schroeder A. Aggregated mobility data could help fight COVID-19. *Science.* 2020 Apr 10;368(6487):145-146. doi: 10.1126/science.abb8021. Epub 2020 Mar 23. PubMed PMID: 32205458.
54. Lipsitch M. Estimating case fatality rates of COVID-19. *Lancet Infect Dis.* 2020 Mar 31. pii: S1473-3099(20)30245-0. doi: 10.1016/S1473-3099(20)30245-0. [Epub ahead of print] PubMed PMID: 32243813.
55. Swerdlow DL, Finelli L, Lipsitch M. Epidemiology of Covid-19. Reply. *N Engl J Med.* 2020 Mar 27;382. pii: 10.1056/NEJMc2005157#sa2. doi: 10.1056/NEJMc2005157. [Epub ahead of print] PubMed PMID: 32220201.
56. Lipsitch M, Kahn R, Mina MJ. Antibody testing will enhance the power and accuracy of COVID-19-prevention trials. *Nat Med.* 2020 Apr 27. doi:10.1038/s41591-020-0887-3. [Epub ahead of print] PubMed PMID: 32341581.
57. Eyal N, Lipsitch M, Smith PG. Response to Cioffi. *J Infect Dis.* 2020 Apr 29. pii: jiaa217. doi: 10.1093/infdis/jiaa217. [Epub ahead of print] PubMed PMID: 32348499.
58. Thaler DS, Lipsitch M. Coronavirus: sampling now for future analysis. *Nature.* 2020 Apr;580(7805):590. doi: 10.1038/d41586-020-01267-y. PubMed PMID: 32346142.
59. Lipsitch M, Perlman S, Waldor MK. Testing COVID-19 Therapies to prevent progression of mild disease. *Lancet Infect Dis.* 2020 May 6. [https://doi.org/10.1016/S1473-3099\(20\)30372-8](https://doi.org/10.1016/S1473-3099(20)30372-8)

(c) Popular Articles

1. Lipsitch, M. Genetic Tug-of-War May Explain Many of the Troubles of Pregnancy. *New York Times.* 1993 Jul 20. Sect. B:6.
2. Lipsitch, M. Fears Growing over Bacteria Resistant to Antibiotics. *New York Times.* 1995 Sept 12. Sect. C:1.
3. Popular articles on the evolution-creationism debate in School Board News, The Forward, and The Emory Report.
4. Lipsitch, M. Prepare Now for the Return of SARS. Project Syndicate, syndicated to Straits Times, Taiwan Times, Daily Times (Pakistan). 2003 Jul 13. Available from: <https://www.project-syndicate.org/commentary/prepare-now-for-the-return-of-sars?barrier=accesspaylog>
5. Lipsitch M and Bloom BR. Avian flu: Preparing for a Pandemic. *Harvard Public Health Review.* 2006. Winter 2006. Available from: http://www.hsph.harvard.edu/review/rvw/winter06_dean.html
6. Lipsitch M. The Risk to Academic Freedom That Lurks in Corporate Consulting Contracts. *The Chronicle of Higher Education.* 2010 Jun 27.

Lipsitch, Marc

7. Lipsitch M, Pavia A, Uyeki T, Beigi R, Bernstein H, Bradley J. Data on Flu Treatment [Letter]. New York Times. 2012 April 19. Available from: <http://www.nytimes.com/2012/04/20/opinion/data-on-flu-treatment.html>
8. Lipsitch M. Exceptional Risks, Exceptional Precautions. The European 2013 Apr 23. Available from: <http://www.theeuropean-magazine.com/marc-lipsitch--2/6691-the-risk-from-super-viruses>
9. Lipsitch M. Keine Experimente! Das Züchten neuer Krankheitserreger ist sinnlos und gefährlich. Es muss aufhören. IPG. 2014 June 10. Available from: <http://www.ipg-journal.de/kommentar/artikel/keine-experimente-458/>.
10. Lipsitch M. Anthrax? That's Not the Real Worry. New York Times. 2014 Jun 30. Available from: <http://www.nytimes.com/2014/06/30/opinion/anthrax-thats-not-the-real-worry.html?ref=opinion>
11. Lipsitch M. Make the Pause on Risky Pathogen Research Permanent. Scientific American. 2015 Jan 2:312(2).
12. Lipsitch M and Relman DA. New Game, New Rules: Limiting the Risks of Biological Engineering. Foreign Affairs. 2015 Aug 31. Available from: <https://www.foreignaffairs.com/articles/2015-08-31/new-game-new-rules>
13. Lipsitch M. Keeping biological research safe. The Hill. 2016 Aug 18. Available from: <https://thehill.com/blogs/congress-blog/healthcare/291831-keeping-biological-research-safe>
14. Hanage W and Lipsitch M. How to Report on the COVID-19 Outbreak Responsibly. Scientific American. 2020 Feb 23. Available from: <https://blogs.scientificamerican.com/observations/how-to-report-on-the-covid-19-outbreak-responsibly/>
15. Lipsitch M and Inglesby T. The U.S. is funding dangerous experiments it doesn't want you to know about. Washington Post. 2019 Feb 27. Available from: https://www.washingtonpost.com/opinions/the-us-is-funding-dangerous-experiments-it-doesnt-want-you-to-know-about/2019/02/27/5f60e934-38ae-11e9-a2cd-307b06d0257b_story.html
16. Lipsitch M. Why it's so hard to pin down the risk from coronavirus. Washington Post. 2020 Mar 6. Available from: <https://www.washingtonpost.com/opinions/2020/03/06/why-its-so-hard-pin-down-risk-dying-coronavirus/>
17. Gottlieb S and Lipsitch M. Take smart steps to slow spread of the coronavirus. USA Today. 2020 Mar 9. Available from: <https://www.usatoday.com/story/opinion/2020/03/06/former-fda-chief-gottlieb-actions-needed-fight-coronavirus-covid-19-column/4967137002/>
18. Lipsitch M. The interventions we must take to control the coronavirus. Boston Globe. 2020 Mar 11. Available from: <https://www.bostonglobe.com/2020/03/11/opinion/interventions-we-must-take-control-coronavirus/>
19. Lipsitch M and Allen J. Coronavirus reality check: 7 myths about social distancing, busted. USA Today. 2020 Mar 16. Available from: <https://www.usatoday.com/story/opinion/2020/03/16/coronavirus-social-distancing-myths-realities-column/5053696002/>
20. Lipsitch M. We know enough now to act decisively against Covid-19. Social distancing is a good place to start. STAT News. 2020 Mar 18. Available from: <https://www.statnews.com/2020/03/18/we-know-enough-now-to-act-decisively-against-covid-19/>
21. Danzig R and Lipsitch M. Prepare Now for the Long War Against Covid-19. Bloomberg. 2020 Mar 20. Available from:

- <https://www.bloomberg.com/opinion/articles/2020-03-20/prepare-now-for-the-long-war-against-coronavirus>
22. Lipsitch M. Far more people in the U.S. have the coronavirus than you think. Washington Post. 2020 Mar 23. Available from: <https://www.washingtonpost.com/outlook/2020/03/23/coronavirus-count-confirmed-testing/>
 23. Allen J and Lipsitch M. 6 things to know if you're living with someone who has coronavirus, or think you might be. USA Today. 2020 Mar 24. Available from: <https://www.usatoday.com/story/opinion/2020/03/24/coronavirus-testing-shortage-take-precautions-just-in-case-column/2899989001/>
 24. Lipsitch M and Grad Y. Navigating the Covid-19 pandemic: We're just clambering into a life raft. Dry land is far away. STAT News. 2020 Apr 1. Available from: <https://www.statnews.com/2020/04/01/navigating-covid-19-pandemic/>
 25. Lipsitch M. Who Is Immune to the Coronavirus? New York Times. 2020 Apr 13. Available from: <https://www.nytimes.com/2020/04/13/opinion/coronavirus-immunity.html>
 26. Allen J, Friedman W, Lipsitch M. Keep parks open. The benefits of fresh air outweigh the risks of infection. Washington Post. 2020 Apr 13. Available from: <https://www.washingtonpost.com/outlook/2020/04/13/keep-parks-open-benefits-fresh-air-outweigh-risks-infection/>
 27. Lipsitch M. 'Serology' is the new coronavirus buzzword. Here's why it matters. Washington Post. 2020 May 4. Available from: <https://www.washingtonpost.com/opinions/2020/05/04/serology-is-new-coronavirus-buzzword-heres-why-it-matters/>
 28. Lipsitch M. Good Science is Good Science. Boston Review. 2020 May 12. Available from: <http://www.bostonreview.net/science-nature/marc-lipsitch-good-science-good-science>
 29. Malley R and Lipsitch M. Treating Mild Coronavirus Cases Could Help Save Everyone. New York Times. 2020 May 22. Available from: <https://www.nytimes.com/2020/05/22/opinion/coronavirus-treatment-mild-symptoms.html>
 30. Eyal N, Lipsitch M, Angell M. The True Cost of Vaccine Studies. New York Review of Books. 2020 Aug 20. Available from: <https://www.nybooks.com/articles/2020/08/20/true-cost-of-vaccine-studies/>
 31. Grad Y and Lipsitch M. How to fix public health weaknesses before the next pandemic hits. Washington Post. 2020 Sept 24. Available from: <https://www.washingtonpost.com/opinions/2020/09/24/how-fix-public-health-weaknesses-before-next-pandemic-hits/>
 32. Lipsitch M. Americans, we can fight COVID-19 and save lives now. Wear a mask! USA Today. 2020 Oct 8. Available from: <https://www.usatoday.com/story/opinion/2020/10/08/wear-mask-fight-covid-19-and-save-lives-now-medical-experts-column/5907452002/>

(d) Coauthored (group-authored) reports

33. President's Council of Advisors on Science and Technology, H1N1 Working Group [member author]. Report to the President on US Preparations for 2009-H1N1 Influenza. 2010 Aug 9. Available from:

- https://www.globalsecurity.org/security/library/report/2009/pcast_h1n1-report_090807.htm
34. Wellcome Trust/CIDRAP Team B [member author]. Recommendations for Accelerating the Development of Ebola Vaccines, Report & Analysis. 2015 Feb. Available from:
https://www.cidrap.umn.edu/sites/default/files/public/downloads/ebola_virus_team_b_report-final-021615.pdf
 35. Wellcome Trust/CIDRAP Team B [member author]. Plotting the Course of Ebola Vaccines: Challenges and Unanswered Questions. 2016 Mar. Available from:
https://www.cidrap.umn.edu/sites/default/files/public/downloads/ebola_team_b_report_2-033116-final.pdf
 36. Wellcome Trust/CIDRAP Team B [member author]. Completing the Development of Ebola Vaccines: Current Status, Remaining Challenges and Recommendations. 2017 Jan. Available from:
https://www.cidrap.umn.edu/sites/default/files/public/downloads/ebola_team_b_report_3-011717-final_0.pdf
 37. Moore K, Lipsitch M, Barry JM, Osterholm MT. COVID-19: The CIDRAP Viewpoint. CIDRAP, University of Minnesota. 2020 Apr 30. Available from:
https://www.cidrap.umn.edu/sites/default/files/public/downloads/cidrap-covid19-viewpoint-part1_0.pdf

Exhibit 4

May 29, 2020



Federal Aviation Administration

Information for Airport Sponsors Considering COVID-19 Restrictions or Accommodations

PURPOSE

This document addresses common issues that have arisen or may arise for airport sponsors during the response to the COVID-19 public health emergency. The Federal Aviation Administration (FAA) Office of Airports will evaluate specific requests regarding restrictions or accommodations on a case-by-case basis. The FAA retains maximum flexibility to consider unique circumstances during this public health emergency.

The FAA separately has published frequently asked questions (FAQs) related to the approximately \$10 billion in grants for airports under the Coronavirus Aid, Relief, and Economic Security (CARES) Act. Those FAQs are available at www.faa.gov/airports.

BACKGROUND

The FAA has been receiving inquiries from airport operators about their authority to implement a range of restrictions, changes in operations, terminal service consolidations, and other responses to the COVID-19 public health emergency. Many of these inquiries reflect interest in facilitating social distancing or adapting to a reduced level of activity at the airport.

The FAA's primary concern is that federally obligated airports remain safe and open to the traveling public and aircraft. Particularly during this public health emergency, airports play an essential role in transporting medical and emergency equipment and personnel. The FAA continues to expect all airports to operate safely and stay open.

APPLICABILITY

The guidance here is not legally binding in its own right and will not be relied upon by the FAA as a separate basis for affirmative enforcement action or other administrative penalty. Conformity with this guidance, as distinct from existing statutes, regulations, and grant assurances, is voluntary only, and nonconformity will not affect existing rights and obligations.

ISSUES

Closing airports: All proposed closing of airport access (i.e., passenger and aircraft access) must be approved in advance by the FAA. As noted in [Compliance Guidance Letter, 2020-01](#), in general, the FAA does not permit temporary closure or restriction of federally obligated airports for non-aeronautical purposes. An airport sponsor must obtain FAA approval to allow airport closure for a non-aeronautical purpose. (Grant Assurance 19 and 49 U.S.C. § 47107(a)(8)). Grant Assurance 19

May 29, 2020

further requires that airport sponsors will not cause or permit any activity or action on the airport that would interfere with its use for airport purposes. This includes all airport structures and operational areas. If a proposed action suspends or closes an international Port of Entry, then the sponsor may also need approval from U.S. Customs and Border Protection (CBP).

Prohibiting certain flights (e.g., certain locations, types of aircraft, and types of operations): As is normally the case, actions such as these may violate Federal law and the airport's grant assurances, unless approved in advance by the FAA (and, in some cases, the Office of the Secretary of Transportation (OST) as well). To seek such approval, the airport sponsor should contact the applicable [FAA Airports District Office](#) to discuss the matter.

Requiring flights to land at certain airports for screening: All such requests would ordinarily require prior FAA approval under Grant Assurances 19 and 22 and related statutes. Usually, these restrictions would likely constitute an unreasonable restriction on access; however, FAA has discretion to consider such requests and recognizes the exceptional situation presented by this public health emergency. Depending on the circumstances, such requests might be deemed as reasonable restrictions on access. However, even where FAA is amenable to such a temporary condition, the airport will need to coordinate with OST with regard to requiring route changes, and with CBP if the action appears to suspend or close an international Port of Entry.

Closing of sections of the airfield to allow for aircraft parking: Airports should avoid overflow parking of aircraft on runways except as a last resort. If overflow parking of aircraft is needed, airports should first consider using gates, aprons, and non-movement areas. Airports should also consider suggesting that aircraft owners contact other nearby airports where there may be additional aircraft parking capacity. Based on the location(s) selected, the sponsor must be able to respond with aircraft rescue and firefighting (ARFF) capability and provide required notice. In all cases, operators of airports in the National Plan of Integrated Airport Systems should work with local air traffic facilities (if present) to develop a safe and reasonable parking plan and share that information with their servicing FAA Airports District Office, local FAA Air Traffic Manager, and FAA's Flight Standards Service. For part 139 certificated airports, see [Cert Alert 20-02 – Temporary Parking of Overflow Aircraft](#) (updated March 24, 2020).

Closing restaurants or other retail activities in the terminal: The closing of restaurants, retail stores, or other non-aeronautical functions in a terminal is not likely to violate FAA grant assurances if driven by public health measures or reduced clientele, and especially if restrictions are applicable to all business entities within the jurisdiction. However, airports should coordinate with the FAA Office of Civil Rights with regard to Airport Concession Disadvantaged Business Enterprise regulations.

Closing gates or sections of terminals: In coordination with airport sponsors, airlines, the Transportation Security Administration (TSA), and other entities, closing gates or sections of terminals is likely to be acceptable if the closure is executed in response to reduced passenger volumes and operations, is not discriminatory, and does not provide an unfair competitive advantage to one operator. For example, TSA has reduced lanes or consolidated passenger screening checkpoint operations in numerous airports in response to the reduction in originating passenger volume.

May 29, 2020

Allowing terminals to be used for sheltering of people: This is likely to be acceptable if it does not interfere with airport access or impact security for the traveling public and aircraft operations.

Screening or quarantining passengers boarding or exiting planes: State, local, or territorial public health officials may want to screen or quarantine passengers. In most cases, this is likely to be acceptable as long as passengers are not being categorically refused access to air transportation (e.g., through unapproved blanket closures). Airlines may refuse transportation to a passenger because of a communicable disease if the passenger's condition poses a direct threat to the health or safety of others. Care must also be taken in coordinating with airport sponsors, airlines, TSA, airport law enforcement, and other entities on when, where, and how your government conducts this screening and quarantining, with a goal of minimizing burden and maximizing flexibility for operations. Effort also should be made to minimize undesirable queueing or the formation of large groups of passengers.

Rent abatement / minimum annual guarantee: A decision to abate rent (including "minimum annual guarantees" and also encompassing fees) is a local decision. Rent abatement should be tied to the changed circumstances caused by the public health emergency, and done in accordance with Grant Assurances 22 and 24, as well as related statutes. Where abatement results in shifting costs between various classes of airport tenants and users, the airport sponsor is encouraged to consult with all affected parties and implement a consensus approach if possible.

If a sponsor (or airport tenant, whether aeronautical or non-aeronautical) desires to renegotiate rent, a reasonable basis for such an action might be established if the underlying basis for such rent has temporarily declined or materially altered due to COVID-19. In such circumstances, the offer of accommodation in the form of rent abatement is not barred by the grant assurances as long as it is reasonable under the circumstances and reflects the decline in fair market value, loss of services, and/or changes to volume of traffic and economy of collection.

Sponsors considering such relief are encouraged to consider the business situation of the tenant; the changed circumstances created by the public health emergency; the desirability of having solvent tenants that can resume normal operations when the emergency ends; the availability of other governmental or insurance relief that such entities have or may receive; an appropriate term for such relief; and possible subsequent conditions that, if triggered, would end the abatement. Such a condition could be the receipt of other governmental forms of relief; insurance recovery, if any; or an end to the emergency.

As noted above, where sponsors have residual lease arrangements with aeronautical users, the reduction of rent for certain non-aeronautical entities may shift costs to the aeronautical users such as airlines. Achieving the appropriate balance between these users is a local responsibility that should be managed in consultation with all affected parties. If rent abatement to non-aeronautical users results in an increase to aeronautical rates, that is not necessarily an impediment from a grant assurance perspective, but the aeronautical rates must remain reasonable. For any actions that reallocate costs, FAA encourages sponsors to carefully balance and consider the equities between all airport users. Additionally, the sponsor is encouraged to consult with all affected parties before making its decision and reach a consensus where possible.

May 29, 2020

Apart from any Federal obligations, the FAA also recommends that airport sponsors consult their lease agreements to understand their discretion to act, particularly in a residual methodology context. Airport sponsors should also examine any bond covenants to identify any potential restrictions that may exist.

Deferral of rental payments or other fees: In cases where bond restrictions or other conditions may prevent airports from offering rent abatements, the deferrals of rents and/or fees may be possible. The terms and interest rates applied should be reasonable and applied fairly to similarly situated businesses. Deferral of rental payments and or fees, if adequately justified, is not likely to violate FAA's grant assurances. A primary goal of the statutory sustainability principle is to keep the airport solvent to ensure that the airport can remain open and operate safely. If a deferral exceeds an annual reporting period, interest should be charged based on Treasury note interest rates beginning the date of the deferral and reported on FAA Form 127. The deferred rent amount should be reported in the fiscal year when the rent would have been due but for the deferral. In the event that the rent payment is deferred and not abated, the deferred rent amount should be reported as unpaid invoices (accounts receivables) which would be reflected in the amount of revenue reported on the FAA Form 127. Neither airports nor the FAA have the legal authority, however, to allow air carriers to defer the remittance of collected Passenger Facility Charge (PFC) revenues.

Sponsor's request for reducing hours of operation: If contemplated, it is important that any such proposed action be part of implementing a legitimate public health initiative related to COVID-19. At a minimum, to the extent considered, such an action would require FAA to examine whether it would result in an undue hardship on emergency response or otherwise unjustly discriminate against a specific user of the airport. Finally, FAA is unlikely to approve any such reductions that would restrict either government or emergency operations.

Sheltering-in-place impacts on airport personnel: Because airports are essential in transporting emergency and medical supplies and personnel during emergencies, a critical number of airport and Federal employees should be designated as essential to ensure the continuity, safety, and security of airport operations. Also, airport law enforcement should be informed to facilitate their access to airport and airport facilities. This is particularly true for part 139 certificated airports, which require minimum personnel to meet requirements of the regulation. In addition, the Department of Homeland Security's Cybersecurity and Infrastructure Security Agency has issued [guidance](#) that specifically identifies airport operations personnel as part of the "Essential Critical Infrastructure Workforce" who should not be impeded from their efforts to keep airports safe and operational.

Recreational aeronautical restrictions: Certain States have issued COVID-19 restrictions on activities they deem "non-essential," including certain aeronautical activities such as flight schools and sky diving. With the goal of keeping airports open to ensure access for the traveling public, emergency and medical equipment and supplies, and emergency transportation, FAA does not object to *temporarily* limiting recreational aeronautical activities that are covered by such restrictions. However, the activities limited by a sponsor should be limited to those falling within the scope of a public health measure by an authority whose jurisdiction covers the airport's geographic area (e.g., a State or local government).

May 29, 2020

Prohibiting flights from “hotspot” areas: Prohibiting flights from “hotspots” or areas of high levels of contagion generally is not acceptable. However, a jurisdiction may choose to consider its authority to impose public health screening or quarantine for passengers entering the jurisdiction. The FAA has published [guidance](#) for consideration when implementing quarantine, screening, or movement restrictions that impact air transportation.

Sponsor’s Use of Airport Revenue for Public Health Activities: Federal law requires that federally obligated airports must use airport revenue for the capital or operating costs of the airport. CARES Act grants must be used in the same way as airport revenue and for costs that are directly related to the airport.

Under the extraordinary circumstances of the COVID-19 public health emergency, some activities the airport may undertake to minimize the spread of COVID-19 may be legitimate capital or operating costs of the airport. For example, in this exceptional context, the FAA considers the testing and health screening of airport employees to be a legitimate operating cost of an airport to sustain the airport’s workforce, upon which the continuity of airport operations depends. Additionally, airport operating costs may also include the costs of enhanced cleaning of the terminal and other areas of airport property to minimize transmission of COVID-19. These operating costs may include the purchase of incidentals and supplies to accomplish these purposes, such as screening and testing equipment, cloth face covers, and cleaning and disinfection products. In contrast, the use of airport employees for public health screening is generally not considered a proper use of airport revenue. Airports should properly account for and document allowable costs incurred because of the COVID-19 public health emergency. Airports with specific questions regarding allowable costs related to COVID-19 should contact their Airport District Office.

Sponsor’s Use of Airport Space for Public Health Activities: Under the extraordinary circumstances of the COVID-19 public health emergency, airports are permitted to allocate terminal or office space for testing and health screening activities and the related storage of medical equipment and supplies. In this exceptional context, it is also within an airport sponsor’s discretion to allow tenants to have additional space, beyond what their leases include, for testing and health screening and for storage of medical equipment and supplies. Because these uses support the continuity of airport operations, such accommodations can be for no cost as long as they are temporary, necessitated by the public health emergency, and offered in a way that is not unjustly discriminatory.

CONCLUSION

Airports should be cognizant of, and assume the responsibility for, the implications of their proposed actions in response to COVID-19. Considerations include, among others: (1) coordination with the FAA, (2) coordination with other Federal, State, or local agencies as needed, including airport law enforcement or local law enforcement entities serving the airport; (3) understanding of applicable Federal obligations, (4) impacts on aeronautical use and airport infrastructure; (5) impact on the safe and efficient functioning of air traffic and the National Airspace System; (6) communications and notice requirements; (7) evolving safety and security requirements; (8) the need to document actions; (9) plans for following up on or amending actions as the situation evolves; and (10) the impact to emergency services that rely on air transportation.

Exhibit 5



MANDATORY DIRECTIVE:



Travel

Issued: November 28, 2020

sccgov.org/coronavirus

County of Santa Clara
Public Health Department

Health Officer
976 Lenzen Avenue, 2nd Floor
San José, CA 95126
408.792.5040



MANDATORY DIRECTIVE ON TRAVEL

Please confirm compliance with the State Order. Where there is a difference between the local County Order and the State Order, the more restrictive order must be followed. The State also has specific guidance for certain activities that must be followed in addition to this mandatory directive.

Information on the State's Order and State guidance is available at covid19.ca.gov.

Issued: November 28, 2020
Revised and Effective: January 25, 2021
Effective Upon Release

In light of significant increases in COVID-19 cases and associated hospitalizations across the United States, the State of California, and within Santa Clara County, this Mandatory Directive on Travel is in effect until it is rescinded or modified.

This Directive establishes the County Health Officer's rules for quarantine after travel. The risk of COVID-19 transmission increases as people travel in and out of Santa Clara County and have contact with persons from other households and other communities, especially through travel to regions with significant COVID-19 transmission. To reduce this risk, the County Health Officer has established this mandatory directive related to travel. This Directive applies to all travel into Santa Clara County, whether by residents or non-residents.

This Directive is mandatory, and failure to follow it is a violation of the Health Officer's Order issued on October 5, 2020 ("Order").

Travel Is Discouraged

1. *The County Health Officer discourages travel, especially for non-essential purposes.*
 - a. Because travel involves mixing between regions and households, and because so many areas of the State and United States are also currently experiencing significant surges in COVID-19 cases, travel is discouraged and should be minimized.

- b. In particular, non-essential travel (i.e., travel for leisure or for non-essential business) is strongly discouraged and should be postponed until after the current surge in COVID-19 cases and hospitalizations subsides.

Mandatory Quarantine after Long-Distance Travel into Santa Clara County

2. Quarantine Requirements

- a. Except as otherwise provided in this Directive, all persons traveling into Santa Clara County, whether by air, car, train, or any other means, directly or indirectly from a point of origin greater than 150 miles from the county's borders must quarantine for at least 10 days after arrival.
- b. For the purposes of this Directive, "quarantine" means staying at home or another place of temporary shelter without contact with any persons other than members of one's own immediate traveling party or one's household.
- c. Information, resources, and guidance on quarantine, including recommendations on when to get tested and what to do in the case of a positive test result are available at www.sccstayhome.org.

3. Exemptions from Mandatory Quarantine.

- a. [Licensed healthcare professionals](#), as defined by the Order, and all persons working at acute care hospitals, do not need to quarantine following arrival.
- b. Persons traveling solely for the purpose of performing an essential governmental function, as defined by the governmental entity responsible for that function, do not need to quarantine following that travel.
- c. The following persons are required to quarantine, but may leave their home or place of quarantine solely for the activities specified in this section:
 - i. Persons who perform essential governmental functions, as defined by the governmental entity responsible for those functions, whose purpose for travel does not fall within section 3(b), but only to the extent that the governmental entity determines that it would otherwise lack sufficient staffing to fulfill that essential function.
 - ii. Persons traveling solely for the purpose of performing [essential critical infrastructure work](#), as defined by the State Public Health Officer, but

only to the extent that the employer determines that it would otherwise lack sufficient staffing to perform such work.

- iii. Persons traveling solely for the purpose of work or participation in collegiate or professional athletic activities, provided they are in compliance with all applicable directives.
- d. Persons solely transiting through Santa Clara County and not staying overnight are not required to quarantine.
- e. Persons traveling to Santa Clara County to obtain services from a Healthcare Facility, as defined in the Order, are required to quarantine upon arrival, but may leave their household or place of quarantine to obtain those services.
- f. Persons who are otherwise required to quarantine pursuant to this Directive may leave their household or place of quarantine to the extent necessary to comply with a court order or make an appearance in a court of law or administrative proceeding.

Notification Requirement for Transit Facilities

4. Notification Requirements

- a. All transit facilities, including but not limited to airports, train stations, bus stations, and other facilities where persons may be regularly traveling into Santa Clara County must ensure a copy of this Directive is provided to each passenger upon arrival from a point of origin more than 150 miles from the County borders.
- b. All transit facilities must prominently post notices in such a manner that all persons transiting through such facilities will become aware of the requirements of this Directive. Notices are available to print and post [here](#).

Stay Informed

For answers to frequently asked questions about capacity limitations and other topics, please see the [FAQ page](#). **Please note that this Directive may be updated.** For up-to-date information on the Health Officer Order, visit the County Public Health Department's website at www.sccgov.org/coronavirus.