Project 3 - How Infective Does a Disease Need to be for Masks to be Ineffective?

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Question

Our motivating question is how infective does a disease have to be for masks to no longer be effective? This is a design question because we are designing a model with the end goal of having masks not be effective in mind. We decided to choose this question because it would be useful to know when masks would no longer be effective against a disease and other preventive measures might be required. The people who would benefit from this answer are health officials responsible for spreading information and the general public who would want to know what is effective at preventing the spread of a disease. The background information that is necessary to understand this question and our modeling choices is the odds of getting infected when wearing different types of masks vs. no mask and how much of the population wears each type of mask or no mask.

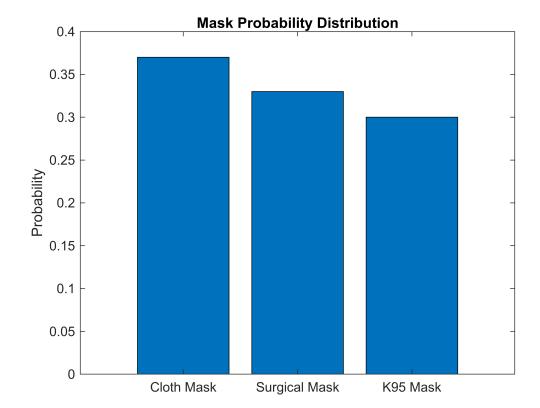
Our literature search is as follows:

	Title	Effectiveness of Face Mask or Respirator Use in Indoor Public Settings for Prevention of SARS-CoV-2 Infection. (Andrejko KL, Pry JM, Myers JF, et al, February 2022) http://dx.doi.org/10.15585/mmwr.mm 7106e1	Evaluation of different types of face masks to limit the spread of SARS-CoV-2: a modeling study (Gurbaxani, Hill, Paul, Prasad, & Slayton, May 2022) https://www.nature.com/articles/s41598-022-11934-x	Factors Associated with Cloth Face Covering Use Among Adults During the COVID-19 Pandemic (Fisher, Barile, Guerin, et al., July 2020) https://www.cdc.gov/mmwr/volumes /69/wr/mm6928e3.htm	https://www.forbes.com/sites /alisondurkee/2022/01/26/m	Introducing the Outbreak Threshold in Epidemiology (Hartfield, Alizon, June 2013) https://www.ncbi.nlm.nih.go v/pmc/articles/PMC3680036
		A study on the most effective methods for staying safe from covid-19, particularlly masks and how effective different types are at defending against the virus.	- ,	The document discusses factors associated with cloth face covering use among adults during the COVID-19 Pandemic in the United States during April and May 2020. Specifically, the document aims to examine the increase in rates of cloth face covering use during that period, the sociodemographic composition of individuals who adopted the practice, and potential barriers to consistent and correct use among those		To introduce the outbreak threshold in epidemiology by providing criteria to determine when a pathogen increases to a risky level, from which it has a large chance of dying out, to when a major outbreak is almost certain. It also discusses the implications of these results for controlling emerging and reemerging pathogens.
		well-fitting face mask or respirator in indoor public settings protects against Covid-19, a respirator offers the best protection.	materials have the potential to substantially reduce transmission of SARS-CoV-2 depending on the type and fit of the mask and the percentage adoption in the population.	Given the promising trend, public health authorities should continue to communicate the importance of cloth face covering use, reinforce positive attitudes, perceived norms, personal agency, and the physical and health benefits of obtaining and wearing face coverings consistently and correctly.	N95 and KN95 masks to be the most effective in protecting against COVID-19, they don't	To introduce the concept of outbreak threshold and how to determine the number of infected individuals needed for a disease to spread in an approximately deterministic manner. The document argues that defining this threshold provides a more accurate and precise method for detecting nascent outbreaks, which is crucial in preventing and controlling outbreaks of
	Important Info	Cloth Masks - 56% lower odds Surgical Masks - 66% lower odds Respirator (N95KN95) - 83% lower odds	• 70% of the population wears a mask	• 36% of people wore a cloth face mask	29% of people wear KN95 masks 33% of people wear surgical masks	If the number of infected people is less than 1log(the infection rate) then there will be no major outbreak of a disease

Model

For our model we used an agent-based model with the transmission rate as our deterministic quantity and mask efficacy as our randomized quantity. To represent the probability that an individual would wear a certain type of mask we used a multinomial distribution as shown below:

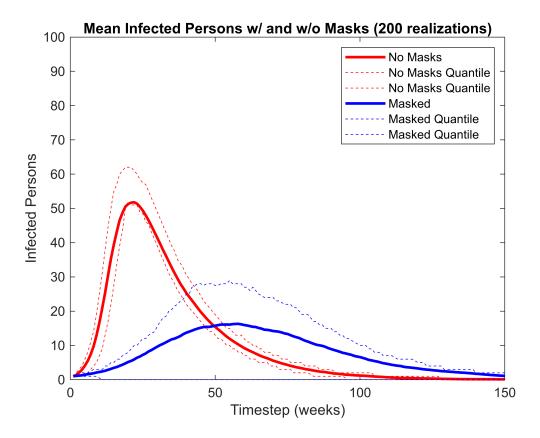
```
% mask efficacy
clothMaskEfficacy = .56;
surgicalMaskEfficacy = .66;
K95MaskEfficacy = .83;
% mask probability distribution (multinomial)
clothMaskProb = 0.37;
surgicalMaskProb = 0.33;
K95MaskProb = 0.3;
pd_mask_probabilities = makedist('Multinomial','Probabilities',[clothMaskProb
surgicalMaskProb K95MaskProb]);
% Visualize
maskOutcomes = categorical({'Cloth Mask', 'Surgical Mask', 'K95 Mask'});
maskOutcomes = reordercats(maskOutcomes,{'Cloth Mask','Surgical Mask','K95 Mask'});
x = 1:3;
pr = pd_mask_probabilities.pdf(x);
figure();
bar(maskOutcomes, pr)
ylabel('Probability')
title('Mask Probability Distribution');
```



We then used simulate_absir() to find run 200 realizations of our model and then graphed the mean and upper and lower quantiles of the max infected with or without masks.

```
mask_matrix_random = random(pd_mask_probabilities, 100, 100);
mask_efficacy = [clothMaskEfficacy surgicalMaskEfficacy K95MaskEfficacy];
mask matrix = mask efficacy(mask matrix random); % converst from 1,2, and 3 to
cloth, surgical, and K95 efficacy values (respectively)
% adjacency matrix 100 ppl
pop = 100;
no mask matrix = ~eye(pop,pop);
% Final Vulnerability Matrix
masked vulnerability matrix = no mask matrix - mask matrix;
masked_vulnerability_matrix = masked_vulnerability_matrix.*-
(eye(height(masked_vulnerability_matrix))-1);
% Simulate absir model w/ and w/o masks
    % Set the number of replications for the agent-based model
n realizations = 200;
    % Run each simulation for 150 timesteps
T simulation = 150;
    % Reserve space for simulation results
I ensemble no mask = zeros(T simulation, n realizations);
I_ensemble_mask = zeros(T_simulation, n_realizations);
    % Set the random seed.
%rng(101)
   % Set parameter values
infection rate = 0.005;
recovery_rate = 0.05;
   % Generate the sample of results
for i = 1 : n realizations
   % Run a single simulation
    Iv1 = zeros(pop, 1);
    Iv1(1) = 1;
    [Sh mask, Ih_mask, Rh_mask] = simulate_absir(masked_vulnerability_matrix, Iv1,
T simulation, infection rate, recovery rate);
    % Check that the number of people stays at 100
    assert(sum(Sh_mask(:, end)) + sum(Ih_mask(:, end)) + sum(Rh_mask(:, end)) ==
pop, 'Number of people in masked realization went is not equal to the population')
    [Sh_no_mask, Ih_no_mask, Rh_no_mask] = simulate_absir(no_mask_matrix, Iv1,
T_simulation, infection_rate, recovery_rate);
    % Check that the number of people stays at 100
    assert(sum(Sh_no_mask(:, end)) + sum(Ih_no_mask(:,end)) + sum(Rh_no_mask(:,
end)) == pop, 'Number of people in no mask realization went is not equal to the
population')
    % Summarize and store the infection history
    I_ensemble_mask(:, i) = sum(Ih_mask, 1);
```

```
I ensemble no mask(:, i) = sum(Ih no mask, 1);
end
% Check that everyone recovered
assert(sum(Ih_mask(:, end)) == 0, 'Not everyone recovered in the masked group')
assert(sum(Ih_no_mask(:,end)) == 0, 'Not everyone recovered in the no masked group')
    % Compute the mean and quantiles
    % Masked
I mean mask = mean(I ensemble mask, 2); % mean
I lo mask = quantile(I ensemble mask, 0.25, 2); % Lower (25%) quantile
I_up_mask = quantile(I_ensemble_mask, 0.75, 2); % Upper (75%) quantile
    % No mask
I mean no mask = mean(I ensemble no mask, 2);
I_lo_no_mask = quantile(I_ensemble_no_mask, 0.25, 2);
I_up_no_mask = quantile(I_ensemble_no_mask, 0.75, 2);
    % Visualize
figure()
% plot w/o masks
plot(I_mean_no_mask, 'r-', 'LineWidth', 2.0, 'DisplayName', 'No Masks'); hold on
plot(I_lo_no_mask, 'r--', 'LineWidth', .5, 'DisplayName', 'No Masks Quantile');
plot(I_up_no_mask, 'r--', 'LineWidth', .5, 'DisplayName', 'No Masks Quantile');
% plot w/ masks
plot(I_mean_mask, 'b-', 'LineWidth', 2.0, 'DisplayName', 'Masked');
plot(I_lo_mask, 'b--', 'LineWidth', .5, 'DisplayName', 'Masked Quantile');
plot(I_up_mask, 'b--', 'LineWidth', .5, 'DisplayName', 'Masked Quantile');hold off
xlabel('Timestep (weeks)')
ylabel('Infected Persons')
ylim([0 pop])
legend()
title('Mean Infected Persons w/ and w/o Masks (200 realizations)')
```



The assumptions we made when creating our model were that all cloth masks were made from the same type of cloth and that every person who was wearing a mask was wearing one properly. These are acceptable for our modeling question because most cloth masks that are known to protect against viruses have very similar effectivity rates, therefore making it redundant to include all of them, and if we assumed some people were not wearing masks properly it would be difficult to know if our results were caused by just the masks or through a mix of masks and human error. We set the parameters of our model based on a mix of our literature research and values from previous worksheets. The mask efficacies and probabilities were determined from our literature research, and the recovery and infection rates were taken from a previous worksheet. The infection rate was swept to determine the answer to our modeling question. The verification facts we came up with are making sure the total number of people stays at 100 and making sure that everyone recovers at some point.

Results

The metric we used to help answer our question was the max number of infected people. This is an appropriate metric because if the max infected for both the population wearing masks and the population not wearing masks is the same that means that masks are ineffective and will answer our modeling question. The results we generated using our model were that a disease must have an infectivity rate of 0.0722 in order for masks to be completely ineffective.

```
% Set infectivity rate values to sweep
infectivity_all = linspace(.005, .1, 100);

% Store metric (mean infection peak)
mean_infection_peak_mask = zeros(1, 100);
```

```
mean infection peak no mask = zeros(1, 100);
% Reserve space for simulation results
I_ensemble_no_mask_sweep = zeros(T_simulation, n_realizations);
I_ensemble_mask_sweep = zeros(T_simulation, n_realizations);
% run simulation for each value of infectivity rate
    for i = 1 : length(infectivity all)
        infectivity rate = infectivity all(i);
       % Run 100 simulations
       for j = 1:n realizations
            Iv1 = zeros(pop, 1);
            Iv1(1) = 1;
            [Sh mask, Ih mask, Rh mask] =
simulate_absir(masked_vulnerability_matrix, Iv1, T_simulation, infectivity_rate,
recovery_rate);
            [Sh no mask, Ih no mask, Rh no mask] = simulate absir(no mask matrix,
Iv1, T_simulation, infectivity_rate, recovery_rate);
           % Summarize and store the infection history
            I_ensemble_mask_sweep(:, j) = sum(Ih_mask, 1);
            I_ensemble_no_mask_sweep(:, j) = sum(Ih_no_mask, 1);
        end
       % Get mean
        I mean mask = mean(I ensemble mask sweep, 2); % mask mean
        I_mean_no_mask = mean(I_ensemble_no_mask_sweep, 2); % no mask mean
       % Get mean peaks and store as metrics
       mean_infection_peak_mask(i) = max(I_mean_mask); % mask
       mean_infection_peak_no_mask(i) = max(I_mean_no_mask); % no mask
    end
```

```
% Determine infectivity value that causes masks to become ineffective based on
determined threshold
percent_threshold = 0.2; % if the masked infection peak is within []% of the no
mask peak, masks are ineffective

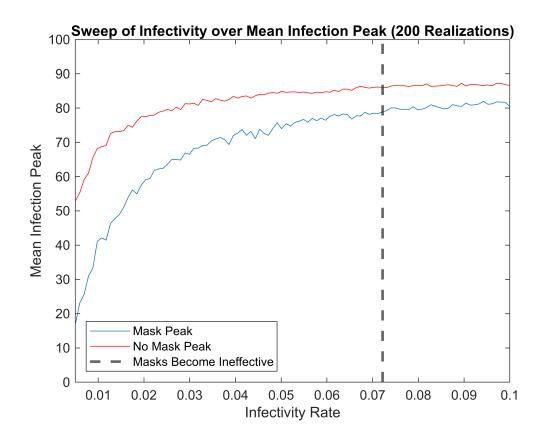
threshhold_check = zeros(1,100);
for i = 1:length(infectivity_all)
    diff_mask_no_mask = mean_infection_peak_no_mask(i) -
mean_infection_peak_mask(i);
    if (diff_mask_no_mask <= percent_threshold * mean_infection_peak_no_mask) %
checks if mask peak is within []% of no mask peak
        threshhold_check(i) = infectivity_all(i);
end
end</pre>
```

threshhold_check % stores the infectivity rate for each value where masks are ineffective (if masks were effective there will be a zero)

point_of_ineffectiveness = threshhold_check(71) % infectivity rate where masks
become ineffective

point of ineffectiveness = 0.0722

```
[mx,my] = polyxpoly(infectivity_all, mean_infection_peak_mask, infectivity_all,
mean_infection_peak_no_mask);
% Plot results
plot(infectivity all, mean infection peak mask, 'DisplayName', 'Mask Peak'); hold
on % mask
plot(mx,my,'g.', 'MarkerSize', 15, 'DisplayName', 'Masks not effective');
xline(point_of_ineffectiveness, 'LineWidth', 2, 'LineStyle', '--', 'DisplayName',
'Masks Become Ineffective') % point where masks become ineffective
plot(infectivity_all, mean_infection_peak_no_mask, 'r', 'DisplayName', 'No Mask
Peak'); hold off % no mask
xlabel('Infectivity Rate')
ylabel('Mean Infection Peak')
title('Sweep of Infectivity over Mean Infection Peak (200 Realizations)')
legend('Location','southwest')
ylim([0 100])
xlim([infectivity_all(1) infectivity_all(end)])
```



The most important trends in our parameter sweep are that both the masked and no masked graphs increase exponentially, with the difference in the mean max infected between them decreasing as they both increased.

Interpretation

Our parameter sweep answers our question of at what infectivity rate are masks no longer effective, which is 0.0722. Some limitations of our model are that it cannot account for any other circumstances which would make someone more or less immune to a disease. It also does not consider the differences in effectivity between the different cloths that can be used to make a cloth mask. If we were to continue to develop our model, we would add more mask types to take into account the different materials used to make cloth masks, we might also take into account that some people do not wear masks the right way.

Validation

According to a 2013 study analyzing what qualifies as an outbreak, in order for the spread of a disease to qualify as an outbreak, at least 57% of people must be infected. According to our results, 75% of people were infected when masks were considered ineffective, meaning that even with everyone wearing masks, a disease with an infection rate of 0.0722 would still cause an outbreak.