## Waning Modeling MEthods and Results

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## Model and parameters

We used a simple, unstructured, susceptible-infectious-removed (=SIR) model to simulate the influenza transmission in a large population. The following parameters were chosen:

| Parameter                    | Symbol                                    | Value              |
|------------------------------|-------------------------------------------|--------------------|
| Total Pop.                   | N                                         | 2.0e + 06          |
| Beginning vaccination uptake | $p_v$                                     | 0.47               |
| vaccination uptake rate      | $\nu$                                     | 0                  |
| Removal rate                 | $\gamma$                                  | 0.25               |
| Basic reprod. No.            | $R_0$                                     | 1.4                |
| Transmission coeff., unvacc. | $\beta = \frac{R_0 \gamma}{N}$            | 1.4                |
| Transmission coeff., vacc.   | $\beta = \frac{R_0 \ \gamma (1-\phi)}{N}$ | 1.4                |
| VE                           | $\phi$                                    | 0.2, 0.3, 0.4, 0.5 |
| Pre-esisting immunity        | $\epsilon$                                | 0                  |

The following initial values were used:

| Parameter                | Symbol                                       | Value                  |
|--------------------------|----------------------------------------------|------------------------|
| No. susceptible, vacc.   | $x_v$                                        | 9.4e + 05              |
| No. susceptible, unvacc. | $x_{nv}$                                     | 1.1e + 06              |
| No. infectious, vacc.    | $y_v = \frac{p_v \phi}{p_v \phi + 1 - \phi}$ | 0.15, 0.21, 0.26, 0.31 |
| No. infectious, unvacc.  | $y_{nv}$                                     | 0.69                   |
| No. removed, vacc.       | $z_v$                                        | 0                      |
| No. removed, unvacc.     | $z_{nv}$                                     | 0                      |

The model used is given by the following system or differential equations:

$$\frac{x_v}{dt} = -\beta (1 - \phi) x_v (y_v + y_{nv}) + \nu x_{nv}$$

$$\frac{x_{nv}}{dt} = -\beta x_{nv} (y_v + y_{nv}) - \nu x_{nv}$$

$$\frac{y_v}{dt} = \beta (1 - \phi) x_v (y_v + y_{nv}) - y_v \gamma$$

$$\frac{y_{nv}}{dt} = \beta x_{nv} (y_v + y_{nv}) - y_{nv} \gamma$$

$$\frac{z_v}{dt} = y_v \gamma$$

$$\frac{z_{nv}}{dt} = y_{nv} \gamma$$

The system is numerically solved using the ode function from the deSolve R package. Test-negative design (TND) studies were simulated by keeping track of the incidence of vaccinated and unvaccinated "cases", i.e. new *infecteds*. The observed VE was calculated based on the ratio of the vaccination odds in the cases to the vaccination odds in the population.

## Results

The trajectories are only shown for the periods of time when there was substantial transmission (more than 10 infectious) and aligned at their "start times".

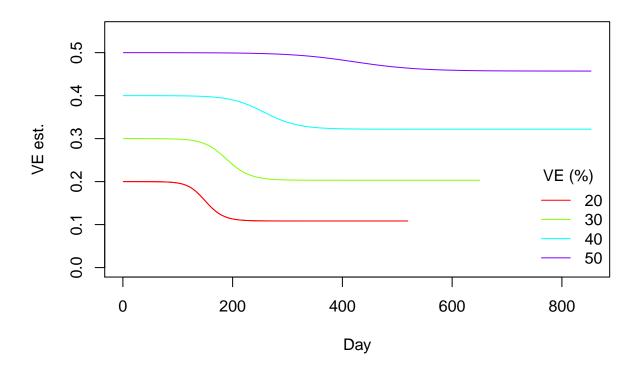
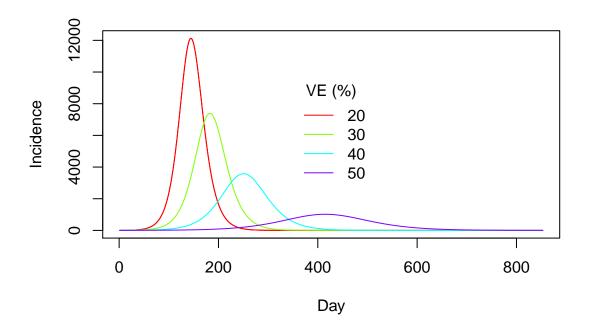


Figure 1: Expected VE estimates over time, by VE.

As Figure 1 shows, the observed VE always declined, The absulute decline was largest with intermediate VE (VE=30%), while the relative decline was most pronounced for low VE (VE=20%)



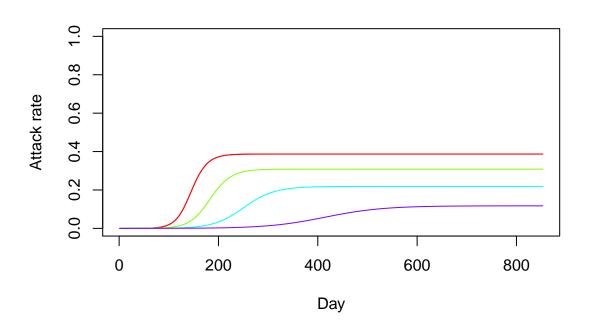


Figure 2: Epi curve (top) and cummulative attack rates (bottom) over time, by VE.