

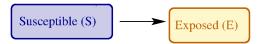


- Different infection phases (compartiments) and their transitions
- ▶ Macroscopic dynamic variables: fractions of the population in the compartiments

Susceptible (S)



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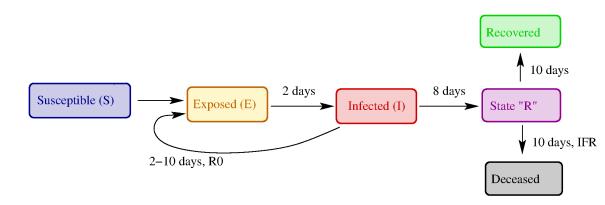




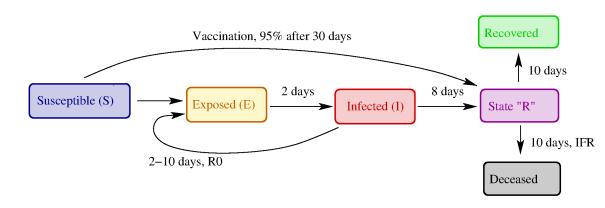
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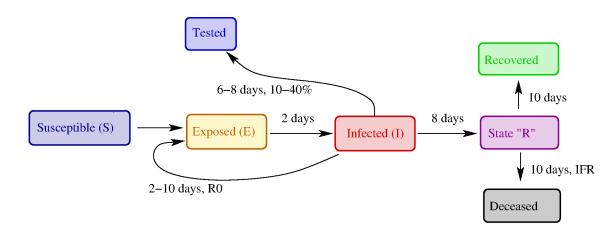
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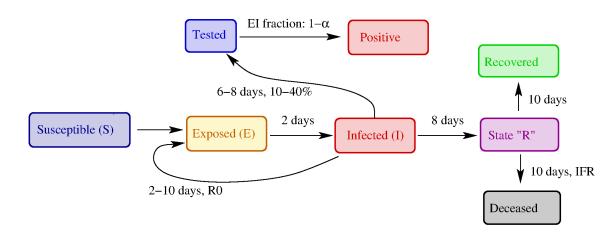


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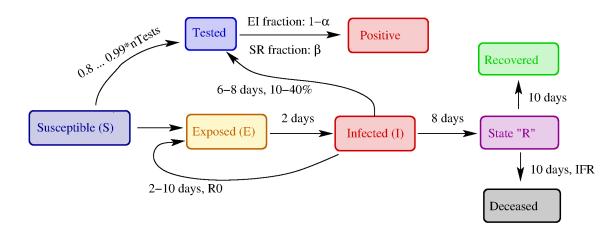




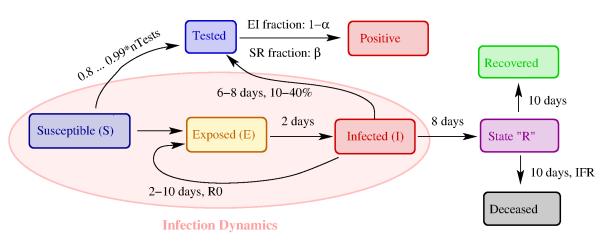
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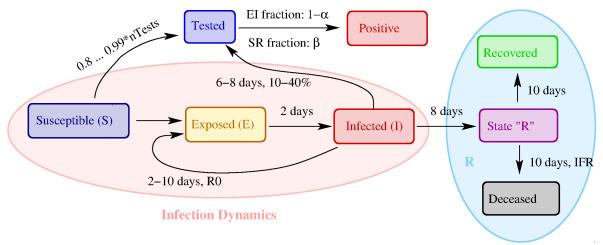
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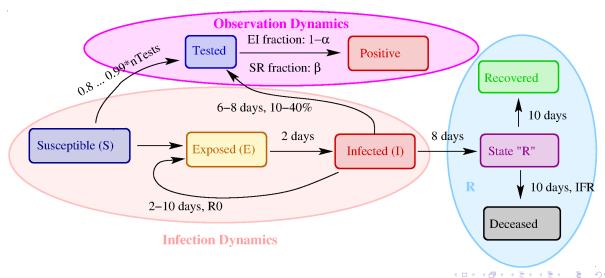
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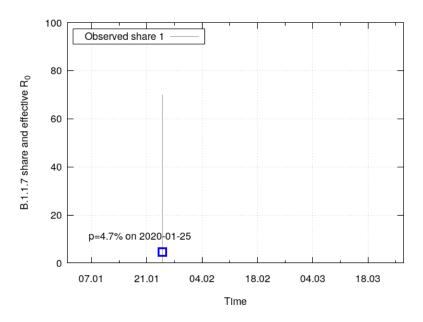


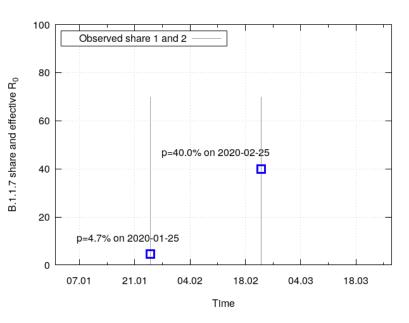
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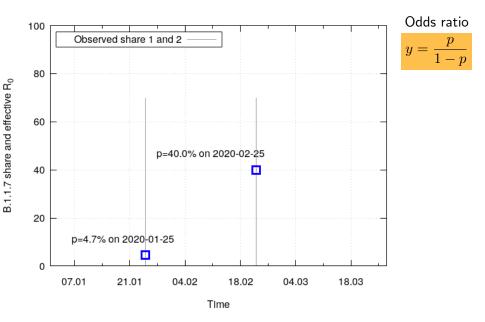


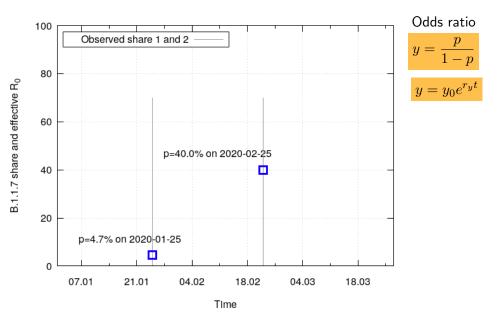
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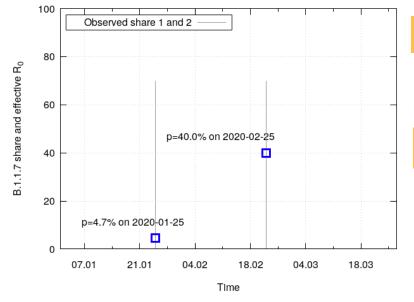












Odds ratio

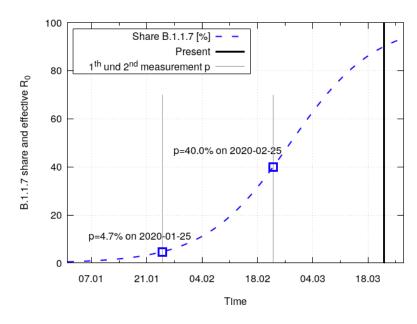
$$y = \frac{p}{1 - p}$$

$$y = y_0 e^{r_y t}$$

Estimation growth rate

$$r_y \approx \frac{\ln y(t_2) - \ln y(t_1)}{t_2 - t_1}$$

#### 1.1 Modelling the mutation dynamics



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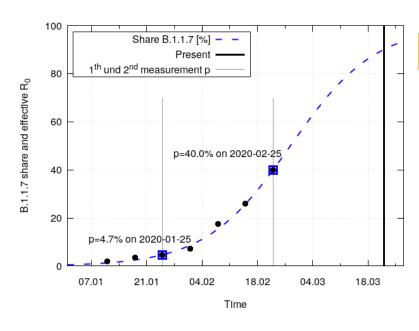
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Estimation B.1.1.7 share

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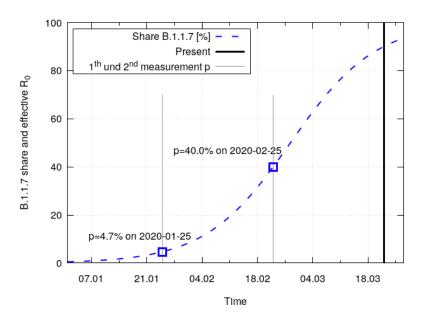
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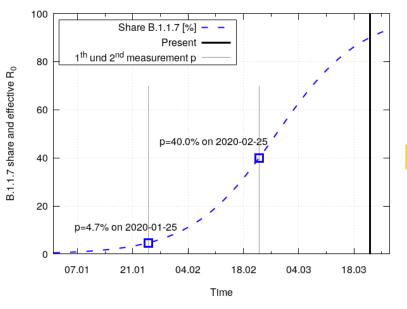
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au: generation time

$$R_0^{\rm mut}/R_0^{\rm wild} = \tau r_y + 1$$

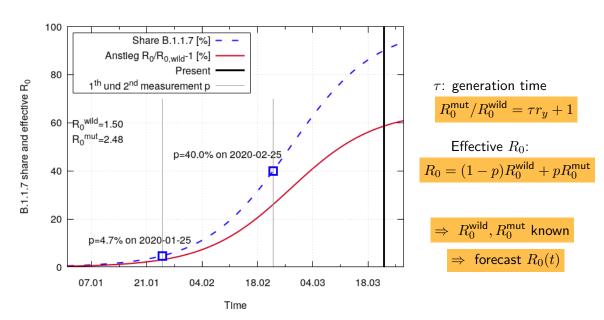


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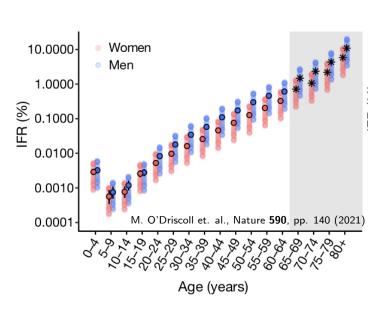
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Effective  $R_0$ :

$$R_0 = (1-p)R_0^{\mathsf{wild}} + pR_0^{\mathsf{mut}}$$

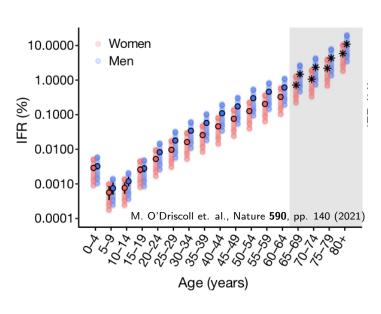


### 1.2 Modelling the age-specific infection fatality rate (IFR)

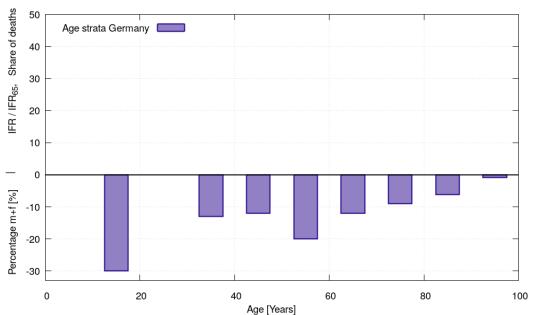


- The IFR exhibits a nearly exponential growth with the age
- The same is true for the case fatality rate (CFR) and for the total one-year fatality
- The gender difference is weak (corresponds to  $\approx$  5 years)
- The infection probability exhibts a much weaker age dependency
- ▶ leave infection spread model unchanged (enables interactive just-in-time calibration) and calculate the IFR in age strata afterwards

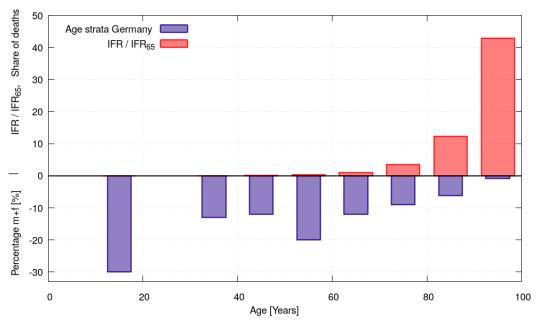
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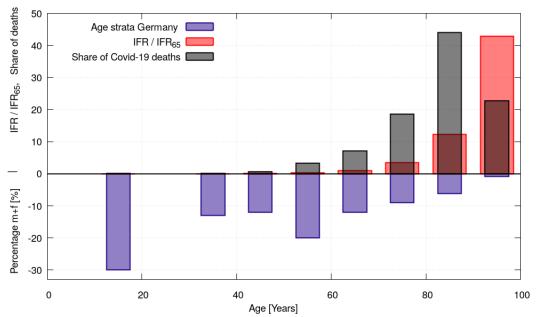


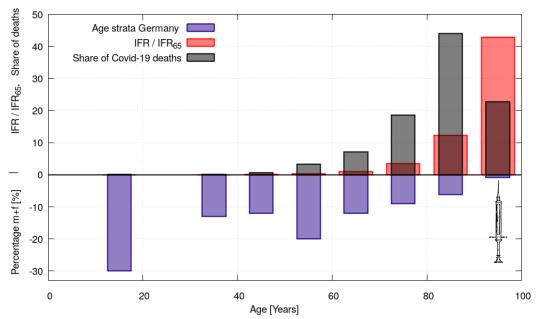
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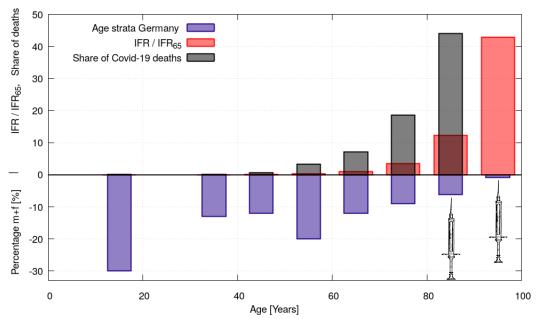


### Calculating the global IFR as multiple of IFR<sub>65</sub>

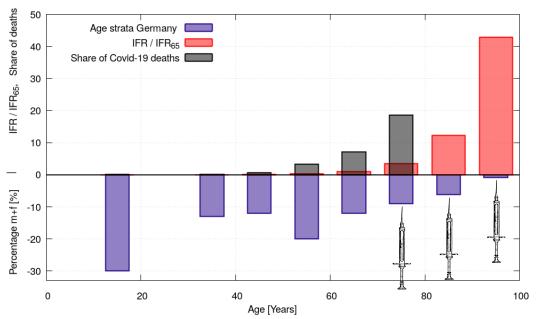


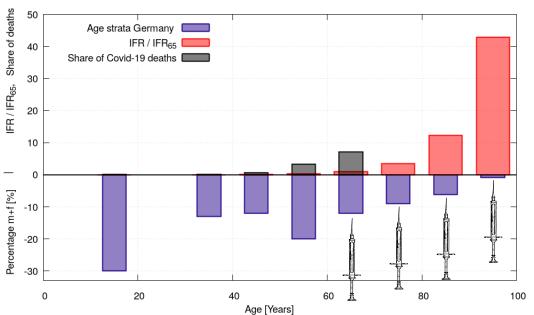




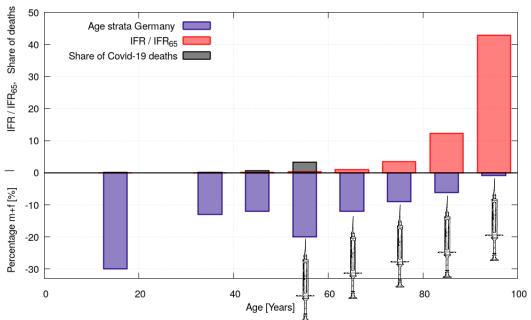


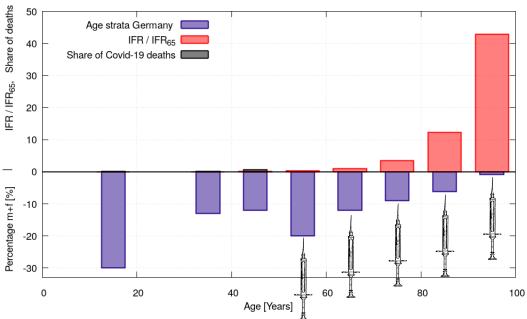




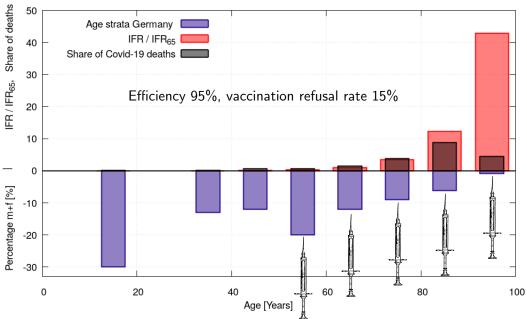












#### TECHNISCHE UNIVERSITÄT

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