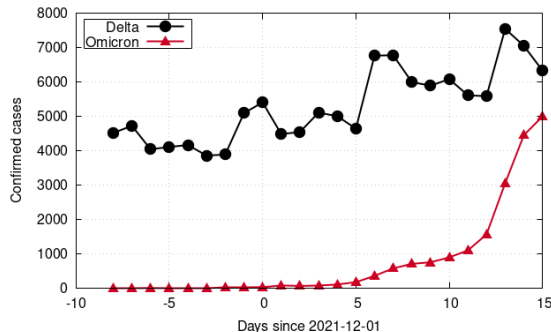
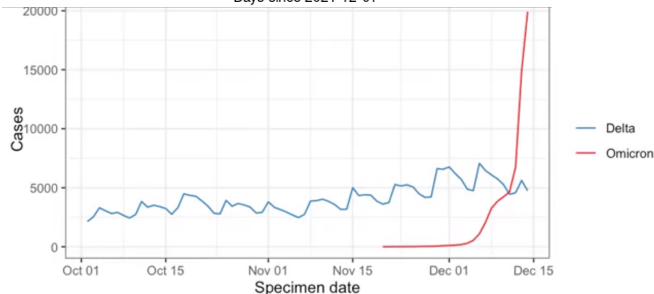


1. Delta and Omicron: Observations

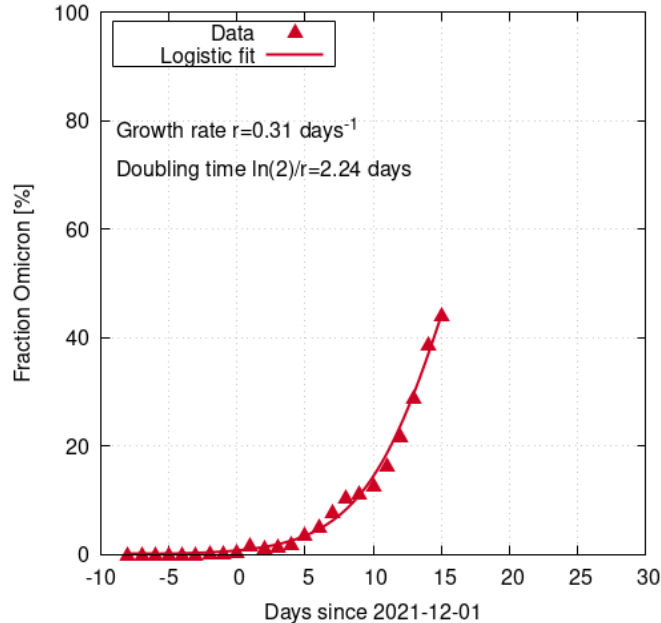


- ▶ Data from Danmark and London
- ▶ The Delta and Omicron variants coexist without directly affecting each other
- ▶ Indirect interaction via competing for common resources, i.e., *first come, first served*



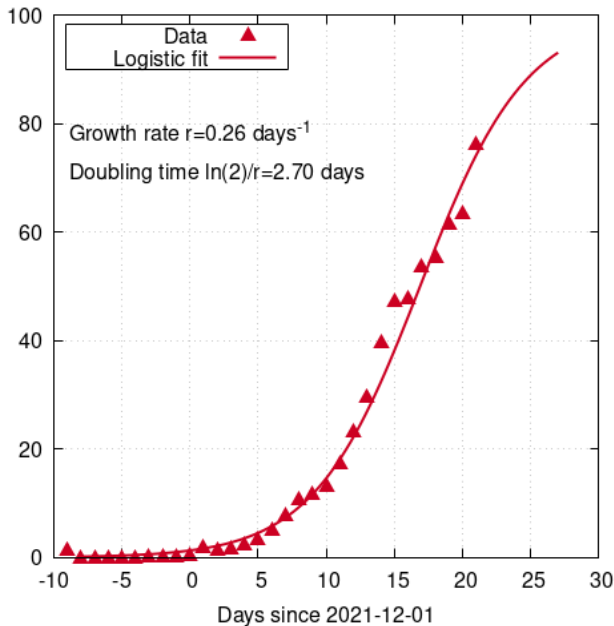
Data: UKHSA. SGTF proportions from daily update combined with case numbers from coronavirus.data.gov.uk.

2. Rate r of the logistic growth of the Omicron share



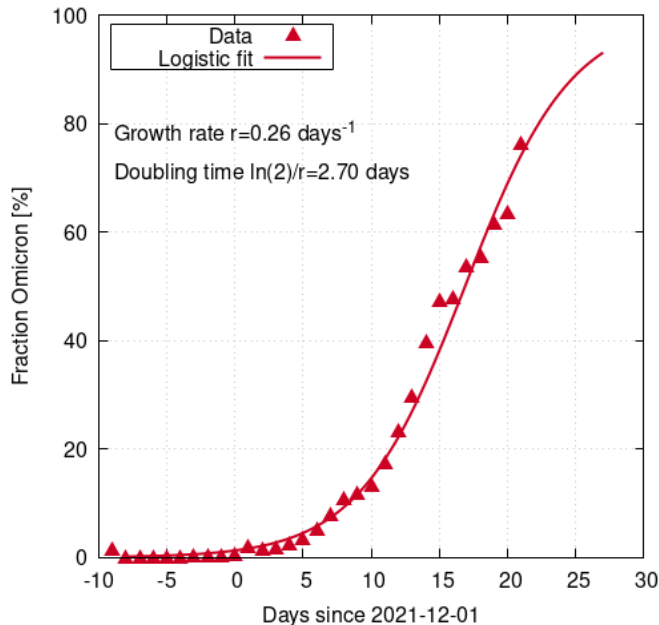
- The share of Omicron can be well described by a logistic function with growth rate r

Update and: how to determine the rate r



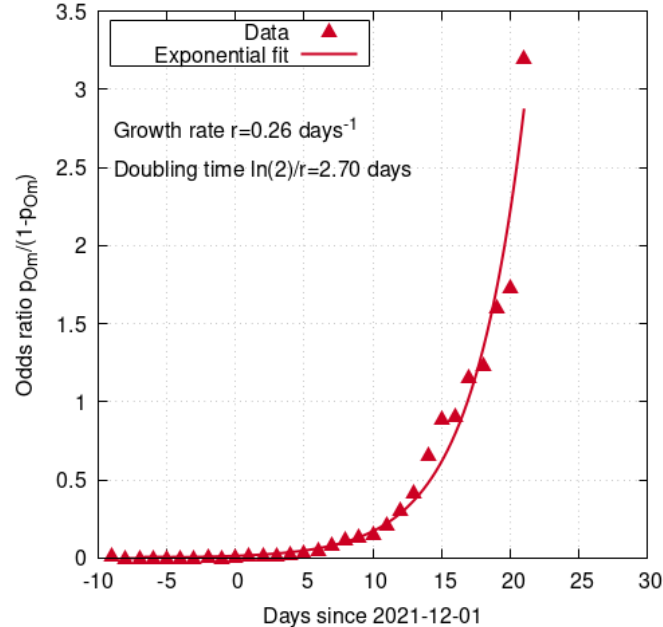
► Data update.

Update and: how to determine the rate r



► Data update. How to get the curve?

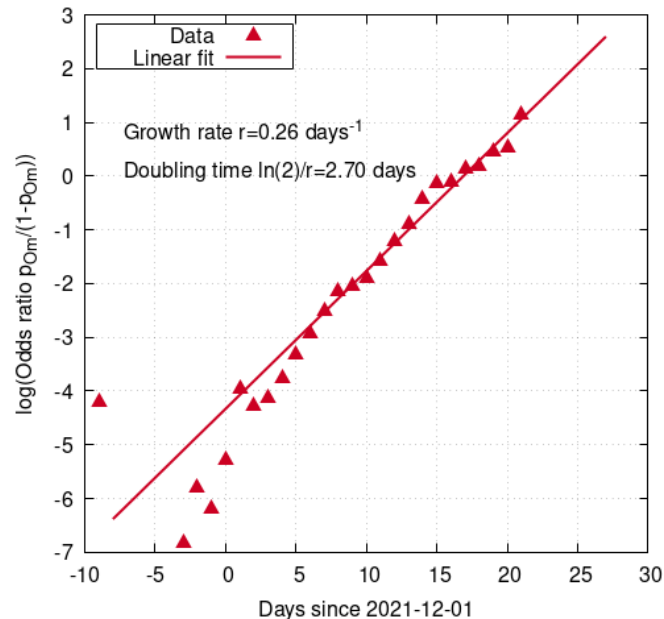
Update and: how to determine the rate r



- Data update. How to get the curve?
- Transform the observed Omicron share p into the **odds ratio** $y = p/(1 - p)$
- From Observation 1 (coexistence), it follows that the odds ratio grows exponentially:

$$y(t) = y_0 e^{rt}$$

Update and: how to determine the rate r



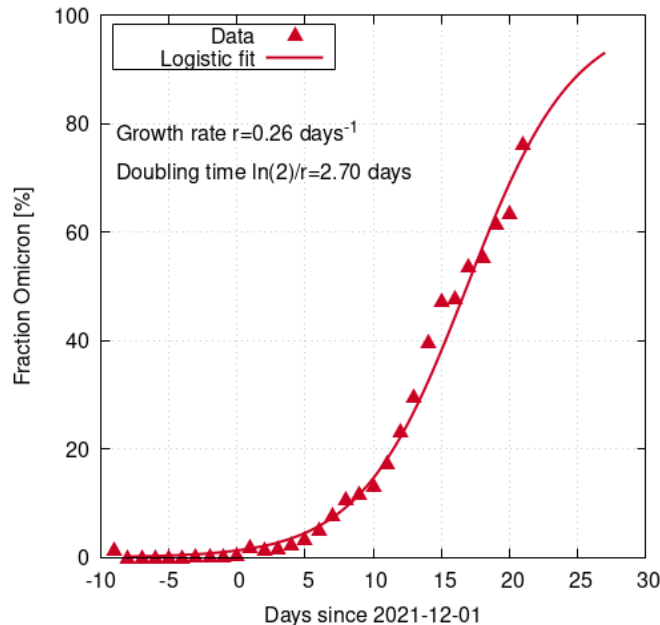
- ▶ Data update. How to get the curve?
- ▶ Transform the observed Omicron share p into the **odds ratio** $y = p/(1-p)$
- ▶ From Observation 1 (coexistence), it follows that the odds ratio grows exponentially:

$$y(t) = y_0 e^{rt}$$

- ▶ This means, the **log-odds** are essentially linear in time:

$$\ln y(t) = \ln y_0 + rt$$

Update and: how to determine the rate r



- Data update. How to get the curve?
- Transform the observed Omicron share p into the **odds ratio** $y = p/(1 - p)$
- From Observation 1 (coexistence), it follows that the odds ratio grows exponentially:

$$y(t) = y_0 e^{rt}$$

- This means, the **log-odds** are essentially linear in time:

$$\ln y(t) = \ln y_0 + rt$$

- Transforming back gives the s-shaped predicted Omicron share (logistic function)

$$p(t) = \frac{y(t)}{1 + y(t)}$$

3. Relation between the logistic growth rate r and the reproduction numbers

Assumptions:

- ▶ Neither positive nor negative **cross effects**: Each variant acts on its own (using common resources of susceptible humans)
- ▶ The Delta and Omicron variants have different **base reproduction numbers** R_{10} and R_{20} and different **generation times** T_1 and T_2 , respectively (e.g., $R_{10} = 5$, $T_1 = 5$ days, $T_2 = 4$ days)
- ▶ The **immunities** I_1 and I_2 (including vaccinations and past infections) against Delta and Omicron are generally different
- ▶ The **reduction factors** f_m by isolation measures and the **seasonal factor** f_s are common
- ▶ All factors influencing the effective reproduction number R are multiplicative

$$\Rightarrow \begin{aligned} x_1(t_0 + T_1) &= R_1 x_1(t_0) = R_{10}(1 - I_1)f_m f_s x_1(t_0), \\ x_2(t_0 + T_2) &= R_2 x_2(t_0) = R_{20}(1 - I_2)f_m f_s x_2(t_0) \end{aligned} \quad (1)$$

3. Relation between the logistic growth rate r and the reproduction numbers

Assumptions:

- ▶ Neither positive nor negative **cross effects**: Each variant acts on its own (using common resources of susceptible humans)
- ▶ The Delta and Omicron variants have different **base reproduction numbers** R_{10} and R_{20} and different **generation times** T_1 and T_2 , respectively (e.g., $R_{10} = 5$, $T_1 = 5$ days, $T_2 = 4$ days)
- ▶ The **immunities** I_1 and I_2 (including vaccinations and past infections) against Delta and Omicron are generally different
- ▶ The **reduction factors** f_m by isolation measures and the **seasonal factor** f_s are common
- ▶ All factors influencing the effective reproduction number R are multiplicative

$$\Rightarrow \begin{aligned} x_1(t_0 + T_1) &= R_1 x_1(t_0) = R_{10}(1 - I_1)f_m f_s x_1(t_0), \\ x_2(t_0 + T_2) &= R_2 x_2(t_0) = R_{20}(1 - I_2)f_m f_s x_2(t_0) \end{aligned} \quad (1)$$

3. Relation between the logistic growth rate r and the reproduction numbers

Assumptions:

- ▶ Neither positive nor negative **cross effects**: Each variant acts on its own (using common resources of susceptible humans)
- ▶ The Delta and Omicron variants have different **base reproduction numbers** R_{10} and R_{20} and different **generation times** T_1 and T_2 , respectively (e.g., $R_{10} = 5$, $T_1 = 5$ days, $T_2 = 4$ days)
- ▶ The **immunities** I_1 and I_2 (including vaccinations and past infections) against Delta and Omicron are generally different
- ▶ The **reduction factors** f_m by isolation measures and the **seasonal factor** f_s are common
- ▶ All factors influencing the effective reproduction number R are multiplicative

$$\Rightarrow \begin{aligned} x_1(t_0 + T_1) &= R_1 x_1(t_0) = R_{10}(1 - I_1)f_m f_s x_1(t_0), \\ x_2(t_0 + T_2) &= R_2 x_2(t_0) = R_{20}(1 - I_2)f_m f_s x_2(t_0) \end{aligned} \quad (1)$$

3. Relation between the logistic growth rate r and the reproduction numbers

Assumptions:

- ▶ Neither positive nor negative **cross effects**: Each variant acts on its own (using common resources of susceptible humans)
- ▶ The Delta and Omicron variants have different **base reproduction numbers** R_{10} and R_{20} and different **generation times** T_1 and T_2 , respectively (e.g., $R_{10} = 5$, $T_1 = 5$ days, $T_2 = 4$ days)
- ▶ The **immunities** I_1 and I_2 (including vaccinations and past infections) against Delta and Omicron are generally different
- ▶ The **reduction factors** f_m by isolation measures and the **seasonal factor** f_s are common
- ▶ All factors influencing the effective reproduction number R are multiplicative

$$\Rightarrow \begin{aligned} x_1(t_0 + T_1) &= R_1 x_1(t_0) = R_{10}(1 - I_1)f_m f_s x_1(t_0), \\ x_2(t_0 + T_2) &= R_2 x_2(t_0) = R_{20}(1 - I_2)f_m f_s x_2(t_0) \end{aligned} \quad (1)$$

3. Relation between the logistic growth rate r and the reproduction numbers

Assumptions:

- ▶ Neither positive nor negative **cross effects**: Each variant acts on its own (using common resources of susceptible humans)
- ▶ The Delta and Omicron variants have different **base reproduction numbers** R_{10} and R_{20} and different **generation times** T_1 and T_2 , respectively (e.g., $R_{10} = 5$, $T_1 = 5$ days, $T_2 = 4$ days)
- ▶ The **immunities** I_1 and I_2 (including vaccinations and past infections) against Delta and Omicron are generally different
- ▶ The **reduction factors** f_m by isolation measures and the **seasonal factor** f_s are common
- ▶ All factors influencing the effective reproduction number R are multiplicative

$$\Rightarrow \begin{aligned} x_1(t_0 + T_1) &= R_1 x_1(t_0) = R_{10}(1 - I_1)f_m f_s x_1(t_0), \\ x_2(t_0 + T_2) &= R_2 x_2(t_0) = R_{20}(1 - I_2)f_m f_s x_2(t_0) \end{aligned} \quad (1)$$

3. Relation between the logistic growth rate r and the reproduction numbers

Assumptions:

- ▶ Neither positive nor negative **cross effects**: Each variant acts on its own (using common resources of susceptible humans)
- ▶ The Delta and Omicron variants have different **base reproduction numbers** R_{10} and R_{20} and different **generation times** T_1 and T_2 , respectively (e.g., $R_{10} = 5$, $T_1 = 5$ days, $T_2 = 4$ days)
- ▶ The **immunities** I_1 and I_2 (including vaccinations and past infections) against Delta and Omicron are generally different
- ▶ The **reduction factors** f_m by isolation measures and the **seasonal factor** f_s are common
- ▶ All factors influencing the effective reproduction number R are multiplicative

$$\Rightarrow \begin{aligned} x_1(t_0 + T_1) &= R_1 x_1(t_0) = R_{10}(1 - I_1) f_m f_s x_1(t_0), \\ x_2(t_0 + T_2) &= R_2 x_2(t_0) = R_{20}(1 - I_2) f_m f_s x_2(t_0) \end{aligned} \quad (1)$$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0)R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0)R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0) R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0) R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0)R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0)R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

Relation between the logistic growth rate r and R_1 and R_2 (ctned)

Assuming continuous infections (slowly varying rates), we can write (1) as

$$x_1(t) = x_1(0)R_1^{t/T_1} = x_1(0) \exp\left(\frac{t}{T_1} \ln R_1\right) \equiv x_1(0) \exp(r_1 t),$$

$x_2(t)$ likewise, or

$$\dot{x}_1 = r_1 x_1, \quad r_1 = \frac{\ln R_1}{T_1}, \quad \dot{x}_2 = r_2 x_2, \quad r_2 = \frac{\ln R_2}{T_2} \quad (2)$$

How does the odds $y = x_2/x_1$ evolve?

$$\begin{aligned} \dot{y} &= \frac{d}{dt} \left(\frac{x_2}{x_1} \right) = \frac{\dot{x}_2}{x_1} - \frac{x_2}{x_1^2} \dot{x}_1 \\ &= \frac{r_2 x_2}{x_1} - \frac{x_2}{x_1^2} r_1 x_1 = (r_2 - r_1) y \end{aligned}$$

$$\Rightarrow \quad r = r_2 - r_1 = \frac{\ln R_2}{T_2} - \frac{\ln R_1}{T_1} \quad (3)$$

Special case $T_2 = T_1 = T$: $R_2 = R_1 \exp(rT)$

4. Determining the Omicron base reproduction rate from the logistic growth rate

Just use Relation (3) and insert the definitions of R_1 and R_2 from (1)

After some manipulations ...

4. Determining the Omicron base reproduction rate from the logistic growth rate

Just use Relation (3) and insert the definitions of R_1 and R_2 from (1)

After some manipulations ...

$$R_{20} = \exp(rT_2) f_m^{\gamma-1} f_s^{\gamma-1} \frac{(R_{10}(1 - I_1))^\gamma}{1 - I_2}, \quad \gamma = \frac{T_2}{T_1} \quad (4)$$

- ▶ For equal generation times $T_1 = T_2 = T$, the measures and the seasonal effects drop out and r depends only on the past infection and vaccination immunities (remains time dependent since the immunities change):

$$T_1 = T_2 = T \quad \Rightarrow \quad R_{20} = e^{rT} \frac{R_{10}(1 - I_1)}{1 - I_2}$$

Example: $e^{rT} = 3$, $R_{10} = 4$, $I_1 = 0.6$, $I_2 = 0.2$, $R_{20} = 3/2 R_{10} = 6$

- ▶ With neither immunities nor measures nor season effects but $T_2 = 0.5T_1$ ($\gamma = 0.5$), we have $R_{20} = e^{rT_2} \sqrt{R_{10}}$, e.g., for $e^{rT_2} = 2$ and $R_{01} = 4$, we have $R_{02} = R_{01} = 4$
- ▶ With measures/season effects and immunities as above, we may have $R_{02} < R_{01}$

4. Determining the Omicron base reproduction rate from the logistic growth rate

Just use Relation (3) and insert the definitions of R_1 and R_2 from (1)

After some manipulations ...

$$R_{20} = \exp(rT_2) f_m^{\gamma-1} f_s^{\gamma-1} \frac{(R_{10}(1 - I_1))^\gamma}{1 - I_2}, \quad \gamma = \frac{T_2}{T_1} \quad (4)$$

- For equal generation times $T_1 = T_2 = T$, the measures and the seasonal effects drop out and r depends only on the past infection and vaccination immunities (remains time dependent since the immunities change):

$$T_1 = T_2 = T \quad \Rightarrow \quad R_{20} = e^{rT} \frac{R_{10}(1 - I_1)}{1 - I_2}$$

Example: $e^{rT} = 3$, $R_{10} = 4$, $I_1 = 0.6$, $I_2 = 0.2$, $R_{20} = 3/2 R_{10} = 6$

- With neither immunities nor measures nor season effects but $T_2 = 0.5T_1$ ($\gamma = 0.5$), we have $R_{20} = e^{rT_2} \sqrt{R_{10}}$, e.g., for $e^{rT_2} = 2$ and $R_{01} = 4$, we have $R_{02} = R_{01} = 4$
- With measures/season effects and immunities as above, we may have $R_{02} < R_{01}$

4. Determining the Omicron base reproduction rate from the logistic growth rate

Just use Relation (3) and insert the definitions of R_1 and R_2 from (1)

After some manipulations ...

$$R_{20} = \exp(rT_2) f_m^{\gamma-1} f_s^{\gamma-1} \frac{(R_{10}(1 - I_1))^\gamma}{1 - I_2}, \quad \gamma = \frac{T_2}{T_1} \quad (4)$$

- For equal generation times $T_1 = T_2 = T$, the measures and the seasonal effects drop out and r depends only on the past infection and vaccination immunities (remains time dependent since the immunities change):

$$T_1 = T_2 = T \quad \Rightarrow \quad R_{20} = e^{rT} \frac{R_{10}(1 - I_1)}{1 - I_2}$$

Example: $e^{rT} = 3$, $R_{10} = 4$, $I_1 = 0.6$, $I_2 = 0.2$, $R_{20} = 3/2 R_{10} = 6$

- With neither immunities nor measures nor season effects but $T_2 = 0.5T_1$ ($\gamma = 0.5$), we have $R_{20} = e^{rT_2} \sqrt{R_{10}}$, e.g., for $e^{rT_2} = 2$ and $R_{01} = 4$, we have $R_{02} = R_{01} = 4$
- With measures/season effects and immunities as above, we may have $R_{02} < R_{01}$

4. Determining the Omicron base reproduction rate from the logistic growth rate

Just use Relation (3) and insert the definitions of R_1 and R_2 from (1)

After some manipulations ...

$$R_{20} = \exp(rT_2) f_m^{\gamma-1} f_s^{\gamma-1} \frac{(R_{10}(1 - I_1))^\gamma}{1 - I_2}, \quad \gamma = \frac{T_2}{T_1} \quad (4)$$

- ▶ For equal generation times $T_1 = T_2 = T$, the measures and the seasonal effects drop out and r depends only on the past infection and vaccination immunities (remains time dependent since the immunities change):

$$T_1 = T_2 = T \quad \Rightarrow \quad R_{20} = e^{rT} \frac{R_{10}(1 - I_1)}{1 - I_2}$$

Example: $e^{rT} = 3$, $R_{10} = 4$, $I_1 = 0.6$, $I_2 = 0.2$, $R_{20} = 3/2 R_{10} = 6$

- ▶ With neither immunities nor measures nor season effects but $T_2 = 0.5T_1$ ($\gamma = 0.5$), we have $R_{20} = e^{rT_2} \sqrt{R_{10}}$, e.g., for $e^{rT_2} = 2$ and $R_{01} = 4$, we have $R_{02} = R_{01} = 4$
- ▶ With measures/season effects and immunities as above, we may have $R_{02} < R_{01}$

5. Effective infection growth rate and reproduction number

The **effective growth rate** r_{eff} of the infection dynamics (not to be confused with the logistic growth rate r of the Omicron shares p) comes directly from (2):

$$\dot{x} = \dot{x}_1 + \dot{x}_2 = r_1 x_1 + r_2 x_2 = [(1-p)r_1 + pr_2]x \equiv r_{\text{eff}}x$$

Effective reproduction number R_{eff} by association $r_{\text{eff}} \equiv \ln R_{\text{eff}}/T_1$:

$$\ln R_{\text{eff}} = (1-p) \ln R_1 + \frac{p}{\gamma} \ln R_2 \quad (5)$$

- ▶ Because $1/\gamma = T_1/T_2 > 1$, influence factors, e.g., measures, have a more sensitive effect on Omicron than on Delta: If $T_1/T_2 = 1/\gamma = 2$ and measures lead to a factor $1/\sqrt{2} \approx 0.7$ on Delta (R_1), they simultaneously lead to a factor $1/2$ on Omicron (R_2)
- ▶ If, at a certain time, the true Omicron share p , the effective reproduction number R_{eff} , and the logistic growth rate r are known (all three can be estimated), and the generation time ratio $\gamma = T_2/T_1$ as well as the total immunities I_1 and I_2 and the effects of the measures and the season at this time can be estimated, the Eqs (1), (4), and (5) allow for a simultaneous estimation of R_{10} and R_{20}

5. Effective infection growth rate and reproduction number

The **effective growth rate** r_{eff} of the infection dynamics (not to be confused with the logistic growth rate r of the Omicron shares p) comes directly from (2):

$$\dot{x} = \dot{x}_1 + \dot{x}_2 = r_1 x_1 + r_2 x_2 = [(1-p)r_1 + pr_2]x \equiv r_{\text{eff}}x$$

Effective reproduction number R_{eff} by association $r_{\text{eff}} \equiv \ln R_{\text{eff}}/T_1$:

$$\ln R_{\text{eff}} = (1-p) \ln R_1 + \frac{p}{\gamma} \ln R_2 \quad (5)$$

- ▶ Because $1/\gamma = T_1/T_2 > 1$, influence factors, e.g., measures, have a more sensitive effect on Omicron than on Delta: If $T_1/T_2 = 1/\gamma = 2$ and measures lead to a factor $1/\sqrt{2} \approx 0.7$ on Delta (R_1), they simultaneously lead to a factor $1/2$ on Omicron (R_2)
- ▶ If, at a certain time, the true Omicron share p , the effective reproduction number R_{eff} , and the logistic growth rate r are known (all three can be estimated), and the generation time ratio $\gamma = T_2/T_1$ as well as the total immunities I_1 and I_2 and the effects of the measures and the season at this time can be estimated, the Eqs (1), (4), and (5) allow for a simultaneous estimation of R_{10} and R_{20}

5. Effective infection growth rate and reproduction number

The **effective growth rate** r_{eff} of the infection dynamics (not to be confused with the logistic growth rate r of the Omicron shares p) comes directly from (2):

$$\dot{x} = \dot{x}_1 + \dot{x}_2 = r_1 x_1 + r_2 x_2 = [(1-p)r_1 + pr_2]x \equiv r_{\text{eff}}x$$

Effective reproduction number R_{eff} by association $r_{\text{eff}} \equiv \ln R_{\text{eff}}/T_1$:

$$\ln R_{\text{eff}} = (1-p) \ln R_1 + \frac{p}{\gamma} \ln R_2 \quad (5)$$

- ▶ Because $1/\gamma = T_1/T_2 > 1$, influence factors, e.g., measures, have a more sensitive effect on Omicron than on Delta: If $T_1/T_2 = 1/\gamma = 2$ and measures lead to a factor $1/\sqrt{2} \approx 0.7$ on Delta (R_1), they simultaneously lead to a factor $1/2$ on Omicron (R_2)
- ▶ If, at a certain time, the true Omicron share p , the effective reproduction number R_{eff} , and the logistic growth rate r are known (all three can be estimated), and the generation time ratio $\gamma = T_2/T_1$ as well as the total immunities I_1 and I_2 and the effects of the measures and the season at this time can be estimated, the Eqs (1), (4), and (5) allow for a simultaneous estimation of R_{10} and R_{20}

5. Effective infection growth rate and reproduction number

The **effective growth rate** r_{eff} of the infection dynamics (not to be confused with the logistic growth rate r of the Omicron shares p) comes directly from (2):

$$\dot{x} = \dot{x}_1 + \dot{x}_2 = r_1 x_1 + r_2 x_2 = [(1-p)r_1 + pr_2]x \equiv r_{\text{eff}}x$$

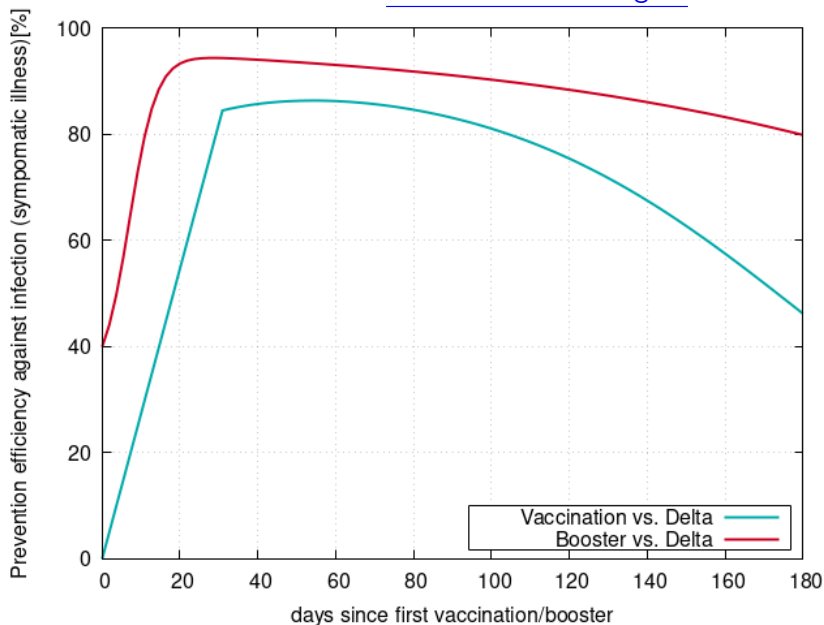
Effective reproduction number R_{eff} by association $r_{\text{eff}} \equiv \ln R_{\text{eff}}/T_1$:

$$\ln R_{\text{eff}} = (1-p) \ln R_1 + \frac{p}{\gamma} \ln R_2 \quad (5)$$

- ▶ Because $1/\gamma = T_1/T_2 > 1$, influence factors, e.g., measures, have a more sensitive effect on Omicron than on Delta: If $T_1/T_2 = 1/\gamma = 2$ and measures lead to a factor $1/\sqrt{2} \approx 0.7$ on Delta (R_1), they simultaneously lead to a factor $1/2$ on Omicron (R_2)
- ▶ If, at a certain time, the true Omicron share p , the effective reproduction number R_{eff} , and the logistic growth rate r are known (all three can be estimated), and the generation time ratio $\gamma = T_2/T_1$ as well as the total immunities I_1 and I_2 and the effects of the measures and the season at this time can be estimated, the Eqs (1), (4), and (5) allow for a simultaneous estimation of R_{10} and R_{20}

6. Immunity I: vaccinations/boosters vs. Delta variant

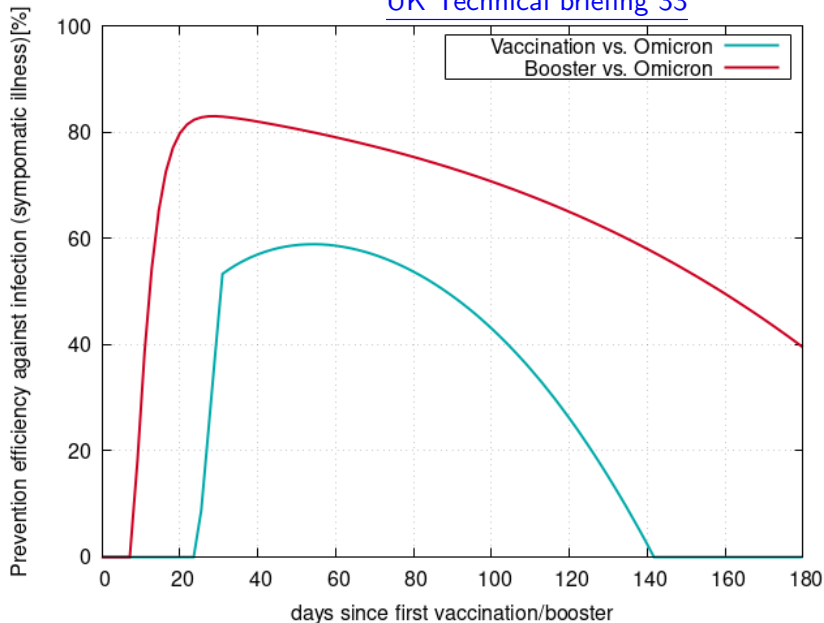
UK Technical briefing 33



“First vaccinated-
first boosted”
principle

6. Immunity II: vaccinations/boosters vs. Omicron variant

[UK Technical briefing 33](#)



Only fresh
full vaccinations
or boosters
help against
Omicron

6. Immunity III: past infections

- ▶ 100 % immunity of Delta against Delta reinfections
- ▶ 100 % immunity of Omicron against Omicron reinfections
- ▶ 100 % no cross immunity (people can get both Delta and Omicron infections)

6. Immunity III: past infections

- ▶ 100 % immunity of Delta against Delta reinfections
- ▶ 100 % immunity of Omicron against Omicron reinfections
- ▶ 100 % no cross immunity (people can get both Delta and Omicron infections)

6. Immunity III: past infections

- ▶ 100 % immunity of Delta against Delta reinfections
- ▶ 100 % immunity of Omicron against Omicron reinfections
- ▶ 100 % no cross immunity (people can get both Delta and Omicron infections)

7. Determining the population immunities in the simulator I: vaccinations

Here, I make following assumptions

- ▶ Vaccination efficiency curves $I_1^v(\tau)$ and $I_2^v(\tau)$ against Delta and Omega as shown,
- ▶ corresponding booster efficiencies $I_1^b(\tau)$ and $I_2^b(\tau)$,
- ▶ *First vaccinated-first boosted*

Since the protection depends on the vaccination times, I sum up the different histories weighted with the past daily vaccination and booster rates $r_{t'}^v$ and $r_{t'}^b$ (fraction of the population per day):

$$I_1^{\text{vacc}}(t) = \sum_{t'=t_v}^t r_{t'}^v I_1^v(t-t') + \sum_{t'=t_b}^t r_{t'}^b I_1^b(t-t')$$

where t_b is the time of the first booster shot, and t_v the oldest time of the first vaccination of any person who is not yet boosted.

The vaccination immunity I_2 is calculated in analogy.

7. Determining the population immunities in the simulator I: vaccinations

Here, I make following assumptions

- ▶ Vaccination efficiency curves $I_1^v(\tau)$ and $I_2^v(\tau)$ against Delta and Omega as shown,
- ▶ corresponding booster efficiencies $I_1^b(\tau)$ and $I_2^b(\tau)$,
- ▶ *First vaccinated-first boosted*

Since the protection depends on the vaccination times, I sum up the different histories weighted with the past daily vaccination and booster rates $r_{t'}^v$ and $r_{t'}^b$ (fraction of the population per day):

$$I_1^{\text{vacc}}(t) = \sum_{t'=t_v}^t r_{t'}^v I_1^v(t-t') + \sum_{t'=t_b}^t r_{t'}^b I_1^b(t-t')$$

where t_b is the time of the first booster shot, and t_v the oldest time of the first vaccination of any person who is not yet boosted.

The vaccination immunity I_2 is calculated in analogy.

7. Determining the population immunities in the simulator I: vaccinations

Here, I make following assumptions

- ▶ Vaccination efficiency curves $I_1^v(\tau)$ and $I_2^v(\tau)$ against Delta and Omega as shown,
- ▶ corresponding booster efficiencies $I_1^b(\tau)$ and $I_2^b(\tau)$,
- ▶ *First vaccinated-first boosted*

Since the protection depends on the vaccination times, I sum up the different histories weighted with the past daily vaccination and booster rates $r_{t'}^v$ and $r_{t'}^b$ (fraction of the population per day):

$$I_1^{\text{vacc}}(t) = \sum_{t'=t_v}^t r_{t'}^v I_1^v(t-t') + \sum_{t'=t_b}^t r_{t'}^b I_1^b(t-t')$$

where t_b is the time of the first booster shot, and t_v the oldest time of the first vaccination of any person who is not yet boosted.

The vaccination immunity I_2 is calculated in analogy.

7. Determining the population immunities in the simulator I: vaccinations

Here, I make following assumptions

- ▶ Vaccination efficiency curves $I_1^v(\tau)$ and $I_2^v(\tau)$ against Delta and Omega as shown,
- ▶ corresponding booster efficiencies $I_1^b(\tau)$ and $I_2^b(\tau)$,
- ▶ *First vaccinated-first boosted*

Since the protection depends on the vaccination times, I sum up the different histories weighted with the past daily vaccination and booster rates $r_{t'}^v$ and $r_{t'}^b$ (fraction of the population per day):

$$I_1^{\text{vacc}}(t) = \sum_{t'=t_v}^t r_{t'}^v I_1^v(t-t') + \sum_{t'=t_b}^t r_{t'}^b I_1^b(t-t')$$

where t_b is the time of the first booster shot, and t_v the oldest time of the first vaccination of any person who is not yet boosted.

The vaccination immunity I_2 is calculated in analogy.

Determining the population immunities in the simulator II: infections and total

- Everybody can only be infected once with any variant but there is no cross immunity, so the immunity is just equal to the total percentage X_1 and X_2 of people infected with either variant:

$$I_1^x = X_1, \quad I_2^x = X_2$$

Notice: X_i is not just the cumulated number of cases divided by the population because any infection, whether detected or not detected, counts

Determining the population immunities in the simulator II: infections and total

- Everybody can only be infected once with any variant but there is no cross immunity, so the immunity is just equal to the total percentage X_1 and X_2 of people infected with either variant:

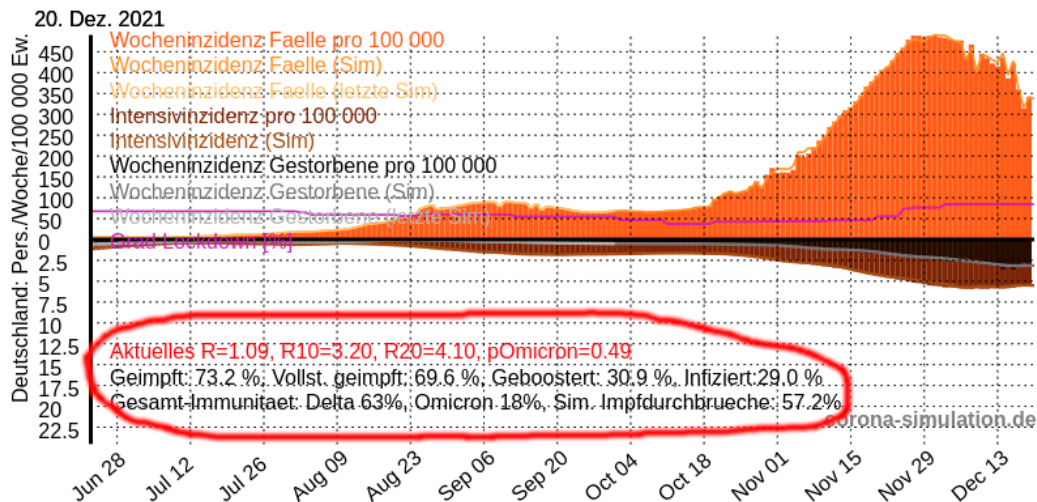
$$I_1^x = X_1, \quad I_2^x = X_2$$

Notice: X_i is not just the cumulated number of cases divided by the population because any infection, whether detected or not detected, counts

- There is no correlation between vaccinations and infections:

$$1 - I_1 = (1 - I_1^v)(1 - I_1^x), \quad 1 - I_2 = (1 - I_2^v)(1 - I_2^x) \quad (6)$$

Simulation



All items I_1 , I_2 , p , R_{10} , R_{20} , f_{season} and $f_{\text{stringency}}$ are displayed in the simulation