

### Contents

Main messages	3
Purpose	4
Methods	4
COVID-19 Omicron variant infectious period	5
The difference in transmission from people with asymptomatic compared with symptomatic COVID-19 Omicron variant	. 28
Inequalities	. 33
Limitations	. 33
Evidence gaps	. 33
Conclusion	. 34
Acknowledgment	.35
Disclaimer	.35
References	. 36
Annexe A. Methods	.43
Protocol	.43
Inclusion and exclusion criteria	.44
Identification of studies	.45
Screening	.45
Data extraction	.45
Risk of bias assessment	.45
Search strategy for infectious period of Omicron variant COVID-19	.46
Search strategy for the difference in transmission of COVID-19 from people with asymptomat and symptomatic Omicron variant COVID-19	

### Main messages

This review (search up to 26 January 2023) identifies and summarises evidence on coronavirus (COVID-19) Omicron variant infectious period (82 studies), and the difference in transmission from people with asymptomatic compared with symptomatic COVID-19 infection (10 studies).

### Infectious period of COVID-19 Omicron variant

- 1. Three studies directly examining transmission suggested that most transmission events happened around symptom onset, with the majority of transmission events occurring up to 5 days after symptom onset (100% in one study, 81% in another study, not reported in the third study).
- 2. Ten studies also reported that viral culture positivity was highest in the first 5 days after symptom onset or diagnosis, and 14 studies reporting on viral load suggested that peak viral loads occurred a median and mean of 2 to 5 days after symptom onset.
- 3. Nine studies measuring serial interval and generation time suggested the median and mean serial interval was between 2 and 4 days (with interquartile ranges between one and 9 days), and generation times of around 3 days.
- 4. Fifty-three studies measuring time to viral clearance suggested that there were substantial differences in viral clearance times between cases and populations. The majority of studies in the general population estimated viral clearance to take around 7 to 11 days, and most studies of hospitalised, immunodeficient, and other high-risk cases estimated viral clearance to take around 10 to 15 days. Whilst detectable viral load does not provide direct evidence on the risk of transmission, it does indicate potential infectivity.
- 5. Overall, the evidence suggests that COVID-19 Omicron variant cases were most infectious up to 5 days after symptom onset but could potentially be infectious for longer, especially if hospitalised, immunocompromised, or otherwise high-risk. Most studies included relatively few cases.

# Asymptomatic compared with symptomatic transmission of COVID-19 Omicron variant

- 6. Three studies compared household secondary attack rates (SAR) of asymptomatic and symptomatic index cases, with 2 studies suggesting more transmission from symptomatic than asymptomatic index cases, and one study suggesting no clear difference.
- 7. Five studies compared viral loads (usually using Ct values) between asymptomatic and symptomatic cases, with 3 studies suggesting similar viral loads, and 2 studies suggesting higher viral loads in symptomatic compared with asymptomatic cases.
- 8. Overall, the evidence on differences in transmission from people with asymptomatic compared with symptomatic COVID-19 Omicron variant was mixed, with some studies suggesting that symptomatic cases were more likely to transmit infection than asymptomatic

cases and others showing no difference. The lack of precision due to small study samples combined with the variability between studies and the small number of studies limit the ability to draw firm conclusions.

### Purpose

To identify and summarise evidence relating to COVID-19 Omicron variant coronavirus infectious period, and the difference in transmission from people with asymptomatic compared with symptomatic COVID-19.

### **Methods**

There were 2 review questions:

- 1. What is the infectious period of COVID-19 Omicron variant?
- 2. What is the difference in transmission from people with asymptomatic compared with symptomatic COVID-19 Omicron variant?

We previously conducted reviews covering the infectious period of COVID-19 (searches to 23 February 2022 and 16 May 2022). However, these could not be used as a source of evidence for this current review (which looked at primary studies) as the previous searches were aimed at identifying existing rapid and systematic reviews. The primary study search for evidence on infectious period therefore started 1 December 2021, shortly after the start of the Omicron wave.

We also conducted a previous review looking at the difference in transmission of COVID-19 between symptomatic and asymptomatic index cases (search to 15 March 2022). This review was used as a source of evidence up to 15 March 2022.

A rapid review was conducted to search for primary studies up to 26 January 2023 for both review questions (from 1 December 2021 for infectious period, and from 15 March 2022 for asymptomatic transmission), following streamlined systematic methodologies to accelerate the review process (<u>1</u>). Only studies where the majority of the participants in the study were stated to have the Omicron variant of COVID-19 were included.

Ten percent of the title and abstract screening was completed in duplicate for each review question, while full text screening and data extraction were performed by one reviewer and checked by another. Risk of bias assessment using the quality criteria checklist (QCC) (<u>2</u>) was performed by one reviewer and checked by another for all analytical studies (not descriptive studies). Full details on the methodology are provided in <u>Annexe A</u>.

The review questions were searched for and reported separately within the same report.

### **COVID-19 Omicron variant infectious period**

### Evidence

In total, 82 observational studies looking at different measures of COVID-19 Omicron variant infectious period were included in this report. As all studies were descriptive rather than analytical, quality was not assessed. Infectious period is difficult to measure directly, so studies reporting on related measures were also included.

In total, 3 studies reported directly on transmission period (3 to 5), 10 studies on culture positivity over time (<u>6 to 15</u>), 8 studies on incubation period (<u>4</u>, <u>5</u>, <u>16 to 21</u>), one study on latent period (<u>5</u>), 9 studies on serial interval (<u>3</u>, <u>4</u>, <u>16 to 18</u>, <u>21 to 24</u>), 6 studies on time to peak viral load (<u>14</u>, <u>25 to 29</u>), 53 studies on time to viral clearance (<u>12</u>, <u>14</u>, <u>15</u>, <u>27</u>, <u>29 to 77</u>), and 9 studies on viral load over time (<u>7</u>, <u>25</u>, <u>78 to 84</u>). Some studies provided evidence for multiple outcomes. Although some studies compared treatments for COVID-19, the effects of treatments on infectious period outcomes were not considered, only the infectious period within different groups in each study.

Studies directly assessing infectious period, large studies (especially those conducted in the UK), and studies of more recent Omicron variants (BA.3 and later) are individually summarised, whereas the remainder of studies are jointly summarised. <u>Table 1</u> gives study characteristics of the 82 included studies.

#### **Transmission period**

There were 3 included studies (3 to 5) (no preprints) that included direct evidence for the transmission period of COVID-19 Omicron variant. These studies identified index cases and known secondary cases and estimated when transmission was likely to have taken place. These studies most directly estimate the period of infectiousness.

Of these studies, 2 studies were conducted in Europe ( $\underline{3}$ ,  $\underline{4}$ ), and one in China ( $\underline{5}$ ). All studies were conducted between December 2021 and May 2022. One study was a prospective cohort study ( $\underline{3}$ ), one study was a retrospective cohort study ( $\underline{4}$ ), and one study was a cross-sectional study ( $\underline{5}$ ). These studies included cases with COVID-19 Omicron variant BA.1 and BA.2 ( $\underline{3}$ ), BA.1 ( $\underline{4}$ ), and BA.1.1 ( $\underline{5}$ ) variant.

All 3 studies are summarised in detail below, with study characteristics in Table 1a.

#### Summaries of individual studies

An der Heiden and others reported on transmission of COVID-19 Omicron variant BA.1 and BA.2 in n=11,512 households in Germany between January 2022 and May 2022 (<u>3</u>). The

results suggested that 81% of transmission occurred by day 5 of symptom onset in the index case, and 95% of transmission occurred by day 10 of symptom onset in the index case.

Del Aguila-Mejia and others reported on transmission of COVID-19 Omicron variant BA.1 between n=622 index cases and n=455 secondary cases in Spain in December 2021 ( $\underline{4}$ ). The median transmission period after symptom onset was 0 days (interquartile range [IQR]: -1 to 2 days).

Xin and others reported on transmission of COVID-19 Omicron variant BA.1.1 between n=113 pairs of cases in China between January and February 2022 (<u>5</u>). The estimated proportion of transmissions occurring 4 or more days before symptom onset was 4.4% (95% credible interval [CrI]: 0.9% to 8.0%), and the estimated proportion of transmissions occurring before symptom onset was 33.6% (95% CrI: 24.8% to 42.5%). Overall, transmission peaked at symptom onset, and all observed transmission events occurred within 5 days of symptom onset.

#### Viral culture positivity over time

There were 10 included studies (6 to 15) (one preprint (8)) that included evidence on viral culture positivity over time for COVID-19 Omicron variant. These studies typically took repeated viral samples from cases over the course of their illness and tested them for the presence of live virus using cultures. The presence of live virus in a sample does not necessarily indicate that the case would infect other people with whom they came into contact, but is a stronger indicator of infectiousness than just the presence of viral RNA as measured using RT-PCR.

Of these studies, 5 were conducted in Asia (9 to 11, 13, 15), 2 in the US (7, 8), one in the UK (6), one in Turkey (12), and one in Brazil (14). Nine studies were prospective cohort studies (6 to 14), and the remaining study was a retrospective cohort study (15). All studies were conducted between November 2021 and May 2022. These studies included cases with COVID-19 Omicron B.1.1.529 (10), BA.1 (6, 14), and BA.1 and BA.2 (13) variant, and some studies did not report the sub-lineage (7, 9, 11, 12, 15), or report the variant at all, though were very likely Omicron variant (8). Study characteristics are given in Table 1b.

The study from the UK is summarised in detail below ( $\underline{6}$ ). The remaining studies did not include more than 100 cases and are jointly summarised below.

#### Summaries of individual studies

Boucau and others reported the time to negative culture in n=32 Omicron BA.1 variant cases in the UK up to January 2022 ( $\underline{6}$ ). The median time from the first positive RT-PCR to negative culture was 5 days (IQR: 3 to 9 days), and the median time from the first positive RT-PCR or symptom onset (whichever was earlier) to negative culture was 8 days (IQR: 5 to 10 days).

#### Joint summary of remaining studies

Three studies measured the median time from symptom onset to negative viral cultures ( $\underline{8}$ ,  $\underline{10}$ ,  $\underline{11}$ ). All 3 studies estimated a median time of 4 days, with interquartile ranges varying between 3 to 5 or 6 days ( $\underline{8}$ ,  $\underline{10}$ ) and one to 7 days ( $\underline{11}$ ).

The remaining 6 studies reported on the proportion of cases (typically mild cases) with positive viral cultures on different days after either symptom onset or diagnosis (7, 9, 12 to 15). Culture positivity was highest in the first 5 days after symptom onset or diagnosis, with different studies reporting 46% at day 5 (14), a peak of around 30% at day 4 dropping to 10% at day 5 (7), 83% at day 5 (12), 51.5% and 86.5% at day one falling to 18.2% and 32.4% at day 5 depending on treatment (13), and 11.8% at days 0 to one rising to 41.7% at days 2 to 5 (15). Beyond 5 days after symptom onset or diagnosis, culture positivity fell, with different studies reporting 20% at day 7 after symptom onset (14), 52% at day 7,13.5% at day 10 and 8% at day 14 (12), and 18.8% at days 6 to 9 (15). The time from symptom onset or diagnosis to no further positive cultures was variable, taking 8 days (9), 10 days (14), 10 to 14 days (15), and 13 days (7) in different studies.

#### Incubation period

There were 8 included studies ( $\underline{4}$ ,  $\underline{5}$ ,  $\underline{16}$  to  $\underline{21}$ ) (2 preprints ( $\underline{18}$ ,  $\underline{21}$ )) that included evidence for the incubation period of COVID-19 Omicron variant. The incubation period is the time from COVID-19 exposure and symptom onset in a case, and studies would typically ask participants about exposure and symptom onset times to measure this, often pairing up or otherwise matching index and secondary cases.

Of these studies, 4 were conducted in Asia ( $\underline{5}$ ,  $\underline{17}$ ,  $\underline{19}$ ,  $\underline{21}$ ) and 4 in Europe ( $\underline{4}$ ,  $\underline{16}$ ,  $\underline{18}$ ,  $\underline{20}$ ), including one in the UK ( $\underline{20}$ ). All studies were conducted between December 2021 and April 2022. Three studies were prospective cohort studies ( $\underline{16}$ ,  $\underline{17}$ ,  $\underline{19}$ ), 3 studies were retrospective cohort studies ( $\underline{4}$ ,  $\underline{18}$ ,  $\underline{20}$ ), and 2 studies were cross-sectional studies ( $\underline{5}$ ,  $\underline{21}$ ). These studies included cases with COVID-19 Omicron variant BA.1 ( $\underline{4}$ ,  $\underline{5}$ ,  $\underline{19}$ ), BA.2 ( $\underline{21}$ ), and BA.1 and BA.2 ( $\underline{17}$ ,  $\underline{20}$ ), and some studies did not report the sub-lineage ( $\underline{16}$ ,  $\underline{18}$ ). Study characteristics are given in Table 1c.

One study from the UK included had a large sample size and is summarised in detail below (20). The remaining studies typically included fewer than 1,000 cases in total and are jointly summarised below.

#### Summaries of individual studies

Ward and others reported on the incubation period of n=124,948 cases (n=116,163 BA.1, n=8,785 BA.2 Omicron variant) in the UK up to February 2022 (<u>20</u>). The estimated mean incubation period for Omicron BA.1 was 3.67 days (95% CrI: 3.61 to 3.72 days) with an

estimated standard deviation (SD) of 3.14 days (95% CrI: 3.06 to 3.22 days). Similarly, the estimated mean incubation period for Omicron BA.2 was 3.48 days (95% CrI: 3.43 to 3.53 days) with an estimated SD of 2.90 days (95% CrI: 2.82 to 2.98 days). The mean incubation period of both BA.1 and BA.2 Omicron did not appear to vary substantially by age group.

#### Joint summary of remaining studies

The remaining 7 studies estimated the median ( $\underline{4}$ ,  $\underline{16}$ ,  $\underline{17}$ ,  $\underline{19}$ ,  $\underline{21}$ ) or mean incubation period ( $\underline{5}$ ,  $\underline{16}$ ,  $\underline{18}$ ). The median incubation period was estimated to be between 2 and 5 days with interquartile ranges between one and 6 days ( $\underline{4}$ ,  $\underline{16}$ ,  $\underline{17}$ ,  $\underline{19}$ ,  $\underline{21}$ ), with one study suggesting little difference between the Omicron BA.1 and BA.2 variants ( $\underline{17}$ ). The mean incubation period was estimated to be between 3 and 5 days ( $\underline{5}$ ,  $\underline{16}$ ,  $\underline{18}$ ), with one study estimating that the 95th percentile to be less than 7 days ( $\underline{5}$ ).

#### Latent period

There was one included study that included evidence for the latent period of COVID-19 Omicron variant (<u>5</u>). The latent period is the time from exposure to COVID-19 and detectable viral levels in a case (potentially indicating the capability to infect other people), and studies would typically ask about COVID-19 exposure and first positive test times to measure this. Study characteristics are given in <u>Table 1d</u>.

Xin and others estimated the latent period of n=114 cases with COVID-19 Omicron BA.1.1 variant between January 2022 and February 2022 in China (<u>5</u>). The estimated mean latent period was 3.1 days (95% credible interval [CrI]: 2.8 to 3.5 days), with 95% of cases developing detectable virus levels 5.9 days after infection (95% CrI: 5.3 to 6.8 days).

#### Serial interval and generation time

There were 9 included studies (<u>3</u>, <u>4</u>, <u>16 to 18</u>, <u>21 to 24</u>) (3 preprints (<u>18</u>, <u>21</u>, <u>22</u>)) that included evidence for the serial interval of COVID-19 Omicron variant. The serial interval is the time from symptom onset of an index case to symptom onset of a secondary case, and studies would typically ensure that transmission happened between individual cases, then ask about symptom onset times to measure this. The generation time is the time from infection of an index case and infection of a secondary case, and studies would typically estimate this in the same way as for serial interval.

Of these studies, 4 were conducted in Europe ( $\underline{3}$ ,  $\underline{4}$ ,  $\underline{16}$ ,  $\underline{18}$ ), 4 in Asia ( $\underline{17}$ ,  $\underline{21}$  to  $\underline{23}$ ), and one in the US ( $\underline{24}$ ). All studies were conducted between November 2021 and April 2022. Six studies were prospective cohort studies ( $\underline{3}$ ,  $\underline{16}$ ,  $\underline{17}$ ,  $\underline{22}$  to  $\underline{24}$ ), 2 studies were retrospective cohort studies ( $\underline{4}$ ,  $\underline{18}$ ), and one study was a cross-sectional study ( $\underline{21}$ ). These studies included cases with Omicron BA.1 ( $\underline{4}$ ,  $\underline{23}$ ), BA.2 ( $\underline{21}$ ), and BA.1 and BA.2 ( $\underline{3}$ ,  $\underline{17}$ ,  $\underline{22}$ ,  $\underline{24}$ ) variant COVID-19, and some studies did not report the sub-lineage of COVID-19 Omicron variant ( $\underline{16}$ ,  $\underline{18}$ ). Study characteristics are given in Table 1e.

One study included over 11,000 households and is summarised in more detail below ( $\underline{3}$ ). The remaining studies typically included fewer than 1,000 cases and are jointly summarised below.

#### Summaries of individual studies

An der Heiden and others reported the serial interval estimated from household transmission of COVID-19 Omicron variant BA.1 and BA.2 in n=11,512 households in Germany between January 2022 and May 2022 (3). The mean serial interval was estimated to be 3.61 days (95% CI: 3.56 to 3.66 days) across all cases, 3.88 days (95% confidence interval [CI]: 3.79 to 3.97 days) for Omicron BA.1 cases, and 3.39 days (95% CI: 3.30 to 3.49 days) for Omicron BA.2 cases.

#### Joint summary of remaining studies

Of the remaining studies, 4 studies reported on estimated median serial interval ( $\underline{4}$ ,  $\underline{17}$ ,  $\underline{21}$ ,  $\underline{24}$ ), 4 studies reported on estimated mean serial interval ( $\underline{16}$ ,  $\underline{18}$ ,  $\underline{22}$ ,  $\underline{23}$ ), and 2 studies reported on the median generation time ( $\underline{17}$ ) or mean forward generation interval ( $\underline{18}$ ). The median serial interval was estimated to be between 2 and 4 days, the interquartile ranges between one and 9 days ( $\underline{4}$ ,  $\underline{17}$ ,  $\underline{21}$ ,  $\underline{24}$ ), with one study suggesting little difference between Omicron BA.1 and BA.2 variants ( $\underline{17}$ ). The mean serial interval was estimated to be between 3 and 4 days ( $\underline{16}$ ,  $\underline{18}$ ,  $\underline{22}$ ,  $\underline{23}$ ), with one study suggesting child index cases may have shorter serial intervals (3.0 days) than adult index cases (5.0 days) ( $\underline{23}$ ). Both the median generation time and mean forward generation interval were estimated to be around 3 days ( $\underline{17}$ ,  $\underline{18}$ ).

#### Time to peak viral load

There were 6 included studies (<u>14</u>, <u>25 to 29</u>) (one preprint (<u>29</u>)) that included evidence for time to peak viral load of COVID-19 Omicron variant cases. The time to peak viral load is the time from exposure, first positive RT-PCR test, or symptom onset to the peak viral load. Studies would typically repeatedly test cases for COVID-19 using RT-PCR, recording the peak viral load as the time of the lowest Ct value to measure this. Higher viral loads, as measured by Ct values (lower Ct values indicate higher viral loads), may be indicative of a higher chance of being infectious, but high viral loads (or low Ct values) do not necessarily indicate infectiousness.

Of these studies, 2 were conducted in the US ( $\underline{27}$ ,  $\underline{29}$ ), and one each in Spain ( $\underline{26}$ ), Canada ( $\underline{28}$ ), Brazil ( $\underline{14}$ ), and South Korea ( $\underline{25}$ ). All studies were conducted between December 2021 and March 2022. Three studies were prospective cohort studies ( $\underline{14}$ ,  $\underline{28}$ ,  $\underline{29}$ ), and 3 studies were retrospective cohort studies ( $\underline{25}$  to  $\underline{27}$ ). These studies included cases with COVID-19 Omicron variant BA.1 ( $\underline{14}$ ,  $\underline{27}$  to  $\underline{29}$ ), and BA.1 and BA.2 ( $\underline{26}$ ), and one study did not report the sub-lineage ( $\underline{25}$ ). Study characteristics are given in <u>Table 1f</u>.

One study included over 5,000 cases and is summarised in more detail below (25). The remaining studies included fewer than 1,000 cases in total and are jointly summarised below.

#### Summaries of individual studies

Choi and others reported time to peak viral load for n=5,187 Omicron variant cases in South Korea in January 2022 ( $\underline{25}$ ). The results suggested the peak viral load occurred 2.4 days (95% CI: 2.2 to 2.5 days) after symptom onset.

#### Joint summary of remaining studies

The remaining 5 studies estimated the mean time to peak viral load (<u>14</u>, <u>26 to 29</u>). The studies suggested the mean time to peak viral load was between 2 and 5 days for the Omicron variant BA.1 (<u>14</u>, <u>26 to 29</u>), with one study suggesting the Omicron variant BA.2 may have a shorter mean time to peak viral load (one day) (<u>26</u>).

#### Time to viral clearance

There were 53 included studies (<u>12</u>, <u>14</u>, <u>15</u>, <u>27</u>, <u>29 to 77</u>) (15 preprints (<u>29 to 31</u>, <u>34</u>, <u>36</u>, <u>37</u>, <u>39</u>, <u>54</u>, <u>55</u>, <u>57</u>, <u>58</u>, <u>60</u>, <u>66</u>, <u>69</u>, <u>77</u>)) that included evidence for time to viral clearance of COVID-19 Omicron variant cases. The time to viral clearance is the time either from exposure, first positive RT-PCR test, or symptom onset to the last positive RT-PCR test or first negative RT-PCR test. These studies typically repeatedly tested cases for COVID-19 using RT-PCR, recording the date at which the last positive or first negative RT-PCR test was conducted, sometimes requiring a repeat negative RT-PCR test to ensure viral clearance. A detectable viral load (as measured using RT-PCR) does not necessarily indicate that a case is infectious.

Of these studies, 33 were conducted in China (33, 34, 39, 40, 43 to 48, 51 to 55, 58, 59, 61 to 72, 74 to 77), 8 in Europe (12, 30, 32, 35, 38, 49, 56, 57), 6 in the US (27, 29, 36, 37, 41, 42), 4 in the rest of Asia (60, 73) (15, 50), and 2 in Brazil (14, 31). Most studies were conducted between March 2022 and July 2022, though some studies were conducted earlier, and some studies were conducted up to August and October 2022 (34, 44). In total, 28 studies were prospective cohort studies (12, 14, 29 to 33, 36 to 38, 41 to 44, 46, 47, 49, 53, 54, 56 to 58, 63, 65, 67, 72, 74, 76) (6 of which were randomised controlled trials [RCTs] or quasi-experimental studies but reported as prospective cohort studies (31, 32, 46, 54, 58, 63)), and 25 studies were retrospective cohort studies (15, 27, 34, 35, 39, 40, 45, 48, 50 to 52, 55, 59 to 62, 64, 66, 68 to 71, 73, 75, 77). These studies included cases with COVID-19 Omicron variant BA.1 (14, 27, 29, 49), BA.2 (39, 40, 45, 46, 51, 54, 55, 57, 60 to 62, 64, 67 to 73, 76, 77), BA.1 and BA.2 (38), and BA.5 (44) , and some studies did not report the sub-lineage (12, 15, 30, 32 to 37, 41 to 43, 47, 48, 50, 52, 53, 56, 58, 59, 63, 65, 66, 74, 75), or report the variant at all, though were very likely Omicron variant (31). Study characteristics are given in Table 1g.

Some studies included both treated and untreated arms, for instance in trials of different drugs compared with standard care or no treatment. In these studies, only the no treatment groups are summarised below, although results for both groups are available in <u>Table 1g</u>. Studies in this section are split into those including cases from the general population, and those that only

included cases who were at high-risk from COVID-19, including cases who were hospitalised (either for COVID-19 or another condition), and cases with immunodeficiency.

One study reported on the Omicron variant BA.5, and 3 studies included over 25,000 cases (with likely substantial overlap in cases between these studies), and all are individually summarised below. The remaining studies typically included fewer than 1,000 cases in total and are jointly summarised below. The jointly summarised studies are split into studies including cases from the general population (including studies of cases quarantining in hospitals, particularly in China) and studies including hospitalised, immunodeficient, or otherwise high-risk cases.

#### Summaries of individual studies

Li and others reported the duration of viral shedding (Ct value of less than 35 on a nucleic acid test) for n=27 cases with COVID-19 Omicron variant BA.5 and n=51 cases with Omicron variant BA.2 in August 2022 in China (44). Of the Omicron BA.5 cases, 96.30% shed virus for at least 7 days, 88.89% shed virus for at least 10 days, and 48.15% shed virus for at least 14 days. Of the Omicron BA.2 cases, 54.38% shed virus for at least 7 days, 21.92% shed virus for at least 10 days, and 4.51% shed virus for at least 14 days.

The following 3 studies reported on cases admitted to shelter hospitals in Shanghai, China between March and May 2022, and therefore likely included a substantial overlap in cases.

Pei and others reported time between first positive and first negative RT-PCR test for n=198,262 asymptomatic and mild cases with likely COVID-19 Omicron variant BA.2 between March and May 2022 in China (51). The median time from first positive to first negative RT-PCR test was 8.29 days (IQR: 5.33 to 11.27 days) in all cases, and 9.29 days (IQR: 6.33 to 12.28 days) in n=33,896 cases with comorbidities.

Yin and others reported the time between first positive and first negative RT-PCR test for n=199,590 asymptomatic or mild cases with Omicron variant BA.2 between March and May 2022 in China (70). The mean time from illness onset to negative RT-PCR test was 7.17 days (SD: 3.42 days).

Ying-Hao and others reported the time between first positive and first negative RT-PCR test for n=25,168 asymptomatic or mild cases with Omicron variant BA.2 between April and May 2022 in China (71). The median time from first positive to first negative RT-PCR test was 6 days (IQR: 4 to 9 days).

#### Joint summary of remaining studies: cases from the general population

In total, 19 studies reported the duration of positivity on a COVID-19 test (typically RT-PCR test, but also rapid antigen tests, and other nucleic acid tests), usually reporting either the mean or median time from either symptom onset or diagnosis to the first of 2 negative tests (<u>27</u>, <u>29 to 31</u>,

<u>36</u>, <u>37</u>, <u>40</u>, <u>41</u>, <u>46</u>, <u>50</u>, <u>57</u>, <u>58</u>, <u>60</u>, <u>61</u>, <u>63 to 65</u>, <u>72</u>, <u>73</u>). Many studies did not specify the outcome, making comparison between studies more difficult. Nonetheless, the time to viral clearance was markedly different across studies, ranging from 4 to 29 days, though most studies estimated viral clearance to take around 7 to 11 days (<u>27</u>, <u>29 to 31</u>, <u>36</u>, <u>37</u>, <u>40</u>, <u>50</u>, <u>57</u>, <u>60</u>, <u>63</u>, <u>72</u>, <u>73</u>).

The remaining 7 studies including cases from the general population reported the proportion of cases who were still positive on a COVID-19 test (typically RT-PCR test, but also rapid antigen tests, and other nucleic acid tests) at certain days after symptom onset or diagnosis (<u>12</u>, <u>14</u>, <u>15</u>, <u>32</u>, <u>42</u>, <u>45</u>, <u>62</u>). As with the studies above, many studies did not specify the outcome and different time periods were reported on, making comparison between studies more difficult. However, all studies reported the proportion of positive tests at day 7, which varied between 20.4% and 100%, which most studies reporting positive test proportions between 87% and 100% (<u>12</u>, <u>14</u>, <u>15</u>, <u>32</u>).

## Joint summary of remaining studies: hospitalised, immunodeficient, and other high-risk cases

In total, 21 studies reported the duration of positivity on a COVID-19 test (typically RT-PCR test, but also rapid antigen tests, and other nucleic acid tests), usually reporting either the mean or median time from either symptom onset, diagnosis, or hospitalisation to the first of 2 negative tests for cases who were hospitalised (for COVID-19 or another condition), immunodeficient, or otherwise at high-risk from COVID-19 (33 to 35, 39, 43, 47 to 49, 52, 53, 55, 56, 59, 66 to 69, 74 to 77). Many studies did not specify the outcome, and cases had variable severity of COVID-19 and other conditions, making comparison between studies more difficult. In general, however, the time to viral clearance was longer than for cases from the general population, ranging from 4 to 21 days across studies, though most studies estimated viral clearance to take around 10 to 15 days (33, 35, 39, 48, 49, 52, 53, 55, 56, 66, 67, 69, 75 to 77). Studies comparing hospitalised cases with and without chronic kidney disease or cases on haemodialysis suggested viral clearance was quicker in cases without chronic kidney disease or on haemodialysis (39, 66, 75).

Two further studies including cases who were hospitalised (for COVID-19 or another condition), immunodeficient, or otherwise at high-risk from COVID-19 reported the proportion of cases who were still positive on a COVID-19 test (typically RT-PCR test, but also rapid antigen tests, and other nucleic acid tests) at certain days after symptom onset or diagnosis (<u>38,54</u>). In one study, 27.9% of immunodeficient cases still showed viral shedding (threshold set at 1,000,000 SARS-CoV-2 RNA copies per ml) 21 days after starting sotrovimab compared with 7.1% for immunocompetent cases (<u>38</u>). In another study, 31.1% of cases had a negative RT-PCR test within 7 days of entry into the study (<u>54</u>).

#### Viral load over time

There were 9 included studies (7, 25, 78 to 84) (4 preprints (78, 80, 83, 84)) that included evidence for the viral load over time for COVID-19 Omicron variant cases. These studies typically repeatedly tested cases for COVID-19 using RT-PCR, and typically expressed the results in a graph indicating mean or median viral loads of participants over a course of infection, either from diagnosis or from symptom onset (or both). For these studies, if no data were presented in the text (and the study reported in a previous section), the graphs were briefly summarised to give an overall impression of how viral load changed over the course of an infection. As with time to viral clearance, a detectable viral load (as measured using RT-PCR) does not necessarily indicate that a case is infectious.

Of these studies, 4 were conducted in Europe (<u>78</u>, <u>80</u>, <u>82</u>, <u>83</u>), including 2 that were conducted in the UK (<u>78</u>, <u>83</u>), 3 were in Asia (<u>25</u>, <u>79</u>, <u>84</u>), and 2 were in the US (<u>7</u>, <u>81</u>). All studies were conducted between November 2021 and May 2022. Five studies were prospective cohort studies (<u>7</u>, <u>80 to 83</u>), 3 studies were retrospective cohort studies (<u>25</u>, <u>79</u>, <u>84</u>), and one study was a cross-sectional study (<u>78</u>). These studies included cases with COVID-19 Omicron variant B.1.1529 (<u>84</u>), BA.1 (<u>81</u>), BA.2 (<u>79</u>), and BA.1 and BA.2 (<u>78</u>, <u>80</u>, <u>82</u>, <u>83</u>), and some studies did not report the sub-lineage (<u>7</u>, <u>25</u>). Study characteristics are given in <u>Table 1h</u>.

One study from England including over a million cases ( $\underline{78}$ ), and a study from South Korea include over 5,000 cases ( $\underline{25}$ ), and both are summarised in detail below. The remaining studies included fewer than 250 cases and are jointly summarised below.

#### Summaries of individual studies

Funk and others (preprint) reported on the Ct values over time for n=1,212,234 cases (n=1,083,976 BA.1, n=128,258 BA.2 Omicron variant) in England between December 2021 and January 2022 (<u>78</u>). Data extracted from a figure suggested Ct values decreased (viral load increased) in the first 2 days after symptom onset, then increased (viral load decreased) from days 2 to 6 (data was truncated at day 6). The study also suggested that BA.2 infections had higher Ct values (lower viral loads) than BA.1 infections, cases with reinfections had higher Ct values (lower viral loads) than cases with no known previous infection, and there was no clear difference in Ct values between people with different doses of vaccine.

Choi and others reported on Ct values over time for n=5,187 Omicron variant cases in South Korea in January 2022 (25). Data extracted from a figure suggested Ct values decreased (viral load increased) up to between 3 and 4 days after symptom onset, increased (viral load decreased) to between 12 and 13 days after symptom onset, then decreased again (viral load increased) up to 17 days after symptom onset (data truncated after this).

#### Joint summary of remaining studies

In the remaining studies, Ct values typically decreased (viral load increased) or remained steady after symptom onset for one (7, 79) or 2 to 3 days (80, 82 to 84), then increased (viral

load decreased), either until the end of testing (7, 79 to 84), or until a slight decrease in Ct values (increase in viral load) between 15 and 17 days for the Omicron BA.2 variant (83). One study also suggested that, for the Omicron BA.2 variant, Ct values increased (viral load decreased) up to one to 3 days after symptom onset, then decreased (viral load increased) up to 7 to 9 days after symptom onset, then increased again (viral load decreased) (82).

### Summary

In total, 82 studies provided evidence on different measures of infectious period.

The 3 studies measuring transmission period directly suggested most transmission events happened around symptom onset, with the vast majority of transmission events happening up to 5 days after symptom onset (100% in one study, 81% in another study, not reported in the third study). However, these studies were reasonably small and therefore the results may be imprecise.

The 10 studies measuring viral culture positivity over time suggested that viral culture positivity was highest in the first 5 days after symptom onset or diagnosis before falling, with variable times for all cases to have negative viral cultures (between 8 and 14 days in different studies). Three of the studies were conducted in healthcare workers, 2 in hospitalised patients or care home residents and the remaining studies in community settings. Whilst viral culture positivity results do not provide direct evidence on the risk of transmission, they indicate potential infectivity.

The 8 studies measuring incubation period (exposure to symptom onset) suggested the median and mean incubation period was between 2 and 5 days, with a large UK study estimating a mean incubation period of between 3 and 4 days.

The one study measuring latent period (exposure to detectable viral levels) suggested the mean latent prior was around 3 days, with 95% of cases developing detectable virus levels around 6 days after infection.

The 9 studies measuring serial interval (symptom onset in an index case to symptom onset in a secondary case) and generation time (exposure in an index case to exposure in a secondary case) suggested the median and mean serial interval was between 2 and 4 days (though this can be variable, with interquartile ranges between one and 9 days), and generation times of around 3 days.

The 6 studies measuring time to peak viral load and 9 studies measuring viral load over time (one study reported on both outcomes) suggested peak viral loads occurred a median and mean of 2 to 5 days after symptom onset, and this may be quicker in Omicron BA.2 compared with BA.1 variant cases. Two of the studies were conducted in healthcare workers, one in hospitalised patients and the remaining studies in community settings.

The 53 studies measuring time to viral clearance (typically positive to negative COVID-19 test) suggested that there were substantial differences in viral clearance times between cases and populations, although the differences in measurement of time to viral clearance between studies may have contributed to these differences. In general, however, most studies of the general population estimated viral clearance to take around 7 to 11 days, and most studies of hospitalised, immunodeficient, and other high-risk cases estimated viral clearance to take around 10 to 15 days. Detectable viral load does not necessarily indicate that a case is infectious.

Overall, the evidence suggests that COVID-19 Omicron variant cases were most infectious up to 5 days after symptom onset, but could potentially be infectious for longer, especially if hospitalised, immunocompromised, or otherwise high-risk cases. However, while some studies included substantial numbers of cases, most studies included relatively few cases, and the majority of studies only included Omicron BA.1 and BA.2 variant cases.

### Table 1. Studies for infectious period

Acronyms: CI = confidence interval, CrI = credible interval, IQR = interquartile range, RT-PCR = reverse transcriptase polymerase chain reaction, SD = standard deviation

#### Table 1a. Transmission period

Study	Country, time period	Study type	Participants	Outcome
An der Heiden ( <u>3</u> )	Germany, January to May 2022	Prospective cohort	n=11,512 households (Omicron BA.1 and BA.2 variants)	81% of transmission occurred by day 5 of symptom onset in occurred by day 10 of symptom onset in the index case
Del Aguila-Mejia ( <u>4</u> )	Spain, December 2021	Retrospective cohort	n=622 index cases, n=455 secondary cases (Omicron BA.1 variant)	Median transmission period after symptom onset: 0 days (IQ
Xin ( <u>5</u> )	China, January to February 2022	Cross- sectional	n=113 pairs of cases (Omicron BA.1.1 variant)	Proportion of transmissions occurring before symptom onset. Transmission peaked at symptom onset, all observed transmisymptom onset

#### Table 1b. Culture positivity over time

Study	Country, time period	Study type	Participants	Outcome
Boucau ( <u>6</u> )	UK, July 2021 to January	Prospective cohort	n=34 cases (Omicron BA.1	Median time from the first positive RT-PCR to negative cu
	2022		variant)	Median time from the first positive RT-PCR or symptom or culture: 8 days (IQR: 5 to 10 days)
Bouton ( <u>7</u> )	US, November 2021	Prospective	n=92 university cases (n=75	Culture positive more than 5 days from diagnosis: n=10 (1
	onwards (end date not stated)	cohort	Omicron, n=17 Delta)	Culture positive more than 5 days from symptom onset: n=
Gilbert ( <u>8</u> ) (Preprint,	US, November 2021 to	Prospective	n=54 university cases (variant	Median time to negative culture: 4 days (IQR: 3 to 5.75 da
conference abstract)	March 2022	cohort	not stated)	No statistically significant association between time to neg COVID-19 vaccine (p=0.34)
Jang ( <u>9</u> )	South Korea, December 2021	Prospective cohort	n=11 hospitalised cases (Omicron variant)	Last positive viral culture after symptom onset: 8 days
Jung ( <u>10</u> )	South Korea, February to March 2022	Prospective cohort	n= 32 healthcare worker asymptomatic and mild cases (Omicron B.1.1.529 variant)	Median time from symptom onset to negative culture: 4 da
				Median time from diagnosis to negative culture: 3 days (95
Kang ( <u>11</u> )	South Korea, February to May 2022	Prospective cohort	n=67 adult cases (Omicron variant)	Median time from symptom onset to negative culture: 4 da
Keske ( <u>12</u> )	Turkey, January to February	Prospective	n=55 healthcare worker non-	Positive viral cultures (day 5): n=44 of 53 (83%)
	2022	cohort	severe cases (Omicron	Positive viral cultures (day 7): n=26 of 50 (52%)
			variant)	Positive viral cultures (day 10): n=7 of 52 (13.5%)
				Positive viral cultures (day 14): n=4 of 50 (8%)
Kim ( <u>13</u> )				Positive viral cultures (day 1, nirmatrelvir and ritanavir grou

n the index case, and 95% of transmission

IQR: -1 to 2 days)

et: 33.6% (95% Crl: 24.8% to 42.5%)

smission events occurring within 5 days after

culture: 5 days (IQR: 3 to 9 days)

onset (whichever was earlier) to negative

(11%)

n=16 of 92 (17%)

days)

egative culture and time since last dose of

days (95% CI: 3 to 5 days) (95% CI: 3 to 4 days)

days (IQR: 1 to 7 days)

roup): 51.5%

Study	Country, time period	Study type	Participants	Outcome
	South Korea, October 2021 to May 2022	Prospective cohort	n=33 mild cases (nirmatrelvir and ritanavir group, Omicron BA.1 and BA.2 variants)	Positive viral cultures (day 5, nirmatrelvir and ritanavir gro
			n=37 mild cases (supportive	Positive viral cultures (day 1, supportive care group): 86.5
			care group, Omicron BA.1 and BA.2 variants)	Positive viral cultures (day 5, supportive care group): 32.4
Luna-Muschi ( <u>14</u> )	Brazil, January 2022	Prospective cohort	n=30 vaccinated healthcare worker mild cases (Omicron BA.1 variant)	Positive viral culture (day 5 after symptom onset): n=11 of
				Positive viral culture (day 7 after symptom onset): n=6 of 3
				Positive viral culture (day 10 after symptom onset): n=0 of
Takahashi ( <u>15</u> )	Japan, November to	Retrospective	n=18 asymptomatic and mild cases (Omicron variant)	Positive viral culture (0 to 1 days after diagnosis): n=2 of 1
	December 2021	cohort		Positive viral culture (2 to 5 days after diagnosis): n=5 of 1
				Positive viral culture (6 to 9 days after diagnosis): n=3 of 1
				Positive viral culture (10 to 14 days after diagnosis): n=0 o
				Positive viral culture (15 days and more after diagnosis): n

#### Table 1c. Incubation period

Study	Country, time period	Study type	Participants	Outcome
Backer ( <u>16</u> )	The Netherlands,	Prospective	n=258 cases (Omicron variant)	Mean incubation period: 3.2 days (95% Crl: 2.9 to 3.6 days),
	December 2021 to January 2022	cohort		Median incubation period: 2.8 days (95% Crl: 2.5 to 3.2 days
Del Aguila-Mejia ( <u>4</u> )	Spain, December 2021	Retrospective cohort	n=622 index cases, n=455 secondary cases (Omicron BA.1 variant)	Median incubation period: 3 days (IQR: 1 to 4 days)
Mefsin ( <u>17</u> ) 2022 to March 2022	Hong Kong, January	Prospective	n=57 cases (Omicron BA.1 variant)	Median incubation period (Omicron BA.1, n=57): 4.38 days (
	cohort	n=23 cases (Omicron BA.2 variant)	Median incubation period (Omicron BA.2, n=23, Gamma dist days)	
Park ( <u>18</u> ) (Preprint)	The Netherlands, November 2021 to January 2022	Retrospective cohort	n=258 cases (Omicron variant)	Mean Incubation period: 4.2 days (95% CI: 3.6 to 4.9 days
Tanaka ( <u>19</u> )	Japan, January 2022	Prospective cohort	n=172 cases (BA.1 Omicron variant)	Median incubation period: 2.6 days (95% CI: 2.4 to 2.8 days) 3.5 days (95% CI: 3.2 to 3.9 days)
Ward ( <u>20</u> )	UK, May 2020 to February 2022	Retrospective cohort	n=124,948 cases (n=116,163 Omicron BA.1, n=8,785 Omicron BA.2 variant)	Mean incubation period (Omicron BA.1): 3.67 days (95% Crl Crl: 3.06 to 3.22 days)
				Mean incubation period (Omicron BA.2): 3.48 days (95% Crl Crl: 2.82 to 2.98 days)
				Mean incubation periods did not appear to vary much by age

roup): 18.2%

5.5% 2.4% of 24 (46%) of 30 (20%) of 30 (0%) f 17 (11.8%) f 12 (41.7%) f 16 (18.8%) 0 of 17 (0%) :: n=0 of 10 (0%)

s), SD: 2.2 days (95% Crl: 1.9 to 2.5 days) hys)

s (95% CI: 3.88 to 4.87 days) istribution): 4.27 days (95% CI: 3.29 to 5.02

ys), IQR: 1.9 days (95% CI: 1.7 to 2.1 days) to

Crl: 3.61 to 3.72 days), SD: 3.14 days (95%)

Crl: 3.43 to 3.53 days), SD: 2.90 days (95%

ge group

Study	Country, time period	Study type	Participants	Outcome
Wei ( <u>21</u> ) (Preprint)	China, April 2022	Cross-sectional	n=52 cases (Omicron BA.2 variant)	Median incubation period: 4.4 days (IQR: 3.1 to 6.0 days, 95t
Xin ( <u>5</u> )	China, January to February 2022	Cross-sectional	n=114 cases (Omicron BA.1.1 variant)	Mean incubation period: 3.8 days (95% Crl: 3.5 to 4.1 days), 6.9 days)

#### Table 1d. Latent period

Study	Country, time period	Study type	Participants	Outcome
Xin ( <u>5</u> )	China, January to	Cross-sectional	n=114 cases (Omicron BA.1.1	Mean latent period: 3.1 days (95% Crl: 2.8 to 3.5 days), 95th
	February 2022		variant)	days)

#### Table 1e. Serial interval and generation time

Study	Country, time period	Study type	Participants	Outcome
Ali ( <u>22</u> ) (Preprint)	Hong Kong, January to February 2022	Prospective cohort	n=229 case pairs (n=204 cases pairs Omicron, n=25 case pairs Delta variant)	Mean serial interval (all cases): 3.6 days (95% Crl: 3.5 to 3.7 days)
			n=30 case pairs (Omicron BA.1 variant)	Mean serial interval (Omicron BA.1): 3.3 days, SD: 2.0 days
			n=174 case pairs (Omicron BA.2 variant)	Mean serial interval (Omicron BA.2): 3.6 days, SD: 1.8 days
An der Heiden ( <u>3</u> )	Germany, January to May	Prospective	n=11,512 households (Omicron	Mean serial interval (Omicron): 3.61 days (95% CI: 3.56 to 3
	2022	cohort	BA.1 and BA.2 variants)	Mean serial interval (Omicron BA.1): 3.88 days (95% CI: 3.79
				Mean serial interval (Omicron BA.2): 3.39 days (95% CI: 3.30
Backer ( <u>16</u> )	The Netherlands, December 2021 to January 2022	Prospective cohort	n=480 household case pairs (Omicron variant)	Mean serial interval (household case pairs): 3.0 days (SD: 2.
Del Aguila-Mejia ( <u>4</u> )	Spain, December 2021	Retrospective cohort	n=622 index cases, n=455 secondary cases (Omicron BA.1 variant)	Median serial Interval: 4 days (IQR: 3 to 6 days)
Kim ( <u>23</u> )	South Korea, November to December 2021	Prospective	n=73 case pairs (Omicron variant)	Mean serial interval: 3.78 days (95% Crl: 3.02 to 4.54 days),
		cohort		Mean serial interval (child index cases): 3.0 days
				Mean serial interval (adult index cases): 5.0 days
Mefsin ( <u>17</u> )	Hong Kong, January 2022	Prospective	n=57 cases (Omicron BA.1 variant)	Median generation time (Omicron BA.1, n=45): 2.38 days (95
	to March 2022	cohort		Median serial Interval (Omicron BA.1, n=30): 3.15 days (95%
			n=23 cases (Omicron BA.2 variant)	Median serial Interval (Omicron BA.2, n=13, Gamma distribu
Park ( <u>18</u> )	The Netherlands,	Retrospective	n=258 cases (Omicron variant)	Mean serial interval: 3.1 days (95% CI: 2.9 to 3.3 days)
(Preprint)	November 2021 to January 2022	cohort		Mean forward generation interval: 3.0 days (95% CI: 2.7 to 3

95th percentile: 8.3 days)
s), 95th percentile: 6.2 days (95% Crl: 5.7 to
th percentile: 5.9 days (95% Crl: 5.3 to 6.8
9.7 days), SD: 3.4 days (95% Crl: 3.3 to 3.5
'S
'S
3.66 days)
.79 to 3.97 days)
.30 to 3.49 days)
2.3 days)
s), SD: 3.33 days (95% Crl: 2.56 to 4.09 days)
95% CI: 2.01 to 2.80 days)
5% CI: 2.49 to 3.92 days)
oution): 2.52 days (95% CI: 1.68 to 3.55 days)
3.2 days)

Study	Country, time period	Study type	Participants	Outcome
Wei ( <u>21</u> ) (Preprint)	China, April 2022		n=234 transmission pairs (Omicron BA.2 variant)	Median serial interval: 4.0 days (IQR: 1.4 to 6.5 days)
Weil ( <u>24</u> )	US, December 2021 to February 2022		n=37 university clusters (Omicron BA.1 and BA.2 variants)	Median serial interval: 2 days (IQR: 1 to 9 days)

#### Table 1f. Time to peak viral load

Study	Country, time period	Study type	Participants	Outcome
Choi ( <u>25</u> )	South Korea, January 2022	Retrospective cohort	n=5,187 (Omicron variant)	Peak viral load after symptom onset: 2.4 days (95% CI: 2.2 to
De Michelena ( <u>26</u> )	Spain, February to March 2022	Retrospective cohort	n=130 cases (Omicron BA.1 variant)	Time to peak viral load (Omicron BA.1): 3 to 5 days
		n=147 cases (Omicron BA.2 variant)	Time to peak viral load (Omicron BA.2): 1 day	
Hay ( <u>27</u> ) US, December 2021 to January 2022		Retrospective cohort	n=878 cases (Omicron BA.1 variant) (number of cases not split by vaccination status)	Mean time to peak viral load (proliferation time, fully or partial to 4.0 days)
				Mean time to peak viral load (proliferation time, booster vacci days)
Kandel ( <u>28</u> )	Canada, December 2021	Prospective cohort	n=41 adult cases (Omicron BA.1 variant)	Mean time from symptom onset to peak viral load: 2.97 days
	to January 2022			Mean time from first positive test to peak viral load: 2.89 days
Luna-Muschi ( <u>14</u> )	Brazil, January 2022	Prospective cohort	n=30 vaccinated healthcare worker mild cases (Omicron BA.1 variant)	Time from symptom onset to lowest Ct value: 5 days
Sikka ( <u>29</u> ) (Preprint)	US, February 2021 to January 2022	Prospective cohort	n=37 cases (Omicron BA.1 variant)	Average time to peak viral load: 1.97 days

#### Table 1g. Time to viral clearance

Study	Country, time period	Study type	Participants	Outcome
Anastasiou ( <u>30</u> )	Germany, January 2022	Prospective	n=72 cases (Omicron variant)	Median time to negative test: 7 days (IQR: 2 to 14 days)
(Preprint)		cohort		Median time to Ct value above 30: 7 days (IQR: 2 to 8 days)
Cabral (31)Brazil, January to May(Preprint)2022	Prospective cohort (RCT)	n=143 mild cases (variant not stated)	Mean time to first negative RT-PCR test (AZVUDINE group): figure)	
			n=138 mild cases (variant not stated)	Mean time to first negative RT-PCR test (placebo group): 8.2
Cegolon ( <u>32</u> )		Prospective cohort (RCT)	n=50 symptomatic or mild cases (Tonimer Lab Panthexyl 800 group, Omicron variant)	Viral shedding time of 7 or more days (Tonimer Lab Panthex)
			n=58 symptomatic or mild cases (control group, Omicron variant)	Viral shedding time of 7 or more days (control group): n=52 c

to 2.5 days)
ially vaccinated cases): 3.6 days (95% Crl: 3.3
ccinated cases): 4.0 days (95% CrI: 3.8 to 4.3
'S
ys
;)
): 5.55 days, SD: 0.45 days (estimated from
.27, SD: 0.59 days (estimated from figure)
exyl 800 group): n=33 of 50 (66%)
of 58 (89%)

Study	Country, time period	Study type	Participants	Outcome	
Chen ( <u>33</u> )	China, March to May	Prospective	n=847 hospitalised cases (Omicron	Median viral shedding time: 13 days (IQR: 10 to 16 days)	
	2022	cohort	variant)	Viral shedding time was longer for older cases ( $p=0.037$ ) and to 5 ( $p<0.001$ ) or heart conditions ( $p=0.030$ ), and shorter for ( $p=0.001$ )	
Chen ( <u>34</u> ) (Preprint)	China, August to October 2022	Retrospective cohort	n=166 hospitalised cases (Azvudine group, Omicron variant)	Median time from treatment to first negative RT-PCR (Azvud	
			n=41 hospitalised cases (control group, Omicron variant)	Median time from treatment to first negative RT-PCR (contro	
Colaneri ( <u>35</u> )	Italy, December 2021 to	Retrospective	n=49 mild and moderate cases with	Median duration of viral load (untreated): 15 days	
	May 2022	cohort	hematologic malignancies	Median duration of viral load (treated with Remdesivir): 21 da	
			(Omicron variant)	Median duration of viral load (treated with Sotrovimab): 17 da	
				Median duration of viral load (treated with Molnupiravir): 17 d	
Cosimi ( <u>36</u> ) (Preprint)	US, January to February 2022	Prospective cohort	n=40 cases (Omicron variant)	Median time from COVID-19 diagnosis or start of symptonic antigen test: 9 days	
Dai ( <u>37</u> ) (Preprint)	US, March to May 2022	Prospective cohort	n=11 cases (Nirmatrelvir-Ritonavir treated, Omicron variant)	Median time from first positive test (diagnosis) to last positive (Nirmatrelvir-Ritonavir treated): 4 days	
			n=25 cases (not treated, Omicron variant)	Median time from first positive test (diagnosis) to last positive treated): 7 days	
Gliga ( <u>38</u> )	Germany, January to February 2022	Prospective cohort	n=43 immunodeficient cases (Omicron BA.1 and BA.2 variants)	Proportion of cases with viral shedding 21 days after starting cases (27.9%)	
			n=14 immunocompetent cases (Omicron BA.1 and BA.2 variants)	Proportion of cases with viral shedding 21 days after starting cases (7.1%)	
Guo ( <u>39</u> ) (Preprint)	China, March to May 2022	Retrospective cohort	n=470 hospitalised cases with chronic kidney disease (Omicron BA.2 variant)	Median time to negative RT-PCR test (cases with chronic kic	
			n=1,508 hospitalised cases without chronic kidney disease (Omicron BA.2 variant)	Median time to negative RT-PCR test (cases without chronic days)	
Hay ( <u>27</u> )	US, December 2021 to January 2022	Retrospective cohort	n=878 cases (Omicron BA.1 variant) (number of cases not split	Mean time to negative RT-PCR test (fully or partially vaccina days)	
			by vaccination status)	Mean time to negative RT-PCR test (booster vaccinated cas	
Hua ( <u>40</u> )	China, July 2022	Retrospective	n=225 adult cases (Omicron	Median duration of viral shedding: 11.0 days (IQR: 9.0 to 13.	
		cohort	BA.2.38 variant)	No statistically significant difference in duration of viral shedo and booster vaccinated cases (p=0.85)	
Keske ( <u>12</u> )	Turkey, January to	Prospective	n=55 healthcare worker non-severe	Positive RT-PCR (day 5): n=53 of 55 (96.4%)	
	February 2022	cohort	cases (Omicron variant)	Positive RT-PCR (day 7): n=48 of 55 (87.3%)	
				Positive RT-PCR (day 10): n=41 of 55 (74.5%)	

and cases with chronic kidney disease stage 4 or cases with full or booster vaccinations

udine group): 5 days (IQR: 1 to 7 days)

trol group): 6 days (IQR: 5 to 7 days)

days

days

days

ns (whichever came first) to first negative rapid

ive RT-PCR (Ct value less than 35)

ive RT-PCR (Ct value less than 35) (not

ng sotrovimab (immunodeficient): n=12 of 43

ng sotrovimab (immunocompetent): n=1 of 14

kidney disease): 13 days (IQR: 8 to 18 days)

nic kidney disease): 10 days (IQR: 7 to 14

nated cases): 6.2 days (95% Crl: 5.8 to 6.6

ases): 8.4 days (95% Crl: 8.0 to 8.7 days)

3.0 days)

dding between unvaccinated, fully vaccinated

Study	Country, time period	Study type	Participants	Outcome	
				Positive RT-PCR (day 14): n=23 of 55 (41.8%)	
Kojima ( <u>41</u> )	US, December 2021	Prospective cohort	n=734 cases (Omicron variant)	Duration of RT-PCR positivity: 14.3 days (SD: 7.0 days)	
_efferts ( <u>42</u> ) US, January to February		Prospective	n=564 symptomatic cases	Positive rapid antigen test (5 days since symptom onset): n=142 of 179 (79.3%)	
	2022	cohort	(Omicron variant)	Positive rapid antigen test (6 days since symptom onset): n=80 of 121 (66.1%)	
				Positive rapid antigen test (7 days since symptom onset): n=74 of 111 (66.7%)	
				Positive rapid antigen test (8 days since symptom onset): n=39 of 93 (41.9%)	
				Positive rapid antigen test (9 days since symptom onset): n=26 of 60 (43.3%)	
			n=165 asymptomatic cases	Positive rapid antigen test (5 days since positive test): n=18 of 58 (31.0%)	
			(Omicron variant)	Positive rapid antigen test (6 days since positive test): n=11 of 45 (24.4%)	
				Positive rapid antigen test (7 days since positive test): n=1 of 33 (3.0%)	
				Positive rapid antigen test (8 days since positive test): n=4 of 19 (21.1%)	
				Positive rapid antigen test (9 days since positive test): n=1 of 10 (10.0%)	
Li ( <u>43</u> )	China, March to April 2022	Prospective cohort	n=175 hospitalised adult cases (nirmatrelvir and ritonavir started 5 or fewer days after symptom onset group, Omicron variant)	Median time from first positive to negative RT-PCR test (nirmatrelvir and ritonavir started 5 or fewer days after symptom onset group): 10 days (IQR: 7 to 12 days)	
			(I n	n=83 hospitalised adult cases (nirmatrelvir and ritonavir started more than 5 days after symptom onset group, Omicron variant)	Median time from first positive to negative RT-PCR test (nirmatrelvir and ritonavir started more than 5 days after symptom onset group): 15 days (IQR: 11 to 21 days)
			n=224 hospitalised adult cases (untreated group, Omicron variant)	Median time from first positive to negative RT-PCR test (untreated group): 17 days (IQR: 12 to 21 days)	
Li ( <u>45</u> )	China, April to May 2022	Retrospective	n=6,134 hospitalised asymptomatic	Time to negative test (7 or fewer days): n=1,249 of 6,134 (20.4%)	
		cohort	and mild cases (Omicron BA.2	Time to negative test (8 to 15 days): n=3,832 of 6,134 (62.4%)	
			variant)	Time to negative test (16 or more days): n=1,059 of 6,134 (17.2%)	
Li ( <u>44</u> )	China, June to August	Prospective	n=51 cases (Omicron BA.2 variant)	Time to negative test (7 days or more, Omicron BA.2): 54.38%	
	2022	cohort		Time to negative test (10 days or more, Omicron BA.2): 21.92%	
				Time to negative test (14 days or more, Omicron BA.2): 4.51%	
			n=27 cases (Omicron BA.5 variant)	Time to negative test (7 days or more, Omicron BA.5): 96.30%	
				Time to negative test (10 days or more, Omicron BA.5): 88.89%	
				Time to negative test (14 days or more, Omicron BA.5): 48.15%	
Liu ( <u>46</u> )	China, April to May 2022	Prospective cohort (quasi- experimental)	n=40 hospitalised asymptomatic, mild, and moderate cases (nasal irrigation group, Omicron BA.2.2 variant)	Time to negative test (nasal irrigation group): 17.58 days (SD: 7.31)	

Study	Country, time period	Study type	Participants	Outcome
			n=40 hospitalised asymptomatic, mild, and moderate cases (conventional treatment group, Omicron BA.2.2 variant)	Time to negative test (conventional treatment group): 29.10
Lu ( <u>47</u> )	China, April to May 2022	Prospective	n=1,337 hospitalised cases aged	Median time from first positive nucleic acid test to first negati
		cohort	over 60 years (Omicron variant)	Viral shedding time was shorter in cases who were fully vaca receiving paxlovid (p=0.003), and cases with mild compared
Luna-Muschi ( <u>14</u> )	Brazil, January 2022	Prospective	n=30 vaccinated healthcare worker	RT-PCR positivity (day 7 after symptom onset): n=30 of 30 (
		cohort	mild cases (Omicron BA.1 variant)	RT-PCR positivity (day 10 after symptom onset): n=29 of 30
				RT-PCR positivity (day 14 after symptom onset): n=17 of 30
Ma ( <u>48</u> )	China, up to June 2022	Retrospective	n=14 liver transplant cases	Time from first positive to first negative RT-PCR test (7 days
	(start date not stated)	cohort	(Omicron variant)	Time from first positive to first negative RT-PCR test (8 to 14
				Time from first positive to first negative RT-PCR test (more the
				Median time from first positive to first negative RT-PCR test:
Martin-Blondel ( <u>49</u> )	France, January to May 2022	Prospective cohort	n=140 mild or moderate cases at high risk of severe COVID-19 (Sotrovimab group, Omicron BA.1 variant)	Median time to negative RT-PCR test (Sotrovimab group, Or days)
			n=10 mild or moderate cases at high risk of severe COVID-19 (Nirmatrelvir group, Omicron BA.1 variant)	Median time to negative RT-PCR test (Nirmatrelvir group, Or days)
			n=43 mild or moderate cases at high risk of severe COVID-19 (Sotrovimab group, Omicron BA.2 variant)	Median time to negative RT-PCR test (Sotrovimab group, Or days)
			n=49 mild or moderate cases at high risk of severe COVID-19 (Nirmatrelvir group, Omicron BA.2 variant)	Median time to negative RT-PCR test (Nirmatrelvir group, Or
Okumura ( <u>50</u> )	Japan, November to	Retrospective	n=11 cases (Omicron variant)	Time for Ct values to become greater than 30: 6.0 days (95%
	December 2021	cohort		Time for Ct values to become greater than 35: 10.6 days (95
				Time for Ct values to become greater than 40: 15.1 days (95
				Time for Ct values to become greater than 45: 19.7 days (95
Pei ( <u>51</u> )	China, March to May 2022	Retrospective cohort	n=198,262 asymptomatic and mild cases (likely Omicron BA.2 variant)	Median time from first positive to first negative RT-PCR test:

0 days (SD: 3.70)

ative test: 9 days (IQR: 6 to 12 days)

ccinated or boosted (p<0.0001), cases ed with severe or critical COVID-19 (p=0.047)

(100%)

80 (97%)

80 (57%)

ys or fewer): n=3 of 14 (21.4%)

14 days): n=4 of 14 (28.6%)

e than 14 days): n=7 of 14 (50.0%)

st: 14 days

Omicron BA.1): 12.5 days (95% CI: 10.5 to 14

Omicron BA.1): 5 days (95% CI: 1 to 12.5

Omicron BA.2): 10.5 days (95% CI: 8 to 12.5

Omicron BA.2): 4 days (95% CI: 4 to 9 days)

5% CI: 4.2 to 7.3 days) 95% CI: 9.5 to 11.9 days) 95% CI: 13.6 to 17.6 days) 95% CI: 17.3 to 23.7 days) st: 8.29 days (IQR: 5.33 to 11.27 days)

Study	Country, time period	Study type	Participants	Outcome
			n=33,896 asymptomatic and mild cases with comorbidities (likely Omicron BA.2 variant)	Median time from first positive to first negative RT-PCR test 6.33 to 12.28 days)
Shao ( <u>52</u> )	China, April to May 2022	Retrospective cohort	n=4 asymptomatic hospitalised cases (Omicron variant)	Median time to negative RT-PCR (asymptomatic cases): 9 d
			n=180 mild hospitalised cases (Omicron variant)	Median time to negative RT-PCR (mild cases): 10 days (IQR
			n=41 moderate hospitalised cases (Omicron variant)	Median time to negative RT-PCR (moderate cases): 13 days
			n=1 severe hospitalised cases (Omicron variant)	Time to negative RT-PCR (n=1 severe case): 15 days
Shen ( <u>53</u> )	China, March 2022	Prospective cohort	n=60 hospitalised non-severe cases (VV116 treated group, Omicron variant)	Time from first positive to first negative nucleic acid test (VV to 10.77 days)
			n=76 hospitalised non-severe cases (untreated group, Omicron variant)	Time from first positive to first negative nucleic acid test (untr 12.04 days)
Shi ( <u>54</u> ) (Preprint)	China, April to May 2022	Prospective cohort (RCT)	n=91 hospitalised asymptomatic and mild cases (Liushen Pill treated group, Omicron BA.2 variant)	Negative RT-PCR test within 7 days of allocation (Liushen P
			n=90 hospitalised asymptomatic and mild cases (control group, Omicron BA.2 variant)	Negative RT-PCR test within 7 days of allocation (control gro
Sikka ( <u>29</u> ) (Preprint)	US, February 2021 to January 2022	Prospective cohort	n=37 cases (Omicron BA.1 variant)	Median time to RT-PCR clearance: 9.5 days
Sun ( <u>55</u> ) (Preprint)	China, April 2022	Retrospective cohort	n=100 hospitalised asymptomatic cases (Omicron BA.2.2.1 variant)	Median time from first positive test to cessation of viral shedo to 11 days)
			n=274 hospitalised mild cases (Omicron BA.2.2.1 variant)	Median time from first positive test to cessation of viral shedd days)
Takahashi ( <u>15</u> )	Japan, November to	Retrospective	n=18 asymptomatic and mild cases	Positive RT-PCR test (0 to 1 days after diagnosis): n=17 of 1
	December 2021	cohort	(Omicron variant)	Positive RT-PCR test (2 to 5 days after diagnosis): n=11 of 1
				Positive RT-PCR test (6 to 9 days after diagnosis): n=16 of 1
				Positive RT-PCR test (10 to 14 days after diagnosis): n=12 o
				Positive RT-PCR test (15 days and more after diagnosis): n=
Tillman ( <u>56</u> )	Germany, study period not stated	Prospective cohort	n=20 vaccinated adult chronic dialysis mild cases (Omicron variant)	Mean time from diagnosis to negative RT-PCR test: 13 days

st (cases with comorbidities): 9.29 days (IQR:

days (IQR: 8.0 to 10.5 days)

QR: 8.0 to 12.5 days)

ys (IQR: 10.0 to 15.0 days)

V116 treated group): 9.92 day (95% CI: 9.06

ntreated group): 11.13 days (95% CI: 10.22 to

Pill treated group): n=44 of 91 (48.35%)

group): n=28 of 90 (31.11%)

edding (asymptomatic cases): 10 days (IQR: 9

edding (mild cases): 10 days (IQR: 9 to 12

f 17 (100%)

f 12 (91.7%)

16 (100%)

2 of 17 (70.6%)

n=3 of 10 (30.0%)

ys

Study	Country, time period	Study type	Participants	Outcome
Van der Veer ( <u>57</u> ) (Preprint)	The Netherlands, November 2021 to	Prospective cohort	n=142 healthcare worker cases (Omicron BA.1 variant)	Median time from first positive to first negative RT-PCR test days)
	February 2022		n=37 healthcare worker cases (Omicron BA.2 variant)	Median time from first positive to first negative RT-PCR test days)
Wei ( <u>58</u> ) (Preprint)	China, May to July 2022	Prospective cohort (RCT)	n=94 asymptomatic or mild adult cases (120mg Cepharanthine group, Omicron variant)	Mean time from randomisation to first negative RT-PCR test (95% CI: 4.11 to 5.30 days)
			n=102 asymptomatic or mild adult cases (60mg Cepharanthine group, Omicron variant)	Mean time from randomisation to first negative RT-PCR test (95% CI: 3.65 to 4.65 days)
			n=85 asymptomatic or mild adult cases (placebo group, Omicron variant)	Mean time from randomisation to first negative RT-PCR test days)
Weng ( <u>59</u> )	China, April to May 2022	Retrospective cohort	n=82 hospitalised mild or moderate cases aged over 60 years (Paxlovid group, Omicron variant)	Median time from first positive test to second negative test (F days)
			n=81 hospitalised mild or moderate cases aged over 60 years (control group, Omicron variant)	Median time from first positive test to second negative test (
Wu ( <u>60</u> ) (Preprint)	Hong Kong, February to July 2022	Retrospective cohort	n=1,084 mild to moderate cases aged 0 to 18 years (Omicron BA.2 variant)	Median time from symptom onset to RT-PCR Ct value of ove cases): 10.4 days (IQR: 8.9 to 11.9 days)
			n=807 mild to moderate cases aged 19 to 64 years (Omicron BA.2 variant)	Median time from symptom onset to RT-PCR Ct value of ove cases): 11.6 days (IQR: 9.3 to 13.7 days)
			n=487 mild to moderate cases aged 65 to 74 years (Omicron BA.2 variant)	Median time from symptom onset to RT-PCR Ct value of ove cases): 13.6 days (IQR: 11.3 to 16.3 days)
			n=682 mild to moderate cases aged 80 years and older (Omicron BA.2 variant)	Median time from symptom onset to RT-PCR Ct value of ove moderate cases): 15.6 days (IQR: 12.7 to 17.9 days)
			n=12,669 symptomatic cases (Omicron BA.2 variant)	The time from symptom onset to RT-PCR Ct value of over 3 (p<0.001), younger cases (p<0.001), and cases with milder i
Wu ( <u>61</u> )	China, March 2022	Retrospective cohort	n=129 vaccinated asymptomatic or mild cases (likely Omicron BA.2 variant)	Mean time from first positive RT-PCR test to RT-PCR Ct val days)
			n=13 unvaccinated asymptomatic or mild cases (likely Omicron BA.2 variant)	Mean time from first positive RT-PCR test to RT-PCR Ct val 4.7 days)

st (Omicron BA.1): 12 days (IQR: 10 to 15

st (Omicron BA.2): 11 days (IQR: 10 to 13

st (120mg Cepharanthine group): 4.70 days

st (60mg Cepharanthine group): 4.15 days

st (placebo group): 4.58 days (3.89 to 5.26

(Paxlovid group): 16.5 days (IQR: 13 to 20

(control group): 20 days (IQR: 17 to 22 days)

over 33 (aged 0 to 18 years, mild to moderate

ver 33 (aged 19 to 64 years, mild to moderate

over 33 (aged 65 to 79 years, mild to moderate

over 33 (age 80 years and over, mild to

33 was shorter for vaccinated cases r infections

alue above 35 (vaccinated): 12.6 days (SD: 3.4

alue above 35 (unvaccinated): 14.8 days (SD:

Study	Country, time period	Study type	Participants	Outcome
Xu ( <u>62</u> )	China, April 2022	Retrospective	n=13,162 asymptomatic or mild	Negative RT-PCR test by 7 days: n=5,437 of 13,162 (41.3%)
		cohort	cases (Omicron BA.2.2 variant)	Negative RT-PCR test by 14 days: n=12,482 of 13,162 (94.8
Xu ( <u>65</u> )	China, April to May 2022	Prospective cohort	n=413 adult asymptomatic or mild cases (Interferon alpha-2b spray group, Omicron variant)	Median time to negative RT-PCR test (interferon alpha-2b sp
			n=458 adult asymptomatic or mild cases (control group, Omicron variant)	Median time to negative RT-PCR test (control group): 12.58
Xu ( <u>66</u> ) (Preprint)	China, April to June 2022	Retrospective cohort	n= 83 hospitalised haemodialysis cases (Omicron variant)	Mean time from first positive to first negative RT-PCR tests ( days)
			n=134 hospitalised non- haemodialysis cases (Omicron variant)	Mean time from first positive to first negative RT-PCR tests ( 3.52 days)
Xu ( <u>63</u> )	China, April to May 2022	Prospective cohort (RCT)	n=1,393 asymptomatic or mild cases (Reyanning group, Omicron variant)	Median time from hospitalisation to negative nucleic acid tes days)
			n=1,407 asymptomatic or mild cases (control group, Omicron variant)	Median time from hospitalisation to negative nucleic acid tes
Xu ( <u>64</u> )	China, April to June 2022	Retrospective cohort	n=346 child asymptomatic or mild cases (Lianhua Qingwen group, likely Omicron BA.2 variant)	Median time from admission to negative RT-PCR test (Lianh 7.0 days)
			n=346 child asymptomatic or mild cases (control group, likely Omicron BA.2 variant)	Median time from admission to negative RT-PCR test (contro
Yan ( <u>67</u> )	China, April to May 2022	Prospective cohort	n=5 hospitalised child mild or moderate cases (Paxlovid group, likely Omicron BA.2 variant)	Median time from first positive RT-PCR test or symptom ons RT-PCR test (Paxlovid group): 9 days (IQR: 9 to 10 days)
			n=30 hospitalised child mild or moderate cases (control group, likely Omicron BA.2 variant)	Median time from first positive RT-PCR test or symptom ons RT-PCR test (control group): 11 days (IQR: 9 to 12 days)
Yang ( <u>68</u> )	China, April to May 2022	Retrospective cohort	n=13 child mild or moderate cases with congenital heart disease (likely Omicron BA.2.2 variant)	Mean duration of viral clearance: 16.4 days (SD: 2.9 days)
Yang ( <u>69</u> )	China, March to May	Retrospective	n=603 hospitalised child cases	Median time from first positive to first negative RT-PCR test:
(Preprint)	2022	cohort	(likely Omicron BA.2.2 variant)	Viral shedding time was longer in cases with abnormal defect in vaccinated cases and cases with higher household vaccin
Yin ( <u>70</u> )	China, March to May 2022	Retrospective cohort	n=199,590 asymptomatic or mild cases (Omicron BA.2 variant)	Mean time from illness onset to negative RT-PCR test: 7.17

%)

1.8%)

spray group): 11.90 days

i8 days

s (haemodialysis cases): 18.15 days (SD: 6.37

s (non- haemodialysis cases): 11.18 days (SD:

est (Reyanning group): 6 days (IQR: 3 to 9

est (control group): 7 days (IQR: 5 to 9 days)

nhua Qingwen group): 5.0 days (IQR: 3.0 to

trol group): 6.0 days (IQR: 5.0 to 8.0 days)

nset (whichever was earlier) to first negative

nset (whichever was earlier) to first negative

#### st: 12 days (IQR: 9 to 14 days

ecation and more severe disease, and shorter cination rates

7 days (SD: 3.42 days)

Study	Country, time period	Study type	Participants	Outcome
Ying-Hao ( <u>71</u> )	China, April to May 2022	Retrospective cohort	n=25,168 asymptomatic or mild cases (Omicron BA.2 variant)	Median time from first positive to first negative RT-PCR test:
Yu ( <u>72</u> )	China, April 2022	Prospective cohort	n=42 asymptomatic cases (Omicron BA.2.2 variant)	Mean time from first positive to first negative nucleic acid tes days)
			n=619 mild cases (Omicron BA.2.2 variant)	Mean time from first positive to first negative nucleic acid tes
Zee ( <u>73</u> )	Hong Kong, February to March 2022	Retrospective cohort	n=422 fully vaccinated healthcare worker cases (likely Omicron BA.2.2 variant)	Mean time to 2 consecutive negative rapid antigen tests: 9.7
Zeng ( <u>74</u> )	China, January 2022	Prospective cohort	n=355 fully vaccinated hospitalised cases (inactive vaccine, Omicron variant)	Median time from first positive to first negative SARS-CoV-2 vaccine cases): 17.0 days (IQR: 12.0 to 22.0 days)
			n=14 fully vaccinated hospitalised cases (recombinant vaccine, Omicron variant)	Median time from first positive to first negative SARS-CoV-2 vaccine cases): 20.5 days (IQR: 17.8 to 26.3 days)
			n=11 partially vaccinated hospitalised cases (Omicron variant)	Median time from first positive to first negative SARS-CoV-2 days (IQR: 9.0 to 25.0 days)
Zhang ( <u>75</u> )	China, March to May 2022	Retrospective cohort	n=33 hospitalised haemodialysis mild cases aged 45 to 99 years (Omicron variant)	Mean time from first positive to first negative nucleic acid tes 5.22 days)
			n=66 hospitalised non- haemodialysis mild cases aged 45 to 99 years (Omicron variant)	Mean time from first positive to first negative nucleic acid tes (SD: 5.43 days)
Zhong ( <u>76</u> )	China, April to May 2022	Prospective cohort	n=106 hospitalised elderly cases (Paxlovid group, likely Omicron BA.2.2 variant)	Mean time from first positive to first negative nucleic acid tes days)
			n=36 hospitalised elderly cases (control group, likely Omicron BA.2.2 variant)	Mean time from first positive to first negative nucleic acid tes
Zhong ( <u>77</u> ) (Preprint)	China, from April 2022 (end date not stated)	Retrospective cohort	n=180 hospitalised elderly cases (Omicron BA.2.2 variant)	Mean time from first positive to first negative nucleic acid tes

#### Table 1h. Viral load over time

Study	Country, time period	Study type	Participants	Outcome
Bouton (7)	US, November 2021	Prospective	n=92 university cases (n=75	Data extracted from figure: Ct values increased from day 0 to d
	onwards (end date not	cohort	Omicron, n=17 Delta variant)	8), Ct values steady from 3 days before to 1 day after symptom
	stated)			after symptom onset (more slowly after day 9), proportion of po
				after diagnosis (to about 30% positivity), then reduced to 0 by 8

st: 6 days (IQR: 4 to 9 days)

est (asymptomatic cases): 9.26 days (SD: 3.16

est (mild cases): 10.62 days (SD: 2.73 days)

.76 days

-2 RNA test (fully vaccinated with inactive

-2 RNA test (fully vaccinated with recombinant

-2 RNA test (partially vaccinated cases): 16.0

est (haemodialysis cases): 16.67 days (SD:

est (non-haemodialysis cases): 14.07 days

est (Paxlovid group): 9.32 days (SD: 2.78

est (control group): 11.11 days (SD: 2.67 days)

est: 10 days (IQR: 8 to 12 days)

day 15 after diagnosis (more slowly after day om onset, increased from day 1 to day 20 positive cultures rose between days 1 and 3 v 8 days after diagnosis (some sporadic

Study	Country, time period	Study type	Participants	Outcome
				positive cultures up to day 15), proportion of positive cultures ro symptom onset (to about 30% positivity), dropped substantially drop until day 13, with no further positive cultures up to day 20
Choi ( <u>25</u> )	South Korea, January 2022	Retrospective cohort	n=5,187 (Omicron variant)	Data extracted from figure: Ct values decreased (viral load increased symptom onset, increased (viral load decreased) to between 12 decreased again (viral load increased) up to 17 days after symptom
Funk ( <u>78</u> ) (Preprint)	England, December 2021 to January 2022	Cross- sectional	n=1,212,234 cases (n=1,083,976 Omicron BA.1, n=128,258 Omicron BA.2 variant)	Data extracted from figure: Ct values for both Omicron BA.1 and symptom onset, then increased up to day 6 since symptom ons infections had higher Ct values than BA.1 infections, reinfection previous infections, and there was no clear difference between
Li ( <u>79</u> )	Hong Kong, January to February 2022	Retrospective cohort	n=104 hospitalised cases (likely Omicron BA.2.2 variant)	Data extracted from figure: The Ct value was lowest (viral load gradually decreased until 10 days after symptom onset (data true
Marking ( <u>80</u> ) (Preprint)	Sweden, January to February 2022	Prospective cohort	n=60 cases (Omicron BA.1, BA.1.1 and BA.2 variants)	Data extracted from figure: Ct values decreased (viral load incre PCR test, then increased (viral load decreased) up to day 15 (d
Tassetto ( <u>81</u> )	US, July 2021 to March 2022	Prospective cohort	n=33 cases (Omicron BA.1 variant)	Data extracted from figure: Ct values increased from days 2 to before day 2), with a decrease in Ct values in days 7 to 9 (data
Teyssou ( <u>82</u> )	France, December 2021 to May 2022	Prospective cohort	n= 84 cases (Omicron BA.1 variant)	Data extracted from figure: For Omicron BA.1, Ct values decrea onset to days 1 to 3 after symptom onset, then increased (viral onset (data truncated at day 10)
			n=60 cases (Omicron BA.2 variant)	Data extracted from figure: For Omicron BA.2, Ct values increations onset to days 1 to 3 after symptom onset, decreased (viral load onset, then increased (viral load decreased) until day 10 after s
Townsley ( <u>83</u> ) (Preprint)	UK, January 2021 to May 2022	Prospective cohort	n=240 cases (Omicron BA.1 and BA.2 variants)	Data extracted from figure: Ct values decreased (viral load incressymptom onset, then increased (viral load decreased) up to day Omicron BA.1 variant cases), then slightly decreased (viral load variant cases (data truncated at day 17 for Omicron BA.2 variant
Young ( <u>84</u> ) (Preprint)	Singapore, December 2021	Retrospective cohort	n=87 cases (Omicron B.1.1529 variant)	Data extracted from figure: Ct values decreased (viral load increased then increased (viral load decreased) until 20 days after illness

rose from 1 day before to 4 days after ly on day 5 (to about 10%), then continued to 0

creased) up to between 3 and 4 days after 12 and 13 days after symptom onset, then mptom onset (data truncated after this)

and BA.2 decreased from days 0 to 2 since nset (data truncated at day 6). BA.2

ons had higher Ct values than no known en people with different doses of vaccine.

d highest) on day 1 after symptom onset, and truncated after this)

creased) up to day 3 from first positive RT-(data truncated at day 15)

o 7 after symptom onset (data truncated ta truncated at day 9)

eased (viral load increased) from symptom al load decreased) until day 10 after symptom

eased (viral load decreased) from symptom ad increased) until days 7 to 9 after symptom <sup>-</sup> symptom onset (data truncated at day 10)

creased) up to days 2 and 3 from first lay 15 (data truncated around day 15 for ad increased) up to day 17 for Omicron BA.2 iant cases)

creased) over the first 2 to 3 days of illness, as onset (data truncated at 20 days)

### The difference in transmission from people with asymptomatic compared with symptomatic COVID-19 Omicron variant

### Evidence

In total, 10 observational studies looking at the difference in COVID-19 transmission between asymptomatic and symptomatic cases were included in this report (5 preprints (55, 85 to 88), one rated as low quality (87), 6 rated as medium quality (55, 86, 88 to 91), and 3 rated as high quality (51, 85, 92)). All studies were identified through the literature search, as no studies of the Omicron variant were found in our previous review (unpublished, search up to 15 March 2022). Three studies reported on the difference in the secondary attack rates (SAR) between asymptomatic and symptomatic index cases (85, 86, 88), while 5 studies reported on differences in viral load (87, 89 to 92) and 2 studies reported on differences in viral shedding time between asymptomatic and symptomatic cases (51, 55).

Five studies were conducted in Asia (51, 55, 88, 90, 91), 3 studies in the US (87, 89, 92), one study in Turkey (85), and one study in Nicaragua (86). While most studies were conducted between December 2021 and June 2022 when the COVID-19 Omicron variant was dominant across the world, other studies started earlier and reported COVID-19 Omicron variant cases separately. All studies were either retrospective (51, 55, 87, 89, 91) or prospective (85, 86, 88, 90, 92) cohort studies.

Table 2 gives study characteristics of the 10 included studies.

### Secondary attack rates

Three prospective cohort studies reported the household SAR of asymptomatic and symptomatic COVID-19 Omicron variant index cases (<u>85</u>, <u>86</u>, <u>88</u>). Study characteristics are given in <u>Table 2a</u>.

Erik and others (preprint, rated as high quality) reported on the household SAR of COVID-19 (variant not stated) from n=42 index cases and n=112 household contacts in Turkey between August 2021 and February 2022 (85). The results suggested that asymptomatic index cases transmitted COVID-19 substantially less than symptomatic index cases, although imprecision was high (odds ratio [OR] of transmission = 0.12, 95% confidence interval [CI]: 0.02 to 0.77, p=0.03).

Frutos and others (preprint, rated as medium quality) reported on the household SAR of COVID-19 (likely Omicron variant) in n=104 households in Nicaragua between January 2022

and June 2022 (86). The results suggested that symptomatic index cases (SAR = 20%, 95% CI: 17% to 25%) transmitted COVID-19 substantially more than asymptomatic index cases (SAR = 2%, 95% CI: 1% to 8%), although imprecision was again high (relative risk [RR] of transmission = 14.77, 95% CI: 3.12 to 70.03).

Wei and others (preprint, rated as medium quality) reported on the household SAR of COVID-19 (likely Omicron BA.2 variant) from n=236 index cases (including n=67 children) and n=546 adult household contacts in China in April 2022 (88). The results suggested that asymptomatic index cases (SAR = 72.7%, 95% CI: 62.0% to 83.5%) transmitted COVID-19 at a similar rate to symptomatic index cases (SAR = 77.7%, 95% CI: 74.0% to 81.4%, p value for difference = 0.46).

### Viral load

Five studies reported the viral loads of asymptomatic and symptomatic COVID-19 Omicron variant cases (87, 89 to 92). Most studies measured Ct (cycle threshold) values with RT-PCR (reverse-transcriptase polymerase chain reaction), which are inversely proportional to viral load, though do not necessarily indicate infectious virus (87, 89 to 91). Study characteristics are given in Table 2b.

Three studies reported similar Ct values or viral loads between asymptomatic and symptomatic cases in the US ( $\underline{87}$ ,  $\underline{89}$ ) and China ( $\underline{91}$ ), although the number of participants was small in all studies. Wu and others (rated as medium quality) reported similar Ct values (Omicron variant COVID-19) between n=22 asymptomatic (median nasopharyngeal Ct value = 27.8, interquartile range [IQR]: 23.4 to 34.5) and n=339 symptomatic cases (median nasopharyngeal Ct value = 30.5, IQR: 24.5 to 35, p value for difference = 0.19) ( $\underline{91}$ ). Kaur and others (conference abstract only, rated as low quality) reported similar viral loads (Omicron variant COVID-19) between n=45 asymptomatic, n=23 pre-symptomatic, and post-infection asymptomatic cases (p value = 0.45) ( $\underline{87}$ ). Laitman and others (rated as medium quality) reported similar Ct values (likely Omicron BA.1 variant COVID-19) between n=272 asymptomatic (median Ct value = 22.98, SD: 4.4) and n=552 symptomatic cases (median Ct value = 21.27, SD: 4.2) ( $\underline{89}$ ).

Conversely, 2 studies reported differences in viral loads or Ct values between asymptomatic and symptomatic cases in Japan (90) and the US (92). Suzuki and others (rated as medium quality) reported proportionately higher Ct values (lower viral loads, Omicron BA.1 and BA.2 variant COVID-19) in n=28 asymptomatic compared with n=382 symptomatic cases, for example, 29% of asymptomatic and 46% of symptomatic cases had Ct values of less than 20, but did not test these differences statistically (90). The Heroes-Recover Network and others (rated as high quality) reported higher viral loads (Omicron B.1.1.529 and BA1 variant COVID-19) in n=628 symptomatic (mean viral load = 3.4 log<sub>10</sub> copies per µL, standard deviation [SD]: 1.6 log<sub>10</sub> copies per µL) compared with n=96 asymptomatic cases (mean viral load = 2.3 log<sub>10</sub> copies per µL, SD: 1.8 log<sub>10</sub> copies per µL), with a mean difference of 1.2 log<sub>10</sub> copies per µL (95% CI: 0.8 log<sub>10</sub> copies per µL to 1.6 log<sub>10</sub> copies per µL).

#### Viral shedding time

Two studies reported the viral shedding time (typically time from symptom onset to negative RT-PCR) of asymptomatic and mildly symptomatic COVID-19 Omicron variant cases in China (51, 55). Study characteristics are given in Table 2c.

Pei and others (rated as high quality) reported that the viral shedding time (likely Omicron BA.2 variant) was longer in cases with a mild COVID-19 infection (n=20,504) compared with asymptomatic cases (n=177,758) (OR for viral shedding time of more than 8.3 days, 1.50, 95% CI: 1.46 to 1.55, p<0.001) (<u>51</u>).

Conversely, Sun and others (preprint, rated as medium quality) reported similar viral shedding times (Omicron BA.2.2.1 variant) between n=100 asymptomatic (median viral shedding time = 10 days, IQR: 9 to 11 days) and n= 274 mildly infected cases (median viral shedding time = 10 days, IQR: 9 to 12 days, p value for difference = 0.35) (<u>55</u>).

### Summary

Ten studies reported on differences in transmission from people with asymptomatic compared with symptomatic COVID-19 Omicron variant. Three studies compared household SAR of asymptomatic and symptomatic index cases, with 2 studies suggesting more transmission from symptomatic than asymptomatic index cases although imprecision was high, and one study suggesting no clear difference. Five studies compared viral loads (usually using Ct values) between symptomatic and asymptomatic cases, with 3 studies suggesting similar viral loads, and 2 studies suggesting higher viral loads in symptomatic compared with asymptomatic cases, with one study suggesting similar viral shedding times, and one study suggesting longer viral shedding times in symptomatic cases.

All studies were relatively small, with only one study having more than a few hundred participants (51), which increases the imprecision of the results. The evidence was also limited due to the small number of studies, the variability between studies and differences in results, which limit our ability to draw conclusions.

### Table 2. Studies for asymptomatic compared with symptomatic transmission

Acronyms: CI = confidence interval, IQR = interquartile range, OR = odds ratio, RT-PCR = reverse transcriptase polymerase chain reaction, SAR = secondary attack rate, SD = standard

#### Table 2a. Secondary attack rates (SARs)

Study	Country, time period	Study type	Participants	Definition of symptomatic	Outcome	
Erik ( <u>85</u> ) (preprint)	Turkey, August 2021 to February 2022	Prospective cohort	n=42 index cases, n=112 household contacts (variant not stated)	Any of: sore throat, runny nose, cough, fever (and other symptoms not listed)	OR for transmission (asymptomatic compared with symptomatic index cases):	0.12 (95% CI: 0.02 to 0.77, p=0.03)
Frutos ( <u>86</u> )	Nicaragua, January to	Prospective	n=104 households	Not stated (daily symptom data collected	SAR (symptomatic index cases):	0.20 (95% CI: 0.17 to 0.25)
(preprint)		(likely Omicron	by staff during visits)	SAR (asymptomatic index cases):	0.02 (95% CI: 0.01 to 0.08)	
			variant)		RR for transmission (symptomatic compared with asymptomatic index cases):	14.77 (95% CI: 3.12 to 70.03)
Wei ( <u>88</u> )	China, April 2022	Prospective	n=236 index cases,	Moderate COVID-19 defined as non-	SAR (asymptomatic index case, n=24):	72.7% (95% CI: 62.0% to 83.5%)
(preprint)	(preprint)	con		severe pneumonia (definitions of mild, severe and critical COVID-19 not stated)	SAR (symptomatic index case, n=212):	77.7% (95% CI: 74.0% to 81.4%)
					p value for difference:	0.46

#### Table 2b. Viral load

Study	Country, time period	Study type	Participants	Definition of symptomatic	Outcome	
Heroes- Recover	US, December 2020 to May 2022	Prospective cohort	n=743 cases (Omicron B.1.1.529	fever, chills, cough, shortness of breath,	Mean viral load (Asymptomatic, n=96):	2.3 log <sub>10</sub> copies per μL (SD: 1.8 log <sub>10</sub> copies per μL)
Network ( <u>92</u> )			and BA1 variants)		Mean viral load (Symptomatic, n=628):	3.4 log <sub>10</sub> copies per $\mu$ L (SD: 1.6 log <sub>10</sub> copies per $\mu$ L)
					Mean difference in viral load (symptomatic compared with asymptomatic cases):	1.2 log <sub>10</sub> copies per μL (95% CI: 0.8 to 1.6 log <sub>10</sub> copies per μL)
Kaur ( <u>87</u> ) (preprint)	US, December 2021	Retrospective cohort	n=68 cases (Omicron variant)	Not stated	No statistically significant difference in the Ct values of asymptomatic (n=45), pre- symptomatic (n=23), and post-infection asymptomatic cases (p=0.45)	
Laitman ( <u>89</u> )	US, December 2021 to January 2022		Not stated	Median Ct value (Roche Cobas) (Asymptomatic, n=272):	22.98 (SD: 4.4)	
	variant)		Median Ct value (Roche Cobas) (Symptomatic, n=552):	21.27 (SD: 4.2)		
Suzuki ( <u>90</u> )	Japan, December 2021 to February 2022	Prospective cohort	n=410 viral samples (Omicron BA.1 and BA.2 variants)	Not stated	Ct values (asymptomatic, n=28):	Less than 20: n=8 (29%); 20 to 24: n=16 (57%); 25 to 29: n=3 (11%); 30 and above: n=1 (4%)

Study	Country, time period	Study type	Participants	Definition of symptomatic	Outcome	
					Ct values (symptomatic, n=382):	Less than 20: n=175 (46%); 20 to 24: n=164 (43%); 25 to 29: n=25 (7%); 30 and above: n=18 (5%)
Wu ( <u>91</u> )	China, January 2022	Retrospective cohort	n=361 cases (Omicron variant)	Patients with one or more of: fever, cough, fatigue, decreased or loss of ability to smell and taste, nasal congestion, runny nose, sore throat, conjunctivitis, myalgia, and diarrhoea	Median nasopharyngeal Ct value (symptomatic, n=339):	30.5 (IQR: 24.5 to 35)
					Median nasopharyngeal Ct value (asymptomatic, n=22):	27.8 (IQR: 23.4 to 34.5)
					p value for difference (nasopharyngeal):	0.19
					Median oropharyngeal Ct value (symptomatic, n=339):	34.5 (IQR: 30 to 37)
					Median oropharyngeal Ct value (asymptomatic, n=22):	33.5 (IQR: 26 to 35)
					p value for difference (oropharyngeal):	0.11

#### Table 2c. Viral shedding time

Study	Country, time period	Study type	Participants	Definition of symptomatic	Outcome	
Pei ( <u>51</u> )	China, March 2022 to May 2022	Retrospective cohort	n=198,262 cases (likely Omicron BA.2 variant)	Not stated	OR of viral shedding time of more than 8.29 days (mild infection [n=20,504] compared with asymptomatic cases [n=177,758]):	1.50 (95% CI: 1.46 to 1.55, p<0.001)
Sun ( <u>55</u> ) (preprint)	China, April 2022	Retrospective cohort	n=382 cases (Omicron BA.2.2.1 variant)	Moderate COVID-19 defined as onset of fever or respiratory symptoms (or both) and radiographic evidence of pneumonia (not stated, but mild COVID-19 likely symptoms less severe than moderate COVID-19)	Median viral shedding time (asymptomatic, n=100):	10 days (IQR: 9 to 11 days)
					Median viral shedding time (mildly infected, n=274):	10 days (IQR: 9 to 12 days)
					p value for difference:	0.35

### Inequalities

There was little evidence available to explore inequalities through variations across populations and subgroups (notably, no included study included cases from more than one country), for example cultural variations or differences between ethnic, social or vulnerable groups. As such, it was not possible to examine inequalities in this report.

### Limitations

The source of evidence in this review included peer-reviewed and preprint articles. We did not conduct an extensive search of other sources (such as websites of public health organisations). As with all reviews, the evidence identified may be subject to publication bias, whereby null or negative results are less likely to have been published by the authors, though descriptive studies may be less susceptible to publication bias than other study types.

For infectious period, 24 of the 82 included studies were preprints, and for asymptomatic compared with symptomatic transmission, 5 of the 10 included studies were preprints. In general, preprints should be treated with caution as they have not been peer reviewed or subject to publishing standards and may be subject to change.

In addition, this rapid review is limited by the fact that we were reviewing evidence for the COVID-19 Omicron variant, which caused outbreaks of COVID-19 in different countries. Studies may have been conducted rapidly, with the aim to provide evidence in a timely manner, and this may have impacted on quality, both in term of design (especially limited statistical analyses) and reporting (insufficient detail).

This review was conducted at pace following streamlined methodology. Quality of the descriptive studies was not assessed and only the main characteristics of the studies were extracted. The narrative summary was conducted by one reviewer and checked by another.

### **Evidence gaps**

For infectious period, there was a reasonable amount of evidence for most of the different measures of infectious period. However, only 3 studies measured infectious period directly, and none from the UK. Additionally, the majority of studies included cases with only COVID-19 Omicron BA.1 and BA.2 variant, which may limit generalisability to the current circulating variants.

For asymptomatic compared with symptomatic transmission of COVID-19, only 3 studies directly compared transmission from asymptomatic and symptomatic cases, again with none from the UK. Additionally, as with infectious period, the vast majority of studies included cases with only Omicron BA.1 and BA.2 variant COVID-19, limiting generalisability to the current circulating variants.

### Conclusion

### Infectious period of Omicron variant COVID-19

In total, 82 studies provided evidence on different measures of the Omicron variant COVID-19 infectious period. The 3 studies directly measuring the transmission period suggested most transmission events happened around symptom onset, with the majority of transmission events happening up to 5 days after symptom onset (100% in one study, 81% in another study, not reported in the third study). The 10 studies measuring viral culture positivity also suggested that viral culture positivity was highest in the first 5 days after symptom onset or diagnosis, and the 14 studies reporting on viral load over suggested that peak viral loads occurred a median and mean of 2 to 5 days after symptom onset.

The 9 studies measuring serial interval and generation time suggested the median and mean serial interval was between 2 and 4 days (though this could be variable, with interquartile ranges between one and 9 days), and generation times of around 3 days. The studies measuring incubation period suggested the median and mean incubation period was between 2 and 5 days.

The 53 studies measuring time to viral clearance suggested that there were substantial differences in viral clearance times between cases and populations, although the differences in measurement of time to viral clearance between studies may have contributed to these differences. In general, however, most studies of the general population estimated viral clearance to take around 7 to 11 days, and most studies of hospitalised, immunodeficient, and other high-risk cases estimated viral clearance to take around 10 to 15 days. Detectable viral load does not necessarily indicate that a case is infectious.

Overall, the evidence suggests that COVID-19 Omicron variant cases were most infectious up to 5 days after symptom onset, but could potentially be infectious for longer, especially if hospitalised, immunocompromised, or otherwise high-risk cases. However, while some studies included substantial numbers of cases, most studies included relatively few cases, and the majority of studies only included Omicron BA.1 and BA.2 variant cases.

# Asymptomatic compared with symptomatic transmission of Omicron variant COVID-19

In total, 10 studies reported on differences in transmission from people with asymptomatic compared with symptomatic Omicron variant COVID-19. Three studies compared household SARs of asymptomatic and symptomatic index cases, with 2 studies suggesting more transmission from symptomatic than asymptomatic index cases (although imprecision was high), and one study suggesting no clear difference. Five studies compared viral loads (usually

using Ct values) between symptomatic and asymptomatic cases, with 3 studies suggesting similar viral loads, and 2 studies suggesting higher viral loads in symptomatic compared with asymptomatic cases. Two studies compared viral shedding times between symptomatic and asymptomatic cases, with one study suggesting similar viral shedding times, and one study suggesting longer viral shedding times in symptomatic compared with asymptomatic cases.

Overall, the evidence on differences in transmission from people with asymptomatic compared with symptomatic Omicron variant COVID-19 was mixed, with some studies suggesting that symptomatic cases were more likely to transmit Omicron variant COVID-19 than asymptomatic cases and others showing no difference. The lack of precision due to small study samples combined with the variability between studies and the small number of studies limit our ability to draw conclusions.

### Acknowledgment

We would like to thank colleagues within the Public Health Clinical Response directorate who either reviewed or input into aspects of this review, particularly Dr Renu Bindra, Dr Colin Brown, Dr Daphne Duval and Dr Katie Kerr.

### Disclaimer

UKHSA's rapid reviews aim to provide the best available evidence to decision makers in a timely and accessible way, based on published peer-reviewed scientific papers, unpublished reports and papers on preprint servers. Please note that the reviews: i) use accelerated methods and may not be representative of the whole body of evidence publicly available; ii) have undergone an internal, but not independent, peer review; and iii) are only valid as of the date stated on the review.

In the event that this review is shared externally, please note additionally, to the greatest extent possible under any applicable law, that UKHSA accepts no liability for any claim, loss or damage arising out of, or connected with the use of, this review by the recipient and/or any third party including that arising or resulting from any reliance placed on, or any conclusions drawn from, the review.

### References

- 1. Tricco A and others. '<u>Rapid reviews to strengthen health policy and systems: a practical guide</u>' World Health Organization, 2017
- 2. Academy of Nutrition and Dietetics. '<u>Evidence Analysis Manual: Steps in the Academy</u> <u>Evidence Analysis Process</u>' 2016
- 3. An Der Heiden M and others. '<u>Serial interval in households infected with SARS-CoV-2</u> variant B.1.1.529 (Omicron) is even shorter compared to Delta' Epidemiology and Infection 2022: volume 150 (no pagination)
- Del Aguila-Mejia J and others. '<u>Secondary Attack Rate, Transmission and Incubation</u> <u>Periods, and Serial Interval of SARS-CoV-2 Omicron Variant, Spain</u>' Emerging Infectious Diseases 2022: volume 28, issue 6, pages 1,224 to 1,228
- 5. Xin H and others. '<u>Transmission dynamics of SARS-CoV-2 Omicron variant infections in</u> <u>Hangzhou, Zhejiang, China, January-February 2022</u>' International Journal of Infectious Diseases 2023: volume 126, pages 132 to 135
- Boucau J and others. '<u>Duration of shedding of culturable virus in SARS-CoV-2 Omicron</u> (<u>BA.1</u>) infection' New England Journal of Medicine 2022: volume 387, issue 3, pages 275 to 277
- 7. Bouton TC and others. '<u>Viral dynamics of Omicron and Delta SARS-CoV-2 variants with</u> <u>implications for timing of release from isolation: a longitudinal cohort study</u>' Clinical Infectious Diseases 2022: volume 23, page 23
- 8. Gilbert M and others. '<u>Time from last COVID-19 vaccination's impact on rapidity of viral</u> <u>culture conversion following SARS-CoV-2 infection: a prospective cohort study</u>'. Open Forum Infectious Diseases 2022: volume 9 (supplement 2), pages S53 to S54
- Jang YR and others. '<u>Clinical Features and Duration of Viral Shedding in Individuals With</u> <u>SARS-CoV-2 Omicron Variant Infection</u>' Open Forum Infectious Diseases 2022: volume 9, issue 7, pages ofac237
- Jung J and others. '<u>Risk of transmission of COVID-19 from healthcare workers returning</u> to work after a 5-day isolation, and kinetics of shedding of viable SARS-CoV-2 variant <u>B.1.1.529 (Omicron)</u>' Journal of Hospital Infection 2023: volume 131, pages 228 to 233
- 11. Kang SW and others. '<u>Comparison of secondary attack rate and viable virus shedding</u> <u>between patients with SARS-CoV-2 Delta and Omicron variants: A prospective cohort</u> <u>study</u>' Journal of Medical Virology 2023: volume 95, issue 1, page e28369
- Keske S and others. '<u>Duration of infectious shedding of SARS-CoV-2 Omicron variant</u> and its relation with symptoms' Clinical Microbiology and Infection 2022: volume 16, page 16
- 13. Kim H and others. '<u>Can nirmatrelvir/ritonavir treatment shorten the duration of COVID-19</u> <u>isolation?</u>' Frontiers in Medicine 2022: volume 9, page 988559
- 14. Luna-Muschi A and others. '<u>Characterization of severe acute respiratory syndrome</u> coronavirus 2 Omicron variant shedding and predictors of viral culture positivity on vaccinated healthcare workers with mild coronavirus disease 2019' Journal of Infectious Diseases 2022: volume 226, issue 10, pages 1,726 to 1,730

- Takahashi K and others. '<u>Duration of infectious virus shedding by SARS-CoV-2 Omicron</u> <u>Variant-Infected vaccinees</u>' Emerging Infectious Diseases 2022: volume 28, issue 5, pages 998 to 1,001
- Backer JA and others. '<u>Shorter serial intervals in SARS-CoV-2 cases with Omicron BA.1</u> variant compared to Delta variant in the Netherlands 13 to 26 December 2021' Eurosurveillance 2022: volume 26
- 17. Mefsin Y and others. Epidemiology of infections with SARS-CoV-2 Omicron BA.2 variant in Hong Kong' January to March 2022. MedRxiv : the Preprint Server for Health Sciences 2022
- Park SW and others. '<u>Inferring the differences in incubation-period and generation-interval distributions of the Delta and Omicron variants of SARS-CoV-2</u>' medRxiv. 2022: volume 5
- Tanaka H and others. <u>Shorter Incubation Period among COVID-19 Cases with the BA.1</u> <u>Omicron Variant</u> International Journal of Environmental Research and Public Health 2022: volume 19, issue 10, page 23
- 20. Ward T and others. '<u>Replacement dynamics and the pathogenesis of the Alpha, Delta,</u> and Omicron variants of SARS-CoV-2' Epidemiology and Infection 2022
- 21. Wei Z and others. <u>Household transmission of SARS-CoV-2 during the Omicron wave in</u> <u>Shanghai, China: a case-ascertained study</u>' MedRxiv 2022
- 22. Ali ST and others. 'Insights into COVID-19 epidemiology and control from temporal changes in serial interval distributions in Hong Kong' MedRxiv 2022
- 23. Kim D and others. '<u>Estimation of serial interval and reproduction number to quantify the</u> <u>transmissibility of SARS-CoV-2 Omicron variant in South Korea</u>' Viruses 2022: volume 14, issue 3
- 24. Weil AA and others. '<u>Genomic surveillance of SARS-CoV-2 Omicron variants on a</u> <u>university campus</u>' Nature communications 2022: volume 13, issue 1, page 5,240
- 25. Choi G and others. '<u>Viral shedding patterns of the symptomatic SARS-CoV-2 infection</u> according to virus-type dominant periods and vaccination status in Gyeonggi Province, <u>Korea</u>' Epidemiology and Health 2022, page e2023008
- 26. de Michelena P and others. '<u>SARS-CoV-2 RNA load in nasopharyngeal specimens from</u> <u>outpatients with breakthrough COVID-19 due to Omicron BA.1 and BA.2</u>' Journal of Medical Virology 2022: volume 94, issue 12, pages 5,836 to 5,840
- Hay JA and others. 'Quantifying the impact of immune history and variant on SARS-CoV-2 viral kinetics and infection rebound: A retrospective cohort study' eLife 2022: volume 11, issue 11, page 16
- 28. Kandel C and others. '<u>Viral dynamics of the SARS-CoV-2 Omicron Variant among</u> household contacts with 2 or 3 COVID-19 vaccine doses'. Journal of Infection 2022: volume 85, issue 6, pages 666 to 670
- 29. Sikka R and others. '<u>COVID testing in the workplace: return to work testing in an</u> occupational cohort' medRxiv 2022: volume 10
- 30. Anastasiou OE and others. <u>'A simple algorithm based on initial Ct values predicts the</u> <u>duration to SARS-CoV-2 negativity and allows more efficient test-to-release and return-</u> <u>to-work schedules</u>' medRxiv 2022: volume 4

- 31. Cabral P and others. 'Serial viral load analysis by Ddpcr to evaluate Fnc efficacy and safety in the treatment of mild cases of COVID-19' Research Square 2022
- 32. Cegolon L and others. '<u>Early negativization of SARS-CoV-2 infection by nasal spray of seawater plus additives: the RENAISSANCE open-label controlled clinical trial</u>' Pharmaceutics 2022: volume 14, issue 11, page 18
- 33. Chen X and others. '<u>Identification of CKD, bedridden history and cancer as higher-risk</u> comorbidities and their impact on prognosis of hospitalized Omicron patients: a multicentre cohort study' Emerging Microbes and Infections 2022: volume 11, issue 1, pages 2,501 to 2,509
- 34. Chen W and others. <u>Oral Azvudine (FNC) tablets in patients infected with SARS-CoV-2</u> <u>Omicron variant: a retrospective cohort study</u>' MedRxiv 2023
- 35. Colaneri M and others. '<u>Assessing the efficacy of early therapies against SARS-CoV-2 in hematological patients: a real-life study from a COVID-19 referral centre in Northern Italy</u>' Journal of Clinical Medicine 2022: volume 11, issue 24 (no pagination)
- 36. Cosimi LA and others. '<u>Evaluation of the role of home rapid antigen testing to determine</u> isolation period after infection with SARS-CoV-2' medRxiv 2022: volume 6
- 37. Dai EY and others. <u>Viral kinetics of severe acute respiratory syndrome coronavirus 2</u> (SARS-CoV-2) Omicron infection in mRNA-vaccinated individuals treated and not treated with Nirmatrelvir-Ritonavir' MedRxiv 2022
- 38. Gliga S and others. '<u>Rapid selection of sotrovimab escape variants in SARS-CoV-2</u> <u>Omicron infected immunocompromised patients</u>'. Clinical infectious diseases : an official publication of the Infectious Diseases Society of America. 2022: volume 03
- 39. Guo Y and others. <u>Clinical characteristics and outcomes in COVID-19 patients with chronic kidney disease in the SARS-CoV-2 Omicron wave: a case-control study</u>. Research Square [Internet]. 2022.
- 40. Hua Q and others. '<u>Effectiveness of Inactivated COVID-19 Vaccines against COVID-19</u> <u>Caused by the SARS-CoV-2 Delta and Omicron Variants: A Retrospective Cohort Study</u>' 2022: volume 10, issue 10, pages 19
- 41. Kojima N and others. '<u>Duration of COVID-19 PCR positivity for Omicron vs earlier</u> variants'. Journal of Clinical Virology Plus 2022: volume 2, issue 3, page 100,085
- 42. Lefferts B and others. '<u>Antigen test positivity after COVID-19 isolation: Yukon-Kuskokwim</u> <u>Delta Region, Alaska, January to February 2022</u>' Morbidity and Mortality Weekly Report 2022: volume 71, issue 8, pages 293 to 298
- 43. Li H and others. '<u>Association of nirmatrelvir/ritonavir treatment on upper respiratory</u> <u>SARS-CoV-2 RT-PCR negative conversion rates among high-risk patients with COVID-</u> <u>19</u>' Clinical Infectious Diseases 2022: volume 23
- 44. Li J and others. '<u>Higher SARS-CoV-2 shedding in exhaled aerosol probably contributed</u> to the enhanced transmissibility of Omicron BA.5 subvariant' Journal of Medical Virology 2022: volume 95, issue 1 (no pagination)
- 45. Li R and others. '<u>Clinical characteristics and risk factors analysis of viral shedding time in</u> mildly symptomatic and asymptomatic patients with SARS-CoV-2 Omicron variant infection in Shanghai' Frontiers in Public Health 2023: volume 10, page 1,073,387

- 46. Liu L and others. '<u>Effect of nasal irrigation in adults infected with Omicron variant of</u> <u>COVID-19: A quasi-experimental study</u>' Frontiers in Public Health 2022: volume 10, page 1,046,112
- 47. Lu G and others. '<u>Geriatric risk and protective factors for serious COVID-19 outcomes</u> <u>among older adults in Shanghai Omicron wave</u>' Emerging Microbes and Infections 2022: volume 11, issue 1, pages 2,045 to 2,054
- 48. Ma E and others. '<u>Omicron infections profile and vaccination status among 1881 liver</u> <u>transplant recipients: a multi-centre retrospective cohort</u>' Emerging Microbes and Infections 2022: volume 11, issue 1, pages 2,636 to 2,644
- 49. Martin-Blondel G and others. '<u>Time to negative PCR conversion amongst high-risk</u> patients with mild-to-moderate Omicron BA.1 and BA.2 COVID-19 treated with sotrovimab or nirmatrelvir' Clinical Microbiology and Infection 2022: volume 28, page 28
- 50. Okumura N and others. '<u>The first 11 cases of SARS-CoV-2 Omicron variant infection in</u> <u>Japan: A focus on viral dynamics</u>' Global Health and Medicine 2022: volume 4, issue 2, pages 133 to 136
- 51. Pei L and others. '<u>Comorbidities prolonged viral shedding of patients infected with SARS-</u> <u>CoV-2 omicron variant in Shanghai: A multi-center, retrospective, observational study</u>' Journal of Infection and Public Health 2023: volume 16, issue 2, pages 182 to 189
- 52. Shao J and others. '<u>Clinical progression and outcome of hospitalized patients infected</u> <u>with SARS-CoV-2 Omicron variant in Shanghai, China</u>' Vaccines (Basel) 2022: volume 10, issue 9
- 53. Shen Y and others. '<u>An open, prospective cohort study of VV116 in Chinese participants</u> <u>infected with SARS-CoV-2 omicron variants</u>' Emerging Microbes and Infections 2022: volume 11, issue 1, pages 1,518 to 1,523
- 54. Shi C and others. '<u>Oral Liushen pill for patients with COVID-19 : a prospective,</u> randomized, controlled trial' Research Square 2022
- 55. Sun W and others. '<u>Duration of viral shedding of the Omicron variant in asymptomatic</u> and mild COVID-19 cases from Shanghai, China' medRxiv 2022: volume 9
- 56. Tillmann FP and others. 'Effect of third and fourth mRNA-Based booster vaccinations on SARS-CoV-2 neutralizing antibody titer formation, risk factors for non-response, and outcome after SARS-CoV-2 Omicron breakthrough infections in patients on chronic hemodialysis: a prospective multicenter cohort study' Journal of Clinical Medicine 2022: volume 11, issue 11 (no pagination)
- 57. Van Der Veer BMJW and others. '<u>Viral load dynamics in healthcare workers with COVID-</u> <u>19 during Delta and Omicron era</u>' Research Square 2022
- 58. Wei J and others. '<u>Safety and efficacy of oral administrated Cepharanthine in non-hospitalized, asymptomatic or mild COVID-19 patients: a double-blind, randomized, placebo-controlled trial</u>' MedRxiv 2023
- 59. Weng C and others. <u>'Safety and efficacy of Paxlovid against Omicron variants of</u> <u>coronavirus disease 2019 in elderly patients</u>' Infectious Diseases and Therapy 2023: volume 25, page 25
- 60. Wu P and others. '<u>Viral shedding among symptomatic COVID-19 cases infected with the</u> ancestral strain and Omicron BA.2' Research Square 2022

- 61. Wu J and others. '<u>Vaccination Is Associated With Shorter Time to Target Cycle</u> <u>Threshold Value in Patients With SARS-CoV-2 Omicron Variant</u>' Frontiers in Cellular and Infection Microbiology 2022: volume 12, page 943,407
- 62. Xu Y and others. 'Using machine learning models to predict the duration of the recovery of COVID-19 patients hospitalized in Fangcang shelter hospital during the Omicron BA. 2.2 pandemic'. Frontiers in Medicine 2022: volume 9, pages1,001,801
- 63. Xu X and others. 'Efficacy and safety of Reyanning mixture in patients infected with SARS-CoV-2 Omicron variant: a prospective, open-label, randomized controlled trial' Phytomedicine 2023: volume 108, page 154,514
- 64. Xu X and others. 'Efficacy of Lianhua Qingwen for children with SARS-CoV-2 Omicron infection: a propensity score-matched retrospective cohort study'. Phytomedicine 2023: volume 111, page 154,665
- 65. Xu N and others. '<u>Interferon alpha-2b spray shortened viral shedding time of SARS-CoV-2 Omicron variant: An open prospective cohort study</u>'. Frontiers in Immunology 2022: volume 13, page 967,716
- 66. Xu L and others. '<u>Risks factors for virus shedding period of COVID-19 in maintenance</u> <u>hemodialysis patients: a retrospective cohort study</u>' Research Square 2022
- 67. Yan G and others. '<u>The feasibility, safety, and efficacy of Paxlovid treatment in SARS-CoV-2-infected children aged 6 to 14 years: a cohort study</u>' Annals of Translational Medicine 2022: volume 10, issue 11, page 619
- 68. Yang Y and others. '<u>Infection with the SARS-CoV-2 Omicron variant in children with</u> <u>congenital heart disease: A case series study during Shanghai epidemic</u>'. Frontiers in Cardiovascular Medicine 2022: volume 9, page 1,001,780
- 69. Yang Y and others. <u>Factors associated with negative conversion of viral RNA in</u> <u>hospitalized children infected with SARS-CoV-2 Omicron Variant in Shanghai, China: a</u> <u>retrospective analysis</u>' Research Square 2022
- 70. Yin Y and others. '<u>The relationship between early isolation and the duration of viral</u> <u>shedding of mild and asymptomatic infection with SARS-CoV-2 Omicron BA.2 variant</u>' Journal of Infection 2022: volume 85, issue 6, pages e184 to e186
- 71. Ying-Hao P and others. '<u>Clinical characteristics and analysis of risk factors for disease</u> progression of patients with SARS-CoV-2 Omicron variant infection: a retrospective study of 25207 cases in a Fangcang hospital' Frontiers in Cellular and Infection Microbiology 2022: volume 12, page 1,009,894
- 72. Yu SY and others. 'Liver test abnormalities in asymptomatic and mild COVID-19 patients and their association with viral shedding time' World Journal of Hepatology 2022: volume 14, issue 11, pages 1,953 to 1,963
- 73. Zee ST and others. '<u>Impact of COVID-19 vaccination on healthcare worker infection rate</u> and outcome during SARS-CoV-2 Omicron variant outbreak in Hong Kong' 2022: volume 10, issue 8, page 15
- 74. Zeng QL and others. '<u>Clinical characteristics of Omicron SARS-CoV-2 variant infection</u> <u>after non-mRNA-based vaccination in China</u>' Frontiers in Microbiology 2022: volume 13, page 901,826
- 75. Zhang X and others. <u>Immunocompromised states caused the prolonged duration of viral</u> shedding in middle-aged and elderly hemodialysis patients infected with the Omicron

<u>variant of COVID-19</u>' Therapeutic Apheresis and Dialysis: Official Peer Reviewed Journal of the International Society for Apheresis, the Japanese Society for Apheresis, the Japanese Society for Dialysis Therapy 2023: volume 23, page 23

- 76. Zhong W and others. '<u>The efficacy of paxlovid in elderly patients infected with SARS-</u> <u>CoV-2 omicron variants: Results of a non-randomized clinical trial</u>' Frontiers in Medicine 2022: volume 9, page 980,002
- 77. Zhong WZ and others. '<u>Factors associated with prolonged viral shedding in older patients</u> infected with Omicron BA.2.2' Research Square 2022
- 78. Funk S and others. '<u>Cycle threshold values in symptomatic COVID-19 cases in England</u>' medRxiv 2022: volume 16
- 79. Li X and others. '<u>Risk factors for slow viral decline in COVID-19 patients during the 2022</u> Omicron wave' Viruses 2022: volume 14, issue 8, page 4
- 80. Marking U and others. '<u>Correlates of protection, viral load trajectories and symptoms in</u> <u>BA.1, BA.1.1 and BA.2 breakthrough infections in triple vaccinated healthcare workers</u>' medRxiv 2022: volume 3
- 81. Tassetto M and others. '<u>Detection of higher cycle threshold values in culturable SARS-</u> <u>CoV-2 Omicron BA.1 sublineage compared with pre-Omicron variant specimens: San</u> <u>Francisco Bay Area, California July 2021 to March 2022</u>' Morbidity and Mortality Weekly Report 2022: volume 71, issue 36, pages 1151 to 1,154</u>
- 82. Teyssou E and others. '<u>Prolonged replication of BA.1 and BA.2 Omicron lineages</u> <u>compared to Delta variant in nasopharyngeal samples from COVID-19 patients</u>' Infectious Diseases Now 2022: volume 30, page 30
- 83. Townsley H and others. '<u>Non-hospitalised</u>, vaccinated adults with COVID-19 caused by Omicron BA.1 and BA.2 present with changing symptom profiles compared to those with Delta despite similar viral kinetics' medRxiv. 2022 volume 10
- 84. Young B and others. '<u>Comparison of the clinical features, viral shedding and immune</u> response in vaccine breakthrough infection by the Omicron and Delta variants' Research Square 2022
- 85. Erik HE and others. '<u>Contamination-related characteristics of COVID-19: a household</u> <u>survey</u>' Research Square 2022
- 86. Frutos AM and others. '<u>Infection-induced immunity is associated with protection against</u> <u>SARS-CoV-2 infection, but not decreased infectivity during household transmission</u>' MedRxiv 2022: volume 11, page 11
- 87. Kaur H and others. <u>'Correlation of SARS-CoV2 viral growth on cultures, Ct values,</u> <u>SARS-CoV-2 variant and vaccination status in asymptomatic, pre-symptomatic and postinfection asymptomatic COVID patients</u>" American Journal of Clinical Pathology 2022: volume 158 (supplement 1), page S130
- 88. Wei Z and others. <u>'Household transmission of SARS-CoV-2 during the Omicron wave in</u> <u>Shanghai, China:a case-ascertained study</u>' medRxiv 2022: volume 27
- 89. Laitman AM and others. '<u>The SARS-CoV-2 Omicron variant does not have higher nasal</u> viral loads compared to the Delta Variant in symptomatic and asymptomatic individuals' Journal of Clinical Microbiology 2022: volume 60, issue 4, page e0013922

- 90. Suzuki H and others. '<u>Analytical performance of the rapid qualitative antigen kit for the</u> <u>detection of SARS-CoV-2 during widespread circulation of the Omicron variant</u>' Journal of Infection and Chemotherapy 2022: volume 20, page 20
- 91. Wu Q and others. '<u>Viral RNA load in symptomatic and asymptomatic COVID-19 Omicron</u> <u>variant-positive patients</u>' Canadian Respiratory Journal 2022: volume 2022, page 5460400
- 92. Heroes-Recover Network and others. '<u>Association of mRNA vaccination with clinical and</u> virologic features of COVID-19 among US essential and frontline workers' Journal of the American Medical Association 2022: volume 328, issue 15, pages 1,523 to 1,533
- 93. Page MJ and others. '<u>The PRISMA 2020 statement: an updated guideline for reporting</u> systematic reviews' British Medical Journal 2021: volume 372, page n71

# **Annexe A. Methods**

This rapid review aimed to answer the following research questions:

- 1. What is the infectious period of Omicron variant COVID-19?
- 2. What is the difference in transmission from people with asymptomatic compared with symptomatic Omicron variant COVID-19?

Only studies where the majority of the participants in the study have the Omicron variant of COVID-19 (any sub-lineage) were included.

We have conducted previous (unpublished) reviews looking for rapid and systematic reviews covering the infectious period of COVID-19 (searches to 23 February 2022 and 16 May 2022). However, these could not be used as source of evidence for primary studies as these searches aimed at identifying existing rapid and systematic reviews, these reviews will not be used as a source of evidence. The infectious period primary study search therefore started 1 December 2021, shortly after the start of the Omicron wave.

We also conducted a previous (unpublished) review looking at the difference in transmission of COVID-19 between symptomatic and asymptomatic index cases (search to 15 March 2022). This review was used as a source of evidence up to 15 March 2022.

The review questions for this current review were searched for and reported separately within the same report.

Our rapid review approach follows streamlined systematic methodologies (<u>1</u>). In particular, 10% of the screening on title and abstract were screened in duplicate; full text screening, data extraction and risk of bias assessment were performed by one reviewer and checked by another. The review has been reported according to PRISMA guidelines (<u>93</u>).

### Protocol

A protocol was produced a priori.

# Inclusion and exclusion criteria

Article eligibility criteria are summarised in Table A.1.

#### Table A.1. Inclusion and exclusion criteria

	Included	Excluded
Population	All	Animals
Settings	All settings	
Context	COVID-19 pandemic	Other infectious diseases
Intervention or exposure	Omicron variant COVID-19 (any sub-lineage)	
Outcomes	<ul> <li>any measure of infectious period of COVID- 19, including live virus culture (for example, from cytopathic effects in cell cultures, and the isolation of live virus from cell cultures), epidemiology and contract tracing, viral RNA shedding, and, if necessary, incubation period</li> <li>transmission of COVID-19 from people with asymptomatic or symptomatic COVID-19, as measured by secondary attack rates</li> <li>viral load comparisons between people with asymptomatic and symptomatic COVID-19</li> </ul>	
Language	English	
Date of publication	1 December 2021 (infectious period search) and 15 March 2022 (asymptomatic transmission search) to 25 January 2023	
Study design	<ul> <li>Interventional studies</li> <li>Observational studies (cohorts, case controls and cross-sectional studies)</li> </ul>	<ul> <li>systematic or narrative reviews</li> <li>case reports (of single cases)</li> <li>guidelines</li> <li>opinion pieces</li> <li>modelling studies</li> <li>laboratory studies</li> <li>ecological studies</li> </ul>
Publication type	Published and preprint	

# Identification of studies

Our previous review on the difference in transmission of COVID-19 between symptomatic and asymptomatic index cases was used to identify studies published up to 15 March 2022 (latest search date).

We searched OVID Medline, OVID Embase, and preprint servers (medRxiv, bioRxiv, aRxiv, and Research Square, via COVID-19 portfolio) for studies published after 1 December 2021 (for the infectious period of Omicron variant COVID-19) and 15 March 2022 (for the difference in asymptomatic and symptomatic Omicron variant COVID-19 transmission).

# Screening

Screening on title and abstract was undertaken in duplicate by 2 reviewers for at least 10% of the eligible studies (separately for each of the searches), with the remainder completed by one reviewer. Disagreement was resolved by discussion.

Screening on full text was undertaken by one reviewer with a second reviewer checking the excluded full texts, and disagreement resolved by discussion.

If the dominant variant was not stated in a study, participants in studies conducted after 1 December 2022 were considered likely to have Omicron variant COVID-19 (meeting the inclusion criteria for COVID-19 variant), whereas studies not reporting the time period of the study were excluded.

# Data extraction

Summary information for each study was extracted and reported in tabular form. Information included country, setting, study design, objective, outcomes measures, participants, study period, results and any relevant contextual data (such as timing or level of community transmission at the time of the study). This was undertaken by one reviewer and checked by a second.

# Risk of bias assessment

Risk of bias for analytical studies (studies comparing asymptomatic and symptomatic transmission of COVID-19) was assessed using the quality criteria checklist (QCC) for primary research which assesses the methodological quality of a study. This tool can be applied quickly to most study designs to consider core areas of potential bias. Risk of bias was assessed by one reviewer and checked by a second.

Risk of bias for descriptive studies (studies measuring the infectious period of Omicron variant COVID-19) were not assessed.

# Search strategy for infectious period of Omicron variant COVID-19

#### Search strategy Ovid Medline

- 1 exp SARS-CoV-2/ (146874)
- 2 exp COVID-19/ (210509)
- 3 (corona\* adj1 (virus\* or viral\*)).tw,kw,kf. (5961)
- 4 (CoV not (Coefficien\* or "co-efficien\*" or covalent\* or Covington\* or covariant\* or covarianc\* or "cut-off value\*" or "cutoff value\*" or "cut-off volume\*" or "cutoff volume\*" or "combined optimi?ation value\*" or "central vessel trunk\*" or CoVR or CoVS)).tw,kw,kf. (113432)
- 5 (coronavirus\* or 2019nCoV\* or 19nCoV\* or "2019 novel\*" or Ncov\* or "n-cov" or "SARS-CoV-2\*" or "SARSCoV-2\*" or SARSCoV2\* or "SARS-CoV2\*" or "severe acute respiratory syndrome\*" or COVID\*2).tw,kw,kf. (344396)
- 6 exp COVID-19 Vaccines/ (19017)
- 7 exp COVID-19 Testing/ (10360)
- 8 or/1-7 (352015)
- 9 ((Transmis\* or transmit\*) adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (13916)
- 10 (Infectious\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (5806)
- 11 (Contagio\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (315)
- 12 (Isolation adj3 (duration\* or time or length\* or period\*)).tw,kw,kf. (3612)
- 13 (shed\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (3487)
- 14 Virus Shedding/ (4172)
- 15 (PCR positiv\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (456)
- 16 (Viral proliferat\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (6)
- 17 cycl\* threshold\*.tw,kw,kf. (2484)
- 18 CT value\*.tw,kw,kf. (4742)
- 19 (peak\* adj1 (vir\* load\* or vir\* concentration)).tw,kw,kf. (384)
- 20 9 or 10 or 11 or 12 or 13 or 14 or 15 or 16 or 17 or 18 or 19 (36921)
- 21 8 and 20 (4990)
- 22 (Viral Load/ or exp Disease Transmission, Infectious/) and exp Time/ (9404)
- 23 COVID-19/tm and exp Time/ (304)
- 24 8 and 22 (291)
- 25 21 or 23 or 24 (5428)
- 26 limit 25 to dt=20211201-20230126 (1822)

#### Search strategy Ovid Embase

- 1 exp severe acute respiratory syndrome coronavirus 2/ (86769)
- 2 coronavirus disease 2019/ (309119)
- 3 experimental coronavirus disease 2019/ (18)
- 4 (corona\* adj1 (virus\* or viral\*)).tw,kw. (6270)

- 5 (CoV not (Coefficien\* or co-efficien\* or covalent\* or covington or covariant\* or covarianc\* or "cut-off value\*" or "cutoff value\*" or "cut-off volume\*" or "cutoff volume\*" or "combined optimi?ation value\*" or "central vessel trunk" or CoVR or CoVS)).tw,kw. (115399)
- 6 (coronavirus\* or 2019nCoV\* or 19nCoV\* or "2019 novel\*" or Ncov\* or "n-cov" or "SARS CoV-2\*" or "SARSCoV-2\*" or SARSCoV2\* or "SARS-CoV2\*" or "severe acute respiratory syndrome\*" or COVID\*2).tw,kw. (388908)
- 7 COVID-19 Testing/ (6418)
- 8 exp SARS-CoV-2 vaccine/ (29066)
- 9 or/1-8 (415212)
- 10 ((Transmis\* or transmit\*) adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (15531)
- 11 (Infectious\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (7489)
- 12 (Contagio\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (313)
- 13 (Isolation adj3 (duration\* or time or length\* or period\*)).tw,kw,kf. (4573)
- 14 (shed\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (3981)
- 15 virus shedding/ (9802)
- 16 (PCR positiv\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (606)
- 17 (Viral proliferat\* adj5 (duration\* or time\* or length\* or period\* or peak\*)).tw,kw,kf. (7)
- 18 cycl\* threshold\*.tw,kw,kf. (3344)
- 19 CT value\*.tw,kw,kf. (7645)
- 20 (peak\* adj1 (vir\* load\* or vir\* concentration)).tw,kw,kf. (606)
- 21 10 or 11 or 12 or 13 or 14 or 15 or 16 or 17 or 18 or 19 or 20 (50254)
- 22 (exp virus load/ or exp disease transmission/) and (time/ or time factor/) (2479)
- 23 9 and 21 (7127)
- 24 22 or 23 (9570)
- 25 limit 24 to dc=20211201-20230126 (3062)

#### Search strategy for COVID-19 portfolio

Search will be carried out on 26 January 2023, date limited from 1 December 2021.

("transmission period"~5 OR "transmission duration"~5 OR "transmission time"~5 "transmission length"~5 OR "transmitted period"~5 OR "transmitted duration"~5 OR "transmitted time"~5 OR "transmitted length"~5 OR "transmissible period"~5 OR "transmissible duration"~5 OR "transmissible time"~5 OR "transmissible length"~5 OR "infectious duration"~5 OR "infectious period"~5 OR "infectious time"~5 OR "infectious time"~5 OR "infectious length"~5 OR "contagious duration"~5 OR "contagious period"~5 OR "contagious time"~5 OR "contagious length"~5 OR "shedding duration"~5 OR "shedding period"~5 OR "shedding time"~5 OR "shedding length"~5 OR "cycle threshold" OR "cycling threshold" OR "CT value")

# Search strategy for the difference in transmission of COVID-19 from people with asymptomatic and symptomatic Omicron variant COVID-19

#### Search strategy Ovid Medline

- 1 exp SARS-CoV-2/ (146325)
- 2 exp COVID-19/ (209172)
- 3 (corona\* adj1 (virus\* or viral\*)).tw,kw,kf. (5942)
- 4 (CoV not (Coefficien\* or "co-efficien\*" or covalent\* or Covington\* or covariant\* or covarianc\* or "cut-off value\*" or "cutoff value\*" or "cutoff volume\*" or "cutoff volume\*" or "combined optimi?ation value\*" or "central vessel trunk\*" or CoVR or CoVS)).tw,kw,kf. (112571)
- 5 (coronavirus\* or 2019nCoV\* or 19nCoV\* or "2019 novel\*" or Ncov\* or "n-cov" or "SARS-CoV-2\*" or "SARSCoV-2\*" or "SARSCoV-2\*" or "SARSCoV2\* or "SARS-CoV2\*" or "severe acute respiratory syndrome\*" or COVID\*2).tw,kw,kf. (341602)
- 6 exp COVID-19 Vaccines/ (18791)
- 7 exp COVID-19 Testing/ (10322)
- 8 or/1-7 (349211)
- 9 transmiss\*.tw,kf. (466722)
- 10 transmit\*.tw,kf. (193714)
- 11 (breakthrough or break through).tw,kf. (27417)
- 12 viral load\*.tw,kf. (39861)
- 13 viral burden.tw,kf. (1148)
- 14 viral level\*.tw,kf. (370)
- 15 (shed\*1 or shedding).tw,kf. (118634)
- 16 cytopath\* effect\*.tw,kf. (8659)
- 17 Viral Load/ (38565)
- 18 exp Disease Transmission, Infectious/ (80110)
- 19 Cytopathogenic Effect, Viral/ (9596)
- 20 Virus Shedding/ (4169)
- 21 9 or 10 or 11 or 12 or 13 or 14 or 15 or 16 or 17 or 18 or 19 or 20 (861934)
- 22 asymptomatic\*.tw,kf. (185094)
- 23 symptomatic\*.tw,kf. (217451)
- 24 pre-symptomatic.tw,kf. (1861)
- 25 non-symptomatic.tw,kf. (1021)
- 26 (symptom free or symptom-free).tw,kf. (9086)
- 27 no symptom\*.tw,kf. (11375)
- 28 with\* symptom\*.tw,kf. (80153)
- 29 symptomless.tw,kf. (3072)
- 30 symptom\* status.tw,kf. (1600)
- 31 exp Asymptomatic Diseases/ (9921)
- 32 22 or 23 or 24 or 25 or 26 or 27 or 28 or 29 or 30 or 31 (431945)

- 33 8 and 21 and 32 (4684)
- 34 COVID-19/tm [Transmission] (4798)
- 35 32 and 34 (559)
- 36 33 or 35 (4770)
- 37 limit 36 to dt=20220315-20230126 (984)

#### Search strategy Ovid Embase

- 1 exp severe acute respiratory syndrome coronavirus 2/ (86533)
- 2 coronavirus disease 2019/ (308026)
- 3 experimental coronavirus disease 2019/ (18)
- 4 (corona\* adj1 (virus\* or viral\*)).tw,kw. (6298)
- 5 (CoV not (Coefficien\* or co-efficien\* or covalent\* or covington or covariant\* or covarianc\* or "cut-off value\*" or "cutoff value\*" or "cut-off volume\*" or "cutoff volume\*" or "combined optimi?ation value\*" or "central vessel trunk" or CoVR or CoVS)).tw,kw. (114978)
- 6 (coronavirus\* or 2019nCoV\* or 19nCoV\* or "2019 novel\*" or Ncov\* or "n-cov" or "SARS CoV-2\*" or "SARSCoV-2\*" or SARSCoV2\* or "SARS-CoV2\*" or "severe acute respiratory syndrome\*" or COVID\*2).tw,kw. (387643)
- 7 COVID-19 Testing/ (6371)
- 8 exp SARS-CoV-2 vaccine/ (28834)
- 9 or/1-8 (413724)
- 10 transmiss\*.tw,kf. (520930)
- 11 transmit\*.tw,kf. (226337)
- 12 (breakthrough or break through).tw,kf. (39022)
- 13 viral load\*.tw,kf. (62319)
- 14 viral burden.tw,kf. (1475)
- 15 viral level\*.tw,kf. (499)
- 16 (shed\*1 or shedding).tw,kf. (138850)
- 17 cytopath\* effect\*.tw,kf. (9405)
- 18 exp virus load/ (103317)
- 19 exp disease transmission/ (235545)
- 20 cytopathogenic effect/ (10696)
- 21 virus shedding/ (9776)
- 22 10 or 11 or 12 or 13 or 14 or 15 or 16 or 17 or 18 or 19 or 20 or 21 (1078321)
- 23 asymptomatic\*.tw,kf. (270074)
- 24 symptomatic\*.tw,kf. (335224)
- 25 pre-symptomatic.tw,kf. (3247)
- 26 non-symptomatic.tw,kf. (1593)
- 27 (symptom free or symptom-free).tw,kf. (12424)
- 28 no symptom\*.tw,kf. (17051)
- 29 with\* symptom\*.tw,kf. (122988)
- 30 symptomless.tw,kf. (2711)
- 31 symptom\* status.tw,kf. (2520)
- 32 exp asymptomatic disease/ (29248)

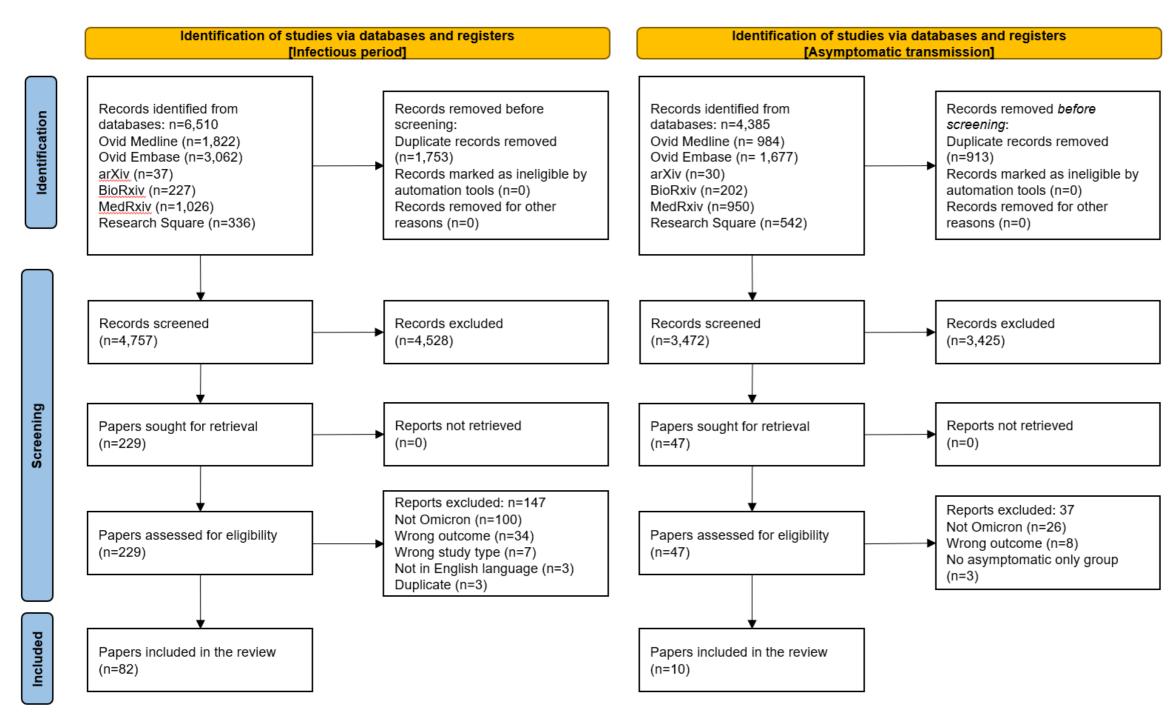
- 33 asymptomatic carrier/ (1267)
- 34 asymptomatic transmission/ (114)
- 35 23 or 24 or 25 or 26 or 27 or 28 or 29 or 30 or 31 or 32 or 33 or 34 (652094)
- 36 asymptomatic coronavirus disease 2019/ (1792)
- 37 22 and 36 (587)
- 38 9 and 22 and 35 (6837)
- 39 37 or 38 (6840)
- 40 limit 39 to dc=20220315-20230126 (1677)

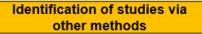
#### Search strategy for COVID-19 portfolio

Search will be carried out on 26 January 2023, date limited from 15 March 2022.

(transmiss\* OR transmit\* OR breakthrough OR "break through" OR "viral load\*" OR "viral burden" OR "viral level\*" OR shed OR sheds OR shedding) AND (asymptomatic\* OR symptomatic\* OR "pre-symptomatic\*" OR "non-symptomatic\*" OR "symptom free" OR "no symptom\*" OR "with\* symptom\*" OR symptomless OR "symptom status")

#### Figure A.1. PRISMA diagram





Records identified from: Expert consultation (n=0)

#### Accessible text version of Figure A.1. PRISMA diagram

A PRISMA diagram showing the flow of studies through this review, ultimately including 82 studies for the research question concerning infectious period, and 10 studies for the research question concerning the difference in transmission of COVID-19 from asymptomatic and symptomatic cases.

From identification of studies via databases and registers for the studies on infectious period, n=6,510 records identified from databases:

- Ovid Medline (n=1,822)
- Ovid Embase (n=3,062)
- arXiv (n=37)
- BioRxiv (n=227)
- MedRxiv (n=1,026)
- Research square (n=336)

From these, records removed before screening:

- duplicate records removed (n=1,753)
- records marked as ineligible by automation tools (n=0)
- records removed for other reasons (n=0)

n=4,757 records screened, of which n=4,528 were excluded, leaving n=229 papers sought for retrieval, all of which were retrieved.

Of the n=229 papers assessed for eligibility, n=147 reports were excluded:

- not Omicron (n=100)
- wrong outcome (n=34)
- wrong study type (n=7)
- not English language (n=3)
- duplicate (n=3)

From identification of studies via other methods, n=0 studies were identified from expert consultation.

Overall, n=82 papers included in the review for the research question concerning infectious period.

From identification of studies via databases and registers for the studies on asymptomatic transmission, n=4,385 records identified from databases:

- Ovid Medline (n=984)
- Ovid Embase (n=1,677)
- arXiv (n=30)
- BioRxiv (n=202)
- MedRxiv (n=950)
- Research square (n=542)

From these, records removed before screening:

- duplicate records removed (n=913)
- records marked as ineligible by automation tools (n=0)
- records removed for other reasons (n=0)

n=3,472 records screened, of which n=3,425 were excluded, leaving n=47 papers sought for retrieval, all of which were retrieved.

Of the n=47 papers assessed for eligibility, n=37 reports were excluded:

- not Omicron (n=26)
- wrong outcome (n=8)
- no asymptomatic only group (n=3)

From identification of studies via other methods, n=0 studies were identified from expert consultation.

Overall, n=10 papers included in the review for the research question concerning asymptomatic transmission.

# About the UK Health Security Agency

UKHSA is responsible for protecting every member of every community from the impact of infectious diseases, chemical, biological, radiological and nuclear incidents and other health threats. We provide intellectual, scientific and operational leadership at national and local level, as well as on the global stage, to make the nation health secure.

<u>UKHSA</u> is an executive agency, sponsored by the <u>Department of Health and Social Care</u>.

© Crown copyright 2023 Version 1

Prepared by: Sean Harrison, Maheen Qureshi, Jennifer Hill For queries relating to this document, please contact: <u>PHCR.Evidence@ukhsa.gov.uk</u>

Published: March 2023 Publishing reference: GOV-14430



You may re-use this information (excluding logos) free of charge in any format or medium, under the terms of the Open Government Licence v3.0. To view this licence, visit <u>OGL</u>. Where we have identified any third party copyright information you will need to obtain permission from the copyright holders concerned.



UKHSA supports the Sustainable Development Goals

