

# SARS-CoV-2, SARS-CoV, and MERS-CoV viral load dynamics, duration of viral shedding, and infectiousness: a systematic review and meta-analysis



Muge Cevik, Matthew Tate, Ollie Lloyd, Alberto Enrico Maraolo, Jenna Schafers, Antonia Ho



## Summary

**Background** Viral load kinetics and duration of viral shedding are important determinants for disease transmission. We aimed to characterise viral load dynamics, duration of viral RNA shedding, and viable virus shedding of severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) in various body fluids, and to compare SARS-CoV-2, SARS-CoV, and Middle East respiratory syndrome coronavirus (MERS-CoV) viral dynamics.

**Methods** In this systematic review and meta-analysis, we searched databases, including MEDLINE, Embase, Europe PubMed Central, *medRxiv*, and *bioRxiv*, and the grey literature, for research articles published between Jan 1, 2003, and June 6, 2020. We included case series (with five or more participants), cohort studies, and randomised controlled trials that reported SARS-CoV-2, SARS-CoV, or MERS-CoV infection, and reported viral load kinetics, duration of viral shedding, or viable virus. Two authors independently extracted data from published studies, or contacted authors to request data, and assessed study quality and risk of bias using the Joanna Briggs Institute Critical Appraisal Checklist tools. We calculated the mean duration of viral shedding and 95% CIs for every study included and applied the random-effects model to estimate a pooled effect size. We used a weighted meta-regression with an unrestricted maximum likelihood model to assess the effect of potential moderators on the pooled effect size. This study is registered with PROSPERO, CRD42020181914.

**Findings** 79 studies (5340 individuals) on SARS-CoV-2, eight studies (1858 individuals) on SARS-CoV, and 11 studies (799 individuals) on MERS-CoV were included. Mean duration of SARS-CoV-2 RNA shedding was 17·0 days (95% CI 15·5–18·6; 43 studies, 3229 individuals) in upper respiratory tract, 14·6 days (9·3–20·0; seven studies, 260 individuals) in lower respiratory tract, 17·2 days (14·4–20·1; 13 studies, 586 individuals) in stool, and 16·6 days (3·6–29·7; two studies, 108 individuals) in serum samples. Maximum shedding duration was 83 days in the upper respiratory tract, 59 days in the lower respiratory tract, 126 days in stools, and 60 days in serum. Pooled mean SARS-CoV-2 shedding duration was positively associated with age (slope 0·304 [95% CI 0·115–0·493];  $p=0\cdot0016$ ). No study detected live virus beyond day 9 of illness, despite persistently high viral loads, which were inferred from cycle threshold values. SARS-CoV-2 viral load in the upper respiratory tract appeared to peak in the first week of illness, whereas that of SARS-CoV peaked at days 10–14 and that of MERS-CoV peaked at days 7–10.

**Interpretation** Although SARS-CoV-2 RNA shedding in respiratory and stool samples can be prolonged, duration of viable virus is relatively short-lived. SARS-CoV-2 titres in the upper respiratory tract peak in the first week of illness. Early case finding and isolation, and public education on the spectrum of illness and period of infectiousness are key to the effective containment of SARS-CoV-2.

**Funding** None.

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## Introduction

Viral load kinetics and the duration of viral shedding are important determinants for disease transmission. They determine the duration of infectiousness, which is a critical parameter to inform effective control measures and disease modelling. Although several studies have evaluated severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) shedding, viral load dynamics and duration of viral shedding reported across studies so far have been heterogeneous.<sup>1</sup> In several case series with serial respiratory sampling, peak viral load was observed just

before, or at the time of, symptom onset.<sup>2–4</sup> Viral RNA shedding was reported to be persistent in the upper respiratory tract and in faeces for more than 1 month after illness onset.<sup>1</sup> However, the duration of SARS-CoV-2 RNA detection has not been well characterised. A comprehensive understanding of viral load dynamics, length of viral shedding, and how these measures relate to other factors, such as age and disease severity, is lacking.

We aimed to characterise the viral load dynamics of SARS-CoV-2, duration of viral RNA shedding by RT-PCR, and viable virus shedding in various body fluids,

*Lancet Microbe* 2021; 2: e13–22

Published Online  
November 19, 2020  
[https://doi.org/10.1016/S2666-5247\(20\)30172-5](https://doi.org/10.1016/S2666-5247(20)30172-5)

Division of Infection and Global Health Research, School of Medicine, University of St Andrews, Fife, UK (M Cevik MRCPP); NHS Lothian Infection Service, Regional Infectious Diseases Unit, Western General Hospital, Edinburgh, UK (M Cevik, O Lloyd MRCPP, J Schafers MRCPP); Respiratory Medicine, Queen Elizabeth University Hospital, Glasgow, UK (M Tate MRCPP); Edinburgh Medical School, College of Medicine and Veterinary Medicine, University of Edinburgh, Edinburgh, UK (O Lloyd); First Division of Infectious Diseases, Cotugno Hospital, AORN dei Colli, Naples, Italy (A E Maraolo MD); and MRC-University of Glasgow Centre for Virus Research, University of Glasgow, Glasgow, UK (A Ho PhD)

Correspondence to:  
Dr Muge Cevik, Division of Infection and Global Health Research, School of Medicine, University of St Andrews, Fife KY16 9TF, UK  
[mc349@st-andrews.ac.uk](mailto:mc349@st-andrews.ac.uk)

### Research in context

#### Evidence before this study

Understanding when patients are most infectious and the duration of infectiousness are of critical importance to controlling the COVID-19 pandemic. The duration of RNA detection across human coronaviruses has not been well characterised, and comprehensive understanding about viral load dynamics and the duration of viral shedding in severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) is lacking. We retrieved all articles reporting the dynamics and the duration of SARS-CoV-2, SARS-CoV, and Middle East respiratory syndrome coronavirus (MERS-CoV) shedding in various specimens through systematic searches of major databases. Our research identified publications that included terms related to viral dynamics and viral shedding. We included case series, cohort studies, and randomised controlled trials in which the viral dynamics or the duration of viral shedding was reported. We excluded case reports, case series with fewer than five patients, and studies that did not have a clear time of symptom onset.

#### Added value of this study

To our knowledge, this is the first systematic review and meta-analysis that has examined and compared the viral dynamics of the three highly pathogenic human coronaviruses: SARS-CoV-2, SARS-CoV, and MERS-CoV. The results provide a comprehensive understanding regarding their viral kinetics and duration of shedding. Mean SARS-CoV-2 RNA shedding duration was 17.0 days (maximum shedding duration 83 days) in upper respiratory tract, 14.6 days (maximum 59 days) in lower respiratory tract, 17.2 days (maximum 35 days) in stool, and 16.6 days (maximum 60 days) in serum samples. Pooled mean SARS-CoV-2 shedding duration was positively associated with

age. No study detected live virus beyond day 9 of illness, despite persistently high viral loads. SARS-CoV-2 viral load in the upper respiratory tract appeared to peak in the first week of illness, whereas SARS-CoV and MERS-CoV peaked later. Several studies reported similar viral loads at the start of infection among asymptomatic and symptomatic patients infected with SARS-CoV-2; however, most studies demonstrated faster viral clearance in asymptomatic individuals, as also seen in MERS-CoV, suggesting a shorter infectious period but with similar potential transmissibility at the onset of infection.

#### Implications of all the available evidence

Our study shows that despite evidence of prolonged SARS-CoV-2 RNA shedding in respiratory and stool samples, viable virus appears to be short-lived. Therefore, RNA detection cannot be used to infer infectiousness. High titres of SARS-CoV-2 are detected early in the disease course, with an early peak observed at the time of symptom onset to day 5 of illness; this finding probably explains the efficient spread of SARS-CoV-2 compared with SARS-CoV and MERS-CoV. This has important implications for SARS-CoV-2 transmission in the community and hospital setting, emphasising the importance of early case finding and prompt isolation as well as public education about the spectrum of illness. Our study shows that isolation practices should be commenced with the start of first symptoms, which can include mild and atypical symptoms, preceding typical symptoms of COVID-19 such as cough and fever. However, given the potential delays in isolation of patients, even the early detection and isolation strategy might not be fully effective in containing SARS-CoV-2.

and to compare SARS-CoV-2 viral dynamics with those of SARS-CoV and Middle East respiratory syndrome coronavirus (MERS-CoV).

## Methods

### Search strategy and selection criteria

We retrieved all English-language research articles reporting viral dynamics or the duration of shedding of SARS-CoV-2, SARS-CoV, or MERS-CoV in various specimens through systematic searches of major databases, including MEDLINE, Embase, Europe PubMed Central, *medRxiv*, and *bioRxiv*, and the grey literature from Jan 1, 2003, to June 6, 2020, using medical subject headings terms (appendix p 14). We also manually screened the references of included original studies to obtain additional studies. Studies published before 2003 were excluded because the first recognised case of SARS-CoV was identified in March, 2003.

Studies were eligible if they met the following inclusion criteria: report on SARS-CoV-2, SARS-CoV, or MERS-CoV

infection, and report viral load kinetics, duration of viral shedding, or viable virus shedding. We excluded review papers; animal studies; studies on environmental sampling; case reports and case series with less than five participants, due to likely reporting bias; papers in which the starting point of viral shedding was not clear or reported from post hospital discharge; and modelling studies with no original data.

### Data extraction

Two authors (MT and OL) screened and retrieved articles according to the eligibility criteria. Four reviewers (MT, OL, JS, and MC) reviewed full-text articles and selected articles to be included. From each study, the following variables were extracted as a minimum: name of first author, year of publication, city and country, sample size, median age, sex ratio, time from symptom onset to viral clearance detected by RT-PCR and culture in different specimens, and longest reported time to viral clearance. If these data were not reported, we also contacted the

See Online for appendix

authors to request the data. If available, we extracted data on peak viral load, clinical outcome, and reported factors associated with duration of viral shedding.

Two authors (OL and JS) independently assessed study quality and risk of bias using the Joanna Briggs Institute Critical Appraisal Checklist tools,<sup>5</sup> which comprise standardised checklists, for the different study designs included in this review. Any disagreements regarding grading of quality were resolved through discussion with a third author (MC).

### Data analysis

For every study included, we calculated the mean duration of viral shedding and 95% CIs. We applied the random-effects model to estimate a pooled effect size. We generated forest plots to show the detailed representation of all studies based on the effect size and 95% CI. If not reported, we derived means and SDs from sample size, and median, IQR, minimum, and maximum values.<sup>6</sup> Heterogeneity between studies was quantified by the  $I^2$  index and Cochran's Q test. We did not assess publication bias because usual appraisal methods are uninformative when meta-analysed studies do not include a test of significance. We used a weighted meta-regression with an unrestricted maximum likelihood model to assess the effect of potential moderators on the pooled effect size ( $p < 0.05$  was considered to be significant). The eligibility criterion for meta-regression was the presence of at least ten studies (referring to one virus) for each covariate. All statistical analyses were done with Comprehensive Meta-Analysis (version 3) software (Biostat, Englewood, NJ, USA).

This systematic review is registered with PROSPERO, CRD42020181914, and will be updated periodically.

### Role of the funding source

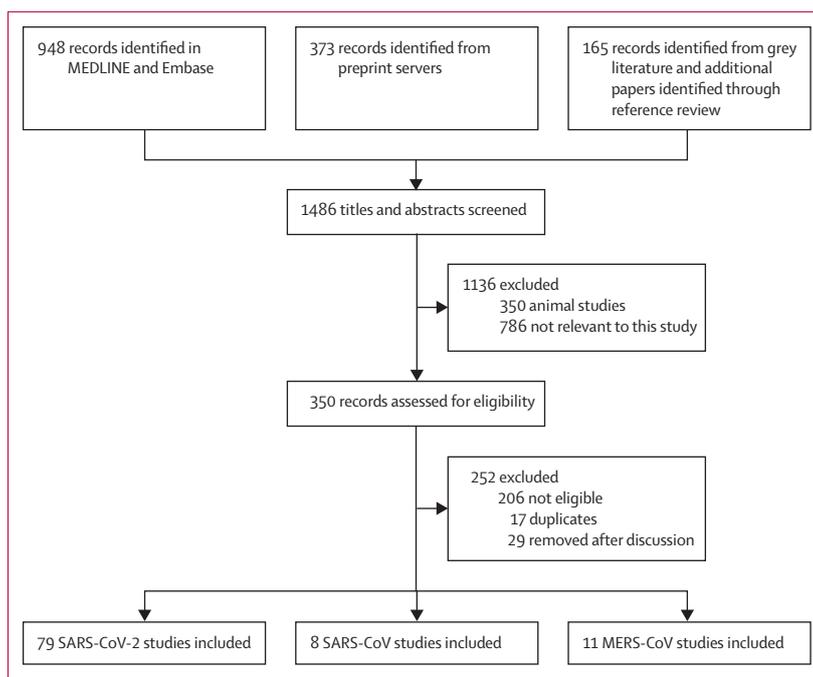
There was no funding source for this study. The corresponding author and senior author (AH) had full access to all the data in the study and had final responsibility for the decision to submit for publication.

### Results

The systematic search identified 1486 potentially relevant articles. 350 articles were retrieved for full-text review. After reviewing the eligibility criteria, 79 studies (5340 individuals) on SARS-CoV-2,<sup>2,4,7-82</sup> eight (1858 individuals) on SARS-CoV,<sup>83-90</sup> and 11 (799 individuals) on MERS-CoV<sup>91-101</sup> were included (figure 1).

Of the 79 papers included, 58 studies were done in China (appendix pp 1-4).<sup>2,10-12,14-17,19-21,23-28,35-41,43,44,48-50,52-63,65-73,75-82</sup> 73 studies included only patients who were admitted to hospital.<sup>3,4,7-29,31,32,34-49,52-54,56-82</sup> Six studies reported viral load dynamics exclusively in children (age younger than 16 years).<sup>7-12</sup> Two additional studies included children, but data on viral load dynamics were presented in aggregate with adults.<sup>13,14</sup>

61 studies reported median or maximum viral RNA shedding in at least one body fluid and were eligible



**Figure 1: Study selection**

MERS-CoV=Middle East respiratory syndrome coronavirus. SARS-CoV=severe acute respiratory syndrome coronavirus. SARS-CoV-2=severe acute respiratory syndrome coronavirus 2.

for quantitative analysis,<sup>3,4,7,9-12,14-17,19-22,24,25,27-30,34-43,45,47,48,50,51,53,54,57-65,</sup>

<sup>67-76,78-80,82</sup> and six studies provided duration of shedding stratified by illness severity only.<sup>13,39,52,55,77,81</sup> Of those studies, 43 (including 3229 individuals) reported duration of shedding in the upper respiratory tract (mean viral shedding duration 17.0 days [95% CI 15.5-18.6]; figure 2), seven (260 individuals) in the lower respiratory tract (14.6 days [9.3-20.0]; appendix p 6), 13 (586 individuals) in stool samples (17.2 days [14.4-20.1]; appendix p 7), and two (108 individuals) in serum samples (16.6 days [3.6-29.7]; appendix p 6). Maximum duration of RNA shedding reported was 83 days in the upper respiratory tract,<sup>35</sup> 59 days in the lower respiratory tract,<sup>27</sup> 126 days in stool samples,<sup>88</sup> and 60 days in serum samples.<sup>78</sup>

Studies reporting duration of viral shedding in upper respiratory tract and stool samples were eligible for meta-regression analysis. Pooled mean viral shedding duration was positively associated with age (slope 0.304 [95% CI 0.115-0.493];  $p=0.0016$ ), but not sex ( $p=0.28$ ; appendix pp 7-8). When adjusted for the proportion of males in a multivariable analysis, mean age was positively associated with the mean duration of viral shedding in upper respiratory tract specimens ( $p=0.0029$ ). There was a positive but non-significant association between mean age and duration of shedding in stool samples ( $p=0.37$ ; appendix p 8).

Eight of 13 studies evaluating SARS-CoV-2 viral load in serial upper respiratory tract samples showed peak viral loads inferred from cycle threshold values within the first week of symptom onset.<sup>2-4,8,15-23</sup> The highest viral loads were

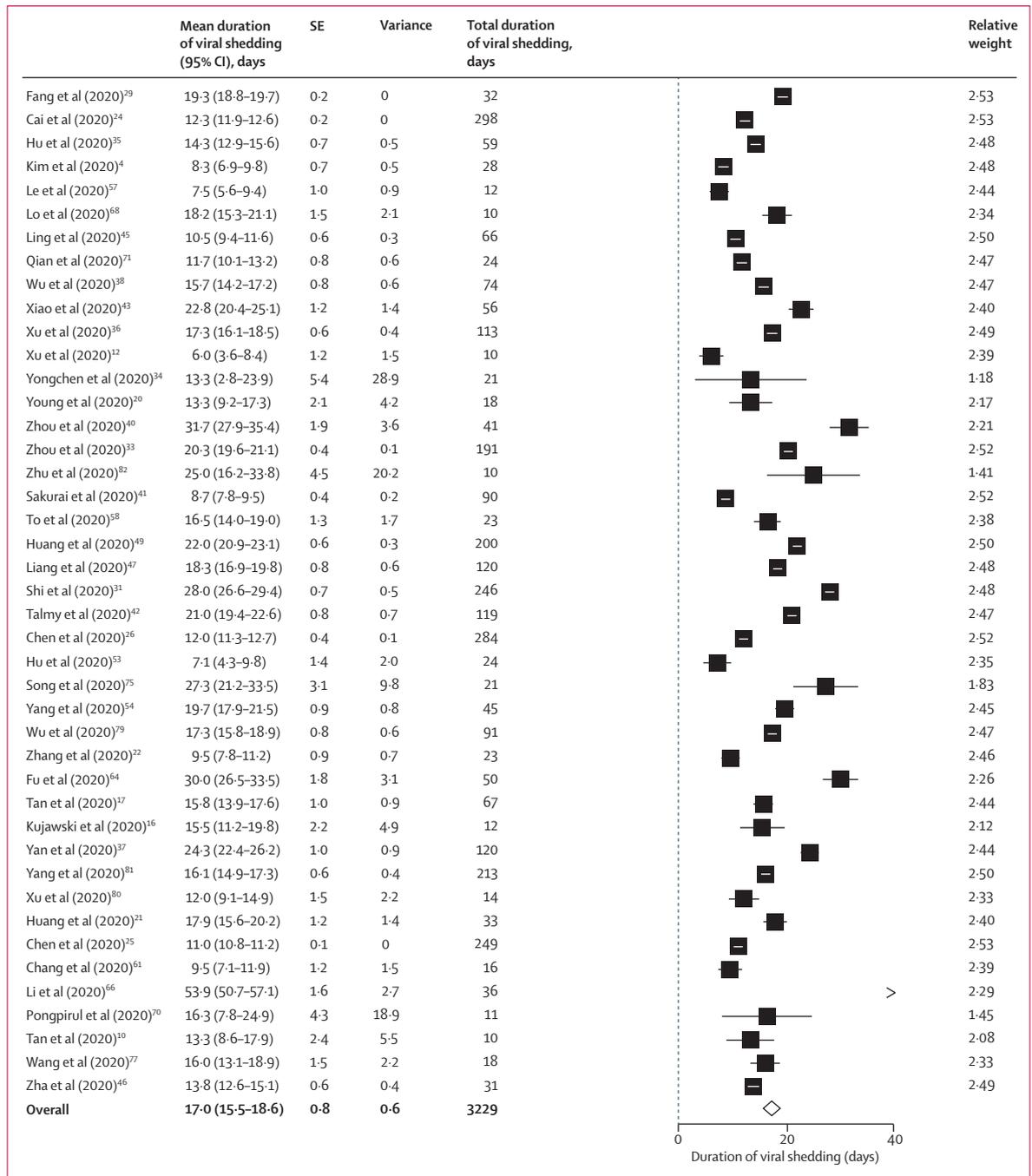


Figure 2: Pooled mean duration (days) of SARS-CoV-2 shedding from the upper respiratory tract (random-effects model) SARS-CoV-2=severe acute respiratory syndrome coronavirus 2.

reported soon after or at the time of symptom onset,<sup>2,8,15,21,23</sup> or at day 3–5 of illness,<sup>3,4,20</sup> followed by a consistent decline.

Five studies that evaluated viral load dynamics in lower respiratory tract samples observed a peak viral load in the second week of illness.<sup>3,4,17,21,23</sup> By contrast, the dynamics of SARS-CoV-2 shedding in stool samples was erratic, with highest viral loads reported on day 7,<sup>17</sup> 2–3 weeks,<sup>22,23</sup> and up

to 5–6 weeks after symptom onset.<sup>21</sup> Although two studies reported significantly higher viral titres in stool samples than in respiratory samples,<sup>8,23</sup> one study reported lower viral load in stool samples than in both lower and upper respiratory tract samples at the time of symptom onset.<sup>21</sup>

20 studies evaluated duration of viral RNA shedding based on disease severity. 13 of these studies reported

longer duration of viral shedding in patients with severe illness than in those with non-severe illness,<sup>17,23–34</sup> whereas five studies in upper respiratory tract samples<sup>16,18,35–37</sup> and one study in stool samples<sup>38</sup> reported similar shedding durations according to disease severity. One study reported shorter duration of viral shedding in moderate to severe illness than in mild to moderate illness.<sup>39</sup> Six studies compared viral shedding among individuals with severe illness versus non-severe illness:<sup>17,23,25,26,36,37</sup> five studies showed significantly longer duration of shedding among those with severe illness than among those with non-severe illness,<sup>17,23,25,26,36</sup> and one study observed no difference<sup>37</sup> (table 1).

All but one study<sup>40</sup> that examined the effect of age on SARS-CoV-2 shedding identified an association between older age (older than 60 years) and prolonged viral RNA shedding.<sup>23,24,26,31,35–37,41–43</sup> Three studies identified age as an independent risk factor for delayed viral clearance.<sup>23,24,36</sup> Male sex was also associated with prolonged shedding,<sup>23,36,44</sup> and the association remained significant even when patients were stratified based on illness severity.<sup>23,36</sup> Corticosteroid treatment was associated with delayed viral clearance in four studies,<sup>31,36,45,46</sup> and one study that recruited 120 patients with critical illness found no difference between corticosteroid and control groups.<sup>47</sup>

A randomised, placebo-controlled trial of remdesivir in adults with severe COVID-19 found a similar decline in viral load over time in remdesivir and control groups, as well as similar proportions of patients with undetectable viral RNA at 28 days.<sup>102</sup> In a phase 2, open-label study evaluating interferon beta-1b, lopinavir–ritonavir, and ribavirin, a shorter duration of viral shedding was seen with combination treatment than with the control.<sup>48</sup> None of the antiviral regimens (chloroquine, oseltamivir, arbidol, and lopinavir–ritonavir) independently improved viral RNA clearance.<sup>26,49</sup> In a retrospective study of 284 patients, lopinavir–ritonavir use was associated with delayed viral clearance even after adjusting for confounders.<sup>26</sup>

12 studies reported viral load dynamics or duration of viral shedding among individuals with asymptomatic SARS-CoV-2 infection (table 2); two demonstrated lower viral loads among asymptomatic individuals than among symptomatic individuals,<sup>8,50</sup> and four found similar initial viral loads.<sup>13,14,51,52</sup> However, Chau and colleagues reported significantly lower viral load in asymptomatic individuals during the follow-up than in symptomatic individuals.<sup>51</sup> Faster viral clearance was observed in asymptomatic individuals in five of six studies.<sup>13,26,51,53,54</sup> The exception, Yongchen and colleagues, found longer shedding duration among asymptomatic cases, but the difference was not significant.<sup>34</sup>

We identified 11 studies that attempted to isolate live virus. All eight studies that attempted virus isolation in respiratory samples successfully cultured viable virus within the first week of illness.<sup>3,9,16,52,55–58</sup> No live virus was

	Classification of illness severity	Median (IQR*) duration of SARS-CoV-2 positivity in cohort, days	Viral dynamics in patients with severe illness vs those with non-severe illness	p value
Chen et al (2020) <sup>25</sup>	ICU vs non-ICU patients	11 (95% CI 10–12)	Median time to viral clearance significantly longer in ICU vs non-ICU patients (HR 3.17, 95% CI 2.29–4.37)	Only HR provided
Chen et al (2020) <sup>36</sup>	China CDC guideline (version 7)	12 (8–16)	Shedding duration varies by severity: asymptomatic 6 days; mild 10 days; moderate 12 days; serious 14 days; critical 32 days	<0.0001
Tan et al (2020) <sup>37</sup>	China CDC guideline (version 6)	Nasopharyngeal swab: 12 (range 3–38); any sample: 22 (range 3–38)	Viral shedding significantly longer in patients with severe illness: any sample 23 days vs 20 days (note that nasopharyngeal swab 14 vs 11 days was non-significant)	0.023 (any sample)
Xu et al (2020) <sup>38</sup>	WHO criteria	17 (13–32)	Higher proportion of patients with severe illness had shedding >15 days (34.2% vs 16.2%)	0.049
Yan et al (2020) <sup>37</sup>	China CDC guideline (version 6)	23 (18–32)	No difference in shedding duration (general illness 23 days vs severe illness 26 days vs critical illness 28 days)	0.51
Zheng et al (2020) <sup>33</sup>	China CDC guideline (version 6)	Respiratory sample: 18 (13–29)	Shedding duration significantly longer in patients with severe illness (21 vs 14 days) in respiratory samples; no difference in shedding duration in stool or serum samples	0.04

CDC=Center for Disease Control and Prevention. HR=hazard ratio. ICU=intensive care unit. SARS-CoV-2=severe acute respiratory syndrome coronavirus 2. \*IQR unless otherwise stated.

**Table 1: Severity of illness and viral dynamics**

isolated from any respiratory samples taken after day 8 of symptoms in three studies,<sup>3,55,56</sup> or beyond day 9 in two studies<sup>16,52</sup> despite persistently high viral RNA loads. One study demonstrated the highest probability of positive culture on day 3 of symptoms.<sup>55</sup> Arons and colleagues cultured viable virus 6 days before typical symptom onset; however, onset of symptoms was unclear.<sup>52</sup>

The success of viral isolation correlated with viral load quantified by RT-PCR. No successful viral culture was obtained from samples with a viral load below  $1 \times 10^6$  copies per mL in one study,<sup>3</sup> cycle threshold values higher than 24 in another study,<sup>55</sup> or higher than 34 in other studies,<sup>52,56</sup> with culture positivity declining with increasing cycle threshold values.<sup>56</sup> Several other studies cultured live virus from RT-PCR-positive specimens; however, they did not correlate these results with viral load titres.<sup>9,57,58</sup>

One study reported the duration of viable virus shedding in respiratory samples; time to clearance from symptom onset was 3–12 days in upper respiratory tract samples and 5–13 days in lower respiratory tract samples, and no positive viral culture was obtained after day 4 in upper respiratory tract infection and day 8 in lower respiratory tract infection.<sup>3</sup> Arons and colleagues cultured viable virus from the respiratory tract in one of three asymptomatic cases.<sup>52</sup>

Viral culture was successful in two of three RT-PCR-positive patients in one study, but the timepoints

	Median (IQR)* duration of SARS-CoV-2 positivity in asymptomatic individuals, days	Viral dynamics in asymptomatic vs symptomatic individuals	p value
Arons et al (2020) <sup>52</sup>	Not reported	No difference in viral load	Not reported
Chau et al (2020) <sup>51</sup>	Not reported	Initial viral load similar; asymptomatic individuals had significantly lower viral load during follow-up and faster viral clearance than symptomatic individuals	0.027
Chen et al (2020) <sup>26</sup>	6 (4–10)	Significantly shorter duration of viral shedding among asymptomatic cases, with increasing shedding duration associated with increasing illness severity	<0.0001
Han et al (2020) <sup>8</sup>	Not reported	Symptomatic children had higher initial RNA load in nasopharyngeal swab specimens than asymptomatic children (9.01 vs 6.32 log <sub>10</sub> copies per mL)	0.048
Hu et al (2020) <sup>53</sup>	6 (2–12)	Asymptomatic individuals had shorter duration of viral shedding compared with pre-symptomatic individuals (median duration of viral shedding was 6 days [2–12] vs 12 days [12–14])	Not reported
Lavezzo et al (2020) <sup>34</sup>	Not reported	No difference in viral load	p=0.62 (E gene); p=0.74 (RdRp gene)
Le et al (2020) <sup>57</sup>	9	Not reported	Not applicable
Sakurai et al (2020) <sup>41</sup>	9 (6–11)	Not reported	Not applicable
Yang et al (2020) <sup>54</sup>	8 (3–12)	Significantly shorter duration of viral shedding from nasopharynx swabs was observed among asymptomatic vs symptomatic individuals	0.001
Yongchen et al (2020) <sup>34</sup>	18 (range 5–28)	Longer shedding duration among asymptomatic cases (median 18 days [range 5–28]) vs non-severe (10 days [2–21]) and severe (14 days [9–33]) cases	Not reported
Zhang et al (2020) <sup>13</sup>	9.6	Initial viral load similar; viral clearance occurred earlier in the asymptomatic (9.6 days) and symptomatic individuals (9.7 days), vs pre-symptomatic group (13.6 days)	<0.05
Zhou et al (2020) <sup>50</sup>	Not reported	Significantly higher viral load in symptomatic (n=22) vs asymptomatic (n=9) individuals (median cycle threshold value 34.5 [IQR 37.5–39.5] vs 39.0 [32.2–37.0]), but duration of shedding was similar	Not reported

SARS-CoV-2=severe acute respiratory syndrome coronavirus 2. \*IQR, if available, unless otherwise stated.

**Table 2: SARS-CoV-2 viral dynamics in asymptomatic individuals compared with symptomatic individuals**

from symptom onset were not reported.<sup>59</sup> Andersson and colleagues were unable to culture virus from 27 RT-PCR-positive serum samples.<sup>60</sup>

Of eight studies on SARS-CoV, none reported mean or median duration of viral shedding and thus were not eligible for quantitative analysis. The maximum duration of viral shedding reported was 8 weeks in upper respiratory tract,<sup>83,84</sup> 52 days in lower respiratory tract,<sup>83,85</sup> 6–7 weeks in serum,<sup>86</sup> and 126 days in stool samples.<sup>83,85,87–89</sup> Studies that evaluated SARS-CoV kinetics found low viral load in the initial days of illness, increasing after the

first week of illness in upper respiratory tract samples, peaking at day 10,<sup>90</sup> or days 12–14,<sup>87</sup> and declining after weeks 3–4.<sup>84</sup> High viral loads correlated with severity of illness and poor survival.<sup>84</sup> Although Chen and colleagues identified an association between younger age and lower viral titres,<sup>84</sup> Leong and colleagues found no difference.<sup>89</sup> Viable SARS-CoV was isolated from stool and respiratory samples up to 4 weeks, and urine specimens up to day 36 from symptom onset.<sup>83,86</sup> All attempts to isolate virus from RT-PCR-positive stool specimens collected more than 6 weeks after disease onset failed.<sup>85</sup> The isolation probability for stool samples was approximately five to ten times lower than for respiratory specimens.<sup>83</sup>

We identified 11 studies on MERS-CoV. Three studies (324 participants) reporting MERS-CoV shedding in the upper respiratory tract<sup>99–101</sup> and four studies (93 participants) reporting MERS-CoV shedding in the lower respiratory tract<sup>91,92,96,101</sup> were included in the quantitative analysis. The mean shedding duration was 15.3 days (95% CI 11.6–19.0) in the upper respiratory tract and 16.3 days (13.8–18.9) in the lower respiratory tract (figures 3, 4). Only one study reported duration of viral shedding in serum with a maximum of 34 days.<sup>91</sup> In a small study, mortality was higher in patients with viraemia (viral RNA in blood).<sup>92</sup> In upper and lower respiratory tract specimens, prolonged shedding was associated with illness severity<sup>93,94</sup> and survival,<sup>95</sup> with the shortest duration observed in asymptomatic individuals.<sup>93</sup> Peak viral loads were observed between days 7 and 10, and higher viral loads were observed among patients with severe illness and fatal outcome.<sup>91,93,94,96,97</sup> Differences in viral loads between survivors and fatal cases was more pronounced in the second week of illness (p=0.0006).<sup>97</sup> The proportion of successful viable culture was 6% in respiratory samples, with a viral load value below 1×10<sup>7</sup> copies per mL.<sup>98</sup>

All but 11 studies (six cohort studies, two cross-sectional studies, and one randomised controlled trial on SARS-CoV-2 and two cohort studies on MERS-CoV) were case series, the majority of which recruited non-consecutive patients and were therefore prone to possible selection bias (appendix pp 9–13).

## Discussion

This systematic review and meta-analysis provides comprehensive data on the viral dynamics of SARS-CoV-2, including the duration of RNA shedding and viable virus isolation. Our findings suggest that, although patients with SARS-CoV-2 infection might have prolonged RNA shedding of up to 83 days in upper respiratory tract infection, no live virus was isolated from culture beyond day 9 of symptoms despite persistently high viral RNA loads. This finding is supported by several studies demonstrating an association between viral load and viability of virus, with no successful culture from samples below a certain viral load threshold. These findings indicate that, in clinical practice, repeat testing might not

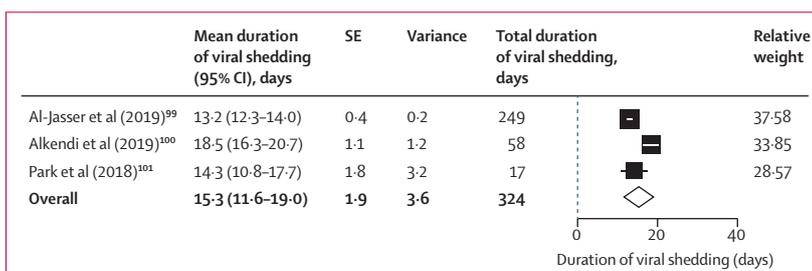
be indicated to deem patients no longer infectious. Duration of infectiousness and subsequent isolation timelines could reflect viral load dynamics and could be counted from symptom onset for 10 days in non-severe cases.

SARS-CoV-2 viral load appeared to peak in the upper respiratory tract within the first week after symptom onset, and later in the lower respiratory tract. By contrast, the viral load of SARS-CoV peaked at days 10–14 of illness and that of MERS-CoV peaked at 7–10 days of illness. Combined with isolation of viable virus in respiratory samples primarily within the first week of illness, patients with SARS-CoV-2 infection are likely to be most infectious in the first week of illness, emphasising the importance of immediate isolation with symptom onset early in the course of illness. Several studies report viral load peaks during the prodromal phase of illness or at the time of symptom onset.<sup>2–4,8,15–21</sup> providing a rationale for the efficient spread of SARS-CoV-2. This finding is supported by the observation in contact-tracing studies that the highest risk of transmission occurs very early in the disease course (a few days before and within the first 5 days after symptom onset).<sup>103,104</sup> Although modelling studies estimated potential viral load peak before symptom onset, we did not identify any study that confirms pre-symptomatic viral load peak.<sup>15</sup>

Similar to SARS-CoV, SARS-CoV-2 can be detected in stool samples for prolonged periods, with high viral loads detected even after 3 weeks of illness. In SARS-CoV, RNA prevalence in stool samples was high, with almost all studies reporting shedding in stools. Although viable SARS-CoV was isolated during up to 4 weeks of illness, faecal–oral transmission was not considered to be a primary driver of infection. By contrast, none of the studies in MERS-CoV reported duration of viral shedding in stool samples and RNA detection was low.<sup>97,105</sup> So far, only a few studies have demonstrated viable SARS-CoV-2 in stool samples.<sup>59,106</sup> Thus, the role of faecal shedding in viral transmission remains unclear.

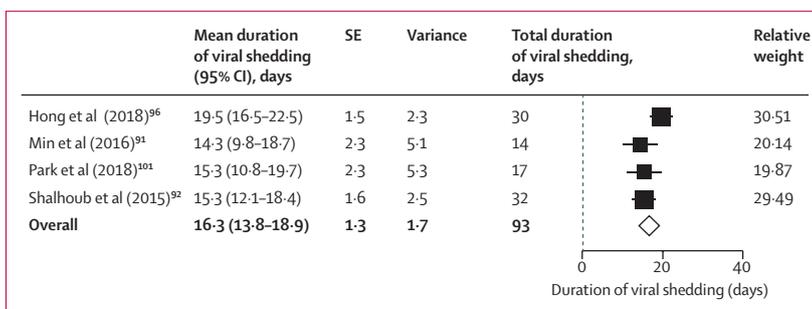
Viral loads appear to be similar between asymptomatic and symptomatic individuals infected with SARS-CoV-2. Nevertheless, most studies demonstrate faster viral clearance among asymptomatic individuals than those who are symptomatic. This finding is in keeping with viral kinetics observed with other respiratory viruses such as influenza and MERS-CoV, in which people with asymptomatic infection have a shorter duration of viral shedding than symptomatic individuals.<sup>93,107</sup> However, data on the shedding of infectious virus in asymptomatic individuals are too scarce to quantify their transmission potential in order to inform policy on quarantine duration in the absence of testing.

To our knowledge, this is the first systematic review to comprehensively examine and compare SARS-CoV-2, SARS-CoV, and MERS-CoV viral dynamics, and the first meta-analysis of viral shedding duration. Our study has limitations. First, almost all patients in the included



**Figure 3:** Pooled mean duration (days) of MERS-CoV shedding from the upper respiratory tract (random-effects model)

MERS-CoV=Middle East respiratory syndrome coronavirus.



**Figure 4:** Pooled mean duration (days) of MERS-CoV shedding from the lower respiratory tract (random-effects model)

MERS-CoV=Middle East respiratory syndrome coronavirus.

studies received a range of treatments, which might have modified the shedding dynamics. Second, our meta-analysis identified substantial study heterogeneity, probably due to differences in study population, follow-up, and management approaches. Furthermore, shedding duration is reported as median with IQR for most studies, but meta-analysis necessitates conversion to mean with SD.<sup>6</sup> The validity of this conversion is based on the assumption that duration of viral shedding is normally distributed, which might not apply to some studies. Last, although there is probably a broad overlap, the true clinical window of infectious shedding might not entirely align with viral culture duration.

We identified a systematic review of SARS-CoV-2 viral load kinetics that included studies published up until May 12, 2020.<sup>108</sup> This review included 26 case reports and 13 case series involving less than five individuals, which did not meet our eligibility criteria; these studies are prone to substantial selection bias, reporting atypical cases with prolonged viral shedding. Additionally, the review included studies that reported viral shedding duration from the time of hospital admission or initial PCR positivity. Furthermore, no meta-analysis of the duration of viral shedding was done.

This review provides detailed understanding about the evidence available so far on viral dynamics of SARS-CoV-2 and has implications for pandemic control strategies and infection control practices. Although SARS-CoV-2 RNA shedding can be prolonged in respiratory and stool

samples, viable virus is short-lived, with culture success associated with viral load levels. Most studies detected the SARS-CoV-2 viral load peak within the first week of illness. These findings highlight that isolation practices should be commenced with the start of first symptoms, including mild and atypical symptoms that precede more typical COVID-19 symptoms. However, given potential delays in the isolation of patients, effective containment of SARS-CoV-2 might be challenging even with an early detection and isolation strategy.<sup>109</sup>

#### Contributors

MC was involved in conceptualisation, methodology, investigation, data curation, and writing of the original draft. MT was involved in investigation, data curation, and writing of the original draft. OL was involved in investigation, data curation, and review and editing of the manuscript. AEM was involved in formal analysis and writing of the original draft. JS was involved in investigation, data curation, and review and editing of the manuscript. AH was involved in conceptualisation, methodology, data curation, writing of the original draft, and supervision of the study.

#### Declaration of interests

We declare no competing interests.

#### Data sharing

Data used in this study are available upon request from the corresponding author.

#### Acknowledgments

We thank Vicki Cormie at the University of St Andrews (Fife, UK) for assistance with the search and obtaining papers not readily accessible.

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