




BMJ Open Ratio of asymptomatic COVID-19 cases among ascertained SARS-CoV-2 infections in different regions and population groups in 2020: a systematic review and meta-analysis including 130 123 infections from 241 studies

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ABSTRACT

Introduction Asymptomatic infection of SARS-CoV-2 may lead to silent community transmission and compromise the COVID-19 pandemic control measures. We aimed to estimate the rate of asymptomatic COVID-19 from published studies and compare this rate among different regions and patient groups.

Methods In this systematic review and meta-analysis, electronic databases including Medline, Embase, PubMed and three Chinese electronic databases (Chinese National Knowledge Infrastructure [CNKI], WanFang Data and China Science, and Technology Journal Database [VIP]) were searched for literature published from 1 November 2019 to 31 December 2020. Original investigations with sample size (or number of subjects) not less than five were included for further analyses. Subgroup analyses were conducted according to different study types, study periods, geographical regions and patient demographics. The STATA (V.14.0) command ‘metaprop’ was implemented to conduct a meta-analysis of the pooled rate estimates of asymptomatic infections with exact binomial and score test-based 95% confidence interval (CI).

Results A total of 130 123 ascertained COVID-19 infections from 241 studies were included in this meta-analysis, including 31 411 asymptomatic infections. The overall rate of asymptomatic infections was 23.6% (18.5%–29.1%) and 21.7% (16.8%–27.0%) before and after excluding presymptomatic cases, respectively. Subgroup analysis showed that significantly higher in pregnant women (48.8%, 28.9%–68.9%), children (32.1%, 24.2%–40.5%), and studies reporting screening programmes (36.0%, 24.6%–48.1%) conducted on or after 1 March 2020 (42.5%, 33.4%–51.9%). In terms of geographical region, the rate was the highest in Africa (64.3%, 56.7%–71.6%), followed by America (40.0%, 27.4%–53.3%), Europe (28.1%, 19.0%–38.1%) and Asia (18.1%, 13.2%–23.5%).

Conclusion We approximated that one-fifth of COVID-19 infections are asymptomatic throughout the course of infection. Public health policies targeting these high-risk groups may be recommended to achieve early

Strengths and limitations of this study

- The comprehensive systematic literature search included a greater number of studies which reported varied rates of asymptomatic infections.
- Several subgroup analyses were conducted by considering different aspects of the study design.
- Heterogeneity in rates of asymptomatic infections shall be noted.
- The cross-sectional design of most reviewed studies may misclassify ‘pre-symptomatic’ as ‘symptomatic’ infections.

identification and more stringent containment of the pandemic.

INTRODUCTION

In 2019, a cluster of severe pneumonia cases of unknown type were reported in Wuhan, China.¹ Later coined as COVID-19, it rapidly resulted in large-scale outbreaks across many regions. On 30 January 2020, the WHO declared COVID-19 as a public health emergency of international concern² and further defined it as a pandemic on 11 March 2020. As of 29 October 2021, a cumulative total of 245 373 039 COVID-19 cases have been confirmed globally, with 4 979 421 associated deaths.

The pathogen of COVID-19, that is, SARS-CoV-2, is highly contagious and could be transmitted from human to human.^{1 3} The viral load of an asymptomatic patient has been found to be similar to that of a symptomatic patient in a study of nine patients,⁴ and this observation was later confirmed in a study involving large samples.⁵ The early

peaking of SARS-CoV-2 viral load during the presymptomatic phase may cause silent community outbreaks.⁶ Further investigations also found that asymptomatic infections may carry SARS-CoV-2 for more than 1 month, indicating the long-lasting risk of secondary infection.^{7–11}

Determining the rate of asymptomatic is important as it may deepen the understanding of the real reproductive number (R_0), as well as the true incidence and mortality rate of COVID-19. The rates may also function as an essential epidemiological parameter to inform disease-combating policies, including the density and range of screening, patient isolation, as well as early intervention.¹² Nevertheless, the significance of the invisibly infected person as the source of infection depends on the distribution in the population and on the amount and duration of the virus excreted.¹³ The reported proportions of asymptomatic individuals in existing literature varied a lot,¹² depending on research setting (e.g., geographical regions, screening vs non-screening studies), demographic characteristics (e.g., age groups, pregnant women, children) and other latent factors. A systematic review and meta-analysis was performed to estimate the rate of asymptomatic infections based on published studies and compare the rate among different patient groups as well as study settings.

METHODS

Search strategy

This systematic review and meta-analysis was conducted strictly following the ‘Preferred Reporting Items for Systematic Reviews and Meta-Analyses’ guideline.¹⁴ XC and ZH searched Medline, Embase, PubMed and the three most commonly used Chinese electronic databases (Chinese National Knowledge Infrastructure [CNKI], WanFang Data, and China Science and Technology Journal Database [VIP]) from 1 November 2019 to 31 December 2020. CNKI, WanFang Data and VIP are affiliates of the Chinese Ministry of Science and Technology, which provides access to peer-reviewed, continuously updated research journal articles in Chinese. The search string related to ‘COVID-19’ and ‘asymptomatic’ was systematically developed in PubMed using Medical Subject Headings (MeSH) terms and was applied to all databases after discussion with an experienced librarian (Maggie Choi). The ‘Text Word’ search field was used to ensure the best possible search evidence (online supplemental table 1). No filters or limitations were applied to retrieve the best possible results and to avoid excluding preindexed materials. Meanwhile, highly relevant references were also searched (by XC and ZH) by reviewing the reference list of the included articles. Two reviewers (XC and ZH) independently determined eligible studies. Consensus was reached by referral to a third reviewer (JW) when there was disagreement. All manuscripts were imported into the EndNote software (V.X8; Thomson Reuters, Carlsbad, California) to store and manage the retrieved citations. Duplicate studies were removed.

Asymptomatic case and asymptomatic ratio

Several similar definitions of ‘asymptomatic case’ were noted in previous studies.¹⁵ We followed the official definition from the State Council of China and the WHO,^{16,17} defining ‘asymptomatic case’ as individuals who (1) have no clinical manifestations of COVID-19, such as fever, cough, sore throat and other self-perceived or clinically identifiable symptoms and signs; (2) have positive result for SARS-CoV-2 pathogen test; and (3) do not develop symptoms until the end of hospital admission or follow-up observations. Asymptomatic COVID-19 infection rate, or asymptomatic ratio, is calculated as the proportion of asymptomatic cases among all COVID-19 infections.

Literature screening and selection criteria

All studies were screened first by title and abstract, followed by full text if the study met the inclusion criteria, which consist of the following: (1) the subjects of the study were diagnosed with SARS-CoV-2 infection; (2) the study was designed as an observational study; and (3) the number of asymptomatic and symptomatic COVID-19 infections was explicitly and exactly reported.

Literature screening was conducted without language or region restrictions. The exclusion criteria were as follows: (1) the study included patients without virological evidence of SARS-CoV-2 infection; (2) the study did not investigate the distribution of asymptomatic COVID-19 infections among all subjects; (3) the study was not classified as original research, such as reviews, comments and case reports; and (4) the study had an overall sample size of less than five.

If the same group of subjects was reported by different articles, only articles with the most updated and detailed information were included for further analysis.

Data extraction and subgrouping schemes

For eligible articles, two types of information were independently extracted by two reviewers (XC and ZH): (1) basic information of individual studies, including name of the first author, investigation period, geographical region, sample size and characteristics of the subjects; and (2) the number of COVID-19 infections, including the total number of test-positive subjects, the number of presymptomatic infections and that of asymptomatic infections.

To ensure accuracy of data, cross-checking was conducted after extraction of the preliminary information. Disagreements were resolved through consensus or by referral to a third reviewer (JW).

To explore the source of heterogeneity among the included studies, several subgroup analyses were performed according to study design and characteristics of subjects in each study. First, we identified three groups based on subjects’ demographic features: the ‘children’ group, consisting of subjects less than 18 years old; the ‘pregnant women’ group, involving expectant mothers as subjects; and the ‘general population’ group. Second, three age groups were identified according to subjects’

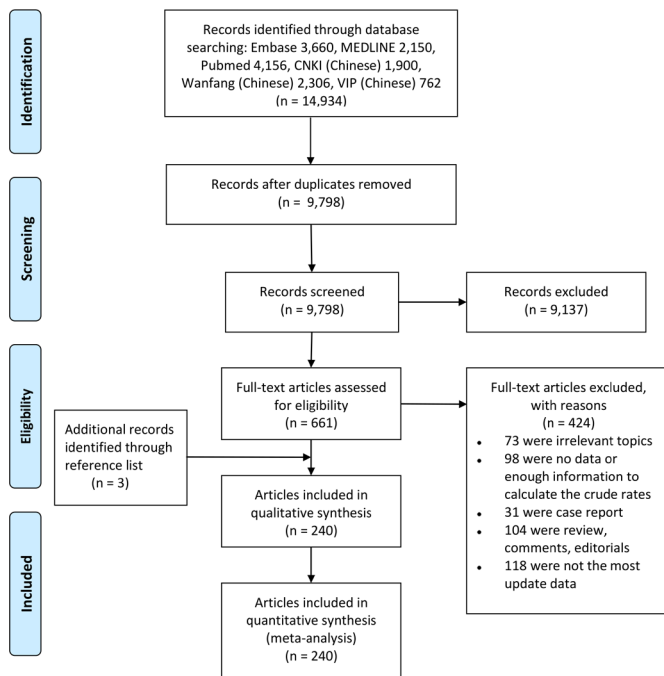


Figure 1 Study selection process for meta-analysis. CNKI, Chinese National Knowledge Infrastructure; PRISMA, Preferred Reporting Items for Systematic Reviews and Meta-Analyses.

mean or median age: ≤ 18 years, 19–45 years and >45 years. The selection of the mean or median age followed the statistics reported in each study. Third, studies were categorised into ‘screening’ or ‘non-screening’ by referring to the positive rate of SARS-CoV-2 pathogen test among the included subjects. In the screening studies, the positive rate was less than 100%, while for non-screening studies all subjects were tested positive. Fourth, we separated studies based on subjects from different geographical regions, including Africa, America, Asia and Europe. Last, subgroup analysis was performed by study period during which the subjects were tested for COVID-19, including ‘before 01 March 2020’ or ‘01 March 2020 and afterwards’.

Quality assessment

The quality of each included study was independently assessed by two researchers (XC and ZH) using the Appraisal tool for Cross-Sectional Studies (AXIS tool),¹⁸ which has 5 components including 20 questions. Seven questions measure the quality of reporting, another seven questions measure the quality of study design, and six questions measure the possible introduction of biases in each study.

Statistical analysis

Accounting for all included studies, pooled estimates of asymptomatic rates were generated with exact binomial-distributed likelihood framework and score test-based 95% CI.¹ The STATA (V.14.0) command ‘metaprop’ was adopted to conduct the meta-analysis. Heterogeneity

across the studies was examined using I^2 statistics, measuring the proportion of total variation contributed by between-study variation. I^2 values $<25\%$, 25% – 75% and $>75\%$ correspond to the thresholds for three ordinal levels of heterogeneity, namely low, moderate and high, respectively.¹⁹

Univariate and multivariate meta-regression analyses were performed to identify any potential effects of modifiers or confounders on the estimated rate. We examined the effects of covariates including study population, age-specific proportion, screening/non-screening study, geographical region and time trend in the multivariate analysis. Sensitivity analysis was conducted by omitting one study at a time, generating pooled estimates and comparing with the original estimates. Potential publication bias was examined by Egger’s test and visualised using a funnel plot. If the tests indicated potential publication bias, the trim and fill’s method, which is based on a modified funnel plot, was adopted to adjust for small-study effects.²⁰

All analyses were performed using STATA statistical software (V.14.0). The figures were generated using R software (V.3.6.3) with the ‘forestplot’ package.

RESULTS

Characteristics of studies and subjects

In total, 9 798 unique citations were identified by the literature search in different databases (figure 1). We screened and assessed the eligibility of 661 full-text articles after 9 137 citations were excluded during title or abstract screening using predetermined criteria. Finally, there were 241 studies in 240 articles included in the meta-analysis, among which 3 studies were additional records identified through reference lists.

The basic characteristics of the included studies are shown in online supplemental table 2. The characteristics include cities, countries and study periods when patients with ascertained COVID-19 infections were recruited; the number of positive tests and asymptomatic infections; the number of presymptomatic infection in patients if available; and the demographics of study subjects. Approximately 72.9% (175 out of 241) of all included studies were conducted in Asia, 11.7% in Europe and 10.42% in America. In terms of study population, 80.5% were performed in general adults, 12.4% in children and 7.1% in pregnant women.

Quality scores of AXIS for the included studies ranged between 11 and 19 points, with 153 studies meeting the criteria for high quality (≥ 16 points) (see online supplemental table 3 for details). Overall, 97.9% (235 out of 240) of all studies met the criteria for both reporting the aims/objectives and quality design, and the risk factors and outcome variables were appropriate for the studies.

Asymptomatic ratio estimates

A total of 130 123 ascertained COVID-19 infections from 241 studies were finally included in the meta-analysis

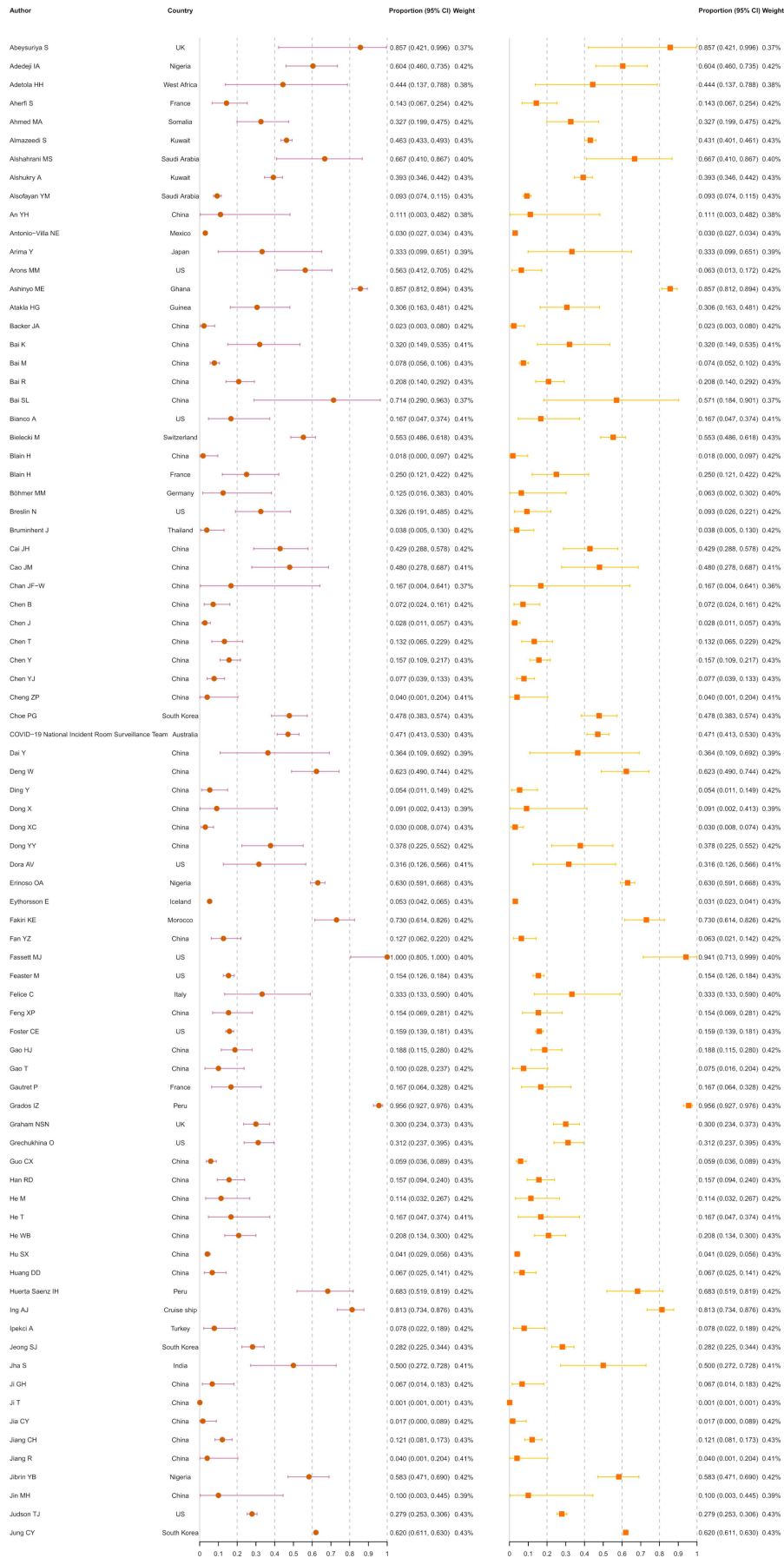


Figure 2 (Continued)

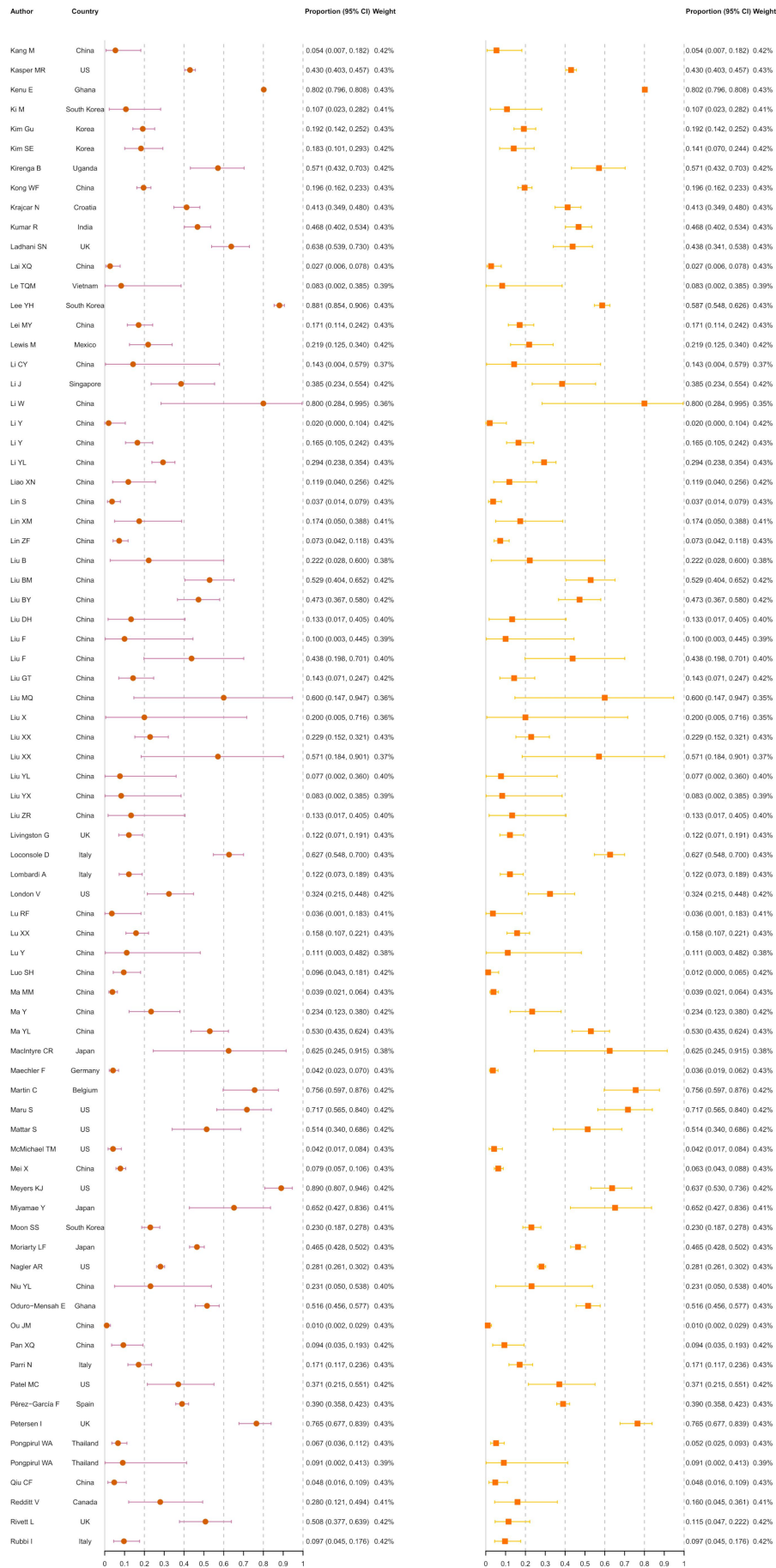


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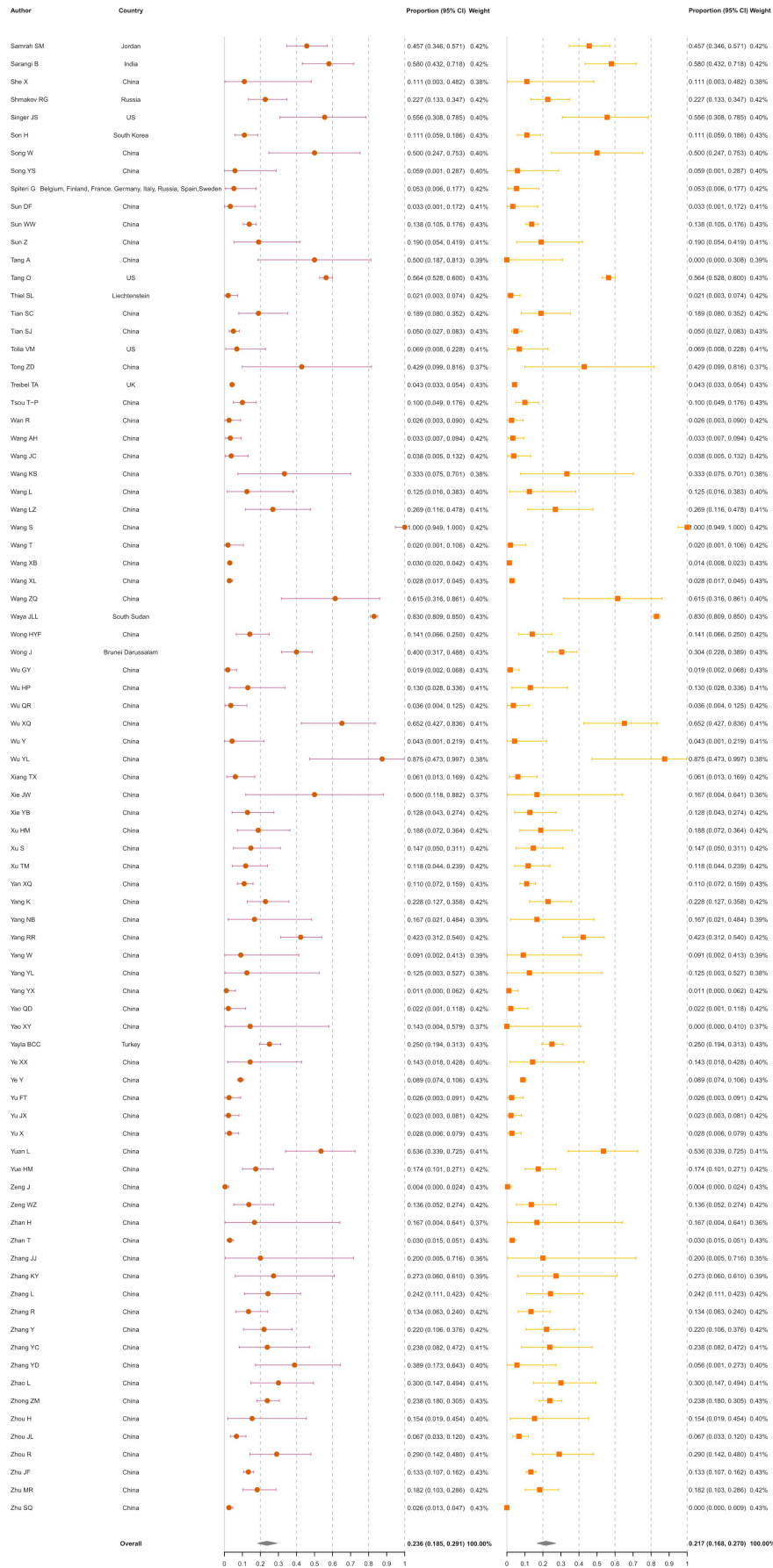


Figure 2 Meta-analysis of the asymptomatic ratio among ascertained COVID-19 infections (including and excluding presymptomatic infections).

(figure 2), including 31 411 asymptomatic infections at the time of diagnosis. The rate of asymptomatic infections varied from 0.1% in general adults in Wuhan, China,²¹ to 95.6% in pregnant women in Peru.²² The overall rate of asymptomatic infections was estimated at 23.6% (18.5%–29.1%). The I^2 equalled 99.7%, indicating high heterogeneity among the studies. Of the 31 411 asymptomatic infections, 448 (1.4%) developed symptoms after admission. After excluding these presymptomatic observations, the pooled rate was 21.7% (16.8%–27.0%) with an I^2 of 99.7%.

In a subgroup analysis, we found that pregnant women (48.8%, 28.9%–68.9%; $I^2=96.9%$) had a significantly higher rate of asymptomatic infections than children (32.1%, 24.2%–40.5%; $I^2=94.4%$), whereas the general non-pregnant adults had the lowest asymptomatic rate (20.4%, 15.0%–26.3%; $I^2=99.8%$). Regarding age-specific proportion, patient groups with a mean/median age of ≤ 18 years (31.1%, 23.5%–39.2%; $I^2=94.4%$) had a significantly higher prevalence of asymptomatic cases than those with a mean/median age of 19–45 years (26.7%, 19.1%–34.9%; $I^2=99.7%$) or >45 years (15.3%, 10.7%–20.6%; $I^2=97.1%$).

The rate of asymptomatic infections was 36.0% (24.6%–48.1%; $I^2=99.8%$) among studies with screening setting, which was almost twice that among studies with non-screening setting (19.4%, 14.3%–24.9%; $I^2=99.6%$). Significantly different asymptomatic rates were observed in different geographical regions ($p<0.001$)—the rate was highest in Africa (64.3%, 56.7%–71.6%; $I^2=96.8%$), followed by America (40.0%, 27.4%–53.3%; $I^2=99.6%$) and Europe (28.1%, 19.0%–38.1%; $I^2=98.5%$). Asia was estimated to have the lowest rate of asymptomatic infections (18.1%, 13.2%–23.5%; $I^2=99.5%$). In the subgroup analysis by time period, we noted that studies conducted before 1 March 2020 reported a pooled asymptomatic infection rate of 13.3% (9.8%–17.1%; $I^2=94.2%$), while the rate increased sharply to 42.5% (33.4%–51.9%; $I^2=99.6%$) among studies conducted on or after 1 March 2020.

The results of all subgroup analyses were shown in figure 3.

We performed both univariate and multivariate meta-regression analyses to investigate study-level factors that may have contributed to the heterogeneity among studies and might have influenced our estimations of the rate of asymptomatic infections. The results of univariate regression analysis (table 1) demonstrated several significant sources of heterogeneity in the estimated rate of asymptomatic infections—studies focused on subjects aged >45 years old ($p=0.015$) and conducted out of Africa ($p\leq 0.001$) were significantly associated with a lower ratio of asymptomatic infections, whereas those using a screening study design ($p<0.001$) and conducted on or after 1 March 2020 ($p<0.001$) tended to report a higher ratio of asymptomatic infections. A screening study design ($p=0.031$), geographical region ($p<0.001$) and time trend ($p<0.001$) remained as significant sources of heterogeneity in the multivariate regression analysis.

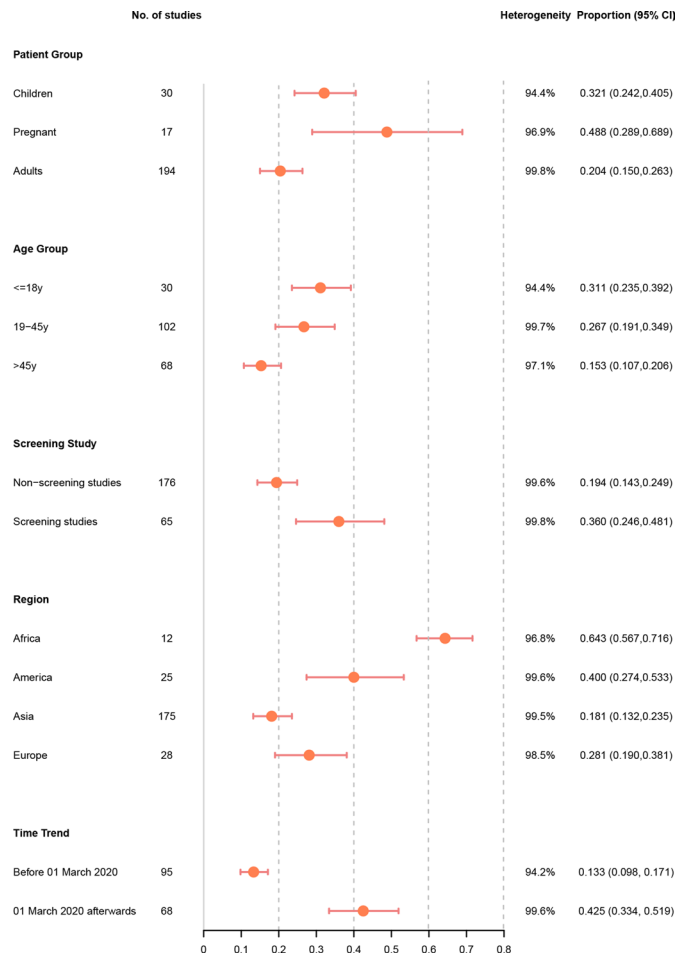


Figure 3 Asymptomatic ratio among ascertained COVID-19 infections by patient group, age group, screening study, region and time trend.

We found no evidence of publication bias among studies by Egger’s test with $p=0.414$. Sensitivity analysis showed that it was unlikely that any individual study significantly influenced the pooled estimates, demonstrating the robustness and reliability of our estimates (online supplemental table 4).

DISCUSSION

Our meta-analysis of 130 123 infections in 241 studies performed in 36 countries provides an up-to-date as well as a comprehensive overview of the asymptomatic infection rate of COVID-19. The estimated rate in our study were 23.6% and 21.7% before and after excluding presymptomatic cases, respectively. There were three key findings in the subgroup analysis.²³ First, African studies reported the highest asymptomatic infection rate, while Asian studies reported the lowest. Second, the proportion of asymptomatic carriers was nearly doubled in screening studies than in non-screening studies. Third, asymptomatic infection rate was more than two times higher in studies conducted on or after 1 March 2020 than those conducted beforehand.

Table 1 Results of univariate and multivariate meta-regression analyses

Variable	Univariate regression analysis			Multivariate regression analysis			
	Studies, n (%)	Crude meta-RR	95% CI	P value	Adjusted meta-RR	95% CI	P value
Patient group							
Children	30 (12.45)	1 (reference)	1 (reference)		1 (reference)	1 (reference)	
Pregnant	17 (7.05)	1.16	0.99 to 1.37	0.074	1.02	0.75 to 1.10	0.901
Adults	194 (80.50)	0.91	0.82 to 1.00	0.057	0.83	0.63 to 1.11	0.209
Age (years)							
≤18	30 (12.45)	1 (reference)	1 (reference)		1 (reference)	1 (reference)	
19–45	102 (42.32)	0.98	0.88 to 1.09	0.682	1.12	0.84 to 1.48	0.446
>45	68 (27.39)	0.87	0.77 to 0.97	0.015	1.09	0.82 to 1.45	0.539
NM	43 (17.84)	0.93	0.83 to 1.05	0.252	1.13	0.85 to 1.49	0.393
Screening							
Non-screening studies	176 (73.03)	1 (reference)	1 (reference)		1 (reference)	1 (reference)	
Screening studies	65 (26.97)	1.16	1.08 to 1.25	<0.001	1.08	1.01 to 1.17	0.031
Region							
Africa	12 (5.00)	1 (reference)	1 (reference)		1 (reference)	1 (reference)	
America	25 (10.42)	0.78	0.67 to 0.90	0.001	0.71	0.62 to 0.82	<0.001
Asia	175 (72.92)	0.64	0.56 to 0.72	<0.001	0.78	0.68 to 0.89	<0.001
Europe	28 (11.67)	0.71	0.62 to 0.82	<0.001	0.69	0.60 to 0.78	<0.001
Time trend							
Before March 2020	95 (39.42)	1 (reference)	1 (reference)		1 (reference)	1 (reference)	
March 2020 afterwards	68 (28.22)	1.33	1.24 to 1.43	<0.001	1.29	1.17 to 1.43	<0.001
NM	78 (32.37)	1.07	1.00 to 1.14	0.092	1.05	0.98 to 1.12	0.158

NM, not mentioned; RR, risk ratio.

Our estimation is higher than some existing meta-analyses, which reported a ratio between 15.6% and 20%.^{24 25} One possible explanation is that the included studies in previous meta-analyses were dominated by that conducted during the early course of the pandemic and in China or the USA, where the asymptomatic ratio was lower than other regions. However, our study included more than 200 articles, resulting in a more precise estimation with significantly narrower CI. With a much larger data set, we conducted more subgroup analyses, which resulted in more implications.

In the subgroup analysis, we noted a large variety of asymptomatic rates among different populations. For instance, children and adults aged ≤45 years old tended to have a significantly higher rate. The inverse correlation between age and symptom severity has also been reported and discussed by a few studies, and the higher prevalence of chronic diseases among elderies was considered as risk factors for symptomatic infection.^{23 26 27} Infection control measures may be targeted on early detection and isolation of asymptomatic youth, as young asymptomatic carriers are more likely to bring in community transmission due

to their more socially active lifestyle habits, with more frequent travelling, compared with people in other age groups.²⁶

We also noted that the rate of asymptomatic infections in pregnant women (48.8%) was more than two times than in general adults (20.4%). A previous study suggested cytotoxic cells may be poised to control virus load and hence affect disease severity during pregnancy.²⁸ Case reports from a New York hospital reported a similarly high rate (14 women, 32.6%) at presentation, yet 71.4% (10 women) of the asymptomatic mothers developed symptoms during their hospitalisation and postpartum course.²⁹ Nevertheless, we noted that 15 out of the 17 included studies on pregnant women were cross-sectional studies, where the high proportion of asymptomatic infections may include both presymptomatic and asymptomatic cases. More follow-up studies among pregnant women are needed before drawing further conclusions. Undetected asymptomatic pregnant women may lead to more severe consequences. Without early detection and proper preventive measures, the delivery of asymptomatic patients brings extra risks to nosocomial infection and

may also result in droplet transmission among women, kids and other family members.²⁹

To our knowledge, this is the first meta-analysis to compare the rate of asymptomatic infections in different continents. A previous meta-analysis only concluded that populations in Asia may have a lower asymptomatic ratio, while not exploring differences among other continents.³⁰ Our results also showed that the estimated ratio in Asia was the lowest among all continents. Possible explanations were summarised in a published meta-analysis, from the perspectives of infection control policies and host characteristics.³⁰ Meanwhile, the rate in Africa (64.3%) was the highest among all continents, although the limited number of subjects (20 271 COVID-19 infections) in African studies reminded us to think twice before generalising the results. The phenomenon may be partly explained by a few factors in African patients, including the generally younger age, the lower proportion of patients and elderlies with chronic diseases living in nursing homes, and the higher serum vitamin D levels due to rich sun exposure.³¹ Despite the higher asymptomatic ratio, it is noted that Africa was the continent with the lowest reverse transcription polymerase chain reaction (RT-PCR) testing rate as well as the lowest vaccination rate. The high rate of asymptomatic infections may further hinder the timely detection and control of COVID-19 outbreaks.

We noted that the rate of asymptomatic infections was nearly doubled in screening studies (36.0%) than in non-screening studies (19.4%). One possible explanation is that people may get tested for varied reasons associated with social, cultural and political factors in each region. For instance, front-line health workers and people with a history of exposure were in general more likely to be reported in screening programmes, hence more likely to be detected during the early stage of their infection. Comparatively, the samples in non-screening studies were likely to be dominated by subjects having severe symptoms requiring hospitalisation.³² If we consider the rate as a constant, the higher asymptomatic infection rate estimates were likely due to higher probabilities of ascertaining asymptomatic COVID-19 infections in the community with screening implemented. This implies the importance of mass screening in detecting infections, which is important in community infection control. Increasing the accessibility and affordability of community testing could be an important surveillance strategy for early containment of diagnosed cases.³³

We also found that the pooled asymptomatic infection rate increased from 13.3% (95% CI 9.8% to 17.1%) in studies conducted before 1 March 2020 to 42.5% (33.4% to 51.9%) in studies conducted on 1 March 2020 or afterwards, although the 95% CI is larger in the latter time period. This timeline was highly consistent with a previous study using publicly released data from the Centre for Health Protection in Hong Kong.³⁴ The increased rate may be due to overlooking the asymptomatic cases, especially presymptomatic cases, during

the early stage of the pandemic, when medical resources were targeted to patients with severe symptoms. Later, with increased public awareness and test accessibility, more COVID-19-infected individuals without symptoms were detected, while more and more studies reported the proportion of asymptomatic patients. Importantly, the SARS-CoV-2 variant with D was replaced by G at the 614th codon in the spike protein, which has dominated the pandemic since late February 2020, and were suggested to raise the asymptomatic cases.^{6 35 36} Future studies may explore whether the proportion of COVID-19 infections with mild or no symptoms is increasing, especially when considering different SARS-CoV-2 variants.

This study has limitations. First, our pooled asymptomatic infection rates were found to have a high level of heterogeneity ($I^2=99.7%$). This could be attributed to the difficulties in generating the exact number of infections and asymptomatic cases during an outbreak. Although we conducted subgroup analysis and meta-regression to figure out the source of heterogeneity, some unobserved factors which have not been included in the original studies may lead to the high heterogeneity, such as changing pandemic control measures in some countries, the diverse definition of asymptomatic infection, varying practices in surveillance and ascertainment of asymptomatic infection, as well as meteorological disparities across time and regions. Nevertheless, previous studies indicated that any amount of heterogeneity is acceptable if both accurate data and predefined eligibility criteria were provided.^{37 38} Second, we followed the WHO definition of COVID-19 infection and only included infections detected by RT-PCR, while the availability of RT-PCR is restricted in many countries.³⁹ An increasing number of studies have reported the results of antibody-based rapid diagnostic tests, yet they were not included in our meta-analysis. Third, our meta-analysis focused on real-world evidence and observational studies, where subjects with more severe and easily recognised symptoms were more likely to be selected. The collider bias caused by non-representative sampling strategies (such as sampling conditional on testing and prognosis conditional on hospitalisation) in existing observational studies on COVID-19 has been deeply discussed in Griffith *et al.*³² Fourth, the cross-sectional studies included in this review might have misclassified presymptomatic cases as 'asymptomatic' and hence resulted in a certain degree of overestimation of the rate of asymptomatic infections. With additional information on the exposure and reporting dates of each case, we remark our estimation can be extended to a right censoring version to further address some potential bias in existing frameworks.¹²

CONCLUSIONS

We estimated the rate of asymptomatic infections at 23.6%, which decreased to 21.7% after excluding presymptomatic cases. Subgroup analysis indicated that pregnant women, children, African residents, screening programmes and studies conducted after 1 March 2020 had higher rates of asymptomatic infections. Our findings



provide further insights on the distribution of asymptomatic COVID-19 infections in different groups of individuals, bearing significance on public health policies that aim to achieve early identification and more stringent containment of the pandemic.

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Contributors XC, ZH, and JW were responsible for the overall study coordination. XC and ZH designed the study and searched the literature. XC, ZH and JW extracted the data and evaluated the quality. XC conducted the statistical analysis. XC, ZH and JW wrote the full manuscript. SZ, MC-SW, KCC, DH and JL provided critical revision to the manuscript for important intellectual content. All authors discussed the results and approved the final version of the manuscript for publication.

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