





Systematic Review of the Prevalence of Long COVID

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Background. Long COVID occurs in those infected with severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) whose symptoms persist or develop beyond the acute phase. We conducted a systematic review to determine the prevalence of persistent symptoms, functional disability, or pathological changes in adults or children at least 12 weeks postinfection.

Methods. We searched key registers and databases from January 1, 2020 to November 2, 2021, limited to publications in English and studies with at least 100 participants. Studies in which all participants were critically ill were excluded. Long COVID was extracted as prevalence of at least 1 symptom or pathology, or prevalence of the most common symptom or pathology, at 12 weeks or later. Heterogeneity was quantified in absolute terms and as a proportion of total variation and explored across predefined subgroups (PROSPERO ID CRD42020218351).

Results. One hundred twenty studies in 130 publications were included. Length of follow-up varied between 12 weeks and 12 months. Few studies had low risk of bias. All complete and subgroup analyses except 1 had $I^2 \ge 90\%$, with prevalence of persistent symptoms range of 0%–93% (pooled estimate [PE], 42.1%; 95% prediction interval [PI], 6.8% to 87.9%). Studies using routine healthcare records tended to report lower prevalence (PE, 13.6%; PI, 1.2% to 68%) of persistent symptoms/pathology than self-report (PE, 43.9%; PI, 8.2% to 87.2%). However, studies systematically investigating pathology in all participants at follow up tended to report the highest estimates of all 3 (PE, 51.7%; PI, 12.3% to 89.1%). Studies of hospitalized cases had generally higher estimates than community-based studies.

Conclusions. The way in which Long COVID is defined and measured affects prevalence estimation. Given the widespread nature of SARS-CoV-2 infection globally, the burden of chronic illness is likely to be substantial even using the most conservative estimates.

Keywords. Long COVID; prevalence; SARS-CoV-2; systematic review.

Long COVID is the state of not fully recovering for many weeks, months, or years after contracting severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection. The World Health Organization (WHO) defines post-COVID-19 condition (Long COVID) as the condition occurring in individuals with a history of probable or confirmed SARS-CoV-2 infection

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3 months after the onset with symptoms that last at least 2 months, cannot be explained by an alternative diagnosis, and generally impacts everyday functioning [1]. These symptoms may be the same as the acute illness or new symptoms developing weeks or months after the acute phase. Clinical guidelines [2, 3] in the United Kingdom and the United States consider Long COVID as symptoms ongoing for 4 weeks or more.

Long COVID can occur across the spectrum of severity of initial infection [4]. A wide range of symptoms have been reported with exhaustion, breathlessness, muscle aches, cognitive dysfunction, headache, palpitations, dizziness, and chest tightness or heaviness among the most common [5, 6]. Patients are still struggling to access adequate recognition, support, medical assessment, and treatment [7, 8].

Studies assessing the prevalence of Long COVID have produced wide-ranging results due to varying settings, case definitions, population denominators, and methods of ascertainment. This is exemplified in the UK Office for National Statistics (ONS) estimates of Long COVID during 2020–2021 where 3 different approaches were used resulting in 3 different

estimates: approach 1 estimated 5.0% prevalence based on respondents reporting any of 12 common symptoms at 12–16 weeks after infection; approach 2 estimated 3.0% prevalence based on respondents reporting any of 12 common continuous symptoms at least 12 weeks after infection; and approach 3 estimated 11.7% prevalence based on respondents describing themselves as having Long COVID [9].

For the purposes of this review, we define Long COVID as persistent (constant, fluctuating or relapsing) symptoms and/ or functional disability and/or the development of new pathology after SARS-CoV-2 infection for equal to or more than 12 weeks from onset of symptoms or from time of diagnosis, in people in whom the infection is self-described, clinically diagnosed, and/or diagnosed through a laboratory test.

We aimed to systematically collate, appraise, and synthesize studies that describe the prevalence of Long COVID and to characterize its typology including patient demographics, symptoms/function disability, and pathology.

METHODS

Search Strategy and Selection Criteria

Included study designs were cohort, cross-sectional, and case control studies with an estimate of the denominator where participants were followed-up/assessed at a minimum of 12 weeks postinfection. Studies were restricted to those published in English between January 1, 2020 and November 2, 2021, including peer-reviewed articles, online reports, letters, and preprints. Only studies with a sample size of 100 or more participants (at the time of follow-up assessment if longitudinal study) were included (50 or more per subgroup).

Studies of adults and children with a confirmed or probable SARS-CoV-2 infection in any age group (as defined by each study) were included. The control group in studies that included one comprised individuals with a confirmed or probable case of SARS-CoV-2 infection (as defined by the study) who had recovered (duration as defined by study as long as under 12 weeks from symptom onset or confirmation of infection) and had no new pathology attributed to SARS-CoV-2 infection. Studies that compared population-based prevalence as the control arm were excluded from the control analysis.

Community-based, hospital-based, and mixed studies were all included, apart from studies that only reported outcomes for critically ill patients admitted to intensive care, because this review did not aim to estimate delayed recovery after intensive care unit (ICU) admission (post-ICU syndrome). Patients who were not hospitalized within 2 weeks of symptom onset but were subsequently hospitalized were counted as nonhospitalized for the purpose of this review.

A systematic search was conducted using MEDLINE (Ovid), Embase (Ovid), the Cochrane COVID-19 Study register (covid-19.cochrane.org; includes Cochrane Central Register of

Controlled Trials [CENTRAL]), WHO International Clinical Trials Registry Platform [ICTRP], medRxiv, Cochrane CENTRAL, MEDLINE [PubMed], ClinicalTrials.gov, and the WHO Global research on coronavirus disease [COVID-19]) database [10]. The initial search was run on November 13, 2020 and updated on November 2, 2021, both by VL. An example of the search strategy applied to Medline is provided in the Supplementary Material; it was adapted for other databases as needed.

The screening management software Covidence was used to screen for eligibility. All articles were screened independently by 2 reviewers at each stage (title, abstract, and full text) with any discrepancies resolved by NAA. This review is reported in line with PRISMA guidelines [11]. The protocol was published on the international prospective register of international reviews, PROSPERO (CRD42020218351): https://www.crd.york.ac.uk/prospero/display_record.php? RecordID=218351.

Data Analysis

Data for each study were extracted independently by 2 of 4 reviewers (MW, DCG, CC, NZ). Any discrepancies were resolved by consensus between the 2 reviewers for each study or by a third reviewer (NAA). In instances in which multiple publications were identified as originating from the same study, all data were extracted but each data point was only used once in the analysis. In addition to excluding duplicate reports, or duplicate results from the same study, several general decisions were made to cope with multiple publications from the same study, either focusing on different lengths of follow-up, different timepoints, or different subgroups. These were guided by the following principles: (1) avoiding double counting individuals; (2) using the most appropriate outcome, for example, general Long COVID definition, in the broadest group such as the widest population, largest sample, most recent update; and (3) unless stratifying by length of follow-up, taking the earliest and/ or most complete follow-up as the main result.

The primary outcome is Long COVID, defined as nonrecovery from COVID-19, according to symptoms, functional ability, or pathology. The SARS-CoV-2 infection can be confirmed, probable, or suspected with prolonged symptoms (including but not limited to those explicitly defined as "new onset"), functional disability, or pathology for equal to or more than 12 weeks from onset of symptoms or positive test date (as defined by the study). Secondary outcomes included the demographics of people with Long COVID in relation to each study's denominator, prevalence of specific persistent or relapsing symptoms, prevalence of functional disability, and the characterization of post-COVID-19 pathology.

A Long COVID-specific risk of bias tool was developed, based on the Newcastle-Ottawa scale, but it was tailored to the relevant sources of bias. The domains used are reported in Supplementary Table 3. Risk of bias was particularly assessed

in relation to the denominator, how the symptoms were assessed (active or passive elicitation of the symptoms), and hospital stay. Subgroup analysis by risk of bias was performed. In studies where follow up was measured posthospital admission or discharge, symptom onset was estimated to have been 7 or 14 days before discharge, respectively, and estimated as 21 days if follow up was measured from a postinfection negative test.

The prevalence was extracted as cumulative incidence. In extracting the prevalence of persistent symptoms, we used either prevalence of at least 1 symptom or pathology, or the prevalence of the most common symptom/pathology, depending on the data reported by the study. Data for each symptom was extracted separately in studies that reported on the prevalence of individual symptoms but did not provide an overall estimate of prevalence of Long COVID. We used the symptom with the highest estimate as our best estimate of overall prevalence, although it is likely to be an underestimate of actual prevalence. In studies with controls, the prevalence of the same symptom was used for comparison. In instances in which length of follow-up varied between study participants, we report a measure of average (eg, mean or median) length of follow-up, or the midpoint of the reported range.

All analysis was conducted in Stata version 17 [12]. The distribution, prevalence estimates, numerators, denominators, and assessment time points in different populations was qualitatively summarized. We used random-effects meta-analysis on the logit of the proportions to ensure estimates and confidence limits did not go below 0% or over 100%, transforming back to the original scale for presentation.

The heterogeneity was quantified both in absolute terms (range of individual study estimates) and as a proportion of total variation (I²), and this was explored across predefined subgroups described below. In a variation to our protocol, we present pooled estimates (PEs) alongside 95% prediction intervals (PIs) to evaluate and incorporate uncertainty in the analysis, as recently recommended for prevalence studies, where true between-study heterogeneity is expected [13, 14]. Heterogeneity was explored by stratifying on predefined subgroups: outcome type (pathology, symptom, functional status), geographical region (China, Europe, North America, Mixed, and other), source of sample (community, healthcare workers, outpatients, hospital inpatients), length of follow-up, study design, confirmed diagnosis, and other risk of bias domains. We also stratified by severity score based on the WHO Clinical Progression Scale (CPS) (Supplementary Methods). Potential small study effects such as publication bias were investigated using contour-enhanced funnel plots and Egger's test of funnel plot asymmetry.

Patient Consent Statement

In this systematic review, we analyzed publicly available data included in published scientific papers. Patient consent and ethical approval were not required.

RESULTS

Literature Search

In our search, we found 11 518 studies in total. After deduplication and title and abstract screening, 457 full-text studies were assessed for eligibility. Using handsearching, we sourced an additional 9 studies and 130 publications in total were included, 120 of these were discrete studies (Figure 1). Twenty-four studies were conducted in China (including Hong Kong), 66 in Europe, 14 in North America, and 16 in various other countries [9, 15–143]. Reasons for exclusion are listed in Supplementary Table 1.

Table 1 summarizes the included studies' key characteristics and primary outcome for the first follow-up. Study design was reported as described by each study or designated based on study description if not explicitly stated. Most studies were in adults and included patients who were hospitalized in the acute phase (24 studies with <10% of the sample hospitalized in the acute phase). However, hospitalization did not always correspond with disease severity, probably due to local diagnostic, treatment, and containment policies. Most studies used polymerase chain reaction (PCR) testing to identify COVID-19 cases at baseline. However, most did not perform COVID-19 diagnostic tests at follow up and therefore did not consider the impact of reinfection on their results. Of the included studies, 21 were community-based studies, 17 were in outpatient settings, 3 were from social media, and 8 were healthcare worker-based studies.

Prevalence Estimates

The prevalence of Long COVID for studies with more than 12 weeks from infection ranged between 0% and 93% (PE, 42.1%; 95% PI, 6.8%–87.9%) (Figure 2). For all complete and subgroup analyses except one, I^2 was >75%. All subgroup analysis results including PEs and PIs can be found in Supplementary Table 4.

Seventy-three included studies had a follow up of 12 weeks to 5 months (PE, 39.8%; PI, 5.1%–89.1%), 49 had a follow up of 6–11 months (PE, 44.9%; PI, 8%–88.4%), and 12 had a follow up of 12 months or more (PE, 48.5%; PI, 12.7%–86%). We recognize that most were not within-study comparisons, but longer follow-up times showed higher pooled estimates (Supplementary Figure 1).

Hospitalization and severity of acute infection were key factors influencing Long COVID prevalence estimates. The prevalence range in analyses in which less than 10% of the participants were hospitalized was 0% to 67% (n = 24) (PE, 26.4%; PI, 2.6%–82.8%), but in studies in which all participants were hospitalized for acute COVID-19 (n = 65), the prevalence range was 5% to 93% (PE, 47.5%; PI, 8.3%–90.0%) (Supplementary Figure 2). Thirty-one studies had 10% or more of their sample admitted to intensive care unit ICU during their acute COVID-19 illness with a Long COVID

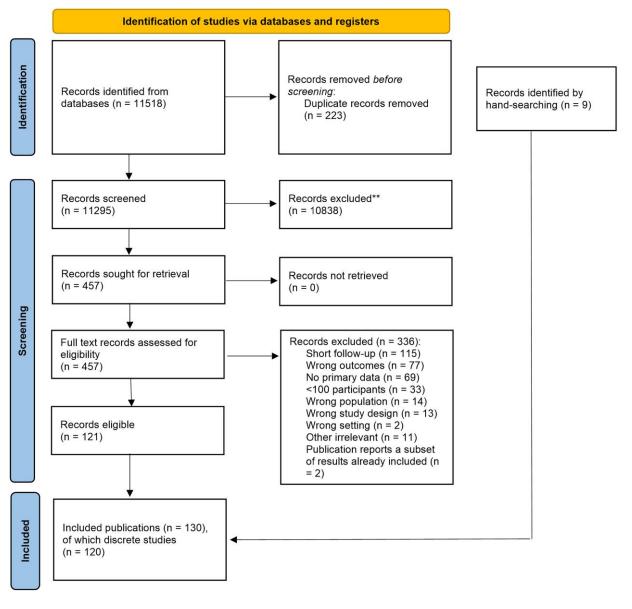


Figure 1. Study selection.

prevalence estimate of 48.8% (PI, 5.7%–93.7%) compared with PE 34.9% (PI, 5.2%–84%, n=48) in studies with <5% of their samples admitted to ICU (Supplementary Figure 3). Studies including more hospitalized participants or more patients in ICU tended to report higher prevalence estimates (Supplementary Table 4). Likewise using the WHO CPS, we found that studies including those with ambulatory mild disease (n=38) generally reported lower prevalence estimates (PE, 23.5%; PI, 1.6%–85.7%) than those with hospitalized severe disease who needed oxygen by noninvasive ventilation or high flow (n=27) (PE, 54.8%; PI, 7.7%–94.7%) (Supplementary Figure 4).

The prevalence of not returning to full health/fitness after at least 12 weeks from infection ranged between 8% and 70% (PE,

34.5%; PI, 4.3%–85.9%; n=10) (Supplementary Figure 5). The prevalence of lower quality of life after at least 12 weeks was 31% (n=2) (Supplementary Figure 6). With regard to individual symptoms, common symptoms reported included fatigue (PE, 21.6%; PI, 2.5%–74.7%; n=72) followed by breathing problems (PE, 14.9%; PI, 1.6%–64.9%; n=78), sleep problems (PE, 13.2%; PI, 1.2%–64.9%; n=42), tingling or itching (PE, 11.3%; PI, 0.7%–69.5%; n=14), and joint/muscle aches and pains (PE, 10.6%; PI, 1.0%–57.5%; n=61) (Figure 3). With regard to pathology, lung pathology was the most common (PE, 38.9%; PI, 3.4%–91.9%, n=26) followed by heart (PE, 6.0%; PI, 0.1%–79.3%; n=12) or neurological pathology (PE, 5.3%; PI, 0.5%–36.5%; n=11) (Figure 3 and Supplementary Figures 7–40).

Table 1. Study characteristics and primary outcome at first follow-up.

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	61.0%	2.9%	9.2%	44.2%	73.6%	27.8%	21.5	32.4%	24.1%	53.8%	52.0%	40.3%
Follow-up Time Days	240–300 (range) after "improvement of acute COVID-19"	126°	150°	243°	₀ 06	131°		124.7 ^e	228 ^b	91–121 ^e	104°	_q 06
Severity	(including 4% ICU)	÷	26.3% ICU	ICU excluded	Mixed	97.5% mild	9:9% ICU	13% ICU	i	27.7% did not require oxygen 11.8% ICU	47% severe	Nonhospitalized
COVID-19 Diagnostic Method	"Tested positive"	"Positive test"	PCR confirmed	PCR confirmed	PCR confirmed or clinico- radiological	PCR confirmed	Laboratory confirmed or clinical diagnosis	"Confirmed diagnosis"	Tested positive or antibody positive	PCR confirmed bronchial swab, serological testing, or suggestive CT	PCR confirmed	PCR confirmed
Age Years) ean/SD Aedian % ((QR) Female	41.8/17.6 65.7	61 (4872) 12.1	1- 5.8	52.7/20.1 52.3	6- 44.0	1- 52.3	64.5/19.2 45.1	57.9/12.8 40.0	8- 63.0	0- 40.3	54.9/10.3 36.0	1.8 70.0 (13.6)
Age (Years) Mean/SD Median (IQR)	41.8/	61 (4	70 (61– 76)	52.7/	60 (46– 73)	43 (31– 54)	64.5/	57.9/	49 (38– 59)	61 (50–71)	54.9/	44.8
Setting	Hospitalized patients and nonhospitalized	Nonhospitalized	Hospitalized patients	Hospitalized patients	Hospitalized patients	Nonhospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients, outpatients and ER attendees	Hospitalized patients	Hospitalized patients	Nonhospitalized patients
Controls N, Type	:	4 526 737 without COVID-19 and not hospitalized	11 868 hospitalized with seasonal influenza	:	:	:	47 780 matched for Hospitalized age, sex patients	:	:	:	:	÷
Denominator ^a	172	60 255	11 800	2839	110	442	47 780	204	740	238	100	129
Study Design (as Described by Study,* If Not Stated)	Prospective cohort	Cohort with controls	Cohort with controls	Retrospective	Prospective cohort	Longitudinal prospective cohort	Observational retrospective matched cohort (with controls)	Cross-sectional	Cross-sectional	Prospective cohort	Prospective	Cohort
Country	Egypt	USA	USA	USA	¥	Germany	ž	Italy	USA	Italy	Spain	Denmark
Author	Abdelrahman et al [15]	Al-Aly et al [16]	Al-Aly et al [16]	Aminian et al [18]	Arnold et al [144]	Augustin et al [20]	Ayoubkhani et al [21]	Baricich et al [22]	Becker et al [23]	Bellan et al [24]	Blanco et al [25]	Bliddal et al [26]
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Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	%9.09	53.0%	66.2%	77.1%	16.5%	10.9%	28.6%	14.8%	%0.08	49.5%	79.3%	53.4%	37.2%
Follow-up Time Days	152–213 (range) after illness	365 ⁵	3 months posthospital discharge	6 months posthospital admission	334–365 (range) after infection	91–150 days posthospital admission	370 ^d	_q 06	107 [†]	6 months posthospital discharge	12 weeks postinfection	7 months postinfection	5.6 months postinfection
Severity	2% asymptomatic,78% symptomatic in community, 21% hospitalized	Mild-to-moderate (home-isolated)	:	Moderate to severe	÷	13% ICU	24% severe	:	89% required at least oxygen support	hospitalized for mild to moderate COVID	Mixed	11.2% asymptomatic	22.3% hospitalized
COVID-19 Diagnostic Method	"Tested positive"	PCR confirmed	PCR confirmed and suspected cases (clinical, imaging and laboratory results)	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	"Hospitalized for COVID-19"	Positive nasal swab Mixed	Laboratory confirmed	"Tested positive"
% Female	51.0	6.09	1.69	53.0	58.0	47.0	51.0	:	43.0	27.7	53.1	62.1	46.2
Age (Years) Mean/SD Median (IQR)	46 (30– 58)	47 (n/a)	88.5/6.7	65/12	25+	63 (50– 76)	65 (59– 70)	:	58.8 (51.6– 66.0)	63.6/12.9	60/13.9	37.2/17.1 62.1	9.2 (10.9– 17.9)
Setting	Hospitalized patients and nonhospitalized	Community	Hospitalized older adult patients	Hospitalized patients with interstitial pneumonia	Community (MoBa: population-based pregnancy cohort study)	Hospitalized patients	Hospitalized cancer and noncancer patients	Community	Hospitalized patients	Hospitalized patients	Hospitalized patients and nonhospitalized	Community	Hospitalized and nonhospitalized children
Controls N, Type	60 seronegative household contacts	:	÷	:	72 953	talized	* * * 	÷	ŧ	÷	E	:	95 randomly selected from non-COVID patients attending the ward
Denominator ^a	312	304	165	118	774	5571	546	357	200	101	111	483	121
Study Design (as Described by Study,* If Not Stated)	Prospective cohort with controls	Prospective	Longitudinal observational	Prospective	Matched cohort	Retrospective cohort	Multicenter ambidirectional cohort	Prospective Iongitudinal	Prospective cohort	Cohort*	Retrospective cohort	Prospective cohort	Cohort
Country	Norway	Italy	Spain	Italy	Norway	NSA	China	USA	Italy	Italy	USA	Spain	Turkey
Author	Blomberg et al [17]	Boscolo-Rizzo et al [27]	Carrillo-Garcia et al [28]	Caruso et al [29]	Caspersen et al [30]	Castro et al [31]	Chai et al [32]	Cirulli et al [33]	Clavario et al [34]	Cristillo et al [35]	Diaz-Fuentes et al [36]	Domenech-Montoliu et al Spain [37]	Erol et al [38]
	12.	13.	4.	15.	16.	17.	18.	19.	20.	21.	22.	23.	24.

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	48.8%	92.6%	81.4%	49.6%	81.2%	75.2%	68.2%	10.1%	44.2%	24.1%	35.3%	91.5%	62.3%	21.4%	76.4%
Follow-up Time Days	365 [†] 4	176 ^f 9	210 ^e 8	210 ^e 4	340 ^e 8	nths tsymptom et	103° 6	6 months posthospital discharge	nths stinfection	180 ^b 2	117.5° 3	3 months gosthospital discharge		122 ^b 2	186° 7
Severity	Mixed	Mixed	7% ICU	7% ICU	6.6% ICU	90.5% required respiratory support	Severe and critical	2.5% ICU	2% hospitalized	14% ICU	Nonhospitalized	Moderate to severe	Severe	mild/moderate (severe excluded)	68% required oxygen therapy 4% ICU
COVID-19 Diagnostic Method	PCR confirmed or clinician diagnosed	Confirmed or clinician-diagnosed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	Not stated	83% PCR confirmed 17% no laboratory confirmation	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	Seropositive	Laboratory confirmed
% Female	39.0	35.7	47.5	47.4	46.9	49.0	41.0	53.3	92.0	39.7	56.0	34.6	30.0	83.0	48.0
Age (Years) Mean/SD Median (IQR)	58.0/12.6	57.9/13	61/17	61/17	61/16	59 (50– 68)	60 (53– 68)	18+	1	64 (54– 76)	49.5/15.3	51/14	54/12	43 (33– 52)	57 (47– 65)
Setting	Hospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients	Not stated	Hospitalized patients	Hospitalized patients	98% nonhospitalized healthcare workers	Hospitalized patients	Community	Hospitalized patients	Hospitalized patients	Health care workers	Hospitalized patients
Controls N, Type	:	ŧ	÷	;	:	÷	:	÷	Ē	i	Norwegian general population norms	ŧ	:	1072 seronegative	÷
Denominator ^a	804	1077	1142	1142	1950	137	107	199	138	116	A47	130	114	323 10	1655
Study Design (as Described by Study,* If Not Stated)	Prospective longitudinal cohort	Prospective longitudinal cohort	Multicenter observational	Multicenter observational	Multicenter cohort	Retrospective	Single-center cohort	Cross-sectional	Cross-sectional	Prospective longitudinal	Cross-sectional survey of a geographical cohort	Prospective Iongitudinal	Prospective Iongitudinal	Cohort with controls	Ambidirectional cohort
Country	¥	¥	Spain	Spain	Spain	France	Belgium	China	¥	Spain	Norway	Mexico	China	Sweden	China
Author	Evans et al (PHOSP-COVID study) [39] (¥)	Evans et al (PHOSP-COVID study) [40] (¥)	Fernandez-de-Las-Penas et al [43] (∞)	Fernandez-de-Las-Penas et al [41] (∞)	Fernandez-de-Las-Penas et al [42] (∞)	Frija-Masson et al [44]	Froidure et al [45]	Fu et al [46]	Gaber et al [47]	Garcia-Abellan et al [48]	Garratt et al [49] (🛮	Gonzalez-Hermosillo et al Mexico [50]	Han et al [51]	Havervall et al [52]	Huang et al [53] (Ω)
	25.	26.	27.	28.	29.	30.	31.	32.	33.	34.	35.	36.	37.	89	39.

Table 1. Continued

Finding:

% With at Least 1 Symptom or Pathology Remaining at Follow up	%0.89	%6.99	41.7%	%2'.29	61.9%	%6.69	37.3%	85.9%	28.7%	64.7%	30.0%	36.4%	20.1%	56.9%
Follow-up Time Days	185°	119.3°	3 months posthospital discharge or visit	195°	93.7%-more than 61.9% 3 months postinfection	90–150 (range) postsymptom onset	395 ^f	90↓	6 months posthospital discharge	111°	169°	6.1 ± 1.1 months postinfection	_p 68	12 months posthospital discharge
Severity	4% ICU	18.6% hospitalized 9.3% ICU	Mild	12% moderate or severe	"Positively tested" Mild and moderate	19.4% severe/critical	62.7% critical/severe	21.1% severe	1.8% ICU	5.9% ICU	6.2% asymptomatic, 84.7% mild illness, 9.0% moderate or severe disease	:	3.9% severe	Mixed
COVID-19 Diagnostic Method	Laboratory confirmed	PCR confirmed	PCR confirmed	PCR confirmed	"Positively tested"	PCR confirmed	"Infected with COVID-19'	PCR confirmed	"Diagnosis of COVID-19"	"Hospitalized for COVID-19"	laboratory- confirmed	"COVID-19 positive patients"	PCR confirmed	Laboratory confirmed
) % Female	47.0	4 46.6	30.6	69.7	9 59.2	4 48.8	80.5	- 48.8	53.3	7 39.9	57.1	3 63.6	53.4	5 49.3
Age (Years) Mean/SD Median (IQR)	59 (49– 67)	43.3/14.4 46.6	18-65	31 (24– 47)	49.8/16.9 59.2	43.6/17.4	39 (33– 48)	47.5 (36– 57)	68 (66-	54.5/16.7	48/15.2	41.4/12.3 63.6	55 (44– 63)	65.1/17.5 49.3
Setting	Hospitalized patients	Hospitalized patients and nonhospitalized	Hospitalized patients and nonhospitalized	Hospitalized patients and nonhospitalized	Community	Hospitalized patients	Hospitalized healthcare workers	Hospitalized patients	Hospitalized patients, elderly	Hospitalized patients	Hospitalized and outpatients	Not stated	Hospitalized patients	Hospitalized patients and ER attendees
Controls N, Type	3383 community dwelling without SARS-CoV-2 infection, 1164 matched pairs	÷	÷	÷	ŧ	:	i	:	466 uninfected spouses who lived together	:	21, "healthy controls recruited via email and flyer advertisements"	÷	:	:
Denominatorª	1227	118	242	006	365	289	303	142	1301 4	153	777	110	204	543
Study Design (as Described by Study,* If Not Stated)	Ambidirectional cohort with controls	Cohort*	Cohort*	Cohort*	Cross-sectional	Cohort	Cohort*	Longitudinal cohort	Cross-sectional	Cohort*	Longitudinal prospective cohort (cross-sectional for controls*)	Observational retrospective	Prospective	Cross-sectional
Country	China	NSA	Pakistan	S Korea	Germany	China	China	China	China	NSA	USA	Italy	China (HK)	Spain
Author	Huang et al [54] (Ω)	Jacobson et al [55]	Kashif et al [56]	Kim et al [57]	Lemhofer et al [58]	Li et al [59]	Liao et al [60]	Liao et al [61]	Liu et al [62]	Liyanage-Don et al [63]	Logue et al [64]	Lucidi et al [65]	Lui et al [66]	Maestre-Muniz et al [67]
	. 04	. 14	42.	43.	4.	45.	46.	47.	48.	49.	. 20	51.	52.	53.

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	26.5%	16.8%	35.8%	82.1%	14.2%	26.5%	24.6%	47.4%	48.0%	21.4%	57.9%	52.0%	39.0%	24.0%	29.0%	39.9%
Follow-up Time Days	168°	180 ^b	90.1⁰	9 months after acute infection	144 [‡]	144 [†]	220°	182 ^e	12 months postdiagnosis	1409	218 ^f	₀ 06	7–9 months postdiagnosis	210 ^b	84°	, 119°
Severity	1.2% hospitalized	27% asymptomatic	78% hospitalized	46.4% hospitalized	Not defined	33.7% severe, 2.6% critical	10.7% asymptomatic, 38.1% severe/very severe	Moderate and severe	:	Not critically ill (ICU/ HDU)	2.6% severe	54% severe	Mild and moderate	Nonsevere	15.5% ICU	29.7% moderate, 1.1% severe
COVID-19 Diagnostic Method	'Positive test'	PCR confirmed	PCR confirmed	Laboratory confirmed	Met relevant clinical criteria	PCR confirmed	PCR confirmed	Not stated	PCR confirmed	PCR confirmed	PCR confirmed and 2.6% severe clinically diagnosed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed
% Female	75.4	:	34.1	34.0	54.1	55.5	49.7	33.0	9.09	19.8	51.1	67.1	67.1	55.0	40.5	74.2
Age (Years) Mean/SD Median (IOR)	Mean range 30–39	9.3 (n/a)	58.5/12.8 34.1	50.6/13.4	59 (47– 68)	59 (47– 68)	47 (33– 58)	59/11	51.5/n/a	37.7/13.7	56 (46– 66)	53.6/13.7	42.7/12.9	36 (27– 48))	62 (n/a)	45/14
Setting	Healthcare workers N	Hospitalized 9 patients and outpatients, pediatric	Hospitalized 5 patients and ER attendees	Hospitalized 5 patients and nonhospitalized	Hospitalized 5 patients	Hospitalized 5 patients	Community 4	Hospitalized 5 patients	Hospitalized 5 patients and outpatients	Hospitalized 3 patients and outpatients	Hospitalized 5 patients	Hospitalized 5 patients	Outpatients 4	Hospitalized 3	Hospitalized 6 patients	Outpatients 4
Controls N, Type	:	:	÷	:	1500, random sample of general population	:	:	÷	:	÷	:	÷	:	:	:	:
Denominator ^a	260	137	226	112	4328	3677	431	135	173	313	2649	173	410	125	200	198
Study Design (as Described by Study,* If Not Stated)	Retrospective cohort	Prospective cohort	Prospective cohort	Single-center cross-sectional	Cohort*	Prospective cohort	Population-based prospective cohort	Prospective cohort	Prospective cohort	Prospective multicenter cross-sectional	Longitudinal cohort	Prospective cross-sectional	Prospective cohort	Cohort*	Prospective cohort	Cross-sectional
Country	Switzerland	France	ltaly	Iraq	China	China	Switzerland	Italy	USA	Bangladesh	Russia	Iran	Switzerland	France	Spain	USA
Author	Martinez et al [68]	Matteudi et al [69]	Mazza et al [70]	Mechi et al [71]	Mei et al [72] (†)	Mei et al [73] (†)	Menges et al [74]	Milanese et al [75]	Millet et al [76]	Mohiuddin Chowdhury et al [77]	Munblit et al [78]	Nabahati et al [79]	Nehme et al [80]	Nguyen et al [81]	Nunez-Fernandez et al [82]	O'Keefe et al [83]
	54.	55.	56.	57.	. 28	29.	.09	61.	62.	63.	. 64.	65.	. 99	. 29	.89	.69

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	11.7%	7.4%	74.3%	24.3%	40.2%	62.2%	52.8%	13.4%	32.6%	3.7%	%2'09	26.7%	19.7%	9.2%
Follow-up Time Days	12 weeks postinfection	°06	At least 3 months postinfection	256 ^f	191°	4 months posttest or first symptoms	125 ^b	3 months posthospital discharge	3 months posthospital discharge	84 ^b	90e ,	6 months postpositive test	84 ^b	₉ 06< 9
Severity	÷	30.1% severe	:	2.7% severe (NIV/IV or PICU)	Mixed	Mixed	4.4% asymptomatic	38% severe	9.4% severe	No hospitalisation	23% severe (ICU), 53% moderate (hospitalized)	4.4% hospitalized	52% hospitalized, 20% ICU	5.8% hospitalized, 2.1% >90 ^b intensive care or ventilation
COVID-19 Diagnostic Method	PCR confirmed	PCR confirmed	Self-report	PCR confirmed	NAAT for confirmed cases; laboratory, imaging or serology for suspected cases	RNA-confirmed	PCR confirmed	PCR confirmed	PCR confirmed	Antibody positive	PCR confirmed	PCR confirmed	PCR confirmed	"Laboratory confirmed"
% Female	52.3	24.6	÷	52.1	53.4	44.0	54.4	56.0	50.0	53.0	39.0	80.0	45.1	60.2
Age (Years) Mean/SD Median (IQR)	2+	44 (33– 56)	÷	10.4 (3.0– 15.2)	53/15.8	48 (37– 57)	39.9/19.4 54.4	58/15	47.5 (37– 57)	6–16	56 (48– 68)	41.6/n/a	56 (45– 66)	:
Setting	Community	Hospitalized patients	Community via social media	Hospitalized children	Hospitalized patients and outpatients	Hospitalized patients and nonhospitalized	96% nonhospitalized patients	Hospitalized patients	Hospitalized patients	Community, children and adolescents	Hospitalized and outpatients	Hospitalized and nonhospitalized healthcare workers	Hospitalized patients and outpatients	Community
Controls N, Type	÷	÷	I	÷	ŧ	÷	:	÷	÷	1246 seronegative	:	125 healthcare workers with negative PCR	÷	Ē
s Denominator ^a	21 374	175	152	518	599	143	180	647	540	109	135	195	421	145 184
Study Design (as Described by Study,* If Not Stated)	Prospective cohort	Prospective longitudinal multicenter cohort	retrospective	Prospective cohort	Bidirectional prospective cohort	Cohort	Longitudinal	Prospective cohort	Multicenter follow-up	Switzerland Longitudinal cohort	Prospective observational cohort	Prospective case-control	Prospective cohort	Matched cohort
Country	¥	Singapore	Italy	Russia	ltaly	USA	Faroe Islands	China	China	Switzerland	Austria	Spain	Italy	Germany
Author	Office for National Statistics [9]	Ong et al [84]	Orru et al [85]	Osmanov et al [86]	Peghin et al [37]	Peluso et al [88]	Petersen et al [89]	Qin et al [90]	Ou et al [91]	Radtke et al [92]	Rass et al [93]	Riestra-Ayora et al [94]		Roessler et al [96] Split cohort (Adults)
	70.	71.	72.	73.	74.	75.	76.	77.	78.	79.	80.	2.	82.	83

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	6.1%	63.9%	23.7%	73.3%	55.4%	57.0%	93.3%	10.0%	54.9%	46.8%	54.9%	51.9%	56.2%	41.0%	46.0%
Follow-up Time Days	q06<	6 months posthospital discharge	₉ 06	140–154 (range) after symptom onset	6 months posthospital discharge	101.5 ^e	222°	183°	3 months after testing positive	104 ^f	103 ^b	258 ^b	126 ^b	117 ^c	117.5°
Severity	1% hospitalized, 0.4% ICU	10.8% ICU	41.6% severe to critical	15.6% mild, 55.2% moderate, 25.0% severe, 4.2% critical	90.8% severe, 9.2% critical	moderate and severe 43% ICU	20.8% no O ₂ , 36.1% supplemental O ₂ , 15.0% noninvasive O ₂ , 28.1% mechanical ventilation	7.5% ICU	14.7% hospitalized	20% ICU	22% ICU	Nonhospitalized, mild	Nonhospitalized	÷	÷
COVID-19 Diagnostic Method	Laboratory confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	Not stated	PCR confirmed or "clinically diagnosed highly suspected"	Spiral chest CT scan or PCR confirmed	PCR confirmed	"Discharge diagnosis of COVID-19"	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed
% Female	48.1	46.3	36.2	57.0	49.2	43.0	41.3	33.3	53.9	38.5	43.0	22	56.8	56.0	56.0
Age (Years) Mean/SD Median (IQR)	:	63/14.4	71.0/5.6	57 (50– 63)	62 (51– 69)	56.1/19.8	59.7 (51.7– 67.7)	54.6/16.9	46.7/n/a	56.2/12.7	57/14	48.6/13.6	48.5/13.5 56.8	49.7/15.2	49.5/15.3
Setting	Community, children	Hospitalized patients	Hospitalized older adult patients	Hospitalized and outpatients	Hospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients and outpatients	Hospitalized patients	Hospitalized and outpatients	Community	Community	Community survey 49.7/15.2 56.0	Community
Controls N, Type	÷	ŧ	:	÷	ŀ	:	1	:	:	:	:	5712 SARS- CoV-2-negative + 3342 randomly selected untested	6006 SARS- COV-2-negative patients	:	ŧ
Denominator ^a	11 950	797	279	146	796	172	327	120	102	126	145	651	672	451	458
Study Design (as Described by Study,* If Not Stated)	Matched cohort	Retrospective longitudinal observational follow-up	Single-center prospective cohort	Prospective cohort	Cohort	Prospective cohort	Prospective cohort	Cohort*	Prospective cohort	Multicenter prospective cohort	Prospective observational	Cohort	Prospective cohort	Cross-sectional	Cross-sectional mixed-mode
Country	Germany	Spain	India	Germany	China	Spain	¥	Iran	Czech Republic	Norway	Austria	Norway	Norway	Norway	Norway
Author	83a. Roessler et al [96] <i>Split</i> (cohort (Children)	84. Romero-Duarte et al [97] 3	85. Sathyamurthy et al [98]	86. Seeβle et al [99]	87. Shang et al [100]	88. Sibila et al [101]	89. Sigfrid et al [102]	90. Simani et al [103]	91. Skala et al [104]	92. Skjorten et al [105]	93. Sonnweber et al [106]	94. Soraas et al [107] (π)	95. Soraas et al [108] (π)	96. Stavem et al [109] (■)	97. Stavem et al [110] (•)

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	96.5%	2.6%	59.1%	47.5%	12.8%	36.5%	29.6%	33.3%	47.9%	53.5%	51.0%
Follow-up Time Days	104°	84 ^b	113	6 months posthospitali zation	180 ^b	g06	3 months posthospital discharge	At least 3 months 33.3% postpositive test	16 weeks posthospital discharge	At least 12 weeks postpositive test	113
Severity	35.4% symptomatic	13.9% visited hospital	87% required oxygen and/or respiratory support, 20% ICU	18.2% ICU	Mixed	Mixed	23% ICU	28.3% moderate, 10.0% severe	÷	1	29.7% ICU, remainder hospitalized
COVID-19 Diagnostic Method	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	"Confirmed diagnosis"	"Confirmed diagnosis", ICD-10 code	"Confirmed COVID-19"	PCR confirmed	"Presumed and confirmed"	PCR confirmed or antibody positive	PCR confirmed or by CT scan
) % Female	63.5	3 57.0	34.3	40.5	55.6	3 55.6	46.0	9 28.0	3 38.2	80.0	1 42.1
Age (Years) Mean/SD Median (IQR)	11–17	46.0/15.8	59.6/14	6.9/14.1	46/19.7	46.3/19.8 55.6	57 (48– 66)	33.7/7.29 58.0	58.6/15.3	20–69	60.9/16.1 42.1
Setting	Community, adolescents	Community	Hospitalized patients	Hospitalized patients	healthcare organisations including hospitals, primary care, and specialist providers	Hospitalized patients and nonhospitalized	Hospitalized patients	Hospitalized and nonhospitalized healthcare workers	Hospitalized patients	Healthcare workers 20–69	Hospitalized patients
Controls N, Type	3739 who tested negative	4182, matched PCR negative***	:	:	105 579 diagnosed with flu, 236 038 with any other RTI including flu	106 578 matched cohort with influenza and without a diagnosis of COVID-19 or positive test	:	÷	÷	1	:
Denominator ^a	3065	4182	127	183	236 379	273 618	115	120	545	217	478
Study Design (as Described by Study,* If Not Stated)	Matched cohort	Prospective observational cohort	Cohort*	Cross-sectional observational	Retrospective cohort with matching	Retrospective cohort	Cohort follow-up	Retrospective cohort	Cohort*	Cross-sectional*	Prospective uncontrolled cohort
Country	¥	UK, USA and Sweden	Y N	Spain	Primarily USA	USA	Italy	Egypt	Y N	Republic of Ireland	France
Author	98. Stephenson et al [111]	99. Sudre et al [112]	100. Sykes et al [113]	101. Taboada et al [114]	102. Taquet et al [116] (<))	103. Taquet et al [115] (<))	104. Tarsitani et al [117]	105. Tawfik et al [118]	106. Taylor et al [119]	107. Tempany et al [120]	108. The Writing Committee for the COMEBAC Study Group [121]

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	1.8%	56.3%	40.2%	5.0%	57.8%	24.4%	43.8%	51.4%	53.8%	44.4%	37.7%	70.4%	49.6%
Follow-up Time Days	3 months after discharge (hospitalized), 4 months postsymptom onset (nonhospitalized)	122 [†]	3 months posthospital discharge	3 months posthospital discharge	72 [†]	194°	6±3 months postpositive test	105°	186 ^f	89.5°	84 ^b	153 [†]	97 [†]
Severity	Mixed	Mixed 30.2% ICU	69.7% severe	25% moderate, 45% severe	55.5% hospitalized	63.7% hospitalized	10.7% hospitalized, 1.6% ICU	88.4% admitted 8.6% ICU	26% severe	28.2% severely ill	0.8% admitted to hospital	100% severe, 5% ICU	5% critical, 33.5% severe
COVID-19 Diagnostic Method	PCR confirmed, or discharge diagnosis of "confirmed or unconfirmed COVID-19"	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	Positive nasopharyngeal swab	PCR confirmed	PCR confirmed	PCR confirmed	Self-reported	"Infected with COVID-19"	"confirmed"
% Female	51.0	23.0	40.2	43.0	53.9	42.0	77.4	32.9	43.0	44.4	57.3	77.0	54.5
Age (Years) Mean/SD Median (IQR)	52.9/15.5	52.5/14.0 23.0	53.6/14.9	55.5/6.2	49.5/15	56 (43– 69)	45/12	63/13.6	74.3/n/a	:	-18+	36 (31– 43)	52 (41– 62)
Setting	Hospitalized patients and nonhospitalized	Hospitalized patients	Hospitalized patients	Hospitalized and outpatients	Hospitalized and nonhospitalized	Hospitalized patients and outpatients	Community via social media	Emergency Department and hospitalized patients	Hospitalized older adult patients	Hospitalized patients	Community	Hospitalized healthcare workers	Hospitalized patients
Controls N, Type	:	:	E	100 randomly recruited from hospital registration system without COVID-19	÷	i	i	÷	÷	÷	ŧ	:	184, volunteers
S Denominator ^a	683	222	239	100	128	168	616	767	106	117	76155	162	538
Study Design (as Described by Study,* If Not Stated)	Multicenter prospective cohort	Prospective cohort	Single-center cohort	Retrospective comparative study with controls	Cross-sectional*	Cross-sectional	Cross-sectional	Cohort*	Cohort	Retrospective	Random community- based survey (REACT-2)	Ambidirectional cohort	Longitudinal with controls
Country	Norway	Saudi Arabia	Brazil	Egypt	Republic of Ireland	Italy	Italy	Italy	Norway	China	¥	China	China
Author	09. Tholin et al [122] (•)	110. Tleyjeh et al [123]	11. Todt et al [124]	12. Tohamy et al [125]	13. Townsend et al [126]	14. Trunfio et al [127]	115. Ursini et al [128]	116. Venturelli et al [129]	17. Walle-Hansen et al [130] Norway	18. Weng et al [131]	119. Whitaker et al [132]	120. Xiong et al [133]	121. Xiong et al [134]
	109. Tho	110. Tle	111. Toc	112. Tot	113. To	114. Tru	115. Urs	116. Vei	117. Wa	118. We	119. W	120. Xio	121. Xio

Table 1. Continued

Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up	%0.0	39.5%	55.8%	35.9%	29.8%	54.9%	72.7%	45.0%	%9.5%
Follow-up Time Days	84°	365°	203 ^b	289 ^b	348°	924	₉ 06	364 [†]	129° (severe cases) 125° (mild)
Severity	asymptomatic/mild symptoms	24% severe	12.8% severe, 3.6% ICU	34.2% hospitalized, 5% ICU	15.7% severe	mild cases excluded, only patients with pulmonary sequelae at discharge included	Nucleic acid testing 9.3% severe/critical	27.9% severe	54.6% severe
COVID-19 Diagnostic Method	"Diagnosed with COVID-19"	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	PCR confirmed	Nucleic acid testing	Laboratory confirmed	PCR and antibody test
% Female	0.0	29.0	49.5	63.0	58.7	50.3	43.8	50.5	56.9
Age (Years) Mean/SD Median (IQR)	35 (30– 49)	53.0/12.2 59.0	53.5/14.8 49.5	49.6/18.7 63.0	49 (40– 57)	51 (31.8–61.0)	43 (33- 54)	60 (49– 68)	:
Setting	Mobile cabin hospital, adult males	Hospitalized patients	Hospitalized patients	Hospitalized patients and outpatients	Hospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients	Hospitalized patients
Controls N, Type	:	i	:	ŧ	:	÷	÷	ŧ	42 healthy controls Hospitalized —negative patients nucleic acid and antibody tests
S Denominator ^a	125	119	337	354	121	122	245	2433	164
Study Design (as Described by Study,* If Not Stated)	Prospective observational	Cohort	Retrospective analysis	Retrospective cohort	Prospective cohort	Retrospective comparative	Cohort*	Retrospective multicenter cohort	Prospective cohort with controls
Country	China	China	China	France	China	China	China	China	China
Author	122. Yan et al [135]	123. Yan et al [136]	124. Yin et al [137]	125. Zayet et al [138]	126. Zhan et al [139]	127. Zhang et al [140]	128. Zhang et al [141]	129. Zhang et al [142]	130. Zhou et al [143]

Abbreviations: COVID, coronavirus disease 2019; CT, computerised tomography; ER, emergency room; HDU, high dependency unit; ICD, intensive care department; ICU, intensive care unit; RTI, respiratory tract infection; SARS-COV-2, severe acute respiratory syndrome coronavirus 2; SD, standard deviation; UK, United Kingdom.

NOTE: Papers coded with the following symbols are different publications from the same study data: Ω , \blacksquare , \diamondsuit , Ψ , \dagger , ∞ , π .

* refers to those studies where study design was not explicitly stated so a design was designated based on the study description.

*** refers to studies where the relevant outcome data was not available for controls.

^a Different denominators specific to each outcome have been used in cases where data are incomplete or where individual symptoms have different denominators.

^bMean number of days postsymptom onset or positive test.

^cMedian number of days postsymptom onset or positive test.

dMedian number of days posthospital admission.

^eMean number of days posthospital discharge.

Median number of days posthospital discharge.

^gMean number of days postnegative test after infection.

	region	mean days since infection	cases	total	% persistent symptoms	12
Abdelrahman et al Al-Aly et al (hospitalized)	Other N America	270 150	105	172	61.0 (53.3, 68.4) 9.2 (8.7, 9.8)	-
Al-Aly et al (hospitalized) Al-Aly et al (non-hospitalized)	N America N America	126	1718	11800 60255	9.2 (8.7, 9.8)	
Aminian et al	N America	243	1255	2839	44.2 (42.4, 46.1)	
Arnold et al	Europe	90	81	110	73.6 (64.4, 81.6)	-
Augustin et al	Europe	131	123	442	27.8 (23.7, 32.3)	• :
Ayoubkhani et al	Europe	154 139	6085 66	28335	21.5 (21.0, 22.0) 32.4 (26.0, 39.2)	•
Baricich et al Becker et al	N America	231	178	204 740	24 1 (21.0, 27.3)	
Bellan et al	Europe	120	128	238	53.8 (47.2, 60.2)	- L
Blanco et al	Europe	104	52	100	52.0 (41.8, 62.1)	-
Bliddal et al	Europe	84	52	129	40.3 (31.8, 49.3)	
Blomberg et al	Europe	183	189	312	60.6 (54.9, 66.0)	
Boscolo-Rizzo et al	Europe	365	161	304	53.0 (47.2, 58.7)	-
COMEBAC study Carrillo-Garcia et al	Europe	127	244	478 151	51.0 (46.5, 55.6)	
Carrillo-Garcia et al Caruso et al	Europe	182	91	118	66.2 (58.1, 73.7) 77.1 (68.5, 84.3)	
Caspersen et al	Europe	350	28	170	16.5 (11.2, 22.9)	
Castro et al	N America	128	721	6619	10.9 (10.2, 11.7)	
Chai et al	China	378	156	546	28.6 (24.8, 32.6)	• 1
Cirulli et al	N America	90	18	122	14.8 (9.0, 22.3)	-
Clavario et al	Europe	121	160	200	80.0 (73.8, 85.3)	
Cristillo et al	Europe	196	50	101	49.5 (39.4, 59.6)	+-
Disz-Fuentes et al Domenech-Montoliu et al	N America	84 213	88 258	111	79.3 (70.5, 86.4) 53.4 (48.9, 57.9)	-
Domeneon-Montoliu et al Erol et al	Europe	170	45	483 121	37 2 (28 6 46 4)	200
Fernandez-de-Las-Penas et al	Europe	227	930	1142	81.4 (79.1, 83.7)	76
Frija-Masson et al	Europe	91	103	137	75.2 (67.1, 82.2)	1
Froidure et al	Europe	95	73	107	68.2 (58.5, 76.9)	-
Fu et al	China	196	20	199	10.1 (6.2, 15.1)	
Gaber et al	Europe	122	61	138	44.2 (35.8, 52.9)	-
Garcia-Abellan et al	Europe	182	28	116	24.1 (16.7, 33.0)	-
Goruzalez-Hermosillo et al	Other	105	119	130	91.5 (85.4, 95.7)	-
Han et al	China	175	71	114	62.3 (52.7, 71.2)	
Havervall et al Huang et al	Europe	122 186	69 1265	323 1655	21.4 (17.0, 26.2) 76.4 (74.3, 78.5)	•
Huang et al Jacobson et al	China N America	186	1265	1655	76.4 (74.3, 78.5) 66.9 (57.7, 75.3)	
Kashif et al	Other	105	101	242	41.7 (35.5, 48.2)	
Kim et al	Other	195	591	900	65.7 (62.5, 68.8)	
Lemhofer et al	Europe	91	226	365	61.9 (56.7, 66.9)	
Li et al	China	120	173	289	59.9 (54.0, 65.6)	-
Liao et al	China	409	113	303	37.3 (31.8, 43.0)	•1
Liao et al	China	104	122	142	85.9 (79.1, 91.2)	
Liu et al Liyanage-Don et al	China N America	196	373	1301 153	28.7 (26.2, 31.2) 64.7 (56.6, 72.3)	•
Liganage-Lion et al Logue et al	N America	169	53	177	30.0 (23.3, 37.3)	
Lucidi et al	Europe	186	40	110	36.4 (27.4, 46.1)	-1
Lui et al	China	96	41	204	20.1 (14.8, 26.3)	•
Maestre-Muniz et al	Europe	379	309	543	56.9 (52.6, 61.1)	
Martinez et al	Europe	168	69	260	26.5 (21.3, 32.3)	•
Matteudi et al	Europe	180	23	137	16.8 (11.0, 24.1)	- 1
Mazza et al	Europe	104	81	226	35.8 (29.6, 42.5)	
Mechi et al	Other	274	92 976	112	82.1 (73.8, 88.7) 26.5 (25.1, 28.0)	-
Mei et al Menges et al	China	158 219	106	3677 431	26.5 (25.1, 28.0) 24.6 (20.6, 28.9)	
Milanoso et al	Europe	196	64	135	47.4 (38.8, 56.2)	
Millet et al	N America	365	83	173	48.0 (40.3, 55.7)	
Mohiuddin et al	Other	161	67	313	21.4 (17.0, 26.4)	
Munblit et al	Europe	232	1534	2649	57.9 (56.0, 59.8)	
Nabahati et al	Other	104	90	173	52.0 (44.3, 59.7)	
Nehme et al	Europe	243	160	410	39.0 (34.3, 43.9)	*
Nguyen et al	Europe	213	30	125	24.0 (16.8, 32.5)	
Nunez-Fernandez et al	Europe	98	58	200	29.0 (22.8, 35.8)	
O'Keefe et all ONS study August 2021	N America Europe	119 84	79 2501	198	39.9 (33.0, 47.1) 11.7 (11.3, 12.1)	
Ong et al	Other	104	13	175	7.4 (4.0, 12.4)	
Orru et al	Europe	91	113	152	74.3 (66.6, 81.1)	- ·
Osmanov et al	Europe	270	126	519	24.3 (20.6, 28.2)	
PHOSP-COVID study	Europe	174	797	861	92.6 (90.6, 94.2)	
Peghin et al	Europe	191	241	599	40.2 (36.3, 44.3)	
Peluso et al	N America	112	89	143	62.2 (53.8, 70.2)	
Petersen et al	Europe	125	95	180	52.8 (45.2, 60.2)	(************************************
Qin et al Qu et al	China	105	87 176	647 540	13.4 (10.9, 16.3)	•
Cou et al Radtke et al	Europe	84	1/6	109	32.6 (28.7, 36.7) 3.7 (1.0, 9.1)	
Rass et al	Europe	90	82	135	60.7 (52.0, 69.0)	
Riestra-Ayora et al	Europe	182	52	195	26.7 (20.6, 33.5)	
Righi et al	Europe	84	83	421	19.7 (16.0, 23.8)	•
Roessier et al (adults)	Europe	91	13396	145184	9.2 (9.1, 9.4)	•
Roessler et al (children)	Europe	91	734	11950	6.1 (5.7, 6.6)	•
Romero-Duarte et al	Europe	196	509	797	63.9 (60.4, 67.2) 23.7 (18.8, 29.1)	_ •
Sathyamurthy et al Seofile et al	Other	104	107	279 146	23.7 (18.8, 29.1)	* 1
Seettle et al. Shang et al.	China	196	441	796	73.3 (65.3, 80.3) 55.4 (51.9, 58.9)	
Sibila et al	Europe	116	98	172	57.0 (49.2, 64.5)	
Sigfrid et al	Europe	222	305	327	93.3 (90.0, 95.7)	
Simani et al	Other	197	12	120	10.0 (5.3, 16.8)	•
Skala et al	Europe	91	56	102	54.9 (44.7, 64.8)	
Skjorten et al	Europe	105	59	126	46.8 (37.9, 55.9)	-
Sonnweber et al	Europe	103	73	133	54.9 (46.0, 63.5)	-
Soraas et al Stavem et al	Europe	126	380	676	56.2 (52.4, 60.0)	
Stavem et al Stephenson et al	Europe Europe	117	185 2038	451 3065	41.0 (36.4, 45.7) 66.5 (64.8, 68.2)	7
Stephenson et al Sudre et al	Other	84	108	4182	06.5 (64.8, 68.2) 2.6 (2.1, 3.1)	
Sykes et al	Europe	127	75	127	59.1 (50.0, 67.7)	-
Taboada et al	Europe	189	87	183	47.5 (40.1, 55.0)	+■-
laquet et al	N America	90	100007	273618	36.5 (36.4, 36.7)	•:
	Europe	105	34	115	29.6 (21.4, 38.8)	
larsitani et al	Other	91	40	120	33.3 (25.0, 42.5)	
fawfik ot al		126	261	545	47.9 (43.6, 52.2) 53.5 (46.6, 60.2)	ff
Tawfik et al Taylor et al	Europe		116	217		-
lawfik et al laylor et al l'empany et al	Europe	84			56.3 (49.5, 62.9)	
lawfik et al laylor et al l'empany et al l'loyjeh et al	Europe Other	136	125		40 2 (33 9 46 7)	
Fawfik et al Faylor et al Fempany et al Floyjieh et al Flodt et al	Europe Other Other	136 105	125 96	239	40.2 (33.9, 46.7)	+
Fawfik et al Faylor et al Fempany et al Floyjeh et al Fock et al Fockmy et al	Europe Other	136	125 96 5		40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5)	-
Fawfik et al Faylor et al Fiempany et al Filoyjeh et al Foodt et al Fohamy et al Fohamy et al	Europe Other Other Europe	136 105 105	125 96	239 100	40.2 (33.9, 46.7) 5.0 (1.6, 11.3)	• •
l'avrifik et al l'aylor et al l'impany et al l'impjeh et al l'importe al l'impany et al l'impany et al l'impany et al	Europe Other Other Europe Europe	136 105 105 86	125 96 5 74	239 100 128	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5)	-
Favelik et al Faylor et al Fempany et al Flogigh et al Fothamy et al Fownsend et al Fransio et al Josini et al	Europe Other Other Europe	136 105 105 86 194	125 96 5 74 41	239 100 128 168	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6)	-
Favefik et al Taylor et al Fempany et al Fempany et al Feogleh et al Footh et al Foothery et al Foomseend et al Frunfio et al Junini et al Vallie-Hanseen et al	Europe Other Other Europe Europe Europe	136 105 105 86 194 182	125 96 5 74 41 270	239 100 128 168 616	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9)	
Favefik et al Taylor et al Fempany et al Fempany et al Feogleh et al Footh et al Foothery et al Foomseend et al Frunfio et al Junini et al Vallie-Hanseen et al	Europe Other Other Other Europe Europe Europe Europe	136 105 105 86 194 182 105	125 96 5 74 41 270 394 57	239 100 128 168 616 767 106 117	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.8 (43.8, 63.5) 44.4 (35.3, 53.9)	-
l'awrific et al l'ayrifor et al l'ayrifor et al l'ord et al l'ord et al l'ord et al l'arrific et al l	Europe Other Other Other Europe Europe Europe Europe Europe China Europe	138 105 105 86 194 182 105 200	125 96 5 74 41 270 394 57	239 100 128 168 616 767 106	40.2 (33.9. 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.8 (43.8, 63.5) 44.4 (36.3, 53.9) 37.7 (37.4, 38.0)	
l'audité o al l'audité o al l'augitor et al l'augitor et al l'otto et al l'otto et al l'audité et al l'	Europe Other Other Other Europe Europe Europe Europe Europe China Europe China	136 105 105 86 194 182 105 200 104 84 167	125 96 5 74 41 270 394 57 52 28713	239 100 128 168 616 767 106 117 76155 162	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.8 (43.8, 63.5) 44.4 (35.3, 53.9) 37.7 (37.4, 38.0) 70.4 (62.7, 77.3)	
l'aurille et al l'Epylor et al l'Epylor et al l'Octo et al l'Octo et al l'Champer et al l'Ocumend et al l'arrier et al l'Aurille et al l'Octo et al l'	Europe Other Other Other Europe Europe Europe Europe Europe China Europe China China	136 105 105 86 194 182 105 200 104 84 167	125 96 5 74 41 270 394 57 52 28713 114 267	239 100 128 168 616 767 106 117 76155 162 538	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.8 (43.8, 63.5) 44.4 (35.3, 53.9) 37.7 (37.4, 38.0) 70.4 (62.7, 77.3) 48.6 (45.3, 53.9)	
l'aurélic et al l'appiro et al l'appiro et al l'appiro et al l'octo et al l'abbanne et al l'abbanne et al l'aurèlic et al l'orige et a	Europe Other Other Other Europe Europe Europe Europe Europe Europe China China China China	136 105 105 86 194 182 105 200 104 84 167 111	125 96 5 74 41 270 394 57 52 28713 114 267	239 100 128 168 616 767 106 117 76155 162 538 125	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 5.7.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.4 (43.8, 63.5) 44.4 (35.3, 53.9) 37.7 (37.4, 38.0) 70.4 (62.7, 77.3) 49.6 (45.3, 53.9) 0.0 (0.0, 2.9)	
Twentik or all Taylor of all Tempany et al Tempany et al Tempany et al Teophe et al Tode et al Todeny et al Townsend et al	Europe Other Other Other Europe Europe Europe China Europe China China China	136 105 105 86 194 182 105 200 104 84 167 111 98 379	125 96 5 74 41 270 394 57 52 28713 114 267 0 47	239 100 128 168 616 767 106 117 76155 162 538 125 119	40.2 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 66.5) 24.4 (18.1, 31.6) 43.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.8 (43.8, 63.5) 44.4 (35.3, 53.9) 37.7 (37.4, 38.0) 70.4 (62.7, 77.3) 49.6 (45.3, 53.9) 93.5 (30.7, 48.9)	
fauntité et al hypotre et al femigrany et al limpoire et al femigrany et al limpoire et al foot et al al limpoire et al foot et al al limpoire et al foot et al al foot et al al foot et al al foot et al f	Europe Other Other Other Europe Europe Europe Europe Europe China Europe China China China China China	136 105 105 86 194 182 105 200 104 84 167 111 98 379 203	125 96 5 74 41 270 394 57 52 28713 114 267 0 47	239 100 128 168 616 767 106 117 76155 162 538 125 119 316	402 (33.9, 46.7) 5.0 (1.6, 11.3) 57.8 (48.8, 68.5) 24.4 (18.1, 31.6) 24.8 (39.9, 47.9) 51.4 (47.8, 55.0) 53.8 (43.8, 63.5) 44.4 (35.3, 53.9) 37.7 (37.4, 38.0) 70.4 (62.7, 77.3) 49.6 (45.3, 53.9) 0.0 (0.0, 2.9) 35.5 (30.7, 48.9) 55.7 (50.0, 61.3)	
Guardie et al Spiyote et al Ferrycaper et al	Europe Other Other Other Other Europe Europe Europe Europe China China China China China China China China China	136 105 105 86 194 182 105 200 104 84 187 111 98 379 200 289	125 96 5 74 41 270 394 57 52 28713 114 267 0 47 176	239 100 128 168 616 767 106 117 78155 162 538 125 119 316 354	40 2 (35, 48.7) 5.0 (1.6, 11.3) 5.7 (4.68, 66.5) 24.4 (181, 31.6) 43.8 (39.8, 47.9) 51.4 (47.6, 55.0) 53.8 (43.8, 63.5) 44.4 (35.3, 53.9) 70.4 (62.7, 77.3) 49.6 (45.3, 53.9) 53.7 (37.4, 38.0) 53.8 (30.7, 48.9) 55.7 (30.0, 61.3) 55.7 (30.0, 61.3)	
l'authic et al l'hyport et al l'implyant et al	Europe Other Other Other Europe Europe Europe Europe China Europe China China China China China China China China	136 105 105 105 86 194 182 105 200 104 84 167 111 98 379 203 289 348	125 96 5 74 41 270 394 57 52 28713 114 267 0 47 176 127 36	239 100 128 168 616 767 106 117 76155 162 538 125 119 316 354 121	40 2 (3) 5, 48.7) 50 (16, 11.3) 57 (2) 48.8 (85.5) 24.4 (181, 31.6) 24.4 (181, 31.6) 51.4 (47.8, 55.0) 53.2 (2) 8, 63.5) 44.4 (35.3, 53.9) 9.7 (37.4, 38.0) 9.0 (00, 2.9) 38.5 (307, 48.9) 55.7 (80.0, 61.3) 29.8 (21.8, 38.7)	
Guardisc et al Stephysic et al Ferregraphy et al	Europe Other Other Other Europe Europe Europe Europe China Europe China China China China Europe China China China China China	138 105 105 86 194 182 105 200 104 84 167 111 98 379 203 289 348	125 96 5 74 41 270 394 57 52 28713 114 267 0 47 176 127 36	239 100 128 168 616 767 106 117 76155 162 538 125 119 316 354 121 2433	40.2(33.4.47) 50.16.1(3) 57.8(48.8.66.5) 24.4(19.3.16) 24.4(19.3.16) 51.4(476.55.0) 53.4(43.6.35) 44.(53.5.35) 70.4(62.7.77.3) 49.6(45.3.53) 00.(00.2.9) 38.5(30.7.48.9) 35.7(37.4.30.0) 70.4(62.7.77.3) 49.6(45.3.53.9) 55.7(50.0.61.3) 35.9(30.8.41.1) 29.8(21.8.38.7)	
Transtant et al Territorio et	Europe Other Other Other Europe Europe Europe Europe China Europe China China China China China China China China	136 105 105 105 86 194 182 105 200 104 84 167 111 98 379 203 289 348	125 96 5 74 41 270 394 57 52 28713 114 267 0 47 176 127 36	239 100 128 168 616 767 106 117 76155 162 538 125 119 316 354 121	40 2 (3) 5, 48.7) 50 (16, 11.3) 57 (2) 48.8 (85.5) 24.4 (181, 31.6) 24.4 (181, 31.6) 51.4 (47.8, 55.0) 53.2 (2) 8, 63.5) 44.4 (35.3, 53.9) 9.7 (37.4, 38.0) 9.0 (00, 2.9) 38.5 (307, 48.9) 55.7 (80.0, 61.3) 29.8 (21.8, 38.7)	

Figure 2. Forest plot of prevalence of Long COVID in the included studies, with 95% prediction intervals.

Pathology tended to be reported in only a small number of studies, with the exception of lung pathology, which was reported in 26 studies.

There were very few studies with a low risk of bias (Supplementary Table 2). Few studies used a sample that was representative of all COVID-19 cases in the population. Approximately half of the studies indicated that symptoms had not been present before infection, whereas the rest did not report ascertaining this. When stratifying by risk of bias, generally lower prevalence estimates were seen in studies with COVID-19 diagnoses confirmed for all participants, studies scored as having a representative sample, studies with an internal or external non-COVID-19 comparator, studies that assessed all participants in the same way, and studies based on community participants (Supplementary Figures 41 and 42).

Comorbidities, ethnicity, and other demographic data were not reported in all studies. Higher prevalence of Long COVID was observed in studies in which study samples had higher proportions of older people (<50 years PE 38.5%, PI 7.9%-82.1%; 50+ years PE 47.7%, PI 7.9%-90.6%), males (<50% female PE 45.6%, PI 5.5%-92.4%; 50%+ female PE 38.7%, PI 8.5%-81.2%), people of non-White ethnicity (<50% White ethnicity PE 56.3%, PI 22.3%-85.2%; 50%+ White ethnicity PE 37.6%, PI 1.7%-95.3%), diabetes (<10% pre-existing diabetes PE 35.4%, PI 5.7%-83.2%; 10%+ pre-existing diabetes PE 51.9%, PI 8.3%-92.8%), hypertension (<30% pre-existing hypertension PE 37.3%, PI 7.0%-82.5%; 30%+ pre-existing hypertension PE 58.5%, PI 16.9%-90.7%), cardiovascular disease (<10% pre-existing CVD PE 38.2%, PI 5.9%-85.9%; 10%+ pre-existing CVD PE 54.7%, PI 9.4%-93.4%), and other comorbidities including obesity, respiratory disease, liver disease, kidney disease, and immunological disorder or allergy (Supplementary Figure 43). Prevalence of Long COVID did not differ substantially with smoking status.

When subgrouping by study design, the range was 0% to 93% (PE, 41.3%; PI, 6.0%-88.6%) in cohort studies and 10% to 82% (PE, 45.9%; PI, 11.2%-85.1%) in cross-sectional studies (Supplementary Figure 50). Prevalence estimates derived from assessing Long COVID as self-reported symptoms and function (n = 93) on the whole tended to report higher prevalence (PE, 43.9%; PI, 8.2%-87.2%) than those that used clinical coding in healthcare records (n = 9) (PE, 13.6%; PI 1.2%–68%). However, studies that had dedicated pathology follow up of COVID-19 patients (for example, pulmonary function tests or scans with pathology discovered at follow up) tended to report the highest prevalence (n = 20) (PE, 51.7%; PI 12.3%– 89.1%) (Figure 4). Studies that defined Long COVID as at least 1 of multiple symptom or pathology domains tended to report a slightly higher prevalence than those that assessed a single symptom/pathology domain (Supplementary Figure 44).

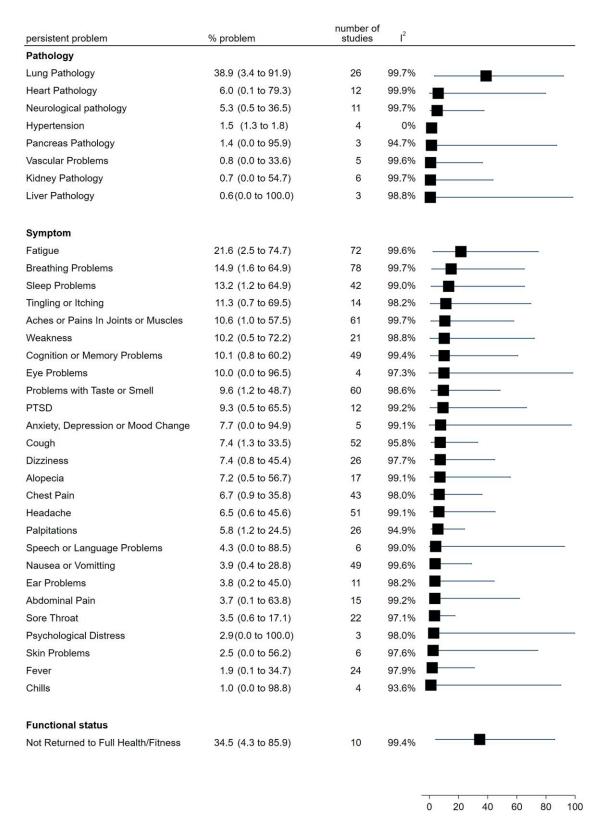


Figure 3. Forest plot of individual symptoms, pathology, and functional disability identified in the included studies, with 95% prediction intervals.

study pathology discovered at follow-up	Dominant source of participants	Breadth of coverage	Mean days since infection	Cases	Total	% per symp	otoms	
lellan et al	hospitalised	multiple domains	120	128	238	53.8	(47.2, 60.2)	•
Sianco et al Frija-Masson et al	hospitalised hospitalised	single domain single domain	104 91	52 103	100	52.0 75.2	(41.8, 62.1) (67.1, 82.2)	•
roidure et al	hospitalised	single domain	95	73	107	68.2	(58.5, 76.9)	-
fan et al	hospitalised	single domain	175	71	114	62.3	(52.7, 71.2)	-
i et al	Localisations	single domain	120	173	289	59.9 85.9	(54.0, 65.6)	•
.iso et al	hospitalised healthcare workers	single domain single domain	104	122	142	37.3	(79.1, 91.2) (31.8, 43.0)	1
Milanese et al	hospitalised	multiple domains	196	64	135	47.4	(38.8, 56.2)	•
Nabahati et al	hospitalised	single domain	104	90	173	52.0	(44.3, 59.7)	-
Nunez-Fernandez et al Din et al	hospitalised hospitalised	single domain single domain	98	58 87	200 647	29.0 13.4	(22.8, 35.8) =- (10.9, 16.3)	
Call et al	hospitalised	single domain	116	98	172	57.0	(49.2, 64.5)	
Sonnweber et al	community	single domain	103	73	133	54.9	(46.0, 63.5)	-
Fohamy et al	outpatients	single domain	105	5	100	5.0	(1.6, 11.3)	00000
Yan et al	hospitalised hospitalised	single domain single domain	379	47 176	119	39.5 55.7	(30.7, 48.9)	- _
Yon et al Zhang et al	hospitalised	single domain	106	67	122	54.9	(50.0, 61.3)	
Zhang et al	hospitalised	single domain	105	178	245	72.7	(66.6, 78.1)	-
Zhou et al	hospitalised	single domain	127	114	164	69.5 51.7	(61.9, 76.5) (12.3, 89.1)	△ ■
						01.7	(12.3, 69.1)	\vee
symptoms & function discovered at follow-up								
Abdelrahman et al	outpatients	multiple domains	270	105	172	61.0	(53.3, 68.4)	-
Amoid et al Augustin et al	hospitalised community	multiple domains multiple domains	90	81 123	110 442	73.6 27.8	(64.4, 81.6) (23.7, 32.3)	-
Saricich et al	hospitalised	single domain	131	66	204	32.4	(26.0, 39.2)	
Becker et al	outpatients	single domain	231	178	740	24.1	(21.0, 27.3)	
Bliddal et al	community	multiple domains	84	52	128	40.3	(31.8, 49.3)	-
Biomberg et al Boscolo-Rizzo et al	outpatients	multiple domains	183 365	189	312 304	60.6 53.0	(54.9, 66.0)	-
SOSCOIO-HIZZO 61 81 COMEBAC study	community hospitalised	multiple domains multiple domains	127	161 244	478	51.0	(47.2, 58.7) (46.5, 55.6)	
Carrillo-Garcia et al	hospitalised	multiple domains	105	100	151	66.2	(58.1, 73.7)	-
Canuso et al	hospitalised	multiple domains	182	91	118	77.1	(68.5, 84.3)	
Chai et al	hospitalised	multiple domains	378	156	546	28.6	(24.8, 32.6)	0.50
Cirulli et al Clavario et al	community	multiple domains	90 121	18 160	122	14.8	(9.0, 22.3) 	-
Clavario et al Cristillo et al	hospitalised hospitalised	single domain single domain	121	160 50	101	80.0 49.5	(73.8, 85.3) (39.4, 59.6)	
Diaz-Fuentes et al	hospitalised	multiple domains	84	88	111	79.3	(70.5, 86.4)	
Domenech-Montoliu et al	community	multiple domains	213	258	483	53.4	(48.9, 57.9)	
Erol et al	hospitalised	multiple domains	170	45	121	37.2	(28.6, 46.4)	F _
Fernandez-de-Las-Penas et al Fu et al	hospitalised hospitalised	single domain single domain	227 196	930	1142	81.4	(79.1, 83.7) (6.2, 15.1)	
ru et al Gaber et al	hospitalised healthcare workers	multiple domains	196	61	199	44.2	(8.2, 15.1)	
Sarcia-Abellan et al	hospitalised	multiple domains	182	28	116	24.1	(16.7, 33.0)	
Gonzalez-Hermosillo et al	hospitalised	multiple domains	105	119	130	91.5	(85.4, 95.7)	
Havervall et al	healthcare workers	multiple domains	122	69 1265	323 1655	21.4 76.4	(17.0, 26.2) (74.3, 78.5)	_
Huang et al Jacobson et al	hospitalised outpatients	multiple domains multiple domains	186	1265	1655	76.4	(74.3, 78.5) (57.7, 75.3)	-
Kashif et al		single domain	105	101	242	41.7	(35.5, 48.2)	-
Kim et al	outpatients	multiple domains	195	591	900	65.7	(62.5, 68.8)	
Lemhofer et al	community	multiple domains	91	226	365	61.9	(56.7, 66.9)	-
Liu et al Livanage-Don et al	hospitalised hospitalised	single domain multiple domains	196	373 99	1301	28.7 64.7	(26.2, 31.2) (56.6, 72.3)	_
.nyanage-bon et al	outpatients	multiple domains	169	53	177	30.0	(23.3, 37.3)	-
ucidi et al		single domain	186	40	110	36.4	(27.4, 46.1)	F
Lui et al	hospitalised	multiple domains	96	41	204	20.1	(14.8, 26.3)	
Maestre-Muniz et al	hospitalised	multiple domains	379	309	543	56.9	(52.6, 61.1) (21.3, 32.3)	•
Martinez et al Matteuck et al	healthcare workers outpatients	multiple domains multiple domains	168 180	69 23	260 137	26.5 16.8	(21.3, 32.3)	
Mazza et al	hospitalised	single domain	104	81	226	35.8	(29.6, 42.5)	F.
Mechi et al	outpatients	multiple domains	274	92	112	82.1	(73.8, 88.7)	-
Mei et al	hospitalised	multiple domains	158	976	3677	26.5	(25.1, 28.0)	
Menges et al Millet et al	community hospitalised	multiple domains multiple domains	219 365	106 83	431 173	24.6 48.0	(20.6, 28.9) (40.3, 55.7)	_
Mohiuddin et al	outpatients	multiple domains	161	67	313	21.4	(17.0, 26.4)	
Munbit et al	hospitalised	multiple domains	232	1534	2649	57.9	(56.0, 59.8)	
Nehme et al		multiple domains	243	160	410	39.0	(34.3, 43.9)	
Nguyen et al O'Kieefe et al	hospitalised outpatients	single domain multiple domains	213 119	30 79	125 198	24.0 39.9	(16.8, 32.5)	
O'Ricete et al ONS study August 2021	community	multiple domains multiple domains	119	2501	21374	11.7	(33.0. 47.1)	
Ong et al	hospitalised	multiple domains	104	13	175	7.4	(4.0, 12.4)	
Omu et al	social media	single domain	91	113	152	74.3	(68.6, 81.1)	-
Osmanov et al	hospitalised	multiple domains	270	126	619	24.3	(20.6, 28.2)	
PHOSP-COVID study Peghin et al	hospitalised outpatients	multiple domains multiple domains	174	797 241	861 599	92.6	(90.6, 94.2)	
Peluso et al	social media	single domain	112	89	143	62.2	(53.8, 70.2)	
Petersen et al	community	multiple domains	125	95	180	52.8	(45.2, 60.2)	-
Du et al	hospitalised	single domain	105	176	540	32.6	(28.7, 36.7)	
Radtke et al	community	multiple domains	84	4	109	3.7	(1.0, 9.1)	
Rass et al Restra-Ayora et al	hospitalised healthcare workers	single domain single domain	90	82 52	135	60.7 26.7	(52.0, 69.0)	-
Restra-Ayora et al Righi et al	healthcare workers hospitalised	multiple domains	84	83	421	19.7	(16.0, 23.8)	
Romero-Duarte et al	hospitalised	multiple domains	196	509	797	63.9	(60.4, 67.2)	-
Sathyamurthy et al	hospitalised	multiple domains	104	66	279	23.7	(18.8, 29.1)	(0.000)
SeeSie et al	outpatients	multiple domains	147	107	146	73.3	(65.3, 80.3)	
Shang et all Sofrid et al	hospitalised hospitalised	multiple domains multiple domains	196 222	441 305	796	55.4 93.3	(51.9, 58.9) (90.0, 95.7)	
Signid et al Simani et al	hospitalised	multiple domains single domain	197	12	120	10.0	(90.0, 95.7)	
Skala et al	outpatients	multiple domains	91	56	102	54.9	(44.7, 64.8)	-
Skjorten et al	hospitalised	single domain	105	59	126	46.8	(37.9, 55.9)	•_
Scraas et al Stavem et al	community	multiple domains multiple domains	126	380 185	676 451	56.2 41.0	(52.4, 60.0) (36.4, 45.7)	
Stavem et al Staphenson et al	community	multiple domains multiple domains	104	2038	3065	66.5	(36.4, 45.7)	
Sudre et al	community	multiple domains	84	108	4182	2.6	(2.1, 3.1)	_
Sykes et al	hospitalised	single domain	127	75	127	59.1	(50.0, 67.7)	_=
Taboada et al	hospitalised	multiple domains	189	87	183	47.5	(40.1, 55.0)	-
Farsitani et al Fawfik et al	hospitalised healthcare workers	single domain single domain	105 91	34 40	115	29.6 33.3	(21.4, 38.8)	20
Taylor et al	hospitalised	single domain	126	261	545	47.9	(43.6, 52.2)	
Tempany et al	healthcare workers	multiple domains	84	116	217	53.5	(46.6, 60.2)	-
Tleyjeh et al	hospitalised	multiple domains	136	125	222	56.3	(49.5, 62.9)	. =
Toott et al Townsend et al	hospitalised outpatients	single domain multiple domains	105	96 74	239 128	40.2 57.8	(33.9, 46.7)	
Trunfio et al	hospitalised	multiple domains multiple domains	194	41	128	24.4	(48.8, 66.5) (18.1, 31.6)	0.75
Ursini et al	social media	single domain	182	270	616	43.8	(39.9, 47.9)	
Venturelli et al	outpatients	multiple domains	105	394	767	51.4	(47.8, 55.0)	•
Walle-Hansen et al	hospitalised	single domain	200	57	105	53.8	(43.8, 63.5)	-
Weng et al Whitaker et al	hospitalised community	single domain multiple domains	104	52 28713	117 76155	44.4 37.7	(35.3, 53.9) - (37.4, 38.0)	-
Anntaker et al	healthcare workers	single domain	167	114	162	70.4	(62.7, 77.3)	
Gong et al	hospitalised	single domain	111	267	538	49.6	(45.3, 53.9)	• -
Yan et al	hospitalised	multiple domains	98	0	125	0.0	(0.0, 2.9)	
Zayet et al	hospitalised	multiple domains	289	127	354	35.9	(30.9, 41.1)	ŀ
Chan et al Chang et al	hospitalised hospitalised	multiple domains multiple domains	348 378	36 1095	121	29.8 45.0	(21.8, 38.7)	
own good on the			5/0	1000	6433	43.9	(8.2, 87.2)	⊼
						5500	Societics	V
							022	
		single domain	150 126	1091	11800	9.2	(8.7, 9.8)	
N-Aly et al (hospitalized)	hospitalised		1.05		2839	44.2	(2.7, 3.0)	
Al-Aly et al (hospitalized) Al-Aly et al (non-hospitalized)	community outpatients	single domain multiple domains	243					
Al-Aly et al (hospitalized) Al-Aly et al (non-hospitalized) Aminian et al	community	single domain multiple domains single domain	243 154	1255 6085	28335	21.5	(21.0, 22.0)	
N-Aly et al (hospitalized) N-Aly et al (non-hospitalized) Aminion et al Nyoushihami et al Caspersen et al	community outpatients hospitalised community	multiple domains single domain single domain	154 350	6085 28	28335 170	21.5 16.5	(21.0, 22.0) (11.2, 22.9) (11.2, 22.9)	
Al-Aly et al (hospitalized) Al-Aly et al (non-hospitalized) Aminian et al Ayubükhani et al Caspersen et al Caspers al	community outpatients hospitalised community hospitalised	multiple domains single domain single domain single domain	154 350 128	6085 28 721	28335 170 6619	21.5 16.5 10.9	(21.0, 22.0) (11.2, 22.9) (10.2, 11.7)	
A4-Ay et al (hospitalized) A4-Ay et al (hospitalized) A4-Ay et al (hospitalized) Ayroubihami et al Caspersen et al Caspersen et al Caspersen et al Caspersen et al	community outpatients hospitalised community hospitalised community	multiple domains single domain single domain single domain multiple domains	154 350 128 91	6085 28 721 13396	28335 170 6619 145184	21.5 16.5 10.9 9.2	(21.0, 22.0) (11.2, 22.9) (10.2, 11.7) (9.1, 9.4)	
contomnes from health record linkage AA-by et al (hospitalized) AA-by et al (non-hospitalized) Annision et al Anysubhanis et al Casprone et al Casprone et al Casprone et al Casprone et al (anutha) Rossister et al (anutha) Rossister et al (anutha)	community outpatients hospitalised community hospitalised	multiple domains single domain single domain single domain	154 350 128	6085 28 721 13396 734	28335 170 6619	21.5 16.5 10.9	(21.0, 22.0) (11.2, 22.9) (10.2, 11.7)	ľ

Figure 4. Forest plot of prevalence of Long COVID in the included studies by method of outcome assessment, with 95% prediction intervals.

Comparison to Controls

Twenty-four of the 130 publications included comparison to at least 1 group of controls (Supplementary Figure 45). The majority of studies used test-negative controls (antigen and antibody, with some matching), but others used untested controls. In community-based studies with controls, the relative risk ranged between 1.0 and 51.4 (pooled relative risk, 2.7; 95% PI, 0.2-39.4) and the absolute risk difference ranged between -1% and 35% (pooled risk difference, 10.1%; 95% PI, -12.7% to 32.8%) (Supplementary Figures 46 and 47). In community-based samples with controls and assessed as having a low risk of bias (n = 4), the pooled relative risk of experiencing symptoms/ill health after COVID-19 was 1.33 compared to controls (95% PI, 1.30. to 1.36; $I^2 = 28.1\%$) (Figure 5) and the absolute risk difference between cases and controls ranged between 1% and 9% (Supplementary Figure 48). There was no evidence of small-study effects such as publication bias (Supplementary Figure 49).

DISCUSSION

This systematic review—which included 120 studies assessing Long COVID symptoms, functional status, or pathology published up to November 2021—demonstrates substantial between-study heterogeneity and wide variation in prevalence estimates. This is due to differences in sources of study samples (community, outpatient clinic, occupational, hospitalized) and number of assessed symptoms and method of assessment (selfreported individual or collective symptoms, healthcare records, clinical investigations at follow up). The only PE with low between-study heterogeneity was a 33% (95% PI, 30%-36%) excess risk of experiencing prolonged symptoms in COVID-19 cases compared to controls in community-based studies with low risk of bias. Although studies that included controls showed, on the whole, lower net prevalence of Long COVID than studies that did not, the evidence from most of these studies is that COVID-19 is associated with a substantially higher risk of being ill 12 weeks after infection than those not infected.

In characterizing Long COVID, the review demonstrated higher prevalence estimates in study samples where a substantial proportion of included individuals were hospitalized during the acute phase of the infection and/or had severe acute disease. It is difficult to comment on prevalence difference by ethnicity, deprivation, or gender because although we conducted subgroup analyses by proportion of participants by gender or ethnicity in included studies, the difference between the prediction estimates may be related to other confounding factors, such as, for example, studies that included more males may indicate that they also include a high proportion of those who had

study	region	type of control	mean days since infection	relative risk (95% CI)				
ONS study August 2021	Europe	negative test result	84	1.47 (1.30, 1.66)		_	-	
Radtke et al	Europe	negative test result	84	1.63 (0.58, 4.57)	•			•
Roessler et al (adults)	Europe	no positive test result	91	1.33 (1.32, 1.35)		ė		
Roessler et al (children)	Europe	no positive test result	91	1.30 (1.25, 1.35)				
95% Prediction Interval				1.33 (1.30, 1.36)		-	_	
					1.0	1.2	1.5 risk (95% CI)	2.0

Figure 5. Forest plot of risk of Long COVID in included studies with community-based samples and controls assessed as having low risk of bias, with 95% prediction intervals.

severe acute illness [145]. Many studies did not report ethnicity or deprivation. These factors will be important to include in future studies if a comprehensive understanding of Long COVID and inequity is to be gained.

Long COVID's proposed pathophysiological mechanisms are multiple and potentially overlapping including persisting viral reservoirs, immune dysfunction, microclotting, and end-organ damage [146]. It is concerning that studies that specifically investigated for pathology tend to report higher prevalence estimates than those depending on healthcare records or even selfreporting of symptoms. The review found that Long COVID presents a significant burden of functional disability, symptoms, and pathology, with a pooled estimate of 34.5% of people not returning to full health/fitness after at least 12 weeks, and estimates of the most common symptoms/pathology including lung pathology (38.9%), fatigue (34.5%), breathing problems (14.9%), sleep problems (13.2%), and tingling or itching (11.3%). The paucity of long-term longitudinal studies after individuals' disease progression means it is difficult to comment on which symptoms are most persistent over time.

The UK's ONS produces population-level Long COVID prevalence estimates where the denominator is the whole population in the specific reported population group, for example, by age, sex, or occupation [147]. These fall out of our inclusion criteria. The ONS also produced prevalence estimates based on following up with those with confirmed SARS-CoV-2 infection, and we used the most recent estimate within the review's search period [9]. This study used multiple approaches including assessing individual symptoms compared to controls and asking participants whether they believe they have Long COVID. The latter approach, in the absence of a standardized method of assessment, may realistically be the best way to assess the presence of Long COVID because most people will take the combination of their symptoms, duration, fluctuation, effect on functional ability, and change from pre-COVID-19 health to shape their responses.

The lack of consensus on the precise definition of Long COVID plays an important part in the wide differences in

prevalence assessments; however, we found that the way the question is specifically asked and the source of retrieved clinical information at follow up are likely to play a crucial role. The ONS study is an example of how different methods of assessment at time of follow up can produce substantially different Long COVID estimates [9]. This was illustrated by our analysis in which studies that asked about multiple symptoms/domains tended to report higher prevalence estimates than single-domain studies. Our analysis indicated higher prevalence estimates with longer follow-up time, although we recognize these were mostly not within-study comparisons. However, in 4 of 10 longitudinal studies, prevalence was higher at the time of the second follow up. These results could be explained by several factors, eg, by the episodic nature of Long COVID, whereby in the early stages people may believe they have recovered from their illness, but with passing time and phases of relapse and remittance, people may be more cautious about reporting they have recovered. People may also be developing new symptoms over time, or perhaps there is more study drop-out by people who believe they have recovered. Overall, however, the results indicate that, over time, prevalence does not substantially reduce.

Studies that used questionnaires/surveys to ask participants about their symptoms, health status, or quality of life tend to report higher prevalence estimates than those that recorded symptoms from healthcare records' clinical coding. This is manifested in the prevalence from Al-Aly et al [16] studies being on the lower side in our analysis because we only included those with symptoms rather than recorded post-COVID-19 pathology, and such symptoms are expected to be severe enough to prompt seeking medical help and being recorded in medical notes. Studies that had dedicated pathology follow up and discovery of COVID-19 patients tended to report the highest prevalence. This is possibly because, in addition to pathology that leads to recognizable signs and symptoms, specific medical investigations as part of the research protocol can pick up latent pathology that may not be accompanied by clinical manifestations.

Studies such as Al-Aly et al [16] that investigated medical diagnoses in the period after COVID-19, report cardiovascular, neurological, and other system-specific clinical sequelae, providing a substantial excess burden in those who survived the acute phase of COVID-19 [13]. However, there is no agreement yet as to whether these outcomes are classified as Long COVID. They are generally not recorded by symptom studies, and the WHO does not yet specifically include such outcomes within its clinical case definition of Post-COVID-19 Condition (also known as Long COVID) [1]. A specific pathology diagnosed after COVID-19 could have been triggered by the infection, but identification as such will depend on the extent of clinical investigations identifying and labeling specific pathology as opposed to differences in the disease manifestation themselves.

Other sources of heterogeneity between studies include study design with some including assessment at 1 point in time, whereas others were longitudinal where assessment of COVID-19 status was conducted before the development of Long COVID. This assessment itself varied in terms of using PCR or antigen testing or self-reporting of history of acute infection.

Ideally, excess absolute risk in comparison to controls is a good measure to estimate the burden of Long COVID. This is likely dependent on the approach to control selection, whether based on self-report of absence of infection history or laboratory results that are not accurate enough to ascertain the state of previous infection (antigen or antibody) and timing of assessment given the predominant episodic nature of Long COVID.

Few studies had a low risk of bias, which suggests there is a gap in the evidence base for strong studies of Long COVID prevalence. In terms of causal inference, many studies were liable to potential collider bias, which presented as selection bias caused by restricting analyses to people who were hospitalized, self-selected for PCR, or lateral flow tests based on symptoms, or simply volunteered their study participation [148]. Similarly, our exploration of potential sources of heterogeneity may be prone to table 2 fallacy in the original studies, where these subgroups do not derive from the focal research question, so these should be interpreted descriptively rather than causally [149].

The strengths of our review include comprehensive electronic searching for relevant studies and comprehensive assessment of risk of bias, data extraction, and checking with each of these processes being done independently by 2 authors. We also adapted the Newcastle-Ottawa scale (Supplementary Table 3) for this prevalence systematic review, which can be used by other researchers for risk assessment and/or to build high-quality study designs. The quality assessment criteria and process were discussed within the study team, which includes 2 authors with lived experience of Long COVID.

Our review was limited by the substantial between-study heterogeneity. We used the most common reported symptom

estimate for studies and did not combine multiple individual symptoms into 1 overall estimate of prevalence of Long COVID. The symptom with the highest prevalence differed from study to study, so this may not be entirely comparable. We did not include more recent studies that assessed the prevalence of Long COVID after infection with different variants of SARS-CoV-2 and/or in double- or triple-vaccinated populations. Recent estimates point to a prevalence of 4%–5% of reporting Long COVID at 12 to 16 weeks after first confirmed SARS-CoV-2 infection depending on variant, with no evidence of difference between variants among those who are triple vaccinated when infected [150]. In those double-vaccinated group, the prevalence of persistent symptoms was approximately 10% compared to 15% of unvaccinated controls [151].

We extracted estimates of "new-onset" Long COVID/ symptoms where possible. In instances in which the proportion is of a symptom-like fatigue, for example, we picked the one quoted as new-onset fatigue if available, or we downgraded quality because it was not possible to ascertain that the symptom is "new" after infection. Because Long COVID is a novel condition, prevalence of the condition is considered equivalent to cumulative incidence. When comparing with controls, we estimated cumulative incidence from reported absolute risk, when appropriate. When reporting risk ratio, we included incidence rate ratio and hazard ratios, but we did not consider the odds ratio an adequate approximation because of the high potential prevalence in some populations.

CONCLUSIONS

We know that significant numbers of people experience ill health after SARS-CoV-2 infection. Long COVID has an impact on society, particularly in places with continuing waves of infection. By reviewing how different research approaches attempted to quantify the population burden of Long COVID, our findings provide insight into how to get more accurate estimates of prevalence and severity. With quantification of prevalence and the associated inequity, we can understand the investment needed for prevention, diagnosis, and treatment as well as the policy decisions needed to resource healthcare and social care services both adequately and equitably, and to mitigate the wider social and economic impact of Long COVID.

Supplementary Data

Supplementary materials are available at *Open Forum Infectious Diseases* online. Consisting of data provided by the authors to benefit the reader, the posted materials are not copyedited and are the sole responsibility of the authors, so questions or comments should be addressed to the corresponding author.

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Author contributions. NAA, DCG, RT, AA, VL, and MW conceptualized and designed the study. MW drafted the protocol and search strategy with input from all coauthors. VL conducted the search. All authors contributed to screening the articles. MW, DCG, NZ, RT, and CC extracted and assessed the data for quality. NAA, MW, DCG, NZ, and CC contributed to the process of checking and verifying the extracted data. DCG planned and conducted the statistical analyses and produced the forest plots. MW, DCG, NZ, and NAA interpreted the data and drafted the manuscript. All authors reviewed the final manuscript. All authors had full access to all the data in the study and had final responsibility for the decision to submit for publication.

Disclaimer. The views and opinions expressed in this review are those of the authors and do not necessarily reflect those of the National Institute for Health Research (NIHR), the Department of Health and Social Care, or the United Kingdom (UK) government's official policies. For the purpose of open access, the author has applied a CC BY public copyright licence to any Author Accepted Manuscript version arising from this submission.

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Potential conflicts of interest. DCG is a coinvestigator on the NIHR-funded LOCOMOTION study. NAA has lived experience of Long COVID, is a coinvestigator on the NIHR-funded STIMULATE-ICP and HI-COVE studies, has contributed in an advisory capacity to World Health Organization (WHO) and the European Union Commission's Expert Panel on effective ways of investing in health meetings in relation to post-COVID-19 condition, and has acted as a collaborator on some of the UK's Office for National Statistics outputs on the prevalence of Long COVID. AA has lived experience of Long COVID, is a co-founder of the Patient-Led Research Collaborative, and has contributed in an advisory capacity to National Institutes of Health, Centers for Disease Control and Prevention, and WHO. All authors: No reported conflicts of interest.

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