



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

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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.

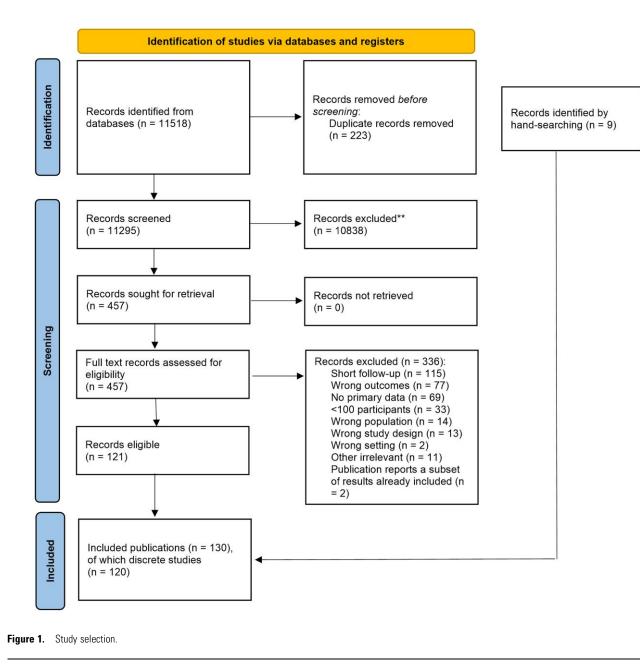
Table 1 summarizes the included studies' key characteristics and primary outcome for the first follow-up. Study design was reported as described by each study or designated based on study description if not explicitly stated. Most studies were in adults and included patients who were hospitalized in 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



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 = 11) (Figure 3 and Supplementary Figures 7–40).

	Author	Country	Study Design (as Described by Study, * If Not Stated)	Denominator ^a	Controls N, Type	Setting	Age (Years) Mean/SD Median (IOR) F	% Female	COVID-19 Diagnostic Method	Severity	Follow-up Time Days	Finding: %With at Least 1 Symptom o or Pathology Remaining at Follow up
<u>-</u>	Abdelrahman et al [15]	Egypt	Prospective cohort	172	÷	Hospitalized patients and nonhospitalized	41.8/17.6	65.7	"Tested positive"	12.8% hospitalized (including 4% ICU)	240–300 (range) after "improvement of acute COVID-19"	61.0%
ci	Al-Aly et al [16]	USA	Cohort with controls	60 255 4	4 526 737 without COVID-19 and not hospitalized	Nonhospitalized	61 (4872) 12.1		"Positive test"	÷	126 ^c	2.9%
2a.	Al-Aly et al [16]	USA	Cohort with controls	11 800 1	11 868 hospitalized with seasonal influenza	Hospitalized patients	70 (61– 5 76)	5.8	PCR confirmed	26.3% ICU	150°	9.2%
က်	Aminian et al [18]	NSA	Retrospective	2839	:	Hospitalized patients	52.7/20.1 5	52.3 F	PCR confirmed	ICU excluded	243°	44.2%
4	Arnold et al [144]	ž	Prospective cohort	110	:	Hospitalized patients	60 (46- 4 73)	44.0 F	PCR confirmed or clinico- radiological	Mixed	°00	73.6%
வ	Augustin et al [20]	Germany	Longitudinal prospective cohort	442	÷	Nonhospitalized patients	43 (31– 5 54)	52.3 F	PCR confirmed	97.5% mild	131 ^c	27.8%
ю́	Ayoubkhani et al [21]	Ъ.	Observational retrospective matched cohort (with controls)	47 780 4	47 780 matched for Hospitalized age, sex patients	Hospitalized patients	64.5/19.2 45.1		Laboratory confirmed or clinical diagnosis	9.9% ICU	140 ^e	21.5
7.	Baricich et al [22]	ltaly	Cross-sectional	204	:	Hospitalized patients	57.9/12.8 4	40.0	"Confirmed diagnosis"	13% ICU	124.7 ^e	32.4%
œ	Becker et al [23]	USA	Cross-sectional	740	÷	Hospitalized patients, outpatients and ER attendees	49 (38– 6 59)	63.0	Tested positive or antibody positive	:	228 ^b	24.1%
ல்	Bellan et al [24]	ltaly	Prospective cohort	238	:	Hospitalized patients	61 (50- 4 71)	40.3 F	PCR confirmed bronchial swab, serological testing, or suggestive CT	27.7% did not require oxygen 11.8% ICU	91–121 ^e	53.8%
10.	Blanco et al [25]	Spain	Prospective	100	:	Hospitalized patients	54.9/10.3 36.0		PCR confirmed	47% severe	104 ^c	52.0%
11.	Bliddal et al [26]	Denmark	Cohort	129	:	Nonhospitalized patients	44.8 (13.6)	70.0 F	PCR confirmed	Nonhospitalized	90 ^b	40.3%

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

Finding: %Writh at Least 1 Symptom or Pathology Remaining at Follow up	60.6%	53.0%	66.2%	77.1%	16.5%	10.9%	28.6%	14.8%	80.0%	49.5%	79.3%	53.4%	37.2%
Follow-up Time Days	152–213 (range) after illness	365 ^b	3 months posthospital discharge	6 months posthospital admission	334–365 (range) after infection	91–150 days posthospital admission	370 ^d	90 ^b	107 ^f	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	60.9	69.1	53.0	58.0	47.0	51.0	÷	43.0	27.7	53.1	62.1	46.2
Age (Years) Mean/SD Median (IOR)	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 a adult patients	vith	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	30 193 hospitalized COVID-19 negative patients	* * 	÷	÷	÷	÷	÷	95 randomly selected from non-COVID patients attending the ward
b Denominator ^a	312	304	165	118	774	5571	546 I	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 longitudinal	Prospective cohort	Cohort*	Retrospective cohort	Prospective cohort	Cohort
Country	Norway	Italy	Spain	Italy	Norway	NSA	China	NSA	ltaly	Italy	NSA	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.	14.	15.	16.	17.	18.	19.	20.	21.	22.	23.	24.

Finding:	% writh 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 ^f	176 ^f	210 ^e	210 ^e	340 ^e	3 months postsymptom onset	103°	6 months posthospital discharge	4 months postinfection	180 ^b	117.5°	3 months posthospital discharge	175 ^b	122 ^b	186°
	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% ICII
	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 (IOR)	58.0/12.6	57.9/13	61/17	61/17	61/16	59 (50– 68)	60 (53– 68)	18+	÷	64 (54– 76)	49.5/15.3 56.0	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	÷	÷	:	:	:	÷	:	÷	÷	:	Norwegian general Community population norms	÷	:	1072 seronegative	:
	Denominator ^a	804	1077	1142	1142	1950	137	107	199	138	116	447 N	130	114	323 1	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 longitudinal	Prospective Iongitudinal	Cohort with controls	Ambidirectional cohort
	Country	Ϋ́	Ϋ́	Spain	Spain	Spain	France	Belgium	China	Х	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. 99.	39.

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Finding: %With at Least 1 Symptom or Pathology Remaining at Follow s up	68.0%	66.9%	41.7% pital je or	65.7%	93.7%-more than 61.9% 3 months postinfection	ange) 59.9% Iptom	37.3%	85.9%	28.7% pital Je	64.7%	30.0%	36.4% hths sction	20.1%	s 56.9% tpital
Follow-up Time Days	185°	119.3°	3 months posthospital discharge or visit	195°	93.7%-more th 3 months postinfection		е 395 ^f	90 ^f	6 months posthospital discharge	111 ^c	169°	6.1 ± 1.1 months postinfection	83q	12 months posthospital
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
b 5D n Female	47.0	43.3/14.4 46.6	30.6	. 69.7	49.8/16.9 59.2	43.6/17.4 48.8	. 80.5	47.5 (36- 48.8 57)	53.3	.7 39.9	57.1	41.4/12.3 63.6	53.4	.5 49.3
Age (Years) Mean/SD Median (IOR)	59 (49– 67)	43.3/14	18–65	31 (24- 47)	49.8/16	43.6/17	39 (33– 48)	47.5 (3(57)	68 (66– 74)	54.5/16.7	48/15.2	41.4/12	55 (44– 63)	65.1/17.5
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
Controls N, Type	3383 community dwelling without SARS-CoV-2 infection, 1164 matched pairs	÷	÷	÷	:	÷	÷	÷	466 uninfected spouses who lived together	÷	21, "healthy controls recruited via email and flyer advertisements"	÷	:	÷
s Denominator ^a	1227	118	242	006	365	289	303	142	1301	153	177	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	USA	Pakistan	S Korea	Germany	China	China	China	China	NSA	USA	ltaly	China (HK)	7] Spain
Author	Huang et al [5 4] (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]
	40.	41.	42.	43.	44.	45.	46.	47.	48.	49.	50.	51.	52.	53.

	Author	Country	Study Design (as Described by Study,* If Not Stated)	Denominator ^a	Controls N, Type	Setting	Age (Years) Mean/SD Median (IOR)	% Female	COVID-19 Diagnostic Method	Severity	Follow-up Time Days	Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up
54.	Martinez et al [68]	Switzerland	Retrospective cohort	260	:	Healthcare workers	Mean range 30–39	75.4	'Positive test'	1.2% hospitalized	168°	26.5%
55.	Matteudi et al [69]	France	Prospective cohort	137	:	Hospitalized patients and outpatients, pediatric	9.3 (n/a)	:	PCR confirmed	27% asymptomatic	180 ^b	16.8%
56.	Mazza et al [70]	Italy	Prospective cohort	226	:	Hospitalized patients and ER attendees	58.5/12.8 34.1	34.1	PCR confirmed	78% hospitalized	90.1 ^e	35.8%
57.	Mechi et al [71]	Iraq	Single-center cross-sectional	112	:	Hospitalized patients and nonhospitalized	50.6/13.4	34.0	Laboratory confirmed	46.4% hospitalized	9 months after acute infection	82.1%
58.	Mei et al [72] (†)	China	Cohort*	4328	1500, random sample of general population	Hospitalized patients	59 (47– 68)	54.1	Met relevant clinical criteria	Not defined	144 ^f	14.2%
59.	Mei et al [73] (†)	China	Prospective cohort	3677	:	Hospitalized patients	59 (47– 68)	55.5	PCR confirmed	33.7% severe, 2.6% critical	144 ^f	26.5%
60.	Menges et al [74]	Switzerland	Switzerland Population-based prospective cohort	431	:	Community	47 (33– 58)	49.7	PCR confirmed	10.7% asymptomatic, 38.1% severe/very severe	220 ^c	24.6%
61.	Milanese et al [75]	ltaly	Prospective cohort	135	:	Hospitalized patients	59/11	33.0	Not stated	Moderate and severe	182 ^e	47.4%
62.	Millet et al [76]	NSA	Prospective cohort	173	:	Hospitalized patients and outpatients	51.5/n/a	50.6	PCR confirmed	:	12 months postdiagnosis	48.0%
63.	Mohiuddin Chowdhury et al [77]	Bangladesh Prospective multicent cross-sec	Prospective multicenter cross-sectional	313	:	Hospitalized patients and outpatients	37.7/13.7 19.8	19.8	PCR confirmed	Not critically ill (ICU/ HDU)	140 ⁹	21.4%
64.	Munblit et al [78]	Russia	Longitudinal cohort	2649	:	Hospitalized patients	56 (46– 66)	51.1	PCR confirmed and 2.6% severe clinically diagnosed	1 2.6% severe	218 ^f	57.9%
65.	Nabahati et al [79]	Iran	Prospective cross-sectional	173	:	Hospitalized patients	53.6/13.7	67.1	PCR confirmed	54% severe	90 ^e	52.0%
.99	Nehme et al [80]	Switzerland	Prospective cohort	410	:	Outpatients	42.7/12.9	67.1	PCR confirmed	Mild and moderate	7–9 months postdiagnosis	39.0%
67.	Nguyen et al [81]	France	Cohort*	125	:	Hospitalized	36 (27– 48))	55.0	PCR confirmed	Nonsevere	210 ^b	24.0%
68.	Nunez-Fernandez et al [82]	Spain	Prospective cohort	200	:	Hospitalized patients	62 (n/a)	40.5	PCR confirmed	15.5% ICU	84 ^e	29.0%
69.	O'Keefe et al [83]	USA	Cross-sectional	198		Outpatients	45/14	74.2	PCR confirmed	29.7% moderate, 1.1% severe	119 ^c	39.9%
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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%	60.7%	26.7%	19.7%	9.2%
Follow-up Time Days	12 weeks postinfection	°06	At least 3 months 74.3% postinfection	256 ^f	191°	4 months posttest or first symptoms	125 ^b	3 months posthospital discharge	3 months posthospital discharge	84 ^b	406	6 months postpositive test	84 ^b	op4 ح
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% 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 (IOR)	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	:	:	:	:	:	:	:	:	:	1246 seronegative	÷	125 healthcare workers with negative PCR	:	÷
Denominator	21 374	175	152	518	599	143	180	647	540	109 1	135	195 1	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	N	Singapore	Italy	Russia	Italy	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 [87]	Peluso et al [88]	Petersen et al [89]	Qin et al [90]	Qu et al [91]	Radtke et al [92]	Rass et al [93]	Riestra-Ayora et al [94]	Righi et al [95]	Roessler et al [96] <i>Split</i> <i>cohort (Adults)</i>
	70.	71.	72.	73.	74.	75.	76.	77.	78.	79.	80.	81.	82.	83.

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Table

	Author	Country	Study Design (as Described by Study.* if Not Stated)	Denominator ^a	Controls N, Type	Setting	Age (Years) Mean/SD Median (IQR)	% Female	COVID-19 Diagnostic Method	Severity	Follow-up Time Days	wwith at Least 1 Symptom or Pathology Remaining at Follow up
83a.	 Roessler et al [96] Split cohort (Children) 	Germany	Matched cohort	11 950	:	Community, children	:	48.1	Laboratory confirmed	1% hospitalized, 0.4% ICU	>06 ⁶	6.1%
84.	Romero-Duarte et al [97]	Spain	Retrospective longitudinal observational follow-up	797	÷	Hospitalized patients	63/14.4	46.3	PCR confirmed	10.8% ICU	6 months posthospital discharge	63.9%
85.	Sathyamurthy et al [98]	India	Single-center prospective cohort	279	÷	Hospitalized older adult patients	71.0/5.6	36.2	PCR confirmed	41.6% severe to critical	- 00 _e	23.7%
86.	Seeβle et al [99]	Germany	Prospective cohort	146	÷	Hospitalized and outpatients	57 (50– 63)	57.0	PCR confirmed	15.6% mild, 55.2% moderate, 25.0% severe, 4.2% critical	140–154 (range) after symptom onset	73.3%
87.	Shang et al [100]	China	Cohort	796	÷	Hospitalized patients	62 (51– 69)	49.2	PCR confirmed	90.8% severe, 9.2% critical	6 months posthospital discharge	55.4%
88.	Sibila et al [101]	Spain	Prospective cohort	172	÷	Hospitalized patients	56.1/19.8	43.0	Not stated	moderate and severe 43% ICU	101.5 ^e	57.0%
80.	Sigfrid et al [102]	ž	Prospective cohort	327	÷	Hospitalized patients	59.7 (51.7– 67.7)	41.3	PCR confirmed or "clinically diagnosed highly suspected"	20.8% no O ₂ , 36.1% supplemental O ₂ , 15.0% noninvasive O ₂ , 28.1% mechanical ventilation	222° al	93.3%
90.	Simani et al [103]	Iran	Cohort*	120	÷	Hospitalized patients	54.6/16.9	33.3	Spiral chest CT scan or PCR confirmed	7.5% ICU	183 ^e	10.0%
91.	Skala et al [104]	Czech Republic	Prospective cohort	102	÷	Hospitalized patients and outpatients	46.7/n/a	53.9	PCR confirmed	14.7% hospitalized	3 months after testing positive	54.9%
92.	Skjorten et al [105]	Norway	Multicenter prospective cohort	126	:	Hospitalized patients	56.2/12.7 38.5	38.5	" Discharge diagnosis of COVID-19"	20% ICU	104 ^f	46.8%
93.	Sonnweber et al [106]	Austria	Prospective observational	145	:	Hospitalized and outpatients	57/14	43.0	PCR confirmed	22% ICU	103 ^b	54.9%
94.	Soraas et al [107] (π)	Norway	Cohort	651 57 (5712 SARS- CoV-2-negative + 3342 randomly selected untested	Community	48.6/13.6	57	PCR confirmed	Nonhospitalized, mild	258 ⁵	51.9%
95.	Soraas et al [108] (π)	Norway	Prospective cohort	672 60 (6006 SARS- COV-2-negative patients	Community	48.5/13.5 56.8	56.8	PCR confirmed	Nonhospitalized	126 ^b	56.2%
96		Norway	Cross-sectional	451	:	Community survey 49.7/15.2	49.7/15.2	56.0	PCR confirmed	:	117 ^c	41.0%
97.	Stavem et al [110] (=)	Norway	Cross-sectional mixed-mode	458	:	Community	49.5/15.3 56.0	56.0	PCR confirmed	:	117.5 ^c	46.0%

Finding: %With at Least 1 Symptom o Pathology Remaining at Follow up	66.5%	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 ^c	84 ^b	113 ^f	6 months posthospitali zation	180 ^b	^d O6	3 months posthospital discharge	At least 3 months 33.3% postpositive test	16 weeks posthospital discharge	At least 12 weeks postpositive test	113 ^f
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	:	:	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	57.0	34.3	40.5	55.6	55.6	46.0	58.0	38.2	80.0	42.1
Age (Years) Mean/SD Median (IQR)	11-17	46.0/15.8 57.0	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	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	÷	÷	÷	:	÷
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	Я	UK, USA and Sweden	NK	Spain	Primarily USA	USA	ltaly	Egypt	UK	Republic of Ireland	France
Author	Stephenson et al [111]	Sudre et al [112]	Sykes et al [113]	101. Taboada et al [114]	102. Taquet et al [116] (众)	103. Taquet et al [115] (◊)	Tarsitani et al [117]	105. Tawfik et al [118]	Taylor et al [119]	107. Tempany et al [120]	108. The Writing Committee for the COMEBAC Study Group [121]
	98.	.66	100.	101.	102.	103.	104.	105.	106.	107.	108.

Finding: %With at Least 1 Symptom o Pathology Remaining Follow-up Time at Follow Days up	months after 1.8% discharge (hospitalized), 4 months postsymptom (nonhospitali cad)	56.3%	s 40.2% ispital rge	s 5.0% spital rge	57.8%	24.4%	nths 43.8% isitive	51.4%	53.8%	44.4%	37.7%	70.4%	49.6%
Follow-L	3 months after discharge (hospitalized) 4 months postsymptor onset (honhospitali zed)	122 ^f	3 months posthospital discharge	5% 3 months posthospital discharge	4 72 [†]	194°	 6 ± 3 months postpositive test 	105°	186 ^f	89.5°	84 ^b	ICU 153 ^f	97 ^f
Severity	r 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%
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 "
0 Female	5 51.0	0 23.0	9 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 (IOR)	52.9/15.5 d	52.5/14.0 23.0	53.6/14.9 40.2	55.5/6.2	49.5/15 d	56 (43– 69)	45/12	63/13.6 hd	ır 74.3/n/a	÷	-18+	36 (31– 43)	52 (41-
Setting	Hospitalized patients and nonhospitalized	Hospitalized patients	Hospitalized patients	Hospitalized and outpatients t	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
a Controls N, Type	:	÷	:	100 randomly recruited from hospital registration system without COVID-19	÷	:	:	:	÷	:	:	:	184, volunteers
Denominator ^a	683	222	239	100	128	168	616	767	106	117	76 155	162	538
Study Design (as Described by Study,* If Not Stated)	Multicenter prospective cohort	Prospective cohort	Single-center cohort	Retrospective comparative study with controls	f Cross-sectional*	Cross-sectional	Cross-sectional	Cohort*	Cohort	Retrospective	Random community- based survey (REACT-2)	Ambidirectional cohort	Longitudinal with
Country	Norway	Saudi Arabia	Brazil	Egypt	Republic of Ireland	ltaly	Italy	Italy	0] Norway	China	¥,	China	China
Author	. Tholin et al [122] (a)	110. Tleyjeh et al [123]	111. Todt et al [124]	112. Tohamy et al [125]	113. Townsend et al [126]	Trunfio et al [127]	Ursini et al [128]	Venturelli et al [129]	117. Walle-Hansen et al [130] Norway	118. Weng et al [131]	119. Whitaker et al [132]	120. Xiong et al [133]	121. Xiong et al [1 34]
	109.	110.	111.	112.	113.	114.	115.	116.	117.	118.	119.	120.	121.

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Author	Country	Study Design (as Described by Study, * If Not Stated)	s Denominator ^a	Controls N, Type	Setting	Age (Years) Mean/SD Median (IQR) F	% Female	COVID-19 Diagnostic Method	Severity	Follow-up Time Days	Finding: %With at Least 1 Symptom or Pathology Remaining at Follow up
122. Yan et al [135]	China	Prospective observational	125	:	Mobile cabin hospital, adult males	35 (30- (49)	. 0.0	" Diagnosed with COVID-19"	asymptomatic/mild symptoms	84 ^e	0.0%
123. Yan et al [136]	China	Cohort	119	:	Hospitalized patients	53.0/12.2 59.0		PCR confirmed	24% severe	365°	39.5%
124. Yin et al [137]	China	Retrospective analysis	337	:	Hospitalized patients	53.5/14.8 49.5		PCR confirmed	12.8% severe, 3.6% ICU	203 ^b	55.8%
125. Zayet et al [138]	France	Retrospective cohort	354	:	Hospitalized patients and outpatients	49.6/18.7 63.0		PCR confirmed	34.2% hospitalized, 5% ICU	289 ^b	35.9%
126. Zhan et al [139]	China	Prospective cohort	121	:	Hospitalized patients	49 (40- 5 57)	58.7 P	PCR confirmed	15.7% severe	348°	29.8%
127. Zhang et al [140]	China	Retrospective comparative	122	÷	Hospitalized patients	51 (31.8– 5 61.0)	50.3 P	PCR confirmed	mild cases excluded, only patients with pulmonary sequelae at discharge included	92 [†]	54.9%
128. Zhang et al [141]	China	Cohort*	245	:	Hospitalized patients	43 (33- 2 54)	43.8 N	Vucleic acid testing	Nucleic acid testing 9.3% severe/critical	90 ^e	72.7%
129. Zhang et al [142]	China	Retrospective multicenter cohort	2433	:	Hospitalized patients	60 (49- 68)	50.5 L	Laboratory confirmed	27.9% severe	364 ^f	45.0%
130. Zhou et al [143]	China	Prospective cohort with controls	164 4.	42 healthy controls Hospitalized —negative patients nucleic acid and antibody tests	Hospitalized patients	:	56.9 P	PCR and antibody test	54.6% severe	129° (severe cases) 125° (mild)	69.5%
						.					

Abbreviations: COVID, coronavirus disease 2019; CT, computerised tomography ; ER, emergency room; HDU, high dependency unit ; ICD, intensive care department ; ICU, intensive care unit; IV, intravenous; IOR, interquartile range; NAAT, nucleic acid amplification test; NIV, noninvasive ventilation; PCR, polymerase chain reaction; PICU, paediatric 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: $\Omega_1 = \langle 0, \Psi, \uparrow, \infty, \pi$. * 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.

^aDifferent 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.

Median number of days posthospital discharge. ^eMean number of days posthospital discharge.

^gMean number of days postnegative test after infection.

Table 1. Continued

study	region	mean days since infection	cases	total	% persistent symptoms	
Abdelrahman et al Al-Aly et al (hospitalized)	Other N America	270 150	105 1091	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	150	1091	11800	9.2 (8.7, 9.8) 2.9 (2.7, 3.0)	
Aminian et al	N America	243	1255	2839	44.2 (42.4, 46.1)	
Arnold et al Augustin et al	Europe Europe	90 131	81 123	110 442	73.6 (64.4, 81.6) 27.8 (23.7, 32.3)	
Augustin et al Ayoubikhani et al	Europe	131	123 6085	442 28335	27.8 (23.7, 32.3) 21.5 (21.0, 22.0)	
Baricich et al	Europe	139	66	204	32.4 (26.0, 39.2)	
Becker et al	N America	231	178	740	24.1 (21.0, 27.3)	•
Bellan et al Blanco et al	Europe Europe	120	128 52	238	53.8 (47.2, 60.2) 52.0 (41.8, 62.1)	
Biddal 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 Europe	127 105	244 100	478 151	51.0 (46.5, 55.6) 66.2 (58.1, 73.7)	
Caruso et al	Europe	182	91	118	77.1 (68.5, 84.3)	
Caspersen et al	Europe	350	28	170	16.5 (11.2, 22.9)	+
Castro et al Chai et al	N America China	128	721	6619 546	10.9 (10.2, 11.7)	•
Charet al Cirulli et al	N America	90	156	122	28.6 (24.8, 32.6) 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)	
Diaz-Fuentes et al Domenech-Montoliu et al	N America Europe	84 213	88 258	111 483	79.3 (70.5, 86.4) 53.4 (48.9, 57.9)	
Erol et al	Other	170	45	121	37.2 (28.6, 46.4)	
Fernandez-de-Las-Penas et al	Europa	227	930	1142	81.4 (79.1, 83.7)	
Frija-Masson et al	Europe	91	103	137	75.2 (67.1, 82.2)	-
Froidure et al Fu et al	Europe China	95 196	73 20	107	68.2 (58.5, 76.9) 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)	
Sonzalez-Hermosillo et al	Other	105	119	130	91.5 (85.4, 95.7)	· · · · ·
Han et al Havervall et al	China Europe	175 122	71 69	114 323	62.3 (52.7, 71.2) 21.4 (17.0, 26.2)	
havervall et al	China	122	1265	1655	21.4 (17.0, 20.2) 76.4 (74.3, 78.5)	
Jacobson et al	N America	119	79	118	66.9 (57.7, 75.3)	
Kashifotal Kimetal	Other	105 195	101 591	242 900	41.7 (35.5, 48.2) 65.7 (62.5, 68.8)	•
Kim et al Lemhofer et al	Other Europe	195	591 226	900	65.7 (62.5, 68.8) 61.9 (56.7, 66.9)	
i 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)	•
Liaoet al	China	104	122 373	142 1301	85.9 (79.1, 91.2) 28.7 (26.2, 31.2)	
Liu et al Liyanage-Don et al	China N America	196	373	1301	28.7 (26.2, 31.2) 64.7 (56.6, 72.3)	1.000
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)	
Lui et al Maestre-Muniz et al	China Europe	96 379	41	204	20.1 (14.8, 26.3) 56.9 (52.6, 61.1)	
Maestre-Muniz et al 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)	-
Mazza et al	Europe	104	81	226	35.8 (29.6, 42.5)	-
Mechietal Meietal	Other	274	92 976	112 3677	82.1 (73.8, 88.7) 26.5 (25.1, 28.0)	
Menges et al	Europe	219	106	431	24.6 (20.6, 28.9)	2.1
Vilanoso et al	Europe	196	64	135	47.4 (38.8, 56.2)	4
Millet et al	N America	365	83	173	48.0 (40.3, 55.7)	÷.
Mohiuddin et al Munblit et al	Other Europe	161 232	67 1534	313 2649	21.4 (17.0, 26.4) 57.9 (56.0, 59.8)	
Nabahati et al	Other	104	90	173	52.0 (44.3, 59.7)	
Vehme 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 O'Keefe et al	Europe N America	98 119	58 79	200 198	29.0 (22.8, 35.8) 39.9 (33.0, 47.1)	-t)
Disaete et al DNS study August 2021	Europe	84	2501	21374	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)	•
Omu et al	Europe	91	113	152	74.3 (66.6, 81.1)	-
Osmanov et al PHOSP-COVID study	Europe	270 174	126 797	519 861	24.3 (20.6, 28.2) 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)	
Din et al Du ot al	China	105	87 176	647 540	13.4 (10.9, 16.3) 32.6 (28.7, 36.7)	100
au et al Radtke et al	Europe	84	1/0	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)	
Restra-Ayora et al	Europe	182	52	195	26.7 (20.6, 33.5)	-
Righi et al Roessker et al (adults)	Europe	84 91	83 13396	421 145184	19.7 (16.0, 23.8)	
Roessler et al (aduits) Roessler et al (children)	Europe	91	13396	145184	9.2 (9.1, 9.4) 6.1 (5.7, 6.6)	- 1
Romero-Duarte et al	Europe	196	509	797	63.9 (60.4, 67.2)	•
Sathyamurthy et al	Other	104	66	279	23.7 (18.8, 29.1)	•
SeefUe et al Shang et al	Europe China	147 196	107 441	146 796	73.3 (65.3, 80.3) 55.4 (51.9, 58.9)	
shang et al Sibila et al	Europe	196	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 Skjorten et al	Europe	91 105	56 59	102 126	54.9 (44.7, 64.8) 46.8 (37.9, 55.9)	
sigorten et al Sonnweber et al	Europe	105	73	126	46.8 (37.9, 55.9) 54.9 (46.0, 63.5)	
Soraas et al	Europe	126	380	676	56.2 (52.4, 60.0)	
Stavem et al	Europe	117	185	451	41.0 (36.4, 45.7)	+
Btephenson et al Sudre et al	Europe Other	104 84	2038 108	3065 4182	66.5 (64.8, 68.2) 2.6 (2.1, 3.1)	
Sucre et al	Europe	127	75	127	59.1 (50.0, 67.7)	-
laboada et al	Europe	189	87	183	47.5 (40.1, 55.0)	
faquet et al	N America	90	100007	273618	36.5 (36.4, 36.7)	
Tarsitani et al Tawfik et al	Europe Other	105 91	34 40	115 120	29.6 (21.4, 38.8) 33.3 (25.0, 42.5)	
faylor et al	Europe	126	261	545	47.9 (43.6, 52.2)	
Tempany et al	Europe	84	116	217	53.5 (46.6, 60.2)	
Fleyjeh et al	Other	136	125	222	56.3 (49.5, 62.9)	
fodt et al fohamy et al	Other	105 105	96 5	239 100	40.2 (33.9, 46.7) 5.0 (1.6, 11.3)	
fonamy et al fownsend et al	Europe	105	74	100	57.8 (48.8, 66.5)	
Fruntio et al	Europe	194	41	168	24.4 (18.1, 31.6)	-
Jrsini et al	Europe	182	270	616	43.8 (39.9, 47.9)	
Venturelli et al	Europe	105	394 57	767	51.4 (47.8, 55.0) 53.8 (43.8, 63.5)	
Walle-Hansen et al Neng et al	Europe China	200	57 52	106	53.8 (43.8, 63.5) 44.4 (35.3, 53.9)	
Weng et al Whitaker et al	Europe	84	52 28713	76155	44.4 (35.3, 53.9) 37.7 (37.4, 38.0)	
Gong et al	China	167	114	162	70.4 (62.7, 77.3)	-
Cong et al	China	111	267	538	49.6 (45.3, 53.9)	•
fan et al	China	98 379	0	125	0.0 (0.0, 2.9)	942
	China	379 203	47	119 316	39.5 (30.7, 48.9) 55.7 (50.0, 61.3)	100
		203	1/6	316	35.9 (30.9, 41.1)	
Yin et al	Europe					100 m
rin et al Zayet et al Zhan et al	Europe China	348	36	121	29.8 (21.8, 38.7)	
Yin et al Zayet et al Zhan et al Zhang et al	China China	348 378	1095	2433	45.0 (43.0, 47.0)	
rin et al Zayet et al Zhan et al Zhang et al Zhang et al	China China China	348 378 105	1095 178	2433 245	45.0 (43.0, 47.0) 72.7 (66.6, 78.1)	
Yan et al Yin et al Zayet et al Zhang et al Zhang et al Zhang et al Zhang et al	China China	348 378	1095	2433	45.0 (43.0, 47.0)	-

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).

persistent problem	% problem	number of studies	I ²	
Pathology	70 problem	3100103		
Lung Pathology	38.9 (3.4 to 91.9)	26	99.7%	_
Heart Pathology	6.0 (0.1 to 79.3)	12	99.9%	
Neurological pathology	5.3 (0.5 to 36.5)	12	99.7%	
Hypertension	1.5 (1.3 to 1.8)	4	0%	
Pancreas Pathology	1.4 (0.0 to 95.9)	3	94.7%	-
Vascular Problems	0.8 (0.0 to 33.6)	5	99.6%	
Kidney Pathology	0.7 (0.0 to 54.7)	6	99.7%	
Liver Pathology	0.6 (0.0 to 100.0)	3	98.8%	
Liver r danology	0.0(0.0 10 100.0)	0	00.070	
Symptom				
Fatigue	21.6 (2.5 to 74.7)	72	99.6%	
Breathing Problems	14.9 (1.6 to 64.9)	78	99.7%	
Sleep Problems	13.2 (1.2 to 64.9)	42	99.0%	
Tingling or Itching	11.3 (0.7 to 69.5)	14	98.2%	
Aches or Pains In Joints or Muscles	10.6 (1.0 to 57.5)	61	99.7%	_
Weakness	10.2 (0.5 to 72.2)	21	98.8%	_
Cognition or Memory Problems	10.1 (0.8 to 60.2)	49	99.4%	
Eye Problems	10.0 (0.0 to 96.5)	4	97.3%	-
Problems with Taste or Smell	9.6 (1.2 to 48.7)	60	98.6%	
PTSD	9.3 (0.5 to 65.5)	12	99.2%	_
Anxiety, Depression or Mood Change	7.7 (0.0 to 94.9)	5	99.1%	-
Cough	7.4 (1.3 to 33.5)	52	95.8%	-
Dizziness	7.4 (0.8 to 45.4)	26	97.7%	
Alopecia	7.2 (0.5 to 56.7)	17	99.1%	-
Chest Pain	6.7 (0.9 to 35.8)	43	98.0%	
Headache	6.5 (0.6 to 45.6)	51	99.1%	-
Palpitations	5.8 (1.2 to 24.5)	26	94.9%	
Speech or Language Problems	4.3 (0.0 to 88.5)	6	99.0%	
Nausea or Vomitting	3.9 (0.4 to 28.8)	49	99.6%	
Ear Problems	3.8 (0.2 to 45.0)	11	98.2%	_
Abdominal Pain	3.7 (0.1 to 63.8)	15	99.2%	
Sore Throat	3.5 (0.6 to 17.1)	22	97.1%	— —
Psychological Distress	2.9(0.0 to 100.0)	3	98.0%	— ——
Skin Problems	2.5 (0.0 to 56.2)	6	97.6%	
Fever	1.9 (0.1 to 34.7)	24	97.9%	
Chills	1.0 (0.0 to 98.8)	4	93.6%	
Functional status				
Not Returned to Full Health/Fitness	34.5 (4.3 to 85.9)	10	99.4%	

0 20 40 60 80 100

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

tudy athology discovered at follow-up	Dominant source of perticipants	Breadth of coverage	Mean days since infection	Cases	Total	% pers sympt	oms	
ellan et al lanco et al	hospitalised hospitalised	multiple domains single domain	120 104	128 52	238 100	53.8 52.0	(47.2, 60.2) (41.8, 62.1)	=
ija-Masson et al	hospitalised	single domain	91	103	137	75.2	(67.1.82.2)	
roidure et al	hospitalised	single domain	95	73	107	68.2	(58.5, 76.9)	-
an et al et al	hospitalised	single domain single domain	175 120	71 173	114 289	62.3 59.9	(52.7, 71.2) (54.0, 65.6)	
ao et al	hospitalised	single domain	104	122	142	85.9	(79.1, 91.2)	
iao et al	healthcare workers	single domain	409	113	303	37.3	(31.8, 43.0)	.
Alanese et al labahati et al	hospitalised hospitalised	multiple domains single domain	196 104	64 90	135	47.4	(38.8, 56.2) (44.3, 59.7)	
lunez-Fernandez et al	hospitalised	single domain	98	58	200	29.0	(22.8, 35.8)	
ân et al	hospitalised	single domain	105	87	647	13.4	(10.9, 16.3)	-
Sibila et al Sonnweber et al	hospitalised community	single domain single domain	116 103	98 73	172 133	57.0 54.9	(49.2, 64.5) (46.0, 63.5)	
ohamy et al	outpatients	single domain	105	5	100	5.0	(1.6, 11.3)	
fan et al	hospitalised	single domain	379	47	119	39.5	(30.7, 48.9)	
fin et al Chang et al	hospitalised hospitalised	single domain single domain	203	176	316	55.7 54.9	(50.0, 61.3) (45.7, 63.9)	
Drang et al	hospitalised	single domain	105	178	245	72.7	(66.6, 78.1)	
hou et al	hospitalised	single domain	127	114	164	69.5 51.7	(61.9, 76.5) (12.3, 89.1)	
ymptoms & function discovered at follow-up								~
bdeirahman et al	outpatients	multiple domains	270	105	172	61.0	(53.3, 68.4)	-
mold et al ucustin et al	hospitalised community	multiple domains multiple domains	90 131	81 123	110	73.6 27.8	(64.4, 81.6) (23.7, 32.3)	
laricich et al	hospitalised	single domain	139	66	204	32.4	(26.0, 39.2)	
lecker et al Nddal et al	outpatients community	single domain multiple domains	231 84	178	740	24.1 40.3	(21.0, 27.3) (31.8, 49.3)	• -
Romberg et al	outpatients	multiple domains	183	189	312	60.6	(51.0, 48.3) (54.9, 66.0)	· ·
loscolo-Rizzo et al	community	multiple domains	365	161	304	53.0	(47.2, 58.7)	-
OMEBAC study	hospitalised	multiple domains	127	244	478	61.0	(46.5, 55.6)	•
amilo-Garcia et al aruso et al	hospitalised hospitalised	multiple domains multiple domains	105 182	100	151 118	66.2 77.1	(58.1, 73.7) (68.5, 84.3)	
hai et al	hospitalised	multiple domains	378	156	546	28.6	(24.8, 32.6)	
iruli et al	community	multiple domains	90	18	122	14.8	(9.0, 22.3)	•
lavario et al ristilio et al	hospitalised hospitalised	single domain single domain	121 196	160 50	200	80.0 49.5	(73.8, 85.3) (39.4, 59.6)	
iaz-Fuentes et al	hospitalised	multiple domains	84	88	111	79.3	(70.5, 86.4)	
omenech-Montoliu et al	community	multiple domains	213	258	483	53.4	(48.9, 57.9)	
rol et al ernandez-de-Las-Penas et al	hospitalised hospitalised	multiple domains single domain	170 227	45 930	121 1142	37.2 81.4	(28.6.46.4) (79.1.83.7)	-
ernandez-de-Las-Penas et al u et al	hospitalised hospitalised	single domain single domain	227	930 20	1142	81.4	(79.1, 83.7) (6.2, 15.1)	
aber et al	healthcare workers	multiple domains	122	61	138	44.2	(35.8, 52.9)	-
iarcia-Abelian et al	hospitalised	multiple domains	182	28	116	24.1	(16.7, 33.0)	
le te officient all levervali et al	hospitalised healthcare workers	multiple domains multiple domains	105 122	119 69	130 323	91.5 21.4	(85.4, 95.7) (17.0, 26.2)	
lavervali et al	hospitalised	multiple domains	122	1265	1655	21.4	(17.0, 26.2) (74.3, 78.5)	
acobson et al	outpatients	multiple domains	119	79	118	66.9	(57.7, 75.3)	
lashif et al lim et al	outpatients	single domain multiple domains	105	101 591	242	41.7	(35.5, 48.2) (62.5, 68.8)	
am et al emboler et al	community	multiple domains multiple domains	195	591 226	365	65.7	(62.5, 68.8) (56.7, 66.9)	
lu et al	hospitalised	single domain	196	373	1301	28.7	(26.2, 31.2)	
lyanage-Don et al	hospitalised	multiple domains	113	99	153	64.7	(56.6, 72.3)	_ +
ogue et al ucidi et al	outpatients	multiple domains single domain	169 186	53 40	177	30.0 36.4	(23.3, 37.3) (27.4, 46.1)	
ui et al	hospitalised	multiple domains	96	41	204	20.1	(14.8, 26.3)	• ⁻
laestre-Muniz et al	hospitalised	multiple domains	379	309	543	56.9	(52.6, 61.1)	
tartinez et al latteuci et al	healthcare workers outpatients	multiple domains multiple domains	168 180	69 23	260	26.5	(21.3, 32.3) (11.0, 24.1)	
taneuci et al tazza et al	hospitalised	single domain	100	23	226	35.8	(11.0, 24.1) (29.6, 42.5)	
fechi et al	outpatients	multiple domains	274	92	112	82.1	(73.8, 88.7)	-
tei et al	hospitalised	multiple domains	158	976	3677	26.5	(25.1, 28.0)	
tenges et al tillet et al	community hospitalised	multiple domains multiple domains	219 365	106 83	431	24.6 48.0	(20.6, 28.9) (40.3, 55.7)	-
lohiuddin et al	outpatients	multiple domains	161	67	313	21.4	(17.0, 26.4)	100
funblit et al	hospitalised	multiple domains	232	1534	2649	67.9	(56.0, 59.8)	
lehme et al	1	multiple domains	243	160	410	39.0	(34.3, 43.9)	-
iguyen et al /Keefe et al	hospitalised outpatients	single domain multiple domains	213 119	30 79	125	24.0 39.9	(16.8, 32.5) (33.0, 47.1)	
WNS study August 2021	community	multiple domains	84	2501	21374	11.7	(11.3, 12.1)	
ing et al	hospitalised	multiple domains	104	13	175	7.4	(4.0, 12.4)	
hmu et al Ismanov et al	social media hospitalised	single domain multiple domains	91 270	113 126	152	74.3 24.3	(66.6, 81.1) (20.6, 28.2)	
HOSP-COVID study	hospitalised	multiple domains	174	797	861	92.6	(90.6, 94.2)	
leghin et al	outpatients	multiple domains	191	241	599	40.2	(36.3, 44.3)	
eluso et al	social media	single domain	112	89	143	62.2	(53.8, 70.2)	
etersen et al u et al	community hospitalised	multiple domains single domain	125 105	95 176	180 540	52.8 32.6	(45.2, 60.2) (28.7, 36.7)	
adiko et al	community	multiple domains	84	4	109	3.7	(1.0, 9.1)	
ass et al	hospitalised	single domain	90	82	135	60.7	(52.0, 69.0)	-
iestra-Ayora et al	healthcare workers	single domain multiple domains	182	62	195	26.7 19.7	(20.6, 33.5) (16.0, 23.8)	
ighi et al omero-Duarte et al	hospitalised hospitalised	multiple domains	84 196	83 509	421 797	63.9	(16.0, 23.8) (60.4, 67.2)	
athyamunthy et al	hospitalised	multiple domains	104	66	279	23.7	(18.8, 29.1)	-
eedle of al	outpatients	multiple domains	147	107	146	73.3	(65.3, 80.3)	=
hang et al Ightid of al	hospitalised hospitalised	multiple domains multiple domains	196 222	441	795	55.4 93.3	(51.9, 58.9) (90.0, 95.7)	
imani et al	hospitalised	single domain	197	12	120	10.0	(5.3, 16.8)	· · · · · · · · ·
ikala et al	outpatients	multiple domains	91	56	102	54.9	(44.7, 64.8)	-
kjorten et al oraas et al	hospitalised community	single domain multiple domains	105 126	59 380	125	46.8	(37.9, 55.9) (52.4, 60.0)	
oraas et al tavem et al	community	multiple domains multiple domains	126	380	451	41.0	(52.4, 60.0) (36.4, 45.7)	
tephenson et al	community	multiple domains	104	2038	3065	66.5	(64.8, 68.2)	
udre et al	community	multiple domains	84	108	4182	2.6	(2.1, 3.1)	• <u>-</u>
iykas et al aboada et al	hospitalised hospitalised	single domain multiple domains	127 189	75 87	127	59.1 47.5	(50.0, 67.7) (40.1, 55.0)	_
arsitani et al	hospitalised	single domain	105	34	163	29.6	(40.1, 55.0) (21.4, 38.8)	- ⁻
awfik et al	healthcare workers	single domain	91	40	120	33.3	(25.0, 42.5)	.
aylor et al empany et al	hospitalised healthcare workers	single domain multiple domains	126 84	261 116	545	47.9 53.5	(43.6, 52.2) (46.6, 60.2)	
empany et al Teyjeh et al	heatthcare workers hospitalised	multiple domains	136	116	217	56.3	(49.5, 62.9)	-
bdt et al	hospitalised	single domain	105	96	239	40.2	(33.9.46.7)	•
ownsend et al runfio et al	outpatients hospitalised	multiple domains	86 194	74 41	128 168	57.8 24.4	(48.8, 66.5) (18.1, 31.6)	
runfio et al rsini et al	hospitalised social media	multiple domains single domain	194	41 270	168 616	24.4 43.8	(18.1.31.6) (39.9.47.9)	
entureli et al	outpatients	multiple domains	105	394	767	51.4	(47.8, 55.0)	
Valle-Hansen et al	hospitalised	single domain	200	57	105	53.8	(43.8, 63.5)	_=
/eng et al /hitaker et al	hospitalised community	single domain multiple domains	104 84	52 28713	117 76155	44.4 37.7	(35.3, 53.9) (37.4, 38.0)	
	healthcare workers	single domain	167	114	162	70.4	(62.7.77.3)	
	hospitalised	single domain	111	267	538	49.6	(45.3, 53.9)	
long et al		multiple domains	98	0	125	0.0	(0.0, 2.9)	• _
long et al long et al lan et al	hospitalised	multiple domains	289 348	127	354 121	35.9 29.8	(30.9, 41.1) (21.8, 38.7)	
llong et al llong et al lan et al layet et al	hospitalised	multiple		36 1095	121 2433	45.0	(43.0, 47.0)	-
iong et al iong et al an et al syste et al tan et al		multiple domains multiple domains	378			43.9	(8.2, 87.2)	()
long et al cong et al an et al systet et al hang of al	hospitalised hospitalised	multiple domains multiple domains	378			41.5		V
cong et al gong et al systet et al sans et al anang et al attornes from health record linkage	hospitalised hospitalised hospitalised	multiple domains			1-0			-
ong et al an al al agrect et al anna et al anna et al anna et al anna et al adorement from health record linkage Ady et al (hospitalized)	hospitalised hospitalised	multiple domains multiple domains single domain single domain	378 150 126	1091 1718	11800	9.2	(8.7, 9.8)	
Jong et al forget al an et al synt et al hang et al outcomes from health record linkage Al-by et al (Dox-Doxplasted) Al-by et al (Dox-Doxplasted)	hospitalised hospitalised hospitalised	multiple domains single domain	150			9.2	(8.7, 9.8) (2.7, 3.0) (42.4, 46.1)	.
Uning 41 at 100 of 41 at 100	hospitalised hospitalised hospitalised community outpatients hospitalised	multiple domains single domain single domain multiple domains single domain	150 126 243 154	1718 1255 6085	60255 2839 28335	9.2 2.9 44.2 21.5	(8.7, 9.8) (2.7, 3.0) (42.4, 46.1) (21.0, 22.0)	÷
Jong et al an et al ser et al	hospitalised hospitalised hospitalised community outpatients hospitalised community	multiple domains single domain single domain multiple domain single domain single domain	150 126 243 154 350	1718 1255 6085 28	60255 2839 28335 170	9.2 2.9 44.2 21.5 16.5	(8.7, 9.8) (2.7, 3.0) (42.4, 46.1) (21.0, 22.0) (11.2, 22.9)	÷
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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

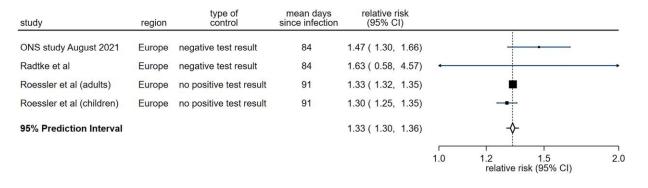


Figure 5. Forest plot of risk of Long COVID in included studies with community-based samples and controls assessed as having low risk of bias, with 95% prediction intervals.

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 sub-groups 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 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.

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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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