

RESEARCH ARTICLE

Open Access



# Risk for newly diagnosed diabetes after COVID-19: a systematic review and meta-analysis

Ting Zhang<sup>1,2</sup>, Qimin Mei<sup>1</sup>, Zhaocai Zhang<sup>3</sup>, Joseph Harold Walline<sup>4</sup>, Yecheng Liu<sup>1\*</sup> , Huadong Zhu<sup>1\*</sup> and Shuyang Zhang<sup>5\*</sup>

## Abstract

**Background:** There is growing evidence that patients recovering after a severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) infection may have a variety of acute sequelae including newly diagnosed diabetes. However, the risk of diabetes in the post-acute phase is unclear. To solve this question, we aimed to determine if there was any association between status post-coronavirus disease (COVID-19) infection and a new diagnosis of diabetes.

**Methods:** We performed a systematic review and meta-analysis of cohort studies assessing new-onset diabetes after COVID-19. PubMed, Embase, Web of Science, and Cochrane databases were all searched from inception to June 10, 2022. Three evaluators independently extracted individual study data and assessed the risk of bias. Random-effects models estimated the pooled incidence and relative risk (RR) of diabetes compared to non-COVID-19 after COVID-19.

**Results:** Nine studies with nearly 40 million participants were included. Overall, the incidence of diabetes after COVID-19 was 15.53 (7.91–25.64) per 1000 person-years, and the relative risk of diabetes after COVID-19 infection was elevated (RR 1.62 [1.45–1.80]). The relative risk of type 1 diabetes was RR=1.48 (1.26–1.75) and type 2 diabetes was RR=1.70 (1.32–2.19), compared to non-COVID-19 patients. At all ages, there was a statistically significant positive association between infection with COVID-19 and the risk of diabetes: <18 years: RR=1.72 (1.19–2.49), ≥18 years: RR=1.63 (1.26–2.11), and >65 years: RR=1.68 (1.22–2.30). The relative risk of diabetes in different gender groups was about 2 (males: RR=2.08 [1.27–3.40]; females: RR=1.99 [1.47–2.80]). The risk of diabetes increased 1.17-fold (1.02–1.34) after COVID-19 infection compared to patients with general upper respiratory tract infections. Patients with severe COVID-19 were at higher risk (RR=1.67 [1.25–2.23]) of diabetes after COVID-19. The risk (RR=1.95 [1.85–2.06]) of diabetes was highest in the first 3 months after COVID-19. These results remained after taking confounding factors into account.

\*Correspondence: ptcaliu@sina.com; drzhuhd@sina.com; ShuyangZhang\_103@sina.com

<sup>1</sup> Emergency Department, State Key Laboratory of Complex Severe and Rare Diseases, Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, Beijing, China

<sup>5</sup> Department of Cardiology, State Key Laboratory of Complex Severe and Rare Diseases, Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, Beijing, China

Full list of author information is available at the end of the article



**Conclusions:** After COVID-19, patients of all ages and genders had an elevated incidence and relative risk for a new diagnosis of diabetes. Particular attention should be paid during the first 3 months of follow-up after COVID-19 for new-onset diabetes.

**Keywords:** SARS-CoV-2 virus, Diabetes mellitus, COVID-19, Unmeasured confounders

## Background

Coronavirus disease 2019 (COVID-19) is a complex clinical syndrome caused by the severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) [1]. Despite many large studies leading to the approval of vaccines and antivirals, the global spread of SARS-CoV-2 continues [2, 3]. As of June 18, 2022, there have been more than 535,863,950 confirmed cases globally, including 6,314,972 deaths (according to the World Health Organization) [4]. Factors associated with poor outcomes, including hospitalization, intensive care unit (ICU) admissions, and mortality in COVID-19 patients, are of considerable interest. More specifically, health comorbidities and baseline physical activity [5] may predispose patients to an increased risk of poor outcomes following COVID-19 infection.

Previous studies have indicated that diabetes mellitus (DM) is associated with an increased risk of severe COVID-19, acute respiratory distress syndrome (ARDS), and in-hospital mortality [6–8]. More intriguingly, a recent meta-analysis has reported that newly diagnosed diabetes is commonly observed in COVID-19 patients [9–11]. The world has raised concerns about a bi-directional relationship between these two health conditions [12].

As the COVID-19 pandemic has progressed, there is growing evidence that after the acute phase of the disease, people with COVID-19 can develop lingering sequelae (called “long COVID”) that may involve pulmonary and extrapulmonary organ system manifestations, such as diabetes [13]. Follow-up of children with COVID-19 has identified that the incidence of type 1 newly diagnosed diabetes has increased [14]. An unregistered meta-analysis [15] in PROSPERO also found an increased risk of diabetes among adults with long COVID-19, but it has some flaws in the study design which limit the interpretation and applicability of the individual studies’ findings. Therefore, there is an urgent need for systematic reviews and meta-analyses of the existing literature, particularly focusing on controlled studies.

This systematic review and meta-analysis was conducted to estimate the prevalence of a new diagnosis of diabetes after COVID-19 compared to non-COVID-19.

## Methods

This review was conducted and reported in accordance with the Meta-analysis of Observational Studies in Epidemiology (MOOSE) [16] and Preferred Reporting Items for Systematic Review and Meta-analysis (PRISMA) [17] guidelines and according to the methods described in the Cochrane Handbook for Systematic Reviews of Interventions (Additional file 2: Table S1 and Table S2). The study protocol was registered in PROSPERO on June 24, 2022 (registration number: CRD42022330723). As all included data was from previously published studies, no institutional review board approval was required.

### Search strategy and eligibility criteria

We systematically searched the following electronic bibliographic databases: PubMed, Embase, Web of Science, and the Cochrane Central Register of Controlled Trials. No time or language restrictions were applied to the search results. Full details of the search strategies used are provided in Additional file 1: Table S1. In brief, combinations of search terms were applied, relating to COVID-19 or SARS-CoV-2, diabetes mellitus, diabetes, or DM. The search was conducted from inception through June 10, 2022.

To be included in this systematic review, prospective or retrospective cohort studies had to meet all of the following criteria: (1) the main exposure of interest was COVID-19, which was defined based on International Classification of Diseases (ICD) codes; (2) to determine relative associations, hospitalized or population controls were utilized as comparators, with priority given to population controls where available; (3) report of newly diagnosed DM that was defined as the new onset of diabetes (no prior history of diabetes with a fasting plasma glucose [FPG]  $\geq 7.0$  mmol/L or a random blood glucose [RBG]  $\geq 11.1$  mmol/L or a HbA1c  $>6.5\%$ ). The following were excluded: case reports or case series, reviews, commentaries, and letters.

### Outcomes

The primary outcome was the incidence of newly diagnosed DM after the diagnosis of SARS-CoV-2 infection, and the relative risk of acquiring DM compared with contemporary or historical controls in the non-SARS-CoV-2 cohort. We performed subgroup analyses according to

age, gender, type of DM, time of onset, and whether the control group was generalized upper respiratory tract infections. We also performed two post hoc subgroup analyses: mild-to-moderate COVID patients versus severe COVID patients, looking at three different follow-up times after COVID-19 (less than 3 months, 3 to 6 months, and greater than 6 months).

### Study selection and data extraction

Data extraction from eligible studies was performed using a standardized spreadsheet. The extracted data included items related to study design and data sources, study participant characteristics, study definition of COVID-19, definition of DM, covariates, and person-years of follow-up (either reported or calculable). The data were extracted by the first reviewer (TZ) and double-checked by a second reviewer (QMM). Disagreement between the two reviewers was resolved by discussion with a third reviewer (YCL).

### Methodological quality

The risk of bias for the included studies was assessed using the Newcastle–Ottawa quality assessment scale for cohort studies [18]. Critical appraisal was carried out by three reviewers (TZ, QMM, and YCL), with discrepancies discussed with the larger authors' group to reach a consensus.

### Statistical analysis

For each included study, the incidence of newly diagnosed DM was calculated using the reported number of newly diagnosed diabetes cases and person-years of follow-up. We first transformed proportions using the Freeman–Tukey double arcsine method [19] and then performed an inverse variance random-effects meta-analysis (DerSimonian and Laird) [20] to calculate the pooled estimates. Diabetes rates between the SARS-CoV-2 patients and control subjects were reported using relative risk (RR) with 95% confidence intervals (CIs).

Heterogeneity between included studies was assessed with the Cochran  $Q$  and  $I^2$  statistics. For the qualitative interpretation of heterogeneity,  $I^2$  values of at least 50% were considered to represent substantial heterogeneity, and values of at least 75% indicated considerable heterogeneity [21]. The significance level for the  $Q$  statistic was set at 0.1. For outcomes reported in 10 or more studies, publication bias was explored by constructing power-enhanced funnel plots (sunset funnel plot) and an Egger's test [22, 23].

To consider how strong uncontrolled confounders in each meta-analyzed study would have to be to negate the observed results, we applied a sensitivity analysis for meta-analyses that are analogous to the  $E$ -value [24,

25]. We calculated an  $E$ -value [26] representing the minimum strengths of associations on the risk ratio scale that uncontrolled confounders would need to jointly have with COVID-19 and with DM across all studies in each meta-analysis to shift the meta-analytic estimate or its 95% CI to the null.

All risk estimates were calculated with the corresponding 95% CIs.  $p$ -values  $<0.05$  were considered statistically significant. All statistical analyses were performed with R software (version 4.1.2, Vienna, Austria) [27, 28], using the packages “metafor,” “EValue,” “confoundedmeta,” “metaviz,” and “metaUtility.”

### Results

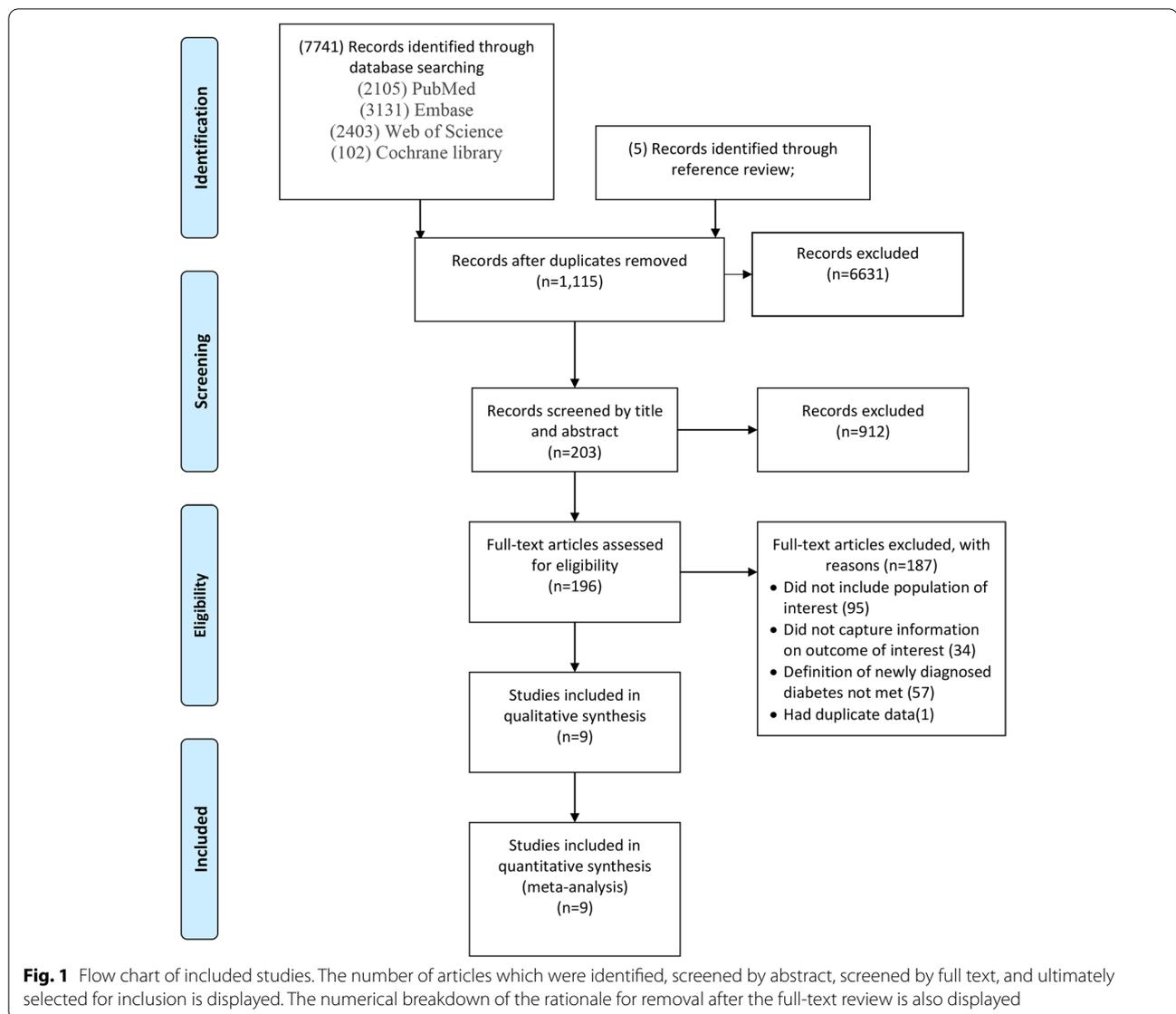
The search yielded 7746 citations. After duplicates were removed and titles and abstracts were reviewed, 7543 articles were excluded. Of the remaining 203 studies, full-text articles of 196 were available. Of the 196, 187 were then excluded after reviewing the full-text manuscripts. After several stages of review, nine eligible studies (with 10 cohorts) were included in the meta-analysis and eight studies were propensity score matching (PSM) cohort studies (Fig. 1) [29–37].

Table 1 and Table S2 show the characteristics of the included studies: six were conducted in the USA [29, 30, 32, 33, 35, 37], two in England [34, 36], and one in Germany [31]. Among the nine studies, seven included only adults, one study included a full population (no restrictions), and one study included only adolescents. Overall, there were 4,002,475 people diagnosed with COVID-19 and 34,717,422 people in the respective control groups.

The risk of bias in the included cohort studies was assessed using the Newcastle–Ottawa Scale and is presented in Table S3. The overall score was 88 of 90 (97.8%), which is considered to be a relatively low risk for bias.

Of the nine included studies, although one study lacked specific follow-up times, the overall incidence of DM was found to be 15.53 cases per 1000 person-years of follow-up ([95% CI 7.91–25.64]; Fig. 2), using a random-effects meta-analysis. The relative risk of diabetes after COVID-19 was 1.62 ([95% CI 1.45–1.80]; Fig. 3) compared with patients not infected with COVID-19. Power-enhanced funnel plots (sunset funnel plot) and Egger's test ( $p = 0.104$ ) did not suggest publication bias (Fig. 4). In the sunset funnel plot, we found that all studies had strong statistical tests.

A subgroup analysis was performed looking at the type of new-onset diabetes. The risk of developing type 1 (insulin-dependent) diabetes was found by meta-analysis to be  $RR=1.48$  ([95% CI 1.26–1.75]; Additional file 1: Fig. S1); type 2 (non-insulin-dependent) diabetes had a  $RR$  of 1.70 ([95% CI 1.32–2.19]; Additional file 1: Fig. S1); and those with an unspecified type of diabetes had a  $RR$  of



1.50 ([95% CI 0.87–2.58]; Additional file 1: Fig. S1) compared to the uninfected population.

Another subgroup analysis was performed according to whether the control group was or was not those with an upper respiratory tract infection. The relative risk of developing diabetes was increased 1.17-fold ([95% CI 1.02–1.34]; Additional file 1: Fig. S2) after COVID-19 compared to patients with upper respiratory tract infections. The relative risk of developing diabetes was increased 1.82-fold ([95% CI 1.47–2.24]; Additional file 1: Fig. S2) after COVID-19 compared to the general population.

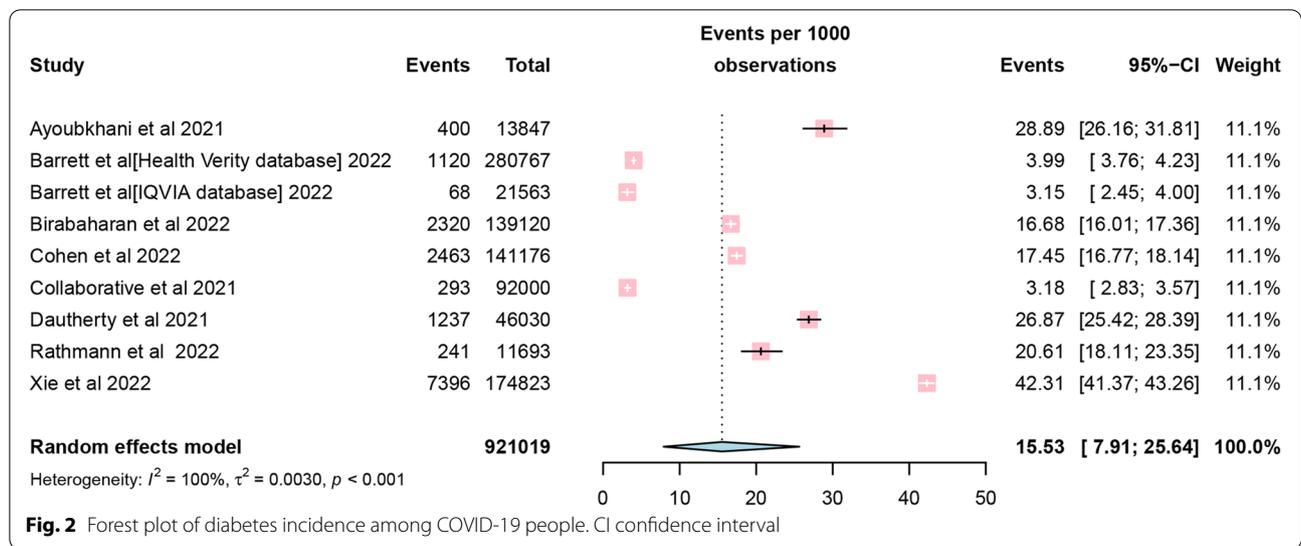
Subgroup analysis was performed again according to the age of onset. The annual incidence rate per 1000 person-years of follow-up was 3.65 (95% CI, 2.91–4.83;

Additional file 1: Fig. S3) in those <18 years; 15.53 ([95% CI 7.91–25.64]; Additional file 1: Fig. S3) in those  $\geq 18$  years; and 17.45 ([95% CI 16.77–18.14]; Additional file 1: Fig. S3) in those >65 years. At all ages, there was a statistically significant positive correlation between COVID-19 infection and the risk of developing diabetes. The positive correlation between COVID-19 infection and the risk of developing diabetes was statistically significant at all ages, <18 years: RR=1.72 ([95% CI 1.19–2.49]; Additional file 1: Fig. S4);  $\geq 18$  years: RR=1.63 ([95% CI 1.26–2.11]; Additional file 1: Fig. S4); >65 years: RR=1.68 ([95% CI 1.22–2.30]; Additional file 1: Fig. S4).

Finally, a subgroup analysis was performed according to gender. The incidence rate was 3.14 ([95% CI 0.75–7.15]; Additional file 1: Fig. S5) per 1000 person-years of

**Table 1** Characteristics of the total study population

First author (year)	Group	Population	Age (years), mean (SD or range), or median (IQR)	Gender (proportion)	Event number	Follow-up time	Levels of COVID-19 severity
Xie, 2022 [35]	COVID-19 group	181,280	60.92 (17.02)	Male: 159,666 (88.08%) Female: 21,614 (11.92%)	7871	352 (IQR 244–406) days	Not hospitalized; hospitalized; intensive care
Rathmann, 2022 [31]	Control group	4,118,441	61.5 (17.08)	Male: 3,655,034 (88.75%) Female: 463,407 (11.25%)	127,002	352 (245–406) days	Not applicable
	COVID-19 group	181,280	42.6 (19.1)	Male: 164,918 (54.4%) Female: 16,362 (45.6%)	241	119 (IQR 0–210) days	Mild COVID-19
Qeadan, 2022 [33]	Control group	4,118,441	42.6 (19.1)	Male: 2,240,432 (54.4%) Female: 16,370 (45.6%)	228	161 (IQR 4–225) days	Not applicable
	COVID-19 group	2,489,266	Not reported	Male: 1,081,608 (46.1%) Female: 1,264,069 (53.9%)	5163	Not reported	Unavailable
Collaborative, 2022 [34]	Control group	24,803,613	Not reported	Male: 10,579,475 (45.9%) Female: 12,491,493 (54.1%)	36,348	Not reported	Not applicable
	COVID-19 group	77,347	68	Male: 39,808 (51.5%) Female: 37,539 (48.5%)	670	274 days	Hospitalized COVID-19
Cohen, 2022 [30]	Control group	386,669	68	Male: 199,013 (51.5%) Female: 187,656 (48.5%)	2763	274 days	Not applicable
	COVID-19 group	87,337	75 (71–82)	Male: 58,110 (44%) Female: 29,227 (56%)	2463	64 (IQR 23–150) days	Not hospitalized: 63,690 Hospitalized: 23,486
Birabharan, 2022 [32]	Control group	87,337	74 (70–80)	Male: 42% Female: 58%	1249	64 (IQR 23–150) days	Not applicable
	COVID-19 group	282,105	Not reported	Not reported	2320	180 days	Mild COVID-19: 276,748 Moderate/severe COVID-19: 5357
Barrett [QVIA database], 2022 [29]	Control group	286,275	Not reported	Not reported	1916	180 days	Not applicable
	COVID-19 group	80,839	1.23 (4.3)	Male: 404,465 (49.9%) Female: 40,517 (50.1%)	68	362 days	Hospitalized: 6473 (0.4)
Barrett [Health Verity database], 2022 [29]	Control group	404,465	1.23 (4.3)	Male: 58,110 (49.9%) Female: 201,880 (50.1%)	132	362 days	Not applicable
	COVID-19 group	439,439	12.7 (3.8)	Male: 219,427 (49.9%) Female: 220,012 (50.1%)	1120	484 days	Hospitalized: 13,118 (3.0)
Ayoubkhani, 2021 [36]	Control group	439,439	12.7 (3.8)	Male: 219,427 (49.9%) Female: 220,012 (50.1%)	853	484 days	Not applicable
	COVID-19 group	36,100	60.9 ± 17.02	Male: 54.9% Female: 45.1%	400	140 days	Intensive unit care (ICU): 4745 Non-ICU: 43,035
Daugherty, 2021 [37]	Control group	36,100	61.5 ± 17.08	Male: 54.9% Female: 45.1%	125	140 days	Not applicable
	COVID-19 group	193,113	41.7 ± 13.9	Male: 47.6% Female: 52.4%	1237	87 (IQR 45–124) days	Mild and hospitalized COVID-19
Control group	193,113	41.6 ± 13.8	Males: 47.5% Females: 52.5%	649	87 (IQR 45–124) days	Not applicable	



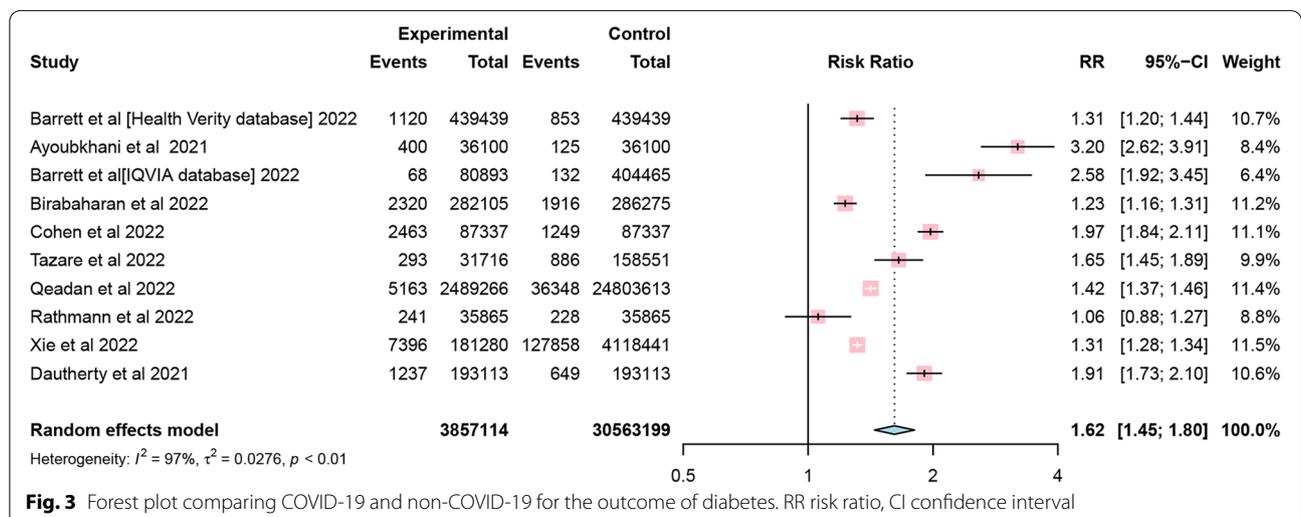
**Fig. 2** Forest plot of diabetes incidence among COVID-19 people. CI confidence interval

follow-up in the male population and 3.00 ([95% CI 0.80–6.59]; Additional file 1: Fig. S5) per 1000 person-years of follow-up in the female population. The relative risk of developing diabetes was increased 2.08-fold in males ([95% CI 1.27–3.40]; Additional file 1: Fig. S6) and 2.15-fold in females ([95% CI 1.26–3.68]; Additional file 1: Fig. S6) after COVID-19 compared to a population without COVID-19 infection.

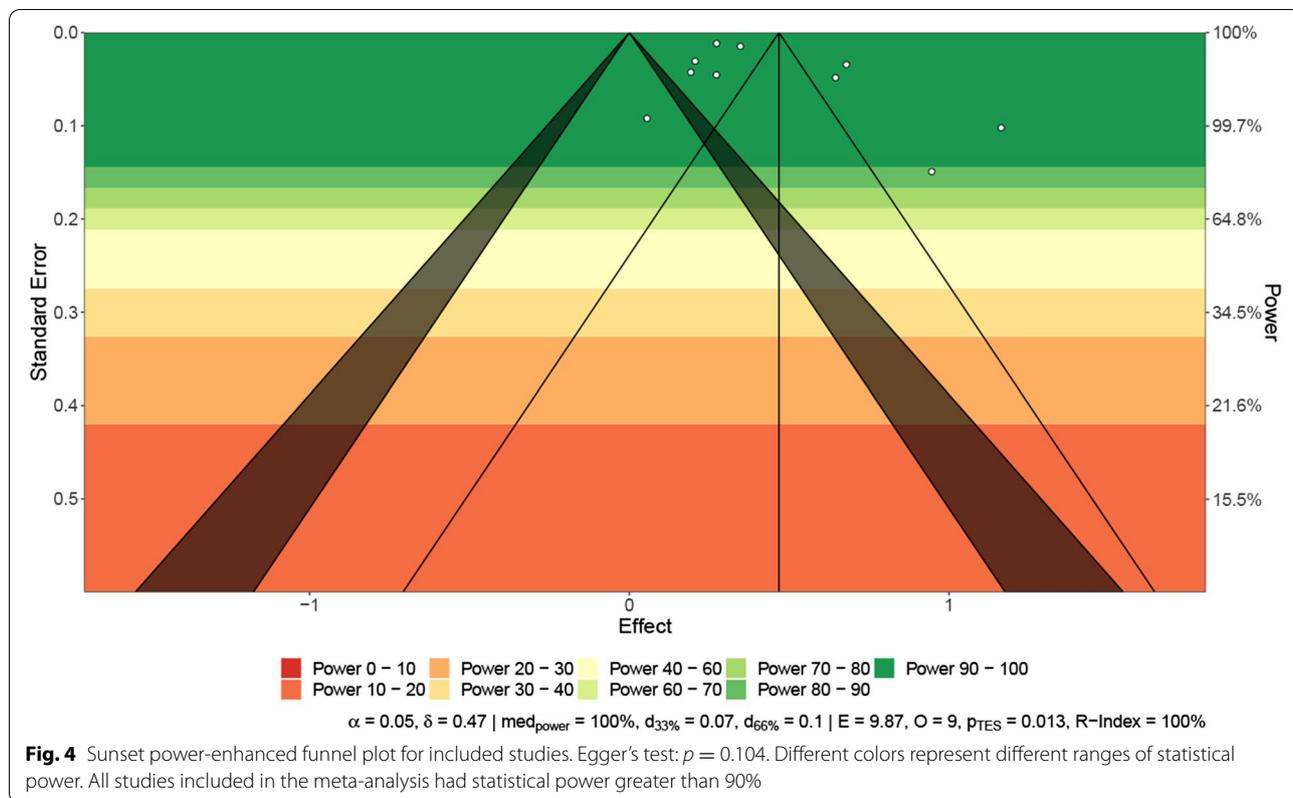
Follow-up time data were available for eight studies (with 9 cohorts) [29–31, 33–37]. The mean follow-up time of participants ranged from 64 to 352 days. The pooled cumulative incidence of diabetes reported at different follow-up time periods (Additional file 1: Fig. S7) was 2.19% (95% CI 1.36–3.21) at less than 3 months, 4.46% (95% CI 0.40–12.53) at 3 to 6 months, and 0.91% (95% CI 0.05–2.82) at more than 6 months. Furthermore,

a significantly higher risk of new-onset diabetes was detected over the reported range of follow-up (Additional file 1: Fig. S8): RR 1.95 at less than 3 months (95% CI 1.85–2.06), RR 1.24 at 3 to 6 months (95% CI 1.12–1.37), and RR 1.38 at more than 6 months (5% CI 1.23–1.55).

Among the included studies, patients in six cohort studies [29–31, 34, 37] suffered from mild-to-moderate COVID-19; patients in three studies [32, 35, 36] included mild, moderate, and severe patients; only one study [33] did not parse the acuity levels of COVID-19 patients included. A subgroup analysis found that, compared with non-COVID-19 patients, the incidence of diabetes in the mild-to-moderate COVID-19 cohort was 2.96% ([95% CI 0.58–7.07]; Additional file 1: Fig. S9), and the risk was 1.48 times ([95% CI 1.25–1.75]; Additional file 1: Fig. S10); the incidence of diabetes in the severe patients was

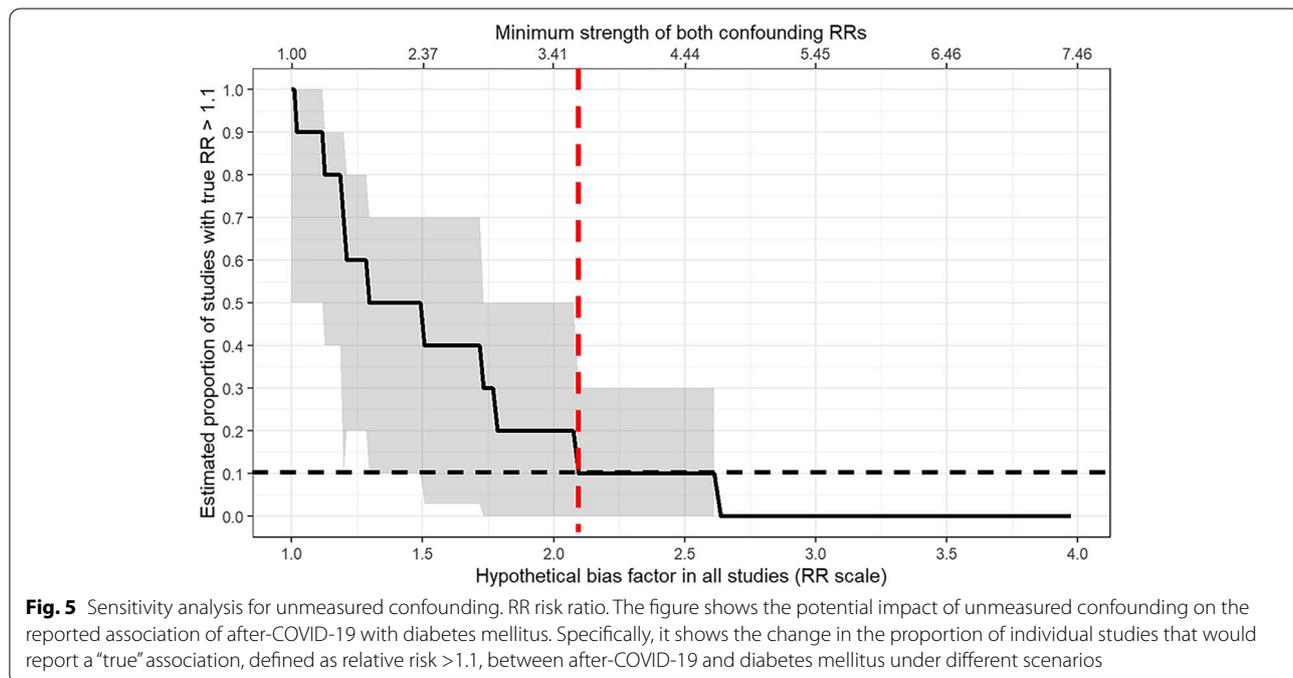


**Fig. 3** Forest plot comparing COVID-19 and non-COVID-19 for the outcome of diabetes. RR risk ratio, CI confidence interval



11.65% ([95% CI 2.59–25.96]; Additional file 1: Fig. S9), and the risk was 1.67 times ([95% CI 1.25–2.23]; Additional file 1: Fig. S10).

The sensitivity analysis for unmeasured confounding showed that to reduce the percentage of RR above 1.1 from 100 to 10%, we estimated a bias factor of at least 2.08 ([95% CI 1.55–2.84]; Fig. 5) would be required in each



study. This means that unmeasured confounding would have needed to shift each study's point estimate away from the null by 2.08-fold on the RR scale. Using the *E*-value metric, this bias factor is equivalent to unmeasured confounders in each study that affected both COVID-19 and diabetes by risk ratios of at least 3.58 each ([95% CI 2.43–5.12]; Additional file 1: Fig. S11 and Fig. S12).

## Discussion

This systematic review and meta-analysis provided comprehensive quantitative estimates of the incidence of diabetes in 10 post-COVID-19 populations. To our knowledge, this is the largest and most wide-ranging analysis of this kind to date. With nearly 40 million participants, and nearly 200,000 cases of diabetes reported, we found a post-COVID-19 incidence of diabetes per 1000 person-years of 15.53, and a relative risk of 1.62 compared to non-COVID-19-infected people. Subgroup analyses suggested that the risk of developing diabetes was also increased regardless of age, gender, type of diabetes, follow-up time, or level of COVID-19 severity, although undifferentiated diabetes did not have a significant relative risk. These results remained significant even after accounting for the possibility of unmeasured confounding.

Similar results have been reported in patients infected with other viruses, with an increased incidence of diabetes compared with those not infected [38–40]. Our subgroup analysis revealed a 1.2-fold increased risk of developing diabetes after COVID-19 compared to patients with other upper respiratory tract infections and a 1.82-fold increased risk of developing diabetes after COVID-19 compared to the general population. This reinforces the need for clinicians to pay attention to patients' glucose metabolism in the post-acute phase of COVID-19.

There is also new evidence regarding the effect of the SARS-CoV-2 virus on pancreatic  $\beta$ -cell function [41]. It has been suggested that SARS-CoV-2 may affect the pancreas by acting on the mRNA of angiotensin-converting enzyme 2 (ACE2) in the endocrine and exocrine glands of the pancreas [11, 42]. The presence of SARS-CoV-2 antigen has recently been reported in the postmortem pancreas of patients who died from COVID-19 [43]. In addition, SARS-CoV-2 can induce a cytokine storm, an exaggerated immune response that produces a broad spectrum of cytokines, thereby establishing a systemic pro-inflammatory environment, which may play a role in promoting insulin resistance and  $\beta$ -cell hyperstimulation, ultimately leading to altered cellular function and the death of  $\beta$ -cells [44–46]. According to our subgroup analysis, there was a 1.48-fold increased risk of developing type 1 diabetes and a 1.7-fold increased risk of

type 2 diabetes compared to patients not infected with COVID-19.

In our analysis, the incidence rate per 1000 person-years of follow-up was 3.65 (95% CI, 2.91 to 4.83), RR=1.72 (95% CI, 1.1 to 2.50) in the <18-year-old population, with similar results in adults and in those >65 years old. Moreover, the relative risk of morbidity was similar across genders. These findings underscore the importance of COVID-19 prevention in all age groups and genders, such as encouraging vaccination of all eligible children and adolescents [47].

Although all of the studies we included reduced confounders by adjusting for the risk of associated factors (propensity score matching) [48], concerns about possible bias due to uncontrolled confounders (e.g., comorbidities, socioeconomic environment, body mass index [BMI], etc.) remain [49]. Our study is the first meta-analysis to consider the *E*-value as a parameter of unmeasured confounders in examining the association between the COVID-19 post-acute phase and diabetes risk, which represents a new methodological contribution to the study of COVID-19 and diabetes [24]. The *E*-value, a sensitivity analysis of unmeasured confounders, is a relatively new method for measuring the association between exposure and outcome robustness and to assess evidence of causality [50]. Our results suggest that an unobserved confounder would need to be associated with a risk ratio of 2.08 for exposure and outcome to fully explain the mean RR of 1.62. In addition, a risk ratio of 3.58 would be required for the confounder to make the risk estimate statistically nonsignificant. Propensity matching was performed in all of the studies we included, and most studies adjusted for at least some clinically important confounding factors, such as patient age, gender, BMI, race, and comorbidities. Therefore, we believe it is implausible that residual confounders exist above and beyond these measured confounders that are sufficient to explain away the above results.

## Limitations

Several potential study limitations need to be considered. First, all included studies used a retrospective design and all studies used the breadth and depth of large electronic healthcare databases [29, 30, 32, 33, 35, 37, 51–53] to construct cohorts and define health characteristics based on validated definitions, which cannot exclude misclassification bias, particularly for diabetes types. Second, some of the studies used contemporary controls, not excluding the possibility that some individuals may be infected with SARS-CoV-2 and have not been tested, which could bias the results toward the null hypothesis if these individuals were present in large numbers in the contemporary control group. Third, because the

included studies were conducted in different countries and in different regions within the same country, differences in national and regional care policies are expected, for which this meta-analysis could not be adapted or adjusted. Fourth, the study designs were heterogeneous (prospective cohort and retrospective cohort studies). As the number of available prospective cohort studies on this topic remains small, more high-quality studies are needed to confirm our results.

## Conclusions

Patients of all ages and genders recovering from COVID-19 had an elevated incidence and relative risk for developing diabetes. Particular attention should be paid to potential new-onset diabetes during the first 3 months of follow-up after COVID-19.

## Abbreviations

ARDS: Acute respiratory distress syndrome; BMI: Body mass index; CI: Confidence interval; COVID-19: Coronavirus disease 2019; DM: Diabetes mellitus;  $I^2$ : Heterogeneity measure; ICD: International Classification of Diseases; ICU: Intensive care unit; PSM: Propensity score matching; RR: Relative risk; SARS-CoV-2: Severe acute respiratory syndrome coronavirus 2.

## Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s12916-022-02656-y>.

**Additional file 1: Table S1.** Search strategies applied in each bibliographic database. **Table S2.** Characteristic of studies included in the present study for analyzing risk of incident diabetes post-COVID-19 versus matched controls. **Table S3.** Quality assessment of included studies with the Newcastle–Ottawa Scale. **Figure S1.** Subgroup analysis of diabetes type. **Figure S2.** Subgroup analysis to whether the control group was or was not upper respiratory tract infections. **Figure S3.** Forest plot showing the incidence of DM among different age groups after COVID-19. **Figure S4.** Subgroup analysis of different age groups. **Figure S5.** Forest plot showing the incidence of DM among different gender after COVID-19. **Figure S6.** Subgroup analysis of gender. **Figure S7.** Forest plot showing the incidence of diabetes among different follow-up time after COVID-19. **Figure S8.** Subgroup analysis of different follow-up time. **Figure S9.** Forest plot showing the incidence of diabetes based on different levels of COVID-19 severity. **Figure S10.** Subgroup analysis of different levels of COVID-19 severity. **Figure S11.** Bias factor = log(1). **Figure S12.** Bias factor = log(2.08).

**Additional file 2: Table S1.** PRISMA checklist. **Table S2.** MOOSE Statement - Reporting Checklist for Authors, Editors, and Reviewers of Meta-analyses of Observational Studies.

## Acknowledgements

The authors thank Dr. Maya B. Mathur from the Quantitative Sciences Unit and Department of Pediatrics, Stanford University, for her assistance in bias analysis on the effect of unmeasured confounding.

## Authors' contributions

TZ, YCL, and QMM conceived and designed this study. TZ, QMM, and YCL conducted the study selection and data extraction. HDZ, TZ, and ZCZ conducted the statistical analyses and participated in the interpretation of the data. JHW and SZ drafted the manuscript. All authors critically reviewed the manuscript. The authors read and approved the final manuscript.

## Funding

The National Key Research and Development Program of China (2021YFC2501800). National High Level Hospital Clinical Research Funding (2022-PUMCH-B-110). Rui E Emergency Medicine Research Special Fund Project (R2021007). 2019 Discipline Development Project of Peking Union Medical College faculty development program in family medicine (No.201920200106).

## Availability of data and materials

All data were collected from publicly available literatures, and all data generated or analyzed during this study are included in this published article and its additional files.

## Declarations

### Ethics approval and consent to participate

Not applicable.

### Consent for publication

Not applicable.

### Competing interests

The authors declare that they have no competing interests.

## Author details

<sup>1</sup>Emergency Department, State Key Laboratory of Complex Severe and Rare Diseases, Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, Beijing, China. <sup>2</sup>Department of Family Medicine & Division of General Internal Medicine, Department of Medicine, Peking Union Medical College Hospital, Chinese Academy of Medical Sciences, State Key Laboratory of Complex Severe and Rare Diseases, Beijing, China. <sup>3</sup>Department of Critical Care Medicine, The Second Affiliated Hospital, Zhejiang University School of Medicine, Hangzhou, China. <sup>4</sup>Department of Emergency Medicine, Penn State Health Milton S. Ebersole Medical Center and Penn State College of Medicine, Hershey, PA, USA. <sup>5</sup>Department of Cardiology, State Key Laboratory of Complex Severe and Rare Diseases, Peking Union Medical College Hospital, Chinese Academy of Medical Science and Peking Union Medical College, Beijing, China.

Received: 7 August 2022 Accepted: 8 November 2022

Published online: 15 November 2022

## References

- Drake TM, Riad AM, Fairfield CJ, Egan C, Knight SR, Pius R, et al. Characterisation of in-hospital complications associated with COVID-19 using the ISARIC WHO Clinical Characterisation Protocol UK: a prospective, multicentre cohort study. *Lancet*. 2021;398:223–37.
- Thomas SJ, Moreira ED, Kitchin N, Absalon J, Gurtman A, Lockhart S, et al. Safety and efficacy of the BNT162b2 mRNA Covid-19 vaccine through 6 months. *N Engl J Med*. 2021;385:1761–73.
- Voysey M, Clemens SAC, Madhi SA, Weckx LY, Folegatti PM, Aley PK, et al. Safety and efficacy of the ChAdOx1 nCoV-19 vaccine (AZD1222) against SARS-CoV-2: an interim analysis of four randomised controlled trials in Brazil, South Africa, and the UK. *Lancet*. 2021;397:99–111.
- WHO coronavirus (COVID-19) dashboard. <https://covid19.who.int>. Accessed 18 June 2022.
- Rahmati M, Shamsi MM, Khoramipour K, Malakoutinia F, Woo W, Park S, et al. Baseline physical activity is associated with reduced mortality and disease outcomes in COVID-19: a systematic review and meta-analysis. *Rev Med Virol*. 2022;32:e2349.
- Huang I, Lim MA, Pranata R. Diabetes mellitus is associated with increased mortality and severity of disease in COVID-19 pneumonia - a systematic review, meta-analysis, and meta-regression. *Diabetes Metab Syndr*. 2020;14:395–403.
- Kumar A, Arora A, Sharma P, Anikhandi SA, Bansal N, Singla V, et al. Is diabetes mellitus associated with mortality and severity of COVID-19? A meta-analysis. *Diabetes Metab Syndr*. 2020;14:535–45.

8. Guo L, Shi Z, Zhang Y, Wang C, Do Vale Moreira NC, Zuo H, et al. Comorbid diabetes and the risk of disease severity or death among 8807 COVID-19 patients in China: a meta-analysis. *Diabetes Res Clin Pract.* 2020;166:108346.
9. Rubino F, Amiel SA, Zimmet P, Alberti G, Bornstein S, Eckel RH, et al. New-onset diabetes in Covid-19. *N Engl J Med.* 2020;383:789–90.
10. Sathish T, Tapp RJ, Cooper ME, Zimmet P. Potential metabolic and inflammatory pathways between COVID-19 and new-onset diabetes. *Diabetes Metab.* 2021;47:101204.
11. Practical recommendations for the management of diabetes in patients with COVID-19 - PubMed. <https://pubmed.ncbi.nlm.nih.gov/32334646/>. Accessed 28 June 2022.
12. Rey-Reñones C, Martínez-Torres S, Martín-Luján FM, Pericas C, Redondo A, Vilaplana-Carnerero C, et al. Type 2 diabetes mellitus and COVID-19: a narrative review. *Biomedicines.* 2022;10:2089.
13. High-dimensional characterization of post-acute sequelae of COVID-19 | Nature. <https://www.nature.com/articles/s41586-021-03553-9>. Accessed 24 June 2022.
14. Rahmati M, Keshvari M, Mirnasuri S, Yon DK, Lee SW, Il Shin J, et al. The global impact of COVID-19 pandemic on the incidence of pediatric new-onset type 1 diabetes and ketoacidosis: a systematic review and meta-analysis. *J Med Virol.* 2022;94:5112–27.
15. Banerjee M, Pal R, Dutta S. Risk of incident diabetes post-COVID-19: a systematic review and meta-analysis. *Prim Care Diabetes.* 2022. <https://doi.org/10.1016/j.pcd.2022.05.009>.
16. Stroup DF, Berlin JA, Morton SC, Olkin I, Williamson GD, Rennie D, et al. Meta-analysis of observational studies in epidemiology: a proposal for reporting. Meta-analysis Of Observational Studies in Epidemiology (MOOSE) group. *JAMA.* 2000;283:2008–12.
17. The PRISMA 2020 statement: an updated guideline for reporting systematic reviews. *BMJ.* <https://www.bmj.com/content/372/bmj.n71>. Accessed 26 June 2022.
18. Ottawa Hospital Research Institute. [https://www.ohri.ca/programs/clinical\\_epidemiology/oxford.asp](https://www.ohri.ca/programs/clinical_epidemiology/oxford.asp). Accessed 19 June 2022.
19. Freeman MF, Tukey JW. Transformations related to the angular and the square root. *Ann Math Stat.* 1950;21:607–11.
20. DerSimonian R, Laird N. Meta-analysis in clinical trials. *Control Clin Trials.* 1986;7:177–88.
21. Measuring inconsistency in meta-analyses. *BMJ.* <https://www.bmj.com/content/327/7414/557.short>. Accessed 19 June 2022.
22. Egger M, Davey Smith G, Schneider M, Minder C. Bias in meta-analysis detected by a simple, graphical test. *BMJ.* 1997;315:629–34.
23. Kossmeier M, Tran US, Voracek M. Power-enhanced funnel plots for meta-analysis: the sunset funnel plot; 2019.
24. Mathur MB, VanderWeele TJ. Sensitivity analysis for unmeasured confounding in meta-analyses. *J Am Stat Assoc.* 2020;115:163–72.
25. Mathur MB, VanderWeele TJ. Methods to address confounding and other biases in meta-analyses: review and recommendations. *Annu Rev Public Health.* 2022;43:19–35.
26. Haneuse S, VanderWeele TJ, Arterburn D. Using the E-value to assess the potential effect of unmeasured confounding in observational studies. *JAMA.* 2019;321:602–3.
27. R: the R project for statistical computing. <https://www.r-project.org/>. Accessed 19 June 2022.
28. Team R. R: a language and environment for statistical computing. undefined. 2014.
29. Barrett CE. Risk for newly diagnosed diabetes 30 days after SARS-CoV-2 infection among persons aged 18 years — United States, March 1, 2020–June 28, 2021. *MMWR Morb Mortal Wkly Rep.* 2022;71.
30. Cohen K, Ren S, Heath K, Dasmariñas MC, Jubilo KG, Guo Y, et al. Risk of persistent and new clinical sequelae among adults aged 65 years and older during the post-acute phase of SARS-CoV-2 infection: retrospective cohort study. *BMJ.* 2022;376:e068414.
31. Rathmann W, Kuss O, Kostev K. Incidence of newly diagnosed diabetes after Covid-19. *Diabetologia.* 2022;65:949–54.
32. Birabaharan M, Kaelber DC, Pettus JH, Smith DM. Risk of new-onset type 2 diabetes in 600 055 people after COVID-19: a cohort study. *Diabetes Obes Metab.* 2022;24:1176–9.
33. Qeadan F, Tingey B, Egbert J, Pezzolesi MG, Burge MR, Peterson KA, et al. The associations between COVID-19 diagnosis, type 1 diabetes, and the risk of diabetic ketoacidosis: a nationwide cohort from the US using the Cerner Real-World Data. *PLoS One.* 2022;17:e0266809.
34. Collaborative, Tazare J, Walker AJ, Tomlinson L, Hickman G, Rentsch CT, et al. Rates of serious clinical outcomes in survivors of hospitalisation with COVID-19 in England: a descriptive cohort study within the OpenSAFELY platform. *Wellcome Open Research;* 2022.
35. Xie Y, Al-Aly Z. Risks and burdens of incident diabetes in long COVID: a cohort study. *Lancet Diabetes Endocrinol.* 2022;10:311–21.
36. Ayoubkhani D, Khunti K, Nafilyan V, Maddox T, Humberstone B, Diamond I, et al. Post-covid syndrome in individuals admitted to hospital with covid-19: retrospective cohort study. *BMJ.* 2021;372:n693.
37. Daugherty SE, Guo Y, Heath K, Dasmariñas MC, Jubilo KG, Samranvedhya J, et al. Risk of clinical sequelae after the acute phase of SARS-CoV-2 infection: retrospective cohort study. *BMJ.* 2021;373:n1098. <https://www.bmj.com/content/373/bmj.n1098.long>. Accessed 8 June 2022.
38. Yang JK, Feng Y, Yuan MY, Yuan SY, Fu HJ, Wu BY, et al. Plasma glucose levels and diabetes are independent predictors for mortality and morbidity in patients with SARS. *Diabet Med.* 2006;23:623–8.
39. Yang J-K, Lin S-S, Ji X-J, Guo L-M. Binding of SARS coronavirus to its receptor damages islets and causes acute diabetes. *Acta Diabetol.* 2010;47:193–9.
40. Apicella M, Campopiano MC, Mantuano M, Mazoni L, Coppelli A, Del Prato S. COVID-19 in people with diabetes: understanding the reasons for worse outcomes. *Lancet Diabetes Endocrinol.* 2020;8:782–92.
41. Kusmartseva I, Wu W, Syed F, Van Der Heide V, Jorgensen M, Joseph P, et al. Expression of SARS-CoV-2 entry factors in the pancreas of normal organ donors and individuals with COVID-19. *Cell Metab.* 2020;32:1041–1051.e6.
42. Dungan KM, Braithwaite SS, Preiser J-C. Stress hyperglycaemia. *Lancet.* 2009;373:1798–807.
43. Müller JA, Groß R, Conzelmann C, Krüger J, Merle U, Steinhart J, et al. SARS-CoV-2 infects and replicates in cells of the human endocrine and exocrine pancreas. *Nat Metab.* 2021;3:149–65.
44. Fabbri A, Marchesini G, Benazzi B, Morelli A, Montesi D, Bini C, et al. Stress hyperglycemia and mortality in subjects with diabetes and sepsis. *Crit Care Explor.* 2020;2:e0152.
45. de Carvalho VF, Guedes Cocate P, Gonçalves Pereira L, de Cássia Gonçalves Alfenas R. The role of hyperglycemia in the induction of oxidative stress and inflammatory process. *Nutr Hosp.* 2012;27:1391–8.
46. Dave GS, Kalia K. Hyperglycemia induced oxidative stress in type-1 and type-2 diabetic patients with and without nephropathy. *Cell Mol Biol (Noisy-le-grand).* 2007;53:68–78.
47. Mariatulqabiah AR, Buttigieg KR. COVID-19 vaccinations for children. *Lancet Infect Dis.* 2022;S1473-3099(22):00414–5.
48. Stephanie. Propensity score matching: definition & overview. *Statistics How To.* 2017. <https://www.statisticshowto.com/propensity-score-matching/>. Accessed 28 June 2022.
49. Deeks JJ, Dinnes J, D'Amico R, Sowden AJ, Sakaravitch C, Song F, et al. Evaluating non-randomised intervention studies. *Health Technol Assess.* 2003;7.
50. Van Der Weele TJ, Ding P. Sensitivity analysis in observational research: introducing the E-value. *Ann Intern Med.* 2017;167:268–74.
51. Rathmann W, Bongaerts B, Carius H-J, Kruppert S, Kostev K. Basic characteristics and representativeness of the German Disease Analyzer database. *Int J Clin Pharmacol Ther.* 2018;56:459–66.
52. NHS England » OpenSAFELY – the coronavirus (COVID-19) research platform. <https://www.england.nhs.uk/contact-us/privacy-notice/how-we-use-your-information/covid-19-response/coronavirus-covid-19-research-platform/>. Accessed 28 Jun 2022.
53. Herbert A, Wijlaars L, Zylbersztejn A, Cromwell D, Hardelid P. Data resource profile: Hospital Episode Statistics Admitted Patient Care (HES APC). *Int J Epidemiol.* 2017;46:1093–1093i.

## Publisher's Note

Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.