

# Transmissibility of COVID-19 among vaccinated individuals

A Rapid Literature Review: Update #2

Date of Literature Search: 8/23/2021 Date of Submission: 9/24/2021

## **Prepared By:**

Charleen Salmon, Jordyn Flanagan, Brenlea Farkas, Liza Mastikhina, Oluwaseun Egunsola, Lynora Saxinger, Becky Skidmore, Fiona Clement, on behalf of the University of Calgary Health Technology Assessment Unit.

### Contact:

**Fiona Clement** 

Email: fclement@ucalgary.ca

**Suggested citation:** Salmon C, Flanagan J, Farkas B, Mastikhina L, Egunsola O, Saxinger L, Skidmore B, Clement FM on behalf of the University of Calgary Health Technology Assessment Unit. Transmissibility of COVID-19 among Vaccinated Individuals: A Rapid Literature Review, Update #2. September 24, 2021







### Funding Acknowledgement(s)

The SPOR Evidence Alliance (SPOR EA) is supported by the Canadian Institutes of Health Research (CIHR) under the Strategy for Patient-Oriented Research (SPOR) initiative.

COVID-19 Evidence Network to support Decision-making (<u>COVID-END</u>) is supported by the Canadian Institutes of Health Research (<u>CIHR</u>) through the Canadian 2019 Novel Coronavirus (COVID-19) Rapid Research Funding opportunity.

### **Project Contributors**

Charleen Salmon<sup>1,2</sup>; Jordyn Flanagan<sup>1,2</sup>; Brenlea Farkas<sup>1,2</sup>; Liza Mastikhina<sup>1,2</sup>; Oluwaseun Egunsola<sup>1,2</sup>; Lynora Saxinger<sup>3</sup>; Becky Skidmore<sup>4</sup>; and Fiona Clement<sup>1,2</sup>

Affiliations

- 1. The Department Community Health Sciences, Teaching Research and Wellness Building, 3280 Hospital Drive NW Calgary Alberta T2N 4N1
- 2. O'Brien Institute for Public Health, Teaching Research and Wellness Building, 3280 Hospital Drive NW Calgary Alberta T2N 4N1
- 3. Department of Medicine University of Alberta, Edmonton, Alberta, Canada
- 4. Ottawa Hospital Research Institute, University of Ottawa, Ottawa, Ontario, Canada

#### **Third-Party Materials**

If you wish to reuse non-textual material from this report that is attributed to a third party, such as tables, figures or images, it is your responsibility to determine whether permission is required for such use and to obtain necessary permission from the copyright holder. The risk of claims resulting from infringement of any third-party-owned material rests solely with the user.

### **General Disclaimer**

This report was prepared by The University of Calgary Health Technology Assessment Unit on behalf of the SPOR Evidence Alliance and COVID-END. It was developed through the analysis, interpretation and synthesis of scientific research and/or health technology assessments published in peer-reviewed journals, institutional websites and other distribution channels. It also incorporates selected information provided by experts and patient partners with lived experience on the subject matter. This document may not fully reflect all the scientific evidence available at the time this report was prepared. Other relevant scientific findings may have been reported since completion of this synthesis report.

SPOR Evidence Alliance, COVID-END and the project team make no warranty, express or implied, nor assume any legal liability or responsibility for the accuracy, completeness, or usefulness of any information, data, product, or process disclosed in this report. Conclusions drawn from, or actions undertaken on the basis of, information included in this report are the sole responsibility of the user.





# **Table of Contents**

Abbreviations and Definitions
Abbreviations3
KEY POINTS
EXECUTIVE SUMMARY
Introduction12
Methods13
Results
Study Characteristics
Risk of Bias Assessment
Vaccine Effectiveness against Infection Transmission
Vaccine Efficacy or Effectiveness Against Asymptomatic Infection
AstraZeneca Vaccine Efficacy in the General Population
AstraZeneca Vaccine Effectiveness in the General Population
Pfizer BioNTech Vaccine Effectiveness in the General Population
mRNA (Pfizer BioNTech and Moderna) Vaccines Effectiveness in the General Population
Moderna Vaccine Efficacy in the General Population43
Moderna Vaccine Effectiveness in the General Population
Janssen Vaccine Efficacy in the General Population
Cycle Threshold (Ct) Values
Viral Load
Discussion
Emerging Evidence
Recommendations
Conclusion
References
Appendix 1: Search Strategy64



# **Abbreviations and Definitions**

## Abbreviations

AZ CDC Ct	AstraZeneca ChAdOx1 nCoV-19 vaccine Centres for Disease Control and Prevention Cycle threshold
COVID-19	Coronavirus Disease 2019
IQR	Interquartile range
J and J	Janssen Ad26.COV2.S
mRNA	Messenger ribonucleic acid
mRNA-1273	Moderna's mRNA vaccine
NR	Not Reported
PCR	Polymerase chain reaction
PfBnT	Pfizer BioNTech's BNT162b2
RCT	Randomized controlled trial
ROBINS-I	Risk of bias for non-randomized studies
SARS-Cov-2	Severe Acute Respiratory Syndrome Coronavirus 2
VOC	Variant of concern
WHO	World Health Organization



# **KEY POINTS**

- Forty-five studies, including six RCTs and 39 observational studies were included in this review.
- COVID-19 vaccination have been demonstrated to be associated with varied degrees of reduced household transmission of SARS-CoV-2, reduced incidence of asymptomatic infection, and a reduction in viral load, although data on the most recently emergent Delta VOC is required.
- Evidence from four large household surveillance studies from the Netherlands, Finland, and Israel suggests that full-dose of AZ, PfBnT, Moderna, or J&J vaccines may prevent significantly reduced household transmission of wild-type or the B.1.1.7 (Alpha) COVID-19 strain after 14 days of vaccination by at least 63%. No studies on vaccine effectiveness against infection transmission of the B.1.617.2 (Delta) strain were found.
- The AZ vaccine trial in the general population suggest that an initial low dose 1 followed with an extended interval standard dose 2 may provide up to 59% protection against asymptomatic or unknown infection. Efficacy against these outcomes was not demonstrated following two standard doses given at a short interval of ?1 mo. The higher efficacy in the low dose study results is felt to be partially explained by the extended interval before dose 2 in that subgroup, which has subsequently been shown to offer higher overall efficacy. A first dose of AZ vaccine was associated with significantly reduced odds (ORs between 0.39 and 0.45) of asymptomatic infection in another observational study.
- Asymptomatic infection is felt to be a risk for post-vaccination transmission, so reduction of asymptomatic infection is a useful end point.
- PfBnT vaccine observational studies in the general population suggest up to 90% effectiveness against asymptomatic infection after seven or more days of full-dose vaccination. For healthcare workers there was up to 75% effectiveness of asymptomatic infection against the wild-type strain after full-dose of the PfBNT vaccine.
- In the general population, vaccine effectiveness of a full dose of PfBNT was reported to be 35.9% effective at reducing asymptomatic infection against the B.1.617.2 (Delta) variant.
- Moderna vaccine observational studies in the general population suggest up to 57.4% effectiveness against asymptomatic infection of the B.1.617.2 (Delta) variant after 14 or more days of full-dose vaccination.
- In a community RCT assessing asymptomatic infection by collection of a RT-PCR swab at the dose 2 visit, a single dose of the Moderna vaccine showed efficacy of 61.4% against asymptomatic infection against wild-type SARS-CoV-2 in adults and 59.5% in adolescents aged 12-17 years. No vaccine efficacy for the full series was reported for adults but data from the adolescent trial of the Moderna vaccine showed two doses two weeks after vaccination had an efficacy of 39.2% against asymptomatic infection of the wild-type strain. The single dose protocol J&J vaccine had an efficacy



of 74% against asymptomatic infection after 28 days of vaccination for adults against the wild-type strain.

- Thirteen of the 20 studies reporting cycle threshold values found significantly increased cycle threshold, suggestive of a lower viral load, in PfBnT, Moderna, or AZ vaccinated individuals compared with unvaccinated. The two included studies reporting viral load found a significant reduction in the viral RNA load in mRNA-based vaccinated individuals.
- Further research is needed to evaluate post-vaccination infectivity and transmission of both the wild type COVID-19 virus and the variants of concern especially the B.1.617.2 (Delta) variant.



# EXECUTIVE SUMMARY

**Objectives:** This is an update of a previous report with a literature search that ended May 4, 2021.<sup>1</sup> A total of 25 additional studies were included in this update for a total of 45 studies. The objective of this report is to identify comparative observational studies and randomized controlled trials (RCTs) evaluating the efficacy and effectiveness of COVID-19 vaccination in reducing forward transmission from vaccinated people, and studies examining the biological plausibility of vaccination induced transmission reduction. There is evolving data around the frequency of asymptomatic COVID-19 and whether the viral load, and therefore infectiousness, is lower among people who develop COVID-19 post-vaccination compared with those who have not been vaccinated. Viral presence is an imperfect proxy of transmissibility although the quantity of virus present does appear to influence risk, as studies document transmission risk is higher with a higher viral load or lower Ct value. Since most COVID-19 vaccine trials use an endpoint of symptomatic infection, there is less data about whether this incurs a risk for viral transmission from vaccinated persons.

Design: Rapid review with grey literature search.

**Method:** A search of databases, MEDLINE, Embase, L-OVE and the Cochrane Central Register of Controlled Trials was conducted to identify RCTs or comparative observational studies evaluating the efficacy and effectiveness of COVID-19 vaccination in the prevention of transmission, asymptomatic infections, and transmissibility of COVID-19 among vaccinated persons. An additional search of grey literature was conducted, including: Clinicaltrials.gov, McMaster Health Forum (COVID-END), MedRxiv, Google, regulatory submissions, and the websites of the Centres for Disease Control and Prevention (CDC) and World Health Organization (WHO). Abstracts were screened by a single reviewer and then reviewed in full text by two independent reviewers. This search is current to August 23, 2021.

A standardized data extraction sheet was used to extract the year of publication, country, study design, patient characteristics including sex, gender and age, variants of COVID-19, seroprevalence, and all the reported outcomes of interest. Quality assessment was conducted based on study design: ROBINS-I for non-randomized studies and Cochrane Risk of Bias for human-subject RCTs. Data were extracted by one reviewer and verified by another. Animal studies were not included in this update.

**Results:** In this update, 25 additional studies were included. Therefore, this review has a total of 45 included studies. Four new studies on COVID-19 transmission to household contacts were included.

**Reduction of household transmission:** A retrospective cohort study in the Netherlands by de Gier et al. of 113,582 confirmed index cases of COVID-19 and 253,168 cohabitating household members or close contacts were assessed for vaccine effectiveness in preventing transmission to the household member or close contact and stratified by vaccination status, vaccine type, and days past date of inoculation.<sup>2</sup> At least one dose of PfBNT, Moderna, AZ, or J&J from past the 14<sup>th</sup> day of vaccination onwards was associated with the reduction of



transmission of COVID-19 to any household contact by 21% (95% CI: 12-28), 23% (95% CI: 14-32) to any unvaccinated household contact, 22% (95% CI: 9-33) to any other close contact, and 22% (95% CI: 8-34) to any unvaccinated close contact.<sup>2</sup> Fully vaccinated individuals with either PfBNT, Moderna, J&J, or AZ from past the 7<sup>th</sup> day of vaccination onwards, was associated with the reduction of transmission of COVID-19 to any household contact by 71% (95% CI: 63-77), 73% (95% CI: 65-79) to any unvaccinated household contact, 22% (95% CI: -5-43) to any other close contact, and 24% (95% CI: -5-43) to any unvaccinated close contact.<sup>2</sup> The low vaccine effectiveness of a fully vaccinated individual against transmission to any close contact or any close unvaccinated contact could be due to the studies being underpowered to detect differences due to the small number of events that occurred in vaccinated individuals compared to unvaccinated index cases.

A similar study was conducted in Finland by Salo et al. This retrospective cohort study investigated the vaccine effectiveness of 95,138 mRNA-based (PfBNT or Moderna) vaccinated healthcare workers against infection transmission to unvaccinated household members compared to unvaccinated healthcare workers and their unvaccinated household members.<sup>3</sup> At least one dose of an mRNA-based vaccine from the 14<sup>th</sup> day of vaccination onward reduced transmission to an unvaccinated spouse by 8.7% (95% CI: -28.9-35.4) and increased to 42.9% (95% CI: 22.3-58.1) reduction in transmission 10 weeks after the first dose.<sup>3</sup> At least one dose of an mRNA-based vaccine from the 14<sup>th</sup> day of vaccination onward increased transmission to an unvaccinated child living in the household between the ages of 3-18 years by 1.0% (95% CI: -53.9-33.7) and decreased transmission to the unvaccinated child by 32.9% (95% CI: 4.1-53.0) 10 weeks after the first dose.<sup>3</sup>

Two studies from Israel found that PfBNT fully vaccinated individuals from past the 7<sup>th</sup> day of vaccination onward had reduced infection transmission to their household contacts.<sup>4,5</sup> A retrospective cohort study using a nationally centralized database investigated the vaccine effectiveness of PfBNT against infection transmission of two-adult households only and one confirmed case of infection during the study period.<sup>4</sup> Of households with a fully vaccinated adult, the PfBNT vaccine was found to reduce infection transmission of the wild-type strain by 80.0% (95% CI: 73.0-85.1) compared to those who were unvaccinated and by 82.0% (95% CI: 75.5-86.7) compared to those who were recently vaccinated with one dose (between 0-7 days after vaccination).<sup>4</sup> A second Israeli study by Layan et al. conducted a case-control study of the PfBNT vaccine's effectiveness on reduction of infection transmission of the wild-type and B.1.1.7 (Alpha) strains in healthcare workers and their households.<sup>5</sup> The risk of transmission from vaccinated cases was 0.22 times (95% CI: 0.06-0.70) the risk of infection transmission compared to unvaccinated cases.<sup>5</sup>

The baseline serology and PCR of household contacts were not reported in any of the studies except for Salo et al. and Gazit et al. who only included seronegative participants.<sup>3,4</sup>

**Reduction of asymptomatic test positive status after vaccination, various populations**: Asymptomatic with lab documented infection data were presented in the UK component of the AstraZeneca ChAdOx1 nCoV-19 (AZ vaccine) vaccine studies. Participants were assessed by weekly self-administered nose and throat swabs for RT-PCR testing. The vaccine demonstrated efficacy against any PCR positive results compared with control in two studies, (67% 95% CI: 49-78)<sup>6</sup> and 46.3% (31.8-57.8)<sup>7</sup>, respectively, after 21 days following the first



dose. However, the AZ vaccine standard dose was reported not to have significant efficacy against asymptomatic or unknown carriage with the wild type virus after 21 days of the first dose (7.8% (95% CI: -46.7-42.1) and after 14 days of the second dose 27.3% (95% CI: -17-54.9)<sup>7</sup> respectively.

Several studies found that a full-dose of PfBNT or Moderna significantly reduced asymptomatic infection from the wild-type strain.<sup>8-12</sup> Tang et al. found a reduction in transmission of asymptomatic infection of fully vaccinated seronegative Qatari healthcare workers between 0-6 days past the date of vaccination (IRR: 0.35 [95% CI: 0.11-1.09]) and from more than 7 days past the date of vaccination (IRR: 0.10 [95% CI: 0.04-0.22]).<sup>8</sup> This finding was supported by Angel et al. who found similar significant reductions in asymptomatic infection.<sup>9</sup> A retrospective cohort study by Andrejko et al. of 525 seronegative California residents found that a full-dose of PfBNT had a 68.3% (95% CI: 27.9-85.7%) reduction of asymptomatic infection of the wild-type strain.<sup>10</sup>

Dagan et al. demonstrated 90% effectiveness (95% CI: 83-94) against asymptomatic infection seven days after the second dose from the wild-type or B.1.17 (Alpha) strain.<sup>13</sup> In an Israeli study, which utilized the national public health surveillance data, Haas et al. reported significantly higher vaccine effectiveness seven or more days after full-dose PfBnT vaccination, 90.4% (95% CI: 89.1-91.5).<sup>14</sup>

Tande et al. evaluated the effectiveness of at least one dose of either Moderna or PfBnT vaccine among people who underwent molecular tests prior to a procedure or surgery.<sup>15</sup> The relative risk for a positive test during asymptomatic pre-procedure screening in vaccinated compared with unvaccinated was significantly lower (0.44 (95% CI: 0.33-0.60)). Ten or more days after the 1st dose, the risk of a positive test was also significantly lower among the vaccinated (0.28 (95% CI: 0.16-0.49; p<.0001)). The risk of test positivity was similarly lower among the vaccinated after the second dose 0.27 (95% CI: 0.12-0.60).<sup>15</sup> Lastly, Chemaitelly et al. found that a full-dose of Moderna was 92.5% effective (95% CI: 84.8-96.9) against asymptomatic carriage 14 days after full vaccination.<sup>11</sup>

There is limited evidence suggesting that mRNA-based vaccines have protection against asymptomatic carriage of the B.1.617.2 (Delta) variant. Tang et al. found that use of at least one dose of an mRNA-based vaccine past the 14<sup>th</sup> day onward from the date of vaccination, reduced asymptomatic infection by 44.3% (95% CI: 0-78.4) against the B.1.617.2 (Delta) variant.<sup>12</sup> The study conducted by Tang et al., stratified by vaccine manufacturer and found that a full dose of PfBNT past the 14<sup>th</sup> day onward from the date of vaccination was found to reduce asymptomatic infection by 35.9% (95% CI: 11.1-53.9) against the B.1.617.2 (Delta) variant<sup>12</sup> and that a full-dose of Moderna had 80.2% vaccine effectiveness (95% CI: 54.2-92.6) against asymptomatic carriage of the Delta strain 14 days after full vaccination<sup>12</sup>

**Possible reduction of viral load / higher Ct values in vaccinated persons, population data:** Twenty studies reported on Ct, an inverse proxy for viral load and two studies reported on viral load.

Results from Phase 2/3 vaccine efficacy studies of AZ vaccine compared with a comparator meningococcal vaccine in the United Kingdom, showed that the Ct values in infected vaccinated participants were statistically significantly higher than the comparator (p<0.0001),



after 14 days of the second dose in baseline seronegative efficacy cohorts.<sup>16</sup> Furthermore, the vaccine recipients were PCR-positive for a significantly shorter period of time (p<0.0001). The Ct values in asymptomatic cases were also significantly higher among vaccine recipients than control (p=0.0040); however, this difference was not significant for primary symptomatic cases (p=0.1534). Vaccine recipients infected with the B.1.1.7 variant also showed significantly higher Ct values than control (p=0.0113).<sup>16</sup>

A longitudinal UK household survey by Pritchard el al. found statistically significant increase in the median Ct values of PfBnT or AZ single or full dose vaccinated individuals compared with unvaccinated individuals at any time point before or after 21 days post-vaccination (p<0.001).<sup>17</sup> Similarly, in another UK study by Shrotri et al., the mean Ct value of unvaccinated individuals within 27 days of vaccination was 26.6 (95% CI: 26-27.1) compared with 26.6 (95% CI: 25.19-26.62) with one dose of PfBnT or AZ, which was not significantly different (p=0.158).<sup>18</sup> However, after 28 days, there was a statistically significant decrease in the mean Ct between vaccinated and unvaccinated persons (mean Ct 26.6 (95% CI: 26-27.1) vs 31.3 (95% CI: 29.6-32.9), p<0.001).<sup>18</sup> Monthly routine PCR testing was conducted in these patients; however, the baseline serology was not reported.<sup>18</sup> In a longitudinal cohort study of HCWs who were offered voluntary nasal and oropharyngeal swab PCR testing every two weeks as well as serological testing, a small study of 49 people vaccinated with either PfBNT or AZ and 96 unvaccinated people in the USA by Mostafa et al. demonstrated non-significant differences in median Ct values (19.26 [Q1, Q3: 16.56-21.96] vs 19.6 [Q1, Q3: 16.28-22.66], respectively).<sup>19</sup> Similar non-significance in the median Ct values of PfBNT or AZ-vaccinated people vs unvaccinated was found in a UK study by Baltas et al. (Median=30.8 [IQR: 25.9-35.4] vs. Median=28.8 [IQR: 25.3-33.7], p=0.053).<sup>20</sup> Lastly, Lumley et al., found vaccination with either PfBnT or AZ to non-significantly increase Ct value by a mean of 2.7.21

A retrospective study of PfBnT mRNA vaccine recipients compared with demographically matched control group of unvaccinated individuals in Israel, found no significant differences in the Ct values for any of the three genes (RdRp, N and E) measured less than 12 days after the first dose in infected persons. However, between 12 and 28 days after the first dose, the Ct values for the three genes were significantly higher among infected vaccinated persons than controls (p<10<sup>-8</sup>).<sup>22</sup> In another UK study of one dose of BNT162b2 vaccine, the median Ct values of infected HCWs were reported to have shown a non-significant trend towards increase between unvaccinated (median=20.3) and vaccinated HCWs after 12 days postvaccination (median=30.3), suggesting that samples from infected vaccinated individuals had lower viral loads.<sup>23</sup> A study by McEllistrem et al. among community living centre residents reported five cases of asymptomatic infections (determined by surveillance nasal swabs every 2-5 days) among baseline PCR negative PfBnT vaccinated and unvaccinated residents. The median Ct values among unvaccinated residents (12.8, IQR: 12.4-14.9) were significantly lower (p=0.009) than vaccinated residents (19.4, IQR: 18.9-25.5).<sup>24</sup> Furthermore, viral load was -2.4 mean log10 lower among the vaccinated cohort (p=0.004).<sup>24</sup> In another large cohort study of HCWs at a large medical centre in Israel by Regev-Yochay et al, the mean Ct values among PfBNT fully vaccinated HCWs (27.3±2.2) was significantly higher (mean difference 5.09, 95% CI: 2.8-7.4, p<0.001) than unvaccinated HCWs (22.2±1.0).<sup>25</sup> A matched casecontrol study by Abu-Raddad et al. from Qatar, evaluating the Ct values of people with two doses of PfBNT with breakthrough infections compared to Ct values of infections in



unvaccinated individuals, found statistically significant higher median Ct values in vaccinated individuals (27.8; IQR: 21.1-32.7) than the median Ct value of unvaccinated individuals (25.8 (IQR: 19.5-31.4; p<0.001).<sup>26</sup> However, studies in France<sup>27</sup> and Greece<sup>28</sup> found no statistically significant differences between PfBNT vaccinated individuals' Ct values and the Ct values of those who were unvaccinated. Bailly et al. found that the Ct values of PfBNT fully vaccinated long-term care residents did not differ from the Ct values of unvaccinated residents (Median=21 [IQR:13-32] vs 15 [IQR: 12-17]; p=0.05).<sup>27</sup> Similarly, Ioannou et al.'s study of fully vaccinated healthcare workers in a Greek hospital amidst an outbreak found no significant differences between the median Ct values of those vaccinated and unvaccinated (18 [15.5-25.5] vs 18.5 [13.5-24]).<sup>28</sup>

A USA study investigating the Ct values of mRNA-based vaccinated healthcare workers (PfBNT or Moderna) from unvaccinated healthcare workers found that there was no statistically significant difference in mean Ct values in the early post-vaccination period defined as less than 14 days post vaccination  $(22.6\pm7 \text{ vs. } 23\pm7.4)$  for partially-vaccinated healthcare workers more than 14 days past first dose but before the second dose  $(27.7\pm8.7 \text{ vs. } 23\pm7.4)$ , or for fully vaccinated healthcare workers at least 14 days past vaccination  $(28.5\pm7.4 \text{ vs. } 23\pm7.4)$ .<sup>29</sup> Another similar study by Duerr et al. reported Ct values for vaccinated individuals in the community but lumped all unvaccinated comparators as under a Ct value equal or less than 30 therefore, no effect size was presented.<sup>30</sup>

Two USA studies reported on viral load, one study is new to this updated version of the report. A prospective cohort study of baseline seronegative vaccinated and unvaccinated healthcare workers across Arizona had their mid-turbinate nasal swabs assessed for viral load.<sup>31</sup> Thompson et al. found that the mean viral RNA load for partially and fully vaccinated healthcare workers, with a mRNA-based vaccine, who were at least 14 days past the date of vaccination had lower presence of virus compared to their unvaccinated counterparts (2.3±1.7 Log<sub>10</sub> copies/mL vs.  $3.8\pm1.7$  Log<sub>10</sub> copies/mL).<sup>31</sup> This represented at least 40.2% lower viral RNA load after at least partial vaccination.<sup>31</sup>

A second retrospective cohort study of five vaccinated and five unvaccinated asymptomatic nursing home residents in a single nursing home evaluated the effectiveness of at least one dose of the PfBNT vaccine on attenuating viral load.<sup>24</sup> Viral load was -2.4 mean log10 lower among the vaccinated cohort (p=0.004).<sup>24</sup>

**Data from vaccine efficacy trials where asymptomatic RT-PCR swabs were collected:** Baden et al. showed that, among participants who received the first dose of the Moderna vaccine while negative for COVID-19 by RT-PCR or antibody testing at baseline, 0.1% had positive swabs but no symptoms at the time of their second dose, compared with 0.27% of the unvaccinated group, which is suggestive of 61.4% efficacy against asymptomatic carriage.<sup>32</sup>

Among participants who were seronegative at baseline (defined as negative RT-PCR and negative serology against SARS-CoV-2 nucleocapsid on day 1), the Ad26.COV2.S vaccine by Janssen Biotech (J&J vaccine), did not show efficacy against asymptomatic infection in the first 28 days of follow-up. However, the vaccine demonstrated 74% (95% CI: 46.8-88.4) efficacy after 28 days. Asymptomatic infection was assessed by lack of symptoms on the day



preceding, the day of, or any time after a positive PCR test. The frequency of swabbing for PCR testing was not reported in this study.<sup>33</sup>

**Emerging Evidence:** Variants of SARS-CoV-2 continue to surface, and the B.617.2 (Delta) variant is currently the one of most concern.<sup>34,35</sup> There has been emerging evidence that indicates that although a full vaccination series might reduce an individual's overall risk of becoming infected, there seems to be a limited difference in the Ct values between vaccinated and unvaccinated.<sup>34,36</sup> Certain outbreaks amongst vaccinated individuals in the USA have led to CDC recommendations for expanded prevention strategies such as universal masking in indoor spaces.<sup>35</sup>

**Conclusion:** Four months since the publication of the previous version of this report, 24 additional relevant studies have been published. Four of these were large household surveillance studies from the Netherlands, Finland, and Israel suggesting that a full dose of PfBNT, Moderna, AZ, or J&J vaccines may prevent household transmission after 14 days of vaccination. Twelve additional studies found that vaccines significantly reduce the risk of asymptomatic infection, with multiple studies finding that vaccines decreased the viral RNA load or increased the cycle threshold, suggestive of reduced viral load. Some studies, such as the RCTs investigating the AZ vaccine, included data on cross sectional prevalence of positive SARS-CoV-2 RT-PCR from routine swabbing, which suggested efficacy against asymptomatic infection, although this was not routinely assessed in a comparable way across studies. Evidence regarding the Ct values for the AZ, PfBnT, and Moderna vaccines suggest their potential to reduce viral load and possibly transmission. Further research is needed to evaluate post-vaccination infectivity and transmission of variants of concern especially the B.1.617.2 (Delta) strain from other jurisdictions.

Protocol/Topic Registration: PROSPERO-CRD42021252485.



# Introduction

Coronavirus disease (COVID-19) is caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2). As of late September 2021, there have been more than 229,903,892 confirmed cases of COVID-19, which have resulted in more than 4,715,485 confirmed deaths worldwide.<sup>37</sup> Since the start of the pandemic, several clinical trials have been conducted to examine the safety and effectiveness of different vaccines to prevent COVID-19. Many of these have found the vaccines to be generally effective against symptomatic COVID-19 infection, with an average efficacy of 85% (95% CI: 71 - 93%) after a full course of vaccination.<sup>38</sup> However, recent evidence suggests that vaccine efficacy/effectiveness may be reduced against certain variants of COVID-19, notably the delta variant, which has been contributing to a recent surge of cases worldwide. Real-world effectiveness of the two-dose vaccine regimen against the delta variant has ranged from 67.0% (95% CI, 61.3 to 71.8) to 88.0% (95% CI, 85.3 to 90.1).<sup>39</sup>

People who have started or finished the COVID19 vaccine series have been documented to have detectable SARSCoV2 by RT-PCR at various time points after vaccination,<sup>6</sup> although demonstration of cultivatable virus and definitive evidence of transmission post vaccination has not been assessed. It is not yet clear whether the current COVID-19 vaccines are as effective at reducing transmission as they are at reducing disease. Moreover, evaluating the ability of vaccinated individuals to transmit the virus after infection is challenging. Therefore, virologic surrogates of possible transmissibility may be a helpful way around this challenge.

Monoclonal antibody studies may provide useful insights into the pathophysiologic plausibility of vaccine induced transmission reduction, since they have been shown to result in circulating neutralizing antibody, with a significant decrease in quantitative viral load.<sup>40</sup> In one study, following quantitative reverse-transcriptase–polymerase-chain-reaction (RT-PCR) testing of nasopharyngeal swabs, an antibody cocktail was found to significantly reduce viral load compared with placebo.<sup>40</sup> The time-weighted average change in viral load in the first 7 days was –0.56 log10 copies per milliliter (95% CI, –1.02 to –0.11) among those who were serum antibody–negative at baseline.<sup>40</sup> Another study reported an elimination of more than 99.97% of viral RNA on day 11 after monoclonal antibody treatment.<sup>41</sup>

There is evolving data around the frequency of asymptomatic COVID-19 and if the viral load, and therefore infectiousness, is lower among people who develop COVID-19 post-vaccination compared with those who have not been vaccinated. Viral presence is an imperfect proxy of transmissibility although the quantity of virus present does appear to influence risk, as studies document transmission risk is higher with a higher viral load or lower Ct value.<sup>42,43</sup> Marks et al. found index viral load to be a major driver of transmission in a Spanish cohort,<sup>43</sup> with only 32% of index cases responsible for transmission, and an attack rate of 12% in contacts of index cases with a viral load <10<sup>6</sup> and 25% in contacts of index cases with a viral load of 10<sup>10</sup>. Similarly, Bjorkman et al. found that higher viral load increased SARS-CoV-2 transmission between asymptomatic residence hall roommates.<sup>44</sup> The index cases who transmitted infection had an average viral load 6.5 log higher than those who did not. Transmission from asymptomatic students to roommates occurred in 20% of rooms with an infected student, with



a lower mean Ct (E gene) of 26.2 in transmission index cases versus 28.9, (median 26.11 in transmission index cases versus 29.32).

However, the risks related to viral presence by RT-PCR may be modulated by individual's immune status, as viral persistence after natural infection has been observed in individuals with neutralizing antibody responses after natural infection, without transmission to close contacts.<sup>45</sup> Although asymptomatic and especially pre-symptomatic transmission of SARSCoV-2 has been well documented, existing studies suggest that transmission risk is lower from asymptomatic individuals than symptomatic individuals.<sup>46</sup>

The evidence for the transmissibility and transmission of COVID-19 infections in vaccinated individuals is rapidly evolving; therefore, the objective of this rapid review was to identify comparative observational studies and randomized controlled trials (RCTs) evaluating the effectiveness or efficacy of COVID-19 vaccination in reducing infection transmission, asymptomatic viral carriage, and other proxies of possible transmission, such as cycle threshold (Ct) values and viral load. This is an update of a previous report with a literature search that ended May 4<sup>th</sup>, 2021.<sup>1</sup>

## **Methods**

An experienced medical information specialist developed and tested the search strategies through an iterative process in consultation with the review team. The MEDLINE strategy was peer reviewed by another senior information specialist prior to execution using the PRESS Checklist.<sup>47</sup>

Using the multifile option and deduplication tool available on the OVID platform, we searched Ovid MEDLINE®, including Epub Ahead of Print, In-Process & Other Non-Indexed Citations, Embase, and EBM Reviews - Cochrane Central Register of Controlled Trials. We also searched for primary studies on the Living Overviews of Evidence (L-OVE) platform. We performed all searches on August 23, 2021.

The strategies utilized a combination of controlled vocabulary (e.g., "COVID-19 Vaccines", "COVID-19/tm [Transmission]", "Disease Transmission, Infectious") and keywords (e.g., "mRNA vaccine", "unvaccinated", "infectiousness"). Vocabulary and syntax were adjusted across the databases. The search strategies are in Appendix 1. No language or date limits were applied. Results were downloaded and deduplicated using EndNote version 9.3.3 (Clarivate Analytics) and uploaded to Word.

A grey literature search was also conducted, including: Clinicaltrials.gov, McMaster Health Forum (CoVID-END), MedRxiv, Google, regulatory submissions, and websites of the Center for Disease Control and Prevention (CDC) and World Health Organization (WHO). This search was limited to studies conducted since May 4, 2021, and current to August 23, 2021. There were no language limitations.

A screening form based on the eligibility criteria was prepared. Citations identified as potentially relevant from the literature search were screened by a reviewer, and subsequently



read in full text by two reviewers and assessed for eligibility based on the criteria outlined below (

Table 1). Discrepancies were resolved by discussion or by a third reviewer. Reference lists of included studies were hand searched to ensure all relevant literature is captured.

Population	Persons who had received COVID-19 vaccination irrespective of age, sex or gender. Animal studies were not included in this update.
Intervention	COVID-19 vaccination
Comparator	Non-vaccinated persons.
Outcome	Ct values, viral load, asymptomatic laboratory confirmed cases by RT-PCR post-vaccination and the number of persons who are infected by someone who has COVID-19 and has had the vaccine. Studies evaluating the transmissibility or infectivity of COVID-19 among vaccinated individuals were included.
Study	Comparative observational studies and RCTs evaluating the efficacy and
Design	effectiveness of COVID-19 vaccination in the prevention of asymptomatic
	viral infections as a proxy of a possible transmission were included. Studies eligible for inclusion had to have a control group of unvaccinated people.

A standardized data extraction sheet was used to extract the year of publication, country, study design, patient characteristics including sex, gender and age, variants of COVID-19, seroprevalence, and all the reported outcomes of interest (e.g., asymptomatic infection, transmission). All reviewers completed a calibration exercise whereby data from two sample studies were extracted by all four reviewers and areas of disagreement were discussed. Data were extracted by one reviewer and verified by another reviewer.

Quality assessment was conducted based on study design: Cochrane risk of bias for non-randomized studies (ROBINS-I) for non-randomized studies<sup>48</sup> and Cochrane Risk of Bias (version 5.1.0) for human-subject RCTs.<sup>49</sup> Quality assessment was conducted by one reviewer and verified by a second reviewer.

## **Results**

Twenty-one studies were included in the previous version of this work published in May 2021.<sup>1</sup> This current search (May 4, 2021 – August 23 2021) yielded 3,340 unique citations, 3,189 of which were excluded after abstract review (Figure 1). A total of 151 studies identified from the database search proceeded to full-text review. An additional 24 studies identified through grey literature search were also reviewed. In total, 150 studies were excluded for the following reasons: outcomes not of interest (n=102), duplicate (n=18), comparator not of interest (n=15), study design not of interest (n=7), intervention not of interest (n=2), population not of interest (n=1), and other (n=5).

Twenty-five new studies were included resulting in a total of 45 studies (Figure 1; 20 studies from the May version and 25 new studies). One study<sup>50</sup> included in the May version was not included in this update because it was a press release from March that could not be linked to

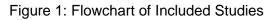


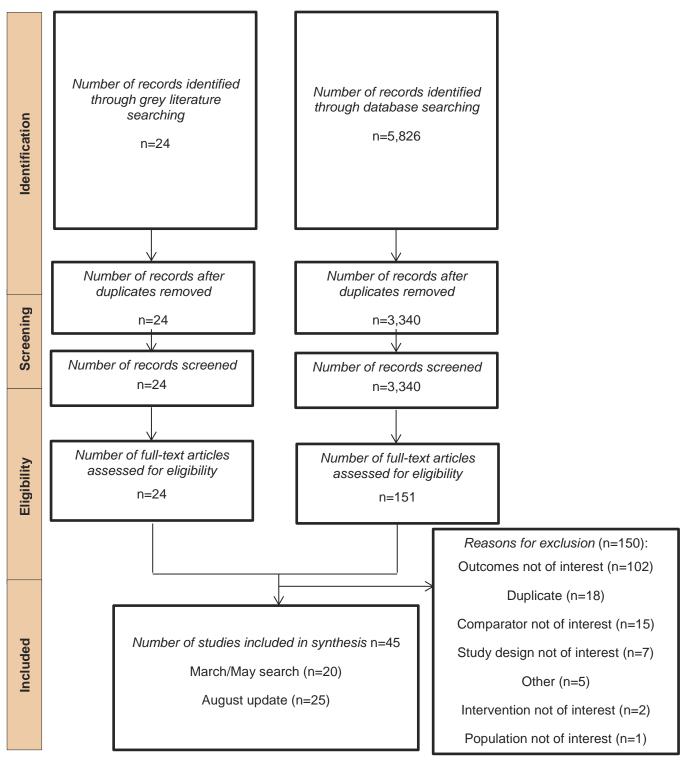
a published study and provided minimal information on the population and outcomes. The update focused on human studies only; therefore, the 12 pre-clinical animal studies included in the March/May report were not included in this update.

## **Study Characteristics**

Across the 45 studies, six were randomized controlled trials (five from the May version,<sup>6,7,16,32,33</sup> one newly identified<sup>51</sup>), 17 were retrospective cohort studies (six from the May version,<sup>15,22-24,52,53</sup> 11 newly identified<sup>2-4,8-10,19,29,36,54,55</sup>, 13 were prospective cohort studies (eight from the May version,<sup>13,14,17,18,21,25,56,57</sup> five newly identified<sup>27,28,31,58,59</sup>), and nine were case control studies (one from the May version,<sup>60</sup> eight newly identified<sup>5,11,12,20,26,30,61,62</sup>).









## Table 2: Characteristics of Included RCTs

Author/Country/Design	Trial Information	Participant Inclusion/Exclusion Criteria	Vaccine Information	Efficacy/Effectiveness Outcomes
Author: Voysey 2021 <sup>7</sup> County: UK, Brazil, S.Africa Date of Recruitment: May-Nov 2020 Trial Phase: 2/3 Design: Single Blind RCT Funding: UK Research and Innovation, National Institutes for Health Research (NIHR), Coalition for Epidemic Preparedness Innovations, Bill & Melinda Gates Foundation, Lemann Foundation, Rede D'Or, Brava and Telles Foundation, NIHR Oxford Biomedical Research Centre, Thames Valley and South Midland's NIHR Clinical Research Network, and AstraZeneca.	Age: NR %Female: Varied Type of comparator: Meningococcal vaccine Sample Size Vaccine: Varied Sample Size Control: Varied Total Sample: Varied VOC: NR	Healthy volunteers aged over 18; at risk of virus, stable pre- existing conditions	Vaccine: ChAdOx1 nCoV-19 Manufacturer: AstraZeneca Dose: Low or Standard Doses Number of Doses: 2	<ul> <li>Symptomatic Infection</li> <li>Severe Cases</li> <li>Asymptomatic infection (weekly self- administered nose and throat swab for NAAT testing from 1 week after first vaccination using kits provided by the UK Department of Health and Social Care)</li> </ul>
Author: Voysey, 2021 <sup>6</sup> County: UK, Brazil, S.Africa Date of Recruitment: May-Dec 2020 Trial Phase: 1/2/3 Design: Single Blind RCT Funding: UKRI, NIHR, CEPI, the Bill & Melinda Gates Foundation, the Lemann Foundation, Rede D'OR, the Brava and Telles Foundation, NIHR Oxford Biomedical Research Centre, Thames Valley and South Midland's NIHR Clinical Research Network, and Astra Zeneca	Age: NR %Female: NR Type of comparator: Meningococcal vaccine Sample Vaccine: 8567 Sample Control: 8580 Total Sample: 17177 VOC: NR	NR	Vaccine: ChAdOx1 nCoV-19 Manufacturer: AstraZeneca Dose: Low or Standard Doses Number of Doses: 2	<ul> <li>Symptomatic Infection</li> <li>Severe Cases</li> <li>Asymptomatic infection (measured by means of weekly self-administered nose and throat swabs using kits provided by the Department of Health and Social Care)</li> </ul>
Author: Emary 2021 <sup>16</sup> County: UK Date of Recruitment: Oct-Jan 2021 Trial Phase: 2/3 Design: RCT	Age: NR %Female: NR Type of comparator: Meningococcal vaccine Sample Vaccine: 4236 Sample Control: 4270 Total Sample: 8506	Aged 18 and over; high- exposure populations eligible for vaccination under the government National Health Service coronavirus vaccine programme.	Vaccine: ChAdOx1 nCoV-19 Manufacturer: AstraZeneca Dose: Low or Standard Doses Number of Doses: 2	<ul> <li>Symptomatic Infection</li> <li>Ct Values (weekly swabs processed. The minimum Ct value across the N and ORF1ab genes from each PCR test was computed)</li> </ul>



Author/Country/Design	Trial Information	Participant Inclusion/Exclusion Criteria	Vaccine Information	Efficacy/Effectiveness Outcomes
Funding: UK Research and Innovation, National Institutes for Health Research (NIHR), Coalition for Epidemic Preparedness Innovations, NIHR Oxford Biomedical Research Centre, Thames Valley and South Midlands NIHR Clinical Research Network, and AstraZeneca. Author: Janssen Biotech, 2021 <sup>33</sup> (Regulatory Submission) County: Argentina, Brazil, Chile, Colombia, Mexico, Peru, South Africa, and the United States Date of Recruitment: Sept 2020-Jan 2021 Trial Phase: 3 Design: Double Blind RCT Funding: Janssen Biotech	VOC: B.1.1.7, Other Age: 51.1 (15.0) %Female: 44.5 Comparator: Placebo Sample Vaccine: 19514 Sample Control:19544 Total Sample: 39058 VOC:NR	Adults 18+ with or without comorbidities.	Vaccine: Ad26.COV2.S Manufacturer: Janssen Biotech Dose: NR Number of Doses: 1	<ul> <li>Asymptomatic Unknown infection (upper airway swabs every week during the trial. Cases were excluded if they occurred before 15 days post the second dose of vaccine or occurred in participants who were not seronegative on a SARS-CoV-2 N protein assay at baseline)</li> <li>Severe cases</li> <li>Moderate to Severe infections</li> <li>Asymptomatic infection (No symptoms on the day preceding, the day of, or any time after the positive PCR test AND has a SARS-CoV- 2 positive RT-PCR test result OR develops a positive serology based on a SARS-CoV-2 N-specific immunoglobulin assay (Elecsys®, Roche) during the study. SARS CoV-2 seropositivity by non-S protein was assessed at Day 1 (pre-vaccination), Day 29 (28 days post-vaccination),</li> </ul>
Author: Baden, 2021 <sup>32</sup> County: USA Date of Recruitment: Jul-Nov, 2020 Trial Phase: 3 Design: Observer Blinded RCT Funding: Biomedical Advanced Research	Age: 51.4 %Female: 47.3 Comparator: saline Sample Vaccine: 14550 Sample Control: 14598 Total Sample: 29148	<b>Include:</b> Eligible participants were persons 18 years of age or older with no known history of SARS-CoV-2 infection and with locations or circumstances that put them at an appreciable risk	Vaccine: Moderna Manufacturer: Moderna Dose: 100mcg Number of Doses: 2	<ul> <li>and Day 71)</li> <li>Symptomatic infection</li> <li>Severe cases</li> <li>Any Positive PCR</li> <li>Asymptomatic infection (Surveillance swab at the second dose visit)</li> </ul>



Author/Country/Design	Trial Information	Participant Inclusion/Exclusion Criteria	Vaccine Information	Efficacy/Effectiveness Outcomes
and Development Authority and the National Institute of Allergy and Infectious Diseases		of SARSCoV-2 infection, a high risk of severe COVID-19, or both. <b>Exclude:</b> Pregnant women and children		
Author: Ali, 2021 <sup>51</sup> County: USA Date of Recruitment: 9 Dec 2020 - 28 Feb 2021 Trial Phase: Phase 2/3 Design: RCT Funding: Moderna and the Biomedical Advanced Research and Development Authority	Age: 14.3 ±1.6 %Female: 49% Comparator: placebo Sample Vaccine: 2139 Sample Control: 1042 Total Sample: 3181	Include: Male and female adolescents between the ages of 12 and 17 years were eligible for enrollment if they were considered to be in good general health by the 26 U.S. investigators Exclude: travel outside of the United States in the 28 days before screening, pregnancy or breast-feeding, acute illness or fever 24 hours before or at screening, previous administration of an investigational vaccine against SARS-CoV-2, or current treatment with investigational agents for prophylaxis against Covid-19	Vaccine: Moderna Manufacturer: Moderna Dose: 100ug Number of Doses: 2	Asymptomatic

IQR: interquartile range, NR: Not Reported, PCR: Polymerase Chain Reaction, RCT: randomized controlled trial, VOC: Variant of Concern, Studies are peer reviewed publications except otherwise stated.

Newly identified RCT in this version is shaded in blue.



## Table 3: Characteristics of Observational Studies

Author/Country/Design	Trial Information	Participant Inclusion/Exclusion Criteria	Vaccine Information		Effectiveness Outcomes
Author: Hall, 2021 <sup>56</sup> County: UK Date of Recruitment: Dec 2020-Feb 2021 Trial Phase: Post Approval Design: Prospective Cohort Funding: Public Health England and the Department of Health and Social Care; NIHR	Age: NR %Female: 84 Type of comparator: Unvaccinated Sample Vaccine: NR Sample Control: NR Total Sample: NR VOC: B.1.1.7	Health care workers at hospital, who could provide informed consent and anticipated remaining engaged in follow-up for 12 months.	Vaccine: BNT162b2 Manufacturer: Pfizer BioNTech Dose: NR Number of Doses: 1 or 2	•	Symptomatic Infection Asymptomatic infection (fortnightly asymptomatic PCR testing (anterior nasal swabs or combined nose and oropharyngeal swabs) and monthly antibody testing) Any positive PCR
Author: Amit, 2021 <sup>52</sup> County: Israel Date of Recruitment: Dec 2020-Jan 2021 Trial Phase: Post Approval Design: Retrospective Cohort Funding: NR	Age: NR %Female: NR Comparator: Unvaccinated Sample Vaccine: NR Sample Control: NR Total Sample: NR VOC: NR	NR	Vaccine: BNT162b2 Manufacturer: Pfizer BioNTech Dose: 1 or 2 Number of Doses:2	•	Symptomatic Infection Any positive PCR
Author: Dagan, 2021 <sup>13</sup> County: Israel Date of Recruitment: Dec 2020-Feb 2021 Trial Phase: Post Approval Design: Prospective Cohort Funding: NR	Age: Unvaccinated: 45 (IQR:35–62), vaccinated: 45 (35–62) %Female: 50 Comparator: Unvaccinated Sample Vaccine: 596618 Sample Control: 596618 Total Sample: 1193236 VOC: B.1.1.7	Include:16 years or older, not having a previously documented positive SARS-CoV-2 PCR test, and being a member of the health care organization during the previous 12 months. Exclude: probability of exposure or the outcomes is high and controlling for the high variability is not feasible.	Vaccine: BNT162b2 Manufacturer: Pfizer BioNTech Dose: NR Number of Doses:2	•	Symptomatic Infection Severe Cases Asymptomatic infection (testing protocol not defined, however SARS- CoV-2 infection without documented symptoms used as proxy)
Author: Levine-Tiefenbrun, 2021 <sup>22</sup> County: Israel Date of Recruitment: Dec 2020-Jan 2021 Trial Phase: Post Approval Design: Retrospective Cohort Funding: NR	Age: NR %Female: NR Comparator: Unvaccinated Sample Vaccine: Varied Sample Control: Varied Total Sample: Varied VOC: NR	Include: All positive post- vaccination samples Exclude: Patients who had a positive sample prior to vaccination; patients age 90 and above	Vaccine: BNT162b2 Manufacturer: Pfizer BioNTech Dose: NR Number of Doses:1	•	Ct values



Author: Jones, 2021 <sup>23</sup> Country: UK Date of Recruitment: Jan 18-31, 2021 Trial Phase: Post Approval Design: Retrospective Cohort Funding: Wellcome Senior Clinical Research Fellowship to MPW (108070/Z/15/Z), a Wellcome Principal Research Fellowship to PJL (210688/Z/18/Z), and an MRC Clinician Scientist Fellowship (MR/P008801/1) and NHSBT workpackage (WPA15-02) to NJM. Funding was also received from Addenbrooke's Charitable Trust and the Cambridge Biomedical Research Centre.	Age: NR %Female: NR Comparator: Unvaccinated Sample Vaccine: 3535 Sample Control: 3252 Total Sample: Varied VOC: B.1.1.7	Include: vaccinated and unvaccinated Health Care Workers Exclude: NR	Vaccine: BNT162b2 Manufacturer: Pfizer BioNTech Number of Doses:1	•	Any positive PCR Ct values Asymptomatic (weekly Screening)
Author: Tande, 2021 <sup>15</sup> Country: USA Date of recruitment: December 2020 to February 2021 Trial Phase: Post approval Design: Retrospective Cohort Funding: Internal funding at the Mayo Clinic	Age: 54.2 (19.7) %Female: 52.5 Comparator: Unvaccinated Sample vaccine: 3006 Sample control: 45,327 Total sample: VOC: NR	Include: 18 or mor years old, underwent preprocedural/presurgical testing within 48-72 hours of procedure Exclude: Patients tested due to symptoms or a known exposure were tested using an alternative ordering process	Vaccine: BNT162b2 or Moderna Manufacturers: Pfizer BioNTech or Moderna Number of Doses: 1 or 2	•	PCR+ among Asymptomatic (consecutive preprocedural molecular screening tests)
Author: McEllistrem, 2021 <sup>24</sup> Country: USA Date of recruitment: December 8, 2020– February 2, 2021 Trial Phase: Post approval Design: Retrospective Cohort Funding: None	Age: NR %Female: NR Comparator: Unvaccinated Sample vaccine: 5 Sample control: 5 Total sample: 10 VOC: NR	Include: A negative baseline nasopharyngeal reverse transcription polymerase chain reaction test (RT-PCR, Palo Alto VA, CA) for SARS-CoV-2 on 12/2/20.	Vaccine: BNT162b2 Manufacturers: Pfizer BioNTech Number of Doses: 1	•	Ct values Viral load Asymptomatic (surveillance nares testing for SARS-CoV-2 with the BD Veritor antigen every 2- 5 days)
Author: Shah, 2021 <sup>53</sup> (Pre-print) Country: UK Date of recruitment: December 8, 2020 – March 3, 2021 Trial Phase: Post approval Design: Retrospective Cohort Funding: British Heart Foundation through an intermediate clinical research fellowship	Age: 44.4(11.4) %Female: 78.7 Comparator: Unvaccinated Sample vaccine: 109,074 Sample control: 144,525 Total sample: VOC: NR	Include: Healthcare workers were included if they were employed by the National Health Service (NHS) in Scotland on or before the 1st of March 2020 (the first positive reported case of COVID-19 in Scotland) and still employed by the NHS on the	Vaccine: BNT162b2 or ChAdOx1 nCoV-19 Manufacturers: Pfizer BioNTech or Oxford AstraZeneca Number of Doses: 1	•	Transmission to contact



**TD** 

(FS/19/17/34172); Wellcome Trust intermediate clinical fellowship and Beit fellowship (201492/Z/16/Z)		1st of November 2020; healthcare worker cohort was restricted to the working-age population (18-65 years of age). The household member cohort included all ages but was restricted to households with no more than one healthcare worker (4% of healthcare workers lived in multiple healthcare worker households)		
Author: Bouton, 2021 <sup>60</sup> (Pre-print) Country: USA Date of recruitment: December 9, 2020- February 23, 2021 Trial Phase: Post approval Design: Case Control Funding:	Age: 40(13) %Female: NR Comparator: Unvaccinated Sample vaccine: 96 Sample control: 329 Total sample: 425 VOC: NR	Include: HCWs had been vaccinated prior to the vaccine initiative and were included in analyses. HCW who received a vaccination following their positive SARS-CoV-2 RT-PCR were included in the unvaccinated group.	Vaccine: BNT162b2 or Moderna Manufacturers: Pfizer BioNTech or Moderna Number of Doses: 1	<ul> <li>Asymptomatic         <ul> <li>(Asymptomatic testing is available to HCWs for workplace exposures, following out-of-state travel, and per request)</li> </ul> </li> <li>All PCR-positive         <ul> <li>(symptomatic and asymptomatic)</li> </ul> </li> </ul>
Author: Regev-Yochay, 2021 <sup>25</sup> (Pre-print) Country: Israel Date of recruitment: December 19, 2020 – March 14, 2021 Trial Phase: Post approval Design: Cohort Funding: Sheba Medical Center, Israel	Age: NR %Female: NR Comparator: Unvaccinated Sample vaccine: Sample control: Total sample: 3578 VOC: NR	Include: HCW at Sheba Medical Center (Israel)	Vaccine: BNT162b2 or Moderna Manufacturers: Pfizer BioNTech or Moderna Number of Doses: 1 or 2	<ul> <li>Asymptomatic (Symptomatic or exposed to confirmed case)</li> <li>Symptomatic</li> <li>Severe cases</li> </ul>
Author: Lumley, 2021 <sup>21</sup> (Pre-print) Country: England Date of recruitment: Through to February 28, 2021 Trial Phase: Post approval Design: Longitudinal Cohort Funding: Supported by the UK Government's Department of Health and Social Care. Also supported by the National Institute for Health Research	Age: 39 (IQR:30-50) %Female: 74.0 Comparator: Unvaccinated seronegative Sample vaccine: NR Sample control: NR Total sample: 13,109 VOC: B.1.1.7	<b>Include:</b> Only those who participated in asymptomatic screening, symptomatic testing or vaccination from 01- September-2020 onwards were included. All staff working for the hospitals were eligible to participate.	Vaccine: BNT162b2 or ChAdOx1 nCoV-19 Manufacturers: Pfizer BioNTech or Oxford AstraZeneca Number of Doses: 1 or 2	<ul> <li>Ct values</li> <li>Symptomatic</li> <li>Asymptomatic (voluntary nasal and oropharyngeal swab PCR testing every two weeks and serological testing every two months)</li> </ul>



Health Protection Research Unit (NIHR HPRU) in Healthcare Associated Infections and Antimicrobial Resistance at Oxford University in partnership with Public Health England (PHE) (NIHR200915), the NIHR Biomedical Research Centre, Oxford, and benefactions from the Huo Family Foundation and Andrew Spokes.				
Author: Pritchard, 2021 <sup>17</sup> (Pre-print) Country: UK Date of recruitment: December 1, 2020 – April 3, 2021 Trial Phase: Post approval Design: Prospective Cohort Funding: Department of Health and Social Care with in-kind support from the Welsh Government, the Department of Health on behalf of the Northern Ireland Government and the Scottish Government.	Age: NR %Female: NR Comparator: Unvaccinated Sample vaccine: Sample control: Total sample: 373,402 VOC: NR	<b>Include:</b> This analysis included participants aged 16 years or over (i.e. those who theoretically could have received vaccination), and all visits with positive or negative swab results from 1 December 2020 to 3 April 2021.	Vaccine: BNT162b2 or ChAdOx1 nCoV-19 Manufacturers: Pfizer BioNTech or Oxford AstraZeneca Number of Doses: 1 or 2	<ul> <li>Asymptomatic (Weekly nose and throat self-swab for first month, then monthly for 12 months from enrolment)</li> <li>Ct values</li> <li>Symptomatic</li> </ul>
Author: Shrotri, 2021. <sup>18</sup> (Pre-print) Country: UK Date of recruitment: December 8, 2020 – March 15, 2021 Trial Phase: Post approval Design: Prospective Cohort Funding: UK Government Department of Health and Social Care.	Age: 86 (IQR: 80-91) %Female: 69.6 Comparator: Unvaccinated Sample vaccine: Sample control: Total sample: 10,412 VOC: NR	Include: At least two PCR test results in total, and $\geq$ 1 PCR result during the analysis period. Residents entered the risk period on 8 December 2020 if they had $\geq$ 1 valid PCR result on or prior to that date; or, if they had no PCR results before 8 December 2020, on the date of their first negative PCR test. Residents with a positive PCR result $\leq$ 90 days before 8 December entered the risk period 90 days after their positive test.	Vaccine: BNT162b2 or ChAdOx1 nCoV-19 Manufacturers: Pfizer BioNTech or Oxford AstraZeneca Number of Doses: 1 or 2	<ul> <li>Ct values</li> <li>Symptomatic</li> </ul>
Author: Haas, 2021 <sup>14</sup> Country: Israel	Age: NR %Female: 50.8 Comparator: Unvaccinated	<b>Include:</b> unvaccinated and vaccinated individuals aged ≥16 years.	Vaccine: BNT162b2 Manufacturers: Pfizer BioNTech	<ul> <li>Asymptomatic (routine testing)</li> <li>Severe cases</li> </ul>



Date of recruitment: Jan 24, 2021–April 3, 2021 Trial Phase: Post approval Design: Prospective Cohort Funding: Israel MoH and Pfizer.	Sample vaccine: NR Sample control: NR Total sample: NR VOC: B.1.1.7		Number of Doses: 1 or 2	•	Symptomatic
Author: Harris, 2021 <sup>57</sup> (Pre-print) Country: UK Date of recruitment: January 4 – February 28, 2021 Trial Phase: Post approval Design: Prospective Cohort Funding: This work was undertaken as part of the core functions of Public Health England in relation to the surveillance of communicable diseases and outbreak response	Age: NR %Female: Unvaccinated index case: 47.6%, Index case vaccinated 21+ day before: 38.3%, Index case vaccinated <21 days before: 40.6% Comparator: Unvaccinated Sample vaccine: Sample control: Total sample: 1,018,842 VOC: NR	<b>Include:</b> Households with an index case occurring between 4 January 2021 to 28 February 2021, with 14 days observable follow up for all contacts; households with a single index case age 16+, and no coprimary cases.	Vaccine: BNT162b2 or ChAdOx1 nCoV-19 Manufacturers: Pfizer BioNTech or Oxford AstraZeneca Number of Doses: 1	•	Transmission to contact
Author: Chemaitelly, 2021 <sup>11</sup> Country: Qatar Date of Recruitment: 1 Feb 2021 - 10 May 2021 Trial Phase: NR Design: Test-negative case control Funding: NR	Age: NR %Female: NR Comparator: no vaccine Sample Vaccine: 1590 Sample Control: 154394 Total Sample: NR VOC: B.1.1.7; B.1.351; B.1.617	Include: This study was conducted in the resident population of Qatar, inclusion: a B.1.1.7 case, a B.1.351 case or a severe or critical or fatal COVID-19 disease case Exclude: All records of vaccination with one or two doses using a vaccine other than Moderna were excluded	Vaccine: Moderna Manufacturer: Moderna Number of Doses: NR	•	Asymptomatic
Author: Ioannou, 2021 <sup>28</sup> Country: Greece Date of Recruitment: 4 Jan 2021 - 14 Apr 2021 Trial Phase: Post Approval Design: Prospective cohort Funding: NR	Age: 42.3 ±9.9 %Female: 74.5% Comparator: no vaccine Sample Vaccine: 21 Sample Control: 31 Total Sample: 52 VOC: B.1.1.7	Include: Vaccinated (with BNT162b2) and non-vaccinated healthcare workers who tested positive for COVID-19 at a single centre in Greece Exclude: NR	Vaccine: BNT162b2 Manufacturer: Pfizer Number of Doses: 2	•	CT values
Author: Thompson, 2021 <sup>31</sup> Country: USA Date of Recruitment: 14 Dec 2020- 10 Apr 2021	Age: NR %Female: 62% Comparator: no vaccine Sample Vaccine: 3179	Include: Eligible participants include Arizona residents aged 18–85 years who currently work at least 20 hours per week in an	Vaccine: mRNA vaccine Manufacturer: Moderna, Pfizer	•	Asymptomatic CT values







Trial Phase: NR Design: Prospective Cohort Funding: National Center for Immunization and Respiratory Diseases, Centers for Disease Control and Prevention under contract numbers 75D30120R68013 awarded to Marshfield Clinic Research Laboratory, 75D30120C08379 to University of Arizona, and 75D30120C08150 awarded to Abt Associates, Inc.	Sample Control: 796 Total Sample: 3975 VOC: B.1.429; B.1.1.7; B.1.427	occupation involving regular direct contact (within three feet) with others, assessed at the participant level <b>Exclude:</b> Exclusion criteria include receipt of a COVID-19 vaccine prior to enrolment, although we continue to follow participants who are vaccinated during the study.	Number of Doses: 1 or 2	
Author: Angel, 2021 <sup>9</sup> Country: Israel Date of Recruitment: 20 Dec 2020- 25 Feb 2021 Trial Phase: Post Approval Design: Retrospective cohort Funding: None	Age: 44.3 ±12.5 %Female: 66.5% Comparator: Non vaccinated HCW Sample Vaccine: 5953 Sample Control: 757 Total Sample: 6710 VOC: NA	Include: HCWs who received at least 1 vaccine dose between December 20, 2020, and February 25, 2021, were as signed to the vaccinated group. The control group was composed of health care workers who did not receive any doses of the BNT162b2 vaccine during this period. Exclude: HCWs who did not undergo at least 1 PCR test during the study period, had incomplete data pertaining to vaccination dates, or contracted SARS CoV-2 infection prior to the study period were excluded from the analysis	Vaccine: BNT162b2 Manufacturer: Pfizer Number of Doses: 1 or 2	<ul> <li>Asymptomatic cases &gt;7 days and &gt;28 days after first and second dose of vaccine</li> </ul>
Author: Tang, 2021 <sup>8</sup> Country: USA Date of Recruitment: 17 Dec 2020 - 20 Mar 2021 Trial Phase: Post Approval Design: Retrospective cohort Funding: American Lebanese Syrian Associated Charities	Age: NR %Female: Vaccinated: 66%, Control: 58.3% Comparator: Unvaccinated individuals Sample Vaccine: 3052 Sample Control: 2165	Include: Vaccine eligible workers that meet state vaccination guidelines Exclude: Individuals with prior COVID-19 exposure were excluded	Vaccine: BNT162b2 Manufacturer: Pfizer Number of Doses: 1 or 2	<ul> <li>Asymptomatic cases 0-11 days and &gt;12 days after first dose</li> <li>Asymptomatic cases 0-6 days and &gt;7 days after second dose</li> </ul>







	Total Sample: 5217 VOC: NA			
Author: Andrejko, 2021 <sup>10</sup> Country: USA Date of Recruitment: 24 Feb 2021- 29 Apr 2021 Trial Phase: Post Approval Design: Retrospective cohort Funding: California Department of Public Health, grant from the ELC program of the US CDC and NIH/NIAID grant	Age: NR %Female: 49.30% Comparator: unvaccinated Sample Vaccine: 20 Sample Control: 454 Total Sample: 525 VOC: NA	Include: California residents with molecular SARS-CoV-2 test results and a telephone number. Controls were persons with negative SARS-CoV-s molecular test results during the same period Exclude: participants who recalled receiving any previous positive test result for SARS- CoV-2 infection or seropositivity, prior to the reported test. Data was excluded from children aged 0-17 years, who were generally ineligible for COVID- 19 vaccination over the study period; and particpants who reported receiving COVID-19 vaccinations other than BNT162b2 or Moderna (due to limited coverage of a third authorized vaccine. or receipt of COVID-19 vaccination without knowledge of vaccination dates	Vaccine: BNT162b2 or Moderna Manufacturer: Pfizer, Moderna Number of Doses: 1 or 2	<ul> <li>Asymptomatic cases &gt;15 days after 2<sup>nd</sup> dose</li> <li>Asymptomatic cases up to 14 days after 1<sup>st</sup> or second dose</li> </ul>
Author: Jacobson, 2021 <sup>29</sup> Country: USA Date of Recruitment: Dec 2020-Apr 2021 Trial Phase: Post Approval Design: Retrospective quality improvement Funding: NR	Age: 37.5 ±10.6 %Female: 69.8% Comparator: Unvaccinated HCP Sample Vaccine: NR Sample Control: NR Total Sample: 283 VOC: B.1.427/B.1.429	Include: include post vaccine SARS CoV-2 cases, defined as HCPs with positive SARS-CoV-2 nucleic acid amplification test after receiving one or more vaccine doses Exclude: NR	Vaccine: BNT162b2 or Moderna Manufacturer: Pfizer, Moderna Number of Doses: 1 or 2	<ul> <li>CT values ≤ 14 after first dose</li> <li>CT values up to 14 days after 1<sup>st</sup> or 2<sup>nd</sup> dose</li> <li>CT values over 14 days after 2<sup>nd</sup> dose</li> </ul>
Author: Bailly, 2021 <sup>27</sup> Country: France Date of Recruitment: 8 Mar 2021 - 29 Mar 2021	Age: Fully vaccinated residents: 87.0 ± 8.2 years %Female:	<b>Include:</b> Residents and staff from a nursing home unit with a positive COVID case	Vaccine: BNT162b2 Manufacturer: Pfizer Number of Doses: 2 doses	<ul> <li>Asymptomatic cases after 2 doses</li> <li>CT values after 2 doses</li> </ul>



Trial Phase: Post Approval Design: Prospective cohort Funding: Ministry of Health	Fully vaccinated residents: 64.5% Comparator: Non- vaccinated residents Sample Vaccine: 13 Sample Control: 5 Total Sample: 18 VOC: 501Y.V2	Exclude: NR		
Author: Salo, 2021 <sup>3</sup> (Pre-print) Country: Finland Date of Recruitment: 27 Dec 2020- 24 Mar 2021 Trial Phase: Post approval Design: Retrospective Funding: InFLAMES and INVEST Flagship Programmes of the Academy of Finland.	Age: Vaccinated: 47.1 ±13.1 Unvaccinated: 43.8 ±14.5 %Female: 86.5% Comparator: unvaccinated HCW Sample Vaccine: 95138; spouses of vaccinated HCW:52,766 Sample Control: 193000; spouses of control: 111,000 Total Sample: 288138 VOC: NA	Include: Vaccinated and unvaccinated HCW. An individual was included in this sample if their spouse is a healthcare worker and they had not been vaccinated during the sample period. Exclude: NR	Vaccine: BNT162b2 or Moderna Manufacturer: Moderna, Pfizer Number of Doses: 1 or 2	<ul> <li>Transmission to unvaccinated spouse 14 days and 10 weeks after 1<sup>st</sup> dose</li> <li>Transmission to unvaccinated child 3-18 years, 14 days and 10 weeks after 1<sup>st</sup> dose</li> <li>Transmission to unvaccinated child 3-12 years, 6 weeks and 10 weeks after 1<sup>st</sup> dose</li> <li>Transmission to unvaccinated child 13-18 years, 6 and 10 weeks after 1<sup>st</sup> dose</li> </ul>
Author: Muhsen, 2021 <sup>58</sup> (Pre-print) Country: Israel Date of Recruitment: Dec 2020-Jan 2021 Trial Phase: Post approval Design: Prospective cohort Funding: No external funding	Age: 46.2 ±11.8 %Female: 79.5% Comparator: Unvaccinated individuals Sample Vaccine: 20 Sample Control: 44 Total Sample: 9162 VOC: NA	Include: 1) adherence to routine screening for SARS-CoV-2 infection by RT-PCR testing. Specifically, they had 12 or more out of the 20 planned screening tests for the period September 2020 through January 2021; 2) working in LCTFs that vaccinated >75% of their employees collectively during three consecutive days; and 3) being RT-PCR negative for SARS-CoV-2 infection by the date of immunization with the second vaccine dose.	Vaccine: BNT162b2 Manufacturer: Pfizer Number of Doses: 2	CT values over 14 days after 2 <sup>nd</sup> dose



Author: de Gier, 2021 <sup>2</sup> (Pre-print) Country: Netherlands Date of Recruitment: 1 Feb 2021- 27 May 2021 Trial Phase: Post Approval Design: Retrospective cohort Funding: Ministry of Health, Welfare and Sports	Age: NR %Female: NR Comparator: Unvaccinated individuals Sample Vaccine: 2032 Sample Control: 139802 Total Sample: Index cases: 113582; contacts: 253168 VOC: B.1.1.7	Unvaccinated HCWs at baseline, who were vaccinated later, were censored upon receiving their first vaccination dose <b>Exclude:</b> Excluded from the primary analysis were HCWs working at institutions that did not have a collective immunization period, partially vaccinated HCWs at baseline (i.e. received one vaccine dose), and those who had a RT-PCR- confirmed SARS-CoV-2 infection before immunization, or between immunization with the second dose until day seven or 14 days post immunization. <b>Include:</b> household members and other close contacts of confirmed cases <b>Exclude:</b> household contacts of an index were excluded if the most likely setting of infection of the index was 'at home' according to the source tracing interview	Vaccine: ChAdOx1-S, BNT162b2, Moderna, Janssen Manufacturer: Pfizer, Moderna, AstraZeneca, Janssen Number of Doses: 1 or 2	<ul> <li>Transmission to contact over 14 days after 1<sup>st</sup> dose: any household contact, unvaccinated household contacts, any other close contact, unvaccinated close contacts, AstraZeneca vaccinated household contact, Pfizer vaccinated household contact, Moderna vaccinated household contact</li> <li>Transmission to contact over 7 days after 2<sup>nd</sup> dose in: any household contact, unvaccinated household contact, any other close contact, unvaccinated</li> </ul>
---	--	---	---	---



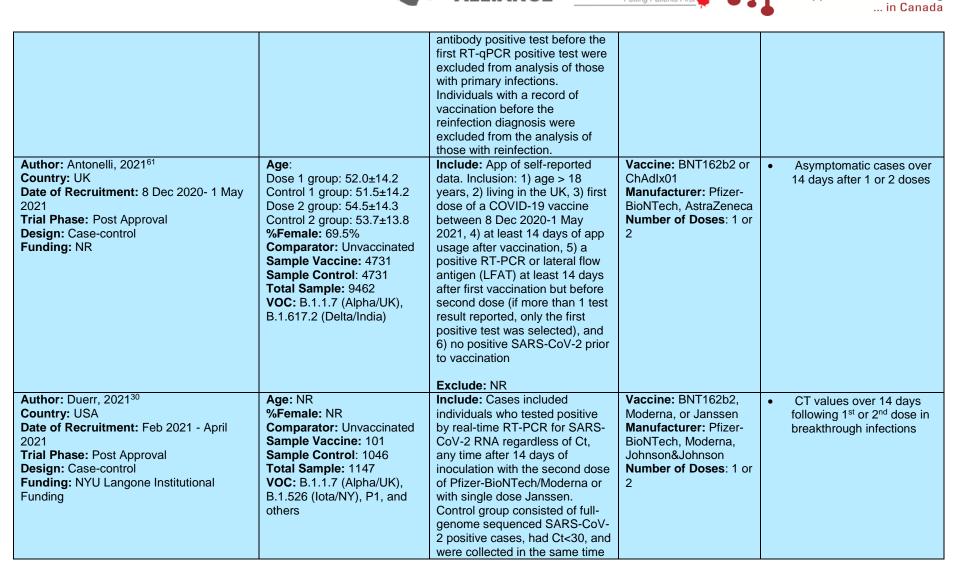
				AstraZeneca vaccinated household contact, Pfizer vaccinated household contact, Moderna vaccinated household contact, Janssen vaccinated household contact
Author: McEllistrem,2021 <sup>55</sup> Country: USA Date of Recruitment: 2 Dec 2020 to 14 May 2021 Trial Phase: Post approval Design: Retrospective Cohort (observational) Funding: NR	Age: 74.5 years (IQR NR) %Female: 7.76% Comparator: unvaccinated Sample Vaccine: 97 Sample Control: 19 Total Sample: NR VOC: NA	Include: residing at the Community Living Center; without a prior history of COVID- 19 who agreed to immunization Exclude: prior history of COVID-19; agreed to immunization after 12/16/21	Vaccine: BNT162b2 Manufacturer: Pfizer- BioNTech Number of Doses: 2	<ul> <li>Asymptomatic cases 1 to 21 days following 1<sup>st</sup> vaccination</li> <li>Asymptomatic cases 14-21 days following 1<sup>st</sup> vaccine</li> <li>Asymptomatic cases 7 days following 2<sup>nd</sup> vaccine</li> </ul>
Author: Souza, 2021 <sup>54</sup> (Pre-print) Country: Brazil Date of Recruitment: 2021-03-01 Trial Phase: Post approval Design: Observational cohort Funding: Sao Paulo Research Foundation, MCTI	Age: 73 (IQR 50-83) %Female: 96.2% Comparator: unvaccinated Sample Vaccine: 23 Sample Control: 3 Total Sample: 26 VOC: B.1.1.7 (UK)	Include: Individuals at least 18 years of age exposed to residents infected with SARS- CoV-2 (from either the convent or LTC facility). Residents and employees from both locations were included in the study Exclude: N/a	Vaccine: ChAdlx01, 1 resident vaxxed with Ad26.COV2.S and CoronaVac, 1 employee vaxxed with ChAdOx1 Manufacturer: AstraZeneca and SinoVac BioTech Number of Doses: 1 or 2	<ul> <li>Asymptomatic cases in individuals &gt;23 days following 1<sup>st</sup> dose</li> <li>Viral load in individuals &gt;23 days following 1<sup>st</sup> dose</li> <li>Asymptomatic cases in individuals 5-27 days following 2<sup>nd</sup> dose</li> <li>Viral load in individuals 5- 27 days following 2<sup>nd</sup> dose</li> </ul>
Author: Abu-Raddad, 2021 <sup>26</sup> (Pre-print) Country: Qatar Date of Recruitment: 28 Feb 2020 - 11 July 2021 Trial Phase: Post approval Design: Matched Case-control 1:1 ratio Funding: NR	Age: 33-35 %Female: 14.90%-21.100% Comparator: Unvaccinated Sample Vaccine: 60-421 Sample Control: 60-421 Total Sample: 120-842 VOC: B.1.1.7 (Alpha/UK), B.1.351 (Beta/South Africa), B.1.617.2 (Delta/India)	Include: All records of RT- qPCR in Qatar but only samples of matched cohorts were included in the analysis. Only breakthrough infections in fully vaccinated individuals were included in the analysis. Being fully vaxxed was defined as >14 days after the second dose Exclude: Individuals with a record of a SARS-CoV-2	Vaccine: BNT162b2, Moderna Manufacturer: Pfizer- BioNTech, Moderna Number of Doses: 2	CT value over 14 days following 2 <sup>nd</sup> dose of vaccine



COVID

COVID-19 Evidence Networ

to support Decision-making









Author: Sansone, 2021 <sup>62</sup> Country: Italy Date of Recruitment: 25 Jan 2021 - 13 Apr 2021 Trial Phase: NR Design: Case-control Funding: NR	Age: NR %Female: NR Comparator: Unvaccinated Sample Vaccine: 40 Sample Control: 52 Total Sample: 92 VOC: B.1.1.7 (Alpha/UK), B1.525	period as the breakthrough infections Exclude: NR Include: Data from health and epidemiological surveillance at the workplaces performed on the ASST "Spedali Civili di Brescia" workforce. Mandatory character of such an activity therefore no ethics committee approval necessary. All workers gave wrwitten consent to the vaccination and data processing. Data was anonymysed. Exclude: NR	Vaccine: BNT162b2 Manufacturer: Pfizer Number of Doses: 2	Asymptomatic cases over 14 days following 2 <sup>nd</sup> vaccine
Author: Layan, 2021 <sup>5</sup> (Pre-print) Country: Israel Date of Recruitment: 31 Dec 2020 to 26 Apr 2021 Trial Phase: Post Approval Design: Case-control (Observational) Funding: Sheba Medical Center. SC acknowledges financial support from the Investissement d'Avenir program, the Laboratoire d'Excellence Integrative Biology of Emerging Infectious Diseases program (grant ANR-10-LABX-62-IBEID), HAS, the INCEPTION project (PIA/ANR-16 CONV-0005), the European Union's Horizon 2020 research and innovation program under grant 101003589 (RECOVER) and 874735 (VEO), AXA and Groupama.	Age: 32±16 %Female: 58% Comparator: unvaccinated Sample Vaccine: 15-124 Sample Control: 200-641 Total Sample: 215-687 VOC: alpha	Include: HCWs employed by Sheba Medical Center with a SARS-CoV-2 case Exclude: missing vaccination status, dates of PCR test and/or symptom onset	Vaccine: BNT162b2 Manufacturer: Pfizer- BioNTech Number of Doses: 2	<ul> <li>Transmission to contact &gt;7 days following 2<sup>nd</sup> dose</li> <li>Infected contacts &gt;7 days following 2<sup>nd</sup> dose</li> </ul>
Author: Mostafa, 2021 <sup>19</sup> (Pre-print) Country: USA Date of Recruitment: Jan 2021- May 2021	Age: 51 (IQR NR) %Female: 63.3% Comparator: unvaccinated	Include: specimens of SARS- CoV-2 positive patients who had received two doses of either	Vaccine: BNT162b2 or ChAdOx1 nCOV-19	CT values 2-100 days following 2 <sup>nd</sup> dose



Trial Phase: Post Approval Design: Retrospective cohort (Observational) Funding: National Institute of Health (The Johns Hopkins Center of Excellence in Influenze Research and Surveillence	Sample Vaccine: 49 Sample Control: 96 Total Sample: NR VOC: P.1, B.1.1.7, B.1.351, B.1.526, and B.1.526.1	Pfizer-BioNTech (BNT162b2) or Moderna (Moderna) vaccines; specimens of a control unvaccinated cohort from a matched time frame	Manufacturer: Pfizer- BioNTech, Moderna Number of Doses: 2	
Influenza Research and Surveillance, HHSN272201400007C), Johns Hopkins University, Maryland Department of Health, Centers for Disease Control and Prevention.		<b>Exclude:</b> samples that had failed sequencing and were thought of as very low viral load or false positives did not yield infectious virus		
Author: Gazit, 2021 <sup>4</sup> (Pre-print) Country: Israel Date of Recruitment: 20 Dec 2020 and 17 Mar 2021 Trial Phase: Post Approval Design: Observational cohort Funding: None	Age: index case: unvaccinated, 56 ±15, recently vaccinated once, 63±12, fully vaccinated 68±9 additional case: unvaccinated, 56±15, recently vaccinated once, 63±12, fully vaccinated once, 63±12, fully vaccinated, 67±9 %Female: 50% Comparator: unvaccinated Sample Vaccine: 381 Sample Control: 2975 Total Sample: 3627 VOC: NA	<ul> <li>Include: a household was defined as having two adults. Only households with two adults were included</li> <li>Exclude: households of only one member, households with varying numbers of children, households in which an infection was recorded before December 20, 2020</li> </ul>	Vaccine: BNT162b2 Manufacturer: Pfizer- BioNTech Number of Doses: 1 and 2	<ul> <li>Transmission to contact 0- 7 days following 1<sup>st</sup> dose</li> <li>Transmsission to contact &gt;7 days following 2<sup>nd</sup> dose</li> </ul>
Author: Pouwels, 2021 <sup>59</sup> (Pre-print) Country: UK Date of Recruitment: 1 Dec 2020 to 16 May 2021 Trial Phase: Post Approval Design: Prospective Cohort Observational Funding: Department of Health and Social Car, Welsh Government, the Department of Health on behalf of the Northern Ireland Government and the Scottish Government.	Age: 28-57 %Female: 53.6-55.8% Comparator: unvaccinated Sample Vaccine: NR Sample Control: 10853 Total Sample: NR VOC: Delta	Include: participants aged 18 years or over (i.e. those who were eligible for vaccination), and all visits with positive or negative swab results from 1 December 2020 to 1 August 2021. Exclude: A small number of visits after reported vaccination with either unknown or vaccines other than	Vaccine: BNT162b2, ChAdOx1, or Moderna Manufacturer: Pfizer- BioNTech, Moderna Number of Doses: 1 or 2	<ul> <li>CT values 0 to 20 days following 1<sup>st</sup> dose</li> <li>CT values &gt;21 days following 1<sup>st</sup> dose, or 0-13 days following 2<sup>nd</sup> dose</li> <li>CT values &gt;14 days following 2<sup>nd</sup> dose</li> </ul>







		ChAdOx1, BNT162b2 or Moderna (for the latter we only included the first dose and only for the period ≥17 May) were excluded as these were too few to provide reliable estimates.		
Author: Tang, 2021 <sup>12</sup> (Pre-print) Country: Qatar Date of Recruitment: 21 Dec 2020 and 21 Jul 2021 Trial Phase: Post Approval Design: matched test-negative, case- control Funding: Unclear	Age: 31-32 years %Female: 18.5-21.9% Comparator: unvaccinated Sample Vaccine: 48-532 Sample Control: 2194-2401 Total Sample: 2318-2862 VOC: Delta	Include: Every case that met the inclusion criteria (a Delta case) and that could be matched to a control Exclude: persons who received mixed vaccines, or who received a vaccine other than BNT162b2 or Moderna	Vaccine: BNT162b2, Moderna Manufacturer: Pfizer- BioNTech, Moderna Number of Doses: 1 or 2	<ul> <li>Asymptomatic cases &gt;14 days following 1<sup>st</sup> and 2<sup>nd</sup> dose</li> </ul>
Author: Baltas, 2021 <sup>20</sup> Country: UK Date of Recruitment: 30th of September 2020 and 15th of March 2021 Trial Phase: Post Approval Design: case-control Funding: COG-UK Hospital-Onset COVID- 19 Infections (HOCI) Wellcome Trust Study through grants from UK Research and Innovation (UKRI); the Wellcome Trust; and the John Black Charitable Foundation	Age: median 79, IQR 65 – 86 %Female: 42.9% Comparator: unvaccinated Sample Vaccine: 112 Sample Control: 399 Total Sample: 511 VOC: B.1.1.7, B.1.525	Include: All SARS CoV-2 first positive cases recruited into the COG-UK-HOCI study between the 30th of September 2020 and 15th of March 2021 Exclude: Positive patients, whose samples were not available for sequencing	Vaccine: BNT162b2 or ChAdOx1 nCOV-19 Manufacturer: Pfizer- BioNTech, Moderna Number of Doses: 1	<ul> <li>CT values &lt;14 and &gt;14 following 1<sup>st</sup> dose</li> </ul>
Author: Riemersma, 2021 <sup>36</sup> (Pre-print) Country: USA Date of Recruitment: 28 June–24 July 2021 Trial Phase: Post Approval Design: Retrospective cohort Funding: NR	Age: median NR %Female: NR Comparator: unvaccinated Sample Vaccine: 79 Sample Control: 212 Total Sample: 291 VOC: B.1.1.7	Include: All samples tested in a Wisconsin commercial laboratory who had Exclude: NR	Vaccine: NR Manufacturer: NR Number of Doses: 2	<ul> <li>CT values &gt;14 following 2<sup>nd</sup> dose</li> </ul>

IQR: interquartile range, NR: Not Reported, PCR: Polymerase Chain Reaction, RCT: randomized controlled trial, VOC: Variant of Concern, NR: Not Reported. Studies are peer-reviewed publications except otherwise stated.

Newly identified observational studies shaded in blue.



## **Risk of Bias Assessment**

Across the six included RCTs, all scored "low" for bias on outcome measurement, and all except for one<sup>16</sup> scored "low" for bias on selection of reported results. For risk of bias on randomization, three scored low,<sup>7,32,51</sup> two were of some concern,<sup>16,33</sup> and one<sup>6</sup> did not report sufficient information. For bias stemming from intended intervention, two scored low,<sup>32,51</sup> and the rest were of some concern. With respect to bias stemming from missing outcome data, only one study was low,<sup>51</sup> one study was high,<sup>16</sup> and the rest were of some concern. On the overall risk of bias domain, only one study scored low,<sup>51</sup> one scored high,<sup>16</sup> and the rest were of some concern.

Across the 39 observational studies, 22 were rated as moderate on risk of bias due to confounding, with nine rated as low, seven rated as high, and one with no information. On risk of bias for participant selection, 19 were rated as low, 14 were rated as serious, and six were rated as moderate. On bias in classification of interventions, all studies were rated as low, with exception of three studies that were rated as moderate. With respect to bias due to deviations from intended interventions, 19 studies scored low, 18 scored moderate, and two were rated as no information. For bias due to missing data, 19 were low, 12 were moderate, three were serious, and five did not have sufficient information. Across the outcome measurement domain, six studies were of moderate risk of bias, one was serious, and the rest were low. On bias in selection of reported results, seven studies scored low, three scored moderate, and the rest did not have sufficient information. On the overall risk of bias domain, 20 studies were rated as moderate risk of bias, 14 were serious, three were critical, one was low, and one did not have sufficient information.



#### Table 4: Risk of Bias Assessment for RCTs

Author	Randomization	Deviation from intended intervention	Missing outcome data	Measurement of outcome	Selection of reported results	Overall Bias
Ali et al.51	Low	Low	Low	Low	Low	Low
Baden et al.32	Low	Low	Some concerns	Low	Low	Some concerns
Emary et al. <sup>16</sup>	Some concerns	Some concerns	High	Low	Some concerns	High
Janssen Biotech33	Some concerns	Some concerns	Some concerns	Low	Low	Some concerns
Voysey et al.7	Low	Some concerns	Some concerns	Low	Low	Some concerns
Voysey et al. <sup>6</sup>	NI	Some concerns	Some concerns	Low	Low	Some concerns

All studies were published in 2021

## Table 5: ROBINS-I Risk of Bias for non-RCTs

Author	Bias due to confounding	Bias in selection of participants into the study	Bias in classification of interventions	Bias due to deviations from intended interventions	Bias due to missing data	Bias in measurement of outcomes	Bias in selection of the reported result	Overall Risk of Bias
Abu-Raddad et al. <sup>26</sup>	Low	Low	Low	Low	Low	Low	NI	Moderate
Amit et al. <sup>52</sup>	NI	Low	Moderate	NI	NI	Moderate	NI	NI
Angel et al. <sup>9</sup>	Moderate	Serious	Low	Moderate	Serious	Low	NI	Serious
Andrejko et al. <sup>10</sup>	Moderate	Low	Low	Low	Moderate	Moderate	NI	Moderate
Antonelli et al. <sup>61</sup>	Serious	Low	Low	Moderate	Serious	Low	NI	Critical
Baltas et al. <sup>20</sup>	Moderate	Serious	Low	Moderate	Low	Low	Low	Serious
Bailly et al. <sup>27</sup>	Serious	Moderate	Low	Moderate	Moderate	Moderate	NI	Serious
Bouton et al. <sup>60</sup>	Low	Low	Low	Low	Low	Low	Low	Low
Chemaitelly et al. <sup>11</sup>	Low	Low	Low	Moderate	Low	Low	NI	Moderate







Dagan et al. <sup>13</sup>	Moderate	Moderate	Low	Low	Low	Low	Low	Moderate
De Gier et al. <sup>2</sup>	Serious	Serious	Low	Low	Low	Moderate	NI	Serious
Duerr et al. <sup>30</sup>	Low	Low	Low	Low	Serious	Serious	NI	Critical
Gazit et al. <sup>4</sup>	Moderate	Serious	Low	Moderate	Low	Low	NI	Serious
Haas et al. <sup>14</sup>	Moderate	Low	Low	Low	Moderate	Low	NI	Moderate
Hall et al. <sup>56</sup>	Moderate	Moderate	Low	Moderate	NI	Low	NI	Moderate
Harris et al. <sup>57</sup>	Moderate	Low	Low	Low	Moderate	Low	NI	Moderate
Ioannou et al. <sup>28</sup>	Moderate	Moderate	Low	Moderate	Low	Low	NI	Moderate
Jacobson et al. <sup>29</sup>	Moderate	Serious	Low	Moderate	Moderate	Moderate	Low	Serious
Jones et al. <sup>23</sup>	Moderate	Low	Low	Low	NI	Low	NI	Moderate
Layan et al. <sup>5</sup>	Serious	Serious	Low	Moderate	Low	Low	NI	Serious
Levine-Tiefenbrun et al. <sup>22</sup>	Moderate	Low	Low	Low	NI	Low	NI	Moderate
Lumley et al. <sup>21</sup>	Low	Low	Low	Low	Moderate	Low	Moderate	Moderate
McEllistrem et al. <sup>24</sup>	Moderate	Low	Low	Low	Low	Low	NI	Moderate
McEllisrem et al.55	Serious	Serious	Low	Moderate	Low	Low	NI	Serious
Mostafa et al. <sup>19</sup>	Moderate	Serious	Low	NI	Low	Low	NI	Serious
Muhsen et al.58	Moderate	Moderate	Low	Moderate	Moderate	Low	Low	Moderate
Pouwels et al. <sup>59</sup>	Moderate	Serious	Low	Moderate	Low	Low	Low	Serious
Pritchard et al. <sup>17</sup>	Low	Low	Low	Low	Moderate	Low	Moderate	Moderate
Regev-Yochay et al. <sup>25</sup>	Low	Low	Low	Low	Low	Low	Moderate	Moderate
Riemersma et al. <sup>36</sup>	Serious	Serious	Low	Moderate	Moderate	Low	NI	Critical
Salo et al. <sup>3</sup>	Moderate	Serious	Moderate	Moderate	Moderate	Moderate	NI	Serious
Sansone et al.62	Low	Low	Low	Low	Low	Low	NI	Moderate
Shah et al. <sup>53</sup>	Moderate	Moderate	Low	Low	Low	Low	NI	Moderate
Shrotri et al. <sup>18</sup>	Moderate	Low	Low	Low	Moderate	Low	NI	Moderate
Souza et al. <sup>54</sup>	Low	Low	Low	Low	Low	Low	NI	Moderate



Tande et al. <sup>15</sup>	Moderate	Low	Low	Low	Low	Low	NI	Moderate
Tang et al. <sup>8</sup>	Moderate	Serious	Low	Moderate	Moderate	Low	NI	Serious
Tang et al. <sup>12</sup>	Moderate	Serious	Low	Moderate	NI	Low	NI	Serious
Thompson et al. <sup>31</sup>	Serious	Serious	Moderate	Moderate	Low	Low	Low	Serious

All studies conducted in 2021, NI, No information



# Vaccine Effectiveness against Infection Transmission

Six studies reported on the effectiveness of the PfBNT, Moderna, J&J, and AZ vaccines against disease transmission; four new studies were included in this update (**Error! Reference source not found.**). Studies included examined vaccine effectiveness against the wild-type or B.1.1.7 (Alpha) strain; none of the studies evaluated vaccine effectiveness against infection transmission against the B.1.617.2 (Delta) strain.

A retrospective cohort study in the Netherlands by de Gier et al. of 113,582 confirmed index cases of COVID-19 and 253,168 cohabitating household members or close contacts were assessed for vaccine effectiveness in preventing transmission to the household member or close contact and stratified by vaccination status, vaccine type, and days past date of inoculation.<sup>2</sup> At least one dose of PfBNT, Moderna, AZ, or J&J from past the 14<sup>th</sup> day of vaccination onwards, was associated with the reduction of transmission of COVID-19 to any household contact by 21% (95% CI: 12-28), 23% (95% CI: 14-32) to any unvaccinated household contact, 22% (95% CI: 9-33) to any other close contact, and 22% (95% CI: 8-34) to any unvaccinated close contact.<sup>2</sup> Fully vaccinated individuals with either PfBNT, Moderna, J&J, or AZ from past the 7<sup>th</sup> day of vaccination onwards, was associated with the reduction of transmission of COVID-19 to any household contact by 71% (95% CI: 63-77), 73% (95% CI: 65-79) to any unvaccinated household contact, 22% (95% CI: -5-43) to any other close contact, and 24% (95% CI: -5-43) to any unvaccinated close contact.<sup>2</sup> The low vaccine effectiveness of a fully vaccinated individual against transmission to any close contact or any close unvaccinated contact could be due to the studies being underpowered to detect differences due to the small number of events that occurred in vaccinated individuals compared to unvaccinated index cases.

de Gier et al. also stratified vaccine effectiveness by vaccine type and found that at least one dose of PfBNT from the 14<sup>th</sup> day of vaccination onward, reduced transmission to any household contact by 26% (95% CI: 12-37) and fully vaccinated individuals reduced transmission to any household contact by 70% (95% CI: 61-77).<sup>2</sup> At least one dose of Moderna from the 14<sup>th</sup> day of vaccination onward, reduced transmission to any household contact by 51% (95% CI: 8-74) and fully vaccinated individuals reduced transmission to any household contact by 88% (95% CI: 50-97).<sup>2</sup> At least one dose of AZ from the 14<sup>th</sup> day of vaccination onward, reduced transmission to any household contact by 48% (95% CI: 50-97).<sup>2</sup> At least one dose of AZ from the 14<sup>th</sup> day of vaccination onward, reduced transmission to any household contact by 15% (95% CI: 4-26) and fully vaccinated individuals reduced transmission to any household contact by 58% (95% CI: 12-84).<sup>2</sup> Fully vaccinated individuals with the J&J vaccine reduced transmission to household contact by 77% (95% CI: 6-94).<sup>2</sup>

A similar study was conducted in Finland by Salo et al. This retrospective cohort study investigated the vaccine effectiveness of 95, 138 mRNA-based (PfBNT or Moderna) vaccinated healthcare workers against infection transmission to unvaccinated household members compared to unvaccinated healthcare workers and their unvaccinated household members.<sup>3</sup> At least one dose of an mRNA-based vaccine from the 14<sup>th</sup> day of vaccination onward, reduced transmission to an unvaccinated spouse by 8.7% (95% CI: -28.9-35.4) and increased to 42.9% (95% CI: 22.3-58.1) reduction in transmission, 10 weeks after the first



dose.<sup>3</sup> At least one dose of an mRNA-based vaccine from the 14<sup>th</sup> day of vaccination onward, increased transmission to an unvaccinated child living in the household between the ages of 3-18 years by 1.0% (95% CI: -53.9-33.7) and decreased transmission to the unvaccinated child by 32.9% (95% CI: 4.1-53.0), 10 weeks after the first dose.<sup>3</sup>

Two studies from Israel found that PfBNT fully vaccinated individuals from past the 7<sup>th</sup> day of vaccination onward, had reduced infection transmission to their household contacts.<sup>4</sup> <sup>5</sup> A retrospective cohort study using a nationally centralized database investigated the vaccine effectiveness of PfBNT against infection transmission of two-adult households only one confirmed case of infection during the study period.<sup>4</sup> Of households with a fully vaccinated adult, the PfBNT vaccine was found to reduce infection transmission of the wild-type strain by 80.0% (95% CI: 73.0-85.1) compared to those who were unvaccinated and by 82.0% (95% CI: 75.5-86.7) compared to those who were recently vaccinated with one dose (between 0-7 days after vaccination).<sup>4</sup> A second Israeli study by Layan et al. conducted a case-control study of the PfBNT vaccine's effectiveness on reduction of infection transmission of the wild-type and B.1.1.7 (Alpha) strains in healthcare workers and their households.<sup>5</sup> The risk of transmission from vaccinated cases was 0.22 times (95% CI: 0.06-0.70) the risk of infection transmission compared to unvaccinated cases.<sup>5</sup>

Shah et al. in a retrospective study of 194,362 household members of 144,525 healthcare workers, who had received at least one dose of the PfBnT or AZ, found that from the 14<sup>th</sup> post-vaccination day onwards, vaccinating a co-habiting healthcare worker was associated with a significantly reduced risk of documented COVID-19 among household members (rate per 100 person-years: 9.40 versus 5.93; HR: 0.70, (95% CI: 0.63-0.78)).<sup>53</sup> The risk of hospitalization was also significantly lower among household contacts of vaccinated HCWs (rate per 100 person-years: 0.51 versus 0.31; HR: 0.77, (95% CI: 0.53-1.10)).<sup>53</sup> Following a second dose, the risks of infection and hospitalization involving a household member were significantly lower, rate per 100 person-years of 9.40 versus 2.98, HR: 0.46 (95% CI: 0.30-0.70) and 0.51 versus 0.22 per 100 person-years, HR: 0.68 (95% CI: 0.17-2.83), respectively).<sup>53</sup> The baseline serology and PCR of household contacts were not reported (**Error! Reference source not found.**).

A study by Harris et al. evaluated the risks of transmission of COVID-19 after one dose of PfBnT and AZ vaccination to unvaccinated household contacts using a retrospective design and a matched case-control method.<sup>57</sup> In the retrospective cohort analysis, there were 96,898 secondary cases among 960,765 household contacts of unvaccinated individuals (10.1%). There were 196 secondary cases in 3,424 contacts (5.72%) where the index case received AZ vaccine more than 21 days before PCR positivity, and 371 secondary cases in 5,939 contacts (6.25%) where the index case received the PfBnT vaccine. Adjusted odds ratio of transmission were 0.53 (95% CI: 0.43-0.63) and 0.51 (95% CI: 0.44-0.59), respectively, which were significantly lower.<sup>57</sup> In the matched case-control method, the odds of secondary infection among contacts of AZ and PfBnT vaccinated individuals were also significantly lower, 0.62 (95% CI: 0.48-0.79) and 0.51 (95% CI: 0.42-0.62) respectively.<sup>57</sup>

The baseline serology and PCR of household contacts were not reported in any of the studies except for Salo et al. and Gazit et al. whom only included seronegative participants (**Error! Reference source not found.**).<sup>3,4</sup>



# Vaccine Efficacy or Effectiveness Against Asymptomatic Infection

Twenty-five studies reported vaccine efficacy or effectiveness against asymptomatic COVID-19 infection, including 11 new observational studies and one new randomized control trial for this update (Table 7 and Table 8). Of the six RCTs included, three studied the AZ vaccine,<sup>6,7,16</sup> one evaluated the J&J vaccine,<sup>33</sup> and two studied the vaccine efficacy of the Moderna vaccine.<sup>32,51</sup>

Of the observational studies included, 12 examined the PfBNT vaccine,<sup>8,9,14,23-25,27,52,55,56,62,63</sup> two studied the AZ vaccine,<sup>21,54</sup> one evaluated the Moderna vaccine,<sup>11</sup> four studies evaluated both Moderna and PfBNT,<sup>10,12,31,60</sup> and two evaluated PfBNT and AZ vaccines.<sup>17,61</sup>

The methods of assessing efficacy or effectiveness against asymptomatic infection used in some of these studies included RT-PCR nasopharyngeal swabs at different time intervals.

## AstraZeneca Vaccine Efficacy in the General Population

No new additional studies investigating vaccine efficacy or effectiveness of the AZ vaccine against asymptomatic infections were included in this update.

#### First Dose AstraZeneca

#### Wild type

Asymptomatic infection data were presented for only the UK component of the AZ vaccine studies. Two AZ vaccine studies reported vaccine efficacy against asymptomatic or unknown infection of 7.8% (-46.7-42.1)<sup>7</sup> and 16% (-88-62)<sup>6</sup>, respectively, after more than 21 days and 22 to 90 days of the first dose. However, vaccine efficacy among participants with positive results, irrespective of symptoms, was 46.3% (31.8-57.8)<sup>7</sup> and 67% (49-78)<sup>6</sup>, respectively, over the same periods (Table 7). These trials implemented weekly self-administered nose and throat swabs for testing on baseline seronegative participants. The PCR status of these participants was not established at baseline.

#### Full Dose AstraZeneca

#### Wild type

After 14 days of the second dose, two AZ vaccine studies did not demonstrate efficacy against asymptomatic or unknown infection with the wild type virus: 22.2% (-9.9-45) and 27.3% (95% CI: -17-54.9), respectively.<sup>6,7</sup> A third study did not show efficacy against asymptomatic infection with the B.1.1.7 variant (26.5% (95% CI: -112-74.5)), following low or standard dose vaccination.<sup>16</sup> All three studies involved baseline seronegative participants. The baseline PCR results of the participants were not reported, therefore, persistent carriage after previous infection was not ruled out. In the subgroup of participants with an initial low dose of the vaccine, followed by a standard dose, two studies reported 49.3%(95% CI: 7.4-72.2)<sup>6</sup> and 58.9%(95% CI: 1-82.9)<sup>7</sup> respective efficacies against asymptomatic and unknown infection 14 days after the second dose (Table 8).



# AstraZeneca Vaccine Effectiveness in the General Population

#### Wild type

### First Dose AstraZeneca

In a large UK household survey with longitudinal follow-up among seronegative or seropositive individuals, Pritchard et al. reported significant reductions in the odds of asymptomatic infections following AZ vaccine 0-7 days, 8-20 days and 21 or more days after the first dose (ORs: 0.45 (95% CI: 0.35-0.57), 0.47 (95% CI: 0.37-0.6) and 0.39 (95% CI: 0.3 -0.51), respectively).<sup>17</sup> Nose and throat self-swabs were conducted every week for a month, and subsequently monthly for 12 months from enrolment.<sup>17</sup>

# Pfizer BioNTech Vaccine Effectiveness in the General Population

Of the 11 studies reporting PfBNT vaccine efficacy or effectiveness, five new additional studies investigating effectiveness of the Pfizer BioNTech vaccine against asymptomatic infections were included in this update.

#### First Dose Pfizer BioNTech Vaccine

## Wild type

An Israeli observational study by Dagan et al., which did not establish baseline seronegativity, showed that one dose of PfBNT significantly reduced asymptomatic infection by 29% (95% CI: 17-39) and 52% (95% CI: 41-60) after 14 to 20 days and 21 to 27 days of follow-up respectively, as assessed by confirmed positive PCR SARS-CoV-2 test without documented symptoms. No routine swabbing was documented for the participants (Table 7). In a large UK household survey with longitudinal follow-up involving participants with unknown baseline serology status, Pritchard et al. reported significant reductions in the odds of asymptomatic infections following PfBNT vaccine 0-7 days, 8-20 days and 21 or more days after the first dose, ORs: 0.48 (95% CI: 0.39-0.6) and 0.54 (95% CI: 0.45-0.65) respectively, compared with unvaccinated previously PCR negative individuals.<sup>17</sup> Nose and throat self-swabs were conducted every week for a month, and subsequently monthly for 12 months from enrolment.<sup>17</sup>

A retrospective study in Qatar by Tang et al. showed that seronegative healthcare workers with at least one dose of PfBNT were 0.58 times (95% CI: 0.3-1.12) less likely than unvaccinated healthcare workers to have an asymptomatic infection of the wild-type virus between 0-11 days past the date of vaccination and 0.58 times (95% CI: 0.3-1.13) less likely than unvaccinated healthcare workers to have an asymptomatic infection between  $\geq$ 12 days after the first dose and second dose.<sup>8</sup> This was a similar nonsignificant finding to an Israeli study by Angel et al. showing that seronegative healthcare workers that were inoculated with at least one dose of PfBNT were 0.48 times (95% CI: 0.19-1.26; p=0.12) less likely than unvaccinated healthcare workers to develop asymptomatic infection against the wild-type of the virus between 7 to 28 days past the vaccination date.<sup>9</sup>

#### Variants of Concern

A study conducted by Tang et al., which did not establish baseline seronegativity nor had a routine swabbing protocol, used a matched test-negative case-control method to investigate the effectiveness of at least one dose of PfBNT against the B.1.617.2 (Delta) variant at



preventing asymptomatic infection.<sup>12</sup> One dose of PfBNT past the 14<sup>th</sup> day onward from the date of vaccination was found to reduce asymptomatic infection by 25.2% (95% CI: 0.0-78.7).<sup>12</sup>

Full Dose Pfizer BioNTech Vaccine

# Wild Type

Several studies found that a full-dose of PfBNT significantly reduced asymptomatic infection from the wild-type strain.<sup>8-10</sup> Tang et al. found a reduction in transmission of asymptomatic infection of fully vaccinated seronegative Qatari healthcare workers between 0-6 days past the date of vaccination (IRR: 0.35 [95% CI: 0.11-1.09]) and from more than 7 days past the date of vaccination of the second dose (IRR: 0.10 [95% CI: 0.04-0.22]).<sup>8</sup> This finding was supported by Angel et al. who found similar significant reductions in asymptomatic infection. PfBNT fully vaccinated individuals were 0.09 times (95% CI: 0.03-0.25; p<0.01) less likely as unvaccinated individuals to have asymptomatic infection after at least 7 days from full vaccination and 0.09 times (95% CI:0.01-0.35; p=0.002) times as likely to have asymptomatic infection 21 days after full vaccination compared to unvaccinated individuals.<sup>9</sup> A retrospective cohort study by Andrejko et al. of 525 seronegative California residents found that a full-dose of PfBNT had a 68.3% (95% CI: 27.9-85.7%) reduction of asymptomatic infection of the wild-type strain.<sup>10</sup>

Dagan et al. also demonstrated 90% effectiveness (95% CI: 83-94) against asymptomatic infection seven days after the second dose from the wild-type or B.1.17 (Alpha) strain. In an Israeli study, which utilized the national public health surveillance data, Haas et al. reported significantly higher vaccine effectiveness seven or more days after full dose PfBnT vaccination, 90.4% (95% CI: 89.1-91.5).<sup>14</sup> The incidence rate per 100 000 person-days among unvaccinated individuals was 54.6 compared with 3.2 in those vaccinated. Vaccine effectiveness after 14 or more days was 93.8% (95% CI: 93.3-94.2).<sup>14</sup> Pritchard et al. also found full dose vaccination with PfBnT vaccine to significantly reduce the odds of asymptomatic infection compared with unvaccinated previously PCR negative UK residents, 0.48 (95% CI: 0.36-0.66).<sup>17</sup>

#### Variants of Concern

The study conducted by Tang et al. found that a full-dose of PfBNT past the 14<sup>th</sup> day onward from the date of vaccination was found to reduce asymptomatic infection by 35.9% (95% CI: 11.1-53.9) against the B.1.617.2 (Delta) variant.<sup>12</sup>

# mRNA (Pfizer BioNTech and Moderna) Vaccines Effectiveness in the General Population

One new study was included in this update that examined the effectiveness of the Pfizer BioNTech or Moderna vaccine against asymptomatic infection.

#### First or second dose of mRNA vaccine

#### Wild type

Tande et al. evaluated the effectiveness of at least one dose of either Moderna or PfBnT vaccine among people who underwent molecular tests prior to a procedure or surgery.<sup>15</sup> The



relative risk for a positive test during asymptomatic pre-procedure screening in vaccinated compared with unvaccinated was significantly lower (0.44 (95% CI: 0.33-0.60)). Ten or more days after the 1st dose, the risk of a positive test was also significantly lower among the vaccinated (0.28 (95% CI: 0.16-0.49; p<.0001)). The risk of test positivity was similarly lower among the vaccinated, after the second dose 0.27 (95% CI: 0.12-0.60).<sup>15</sup>

# Variants of Concern

Similar to the studies by Tande et al. and Bouton et al., Tang et al. found that use of at least one dose of an mRNA-based vaccine past the 14<sup>th</sup> day onward from the date of vaccination reduced asymptomatic infection by 44.3% (95% CI: 0-78.4) against the B.1.617.2 (Delta) variant.<sup>12</sup>

# Moderna Vaccine Efficacy in the General Population

Of the four included studies reporting on Moderna, three new additional studies investigating vaccine efficacy (n= 1) or effectiveness (n=2) of the Moderna vaccine against asymptomatic infections were included in this update.

## First and/or Full Dose Moderna Vaccine

## Wild type

A study of the Moderna vaccine by Baden et al. reported that 0.1% of the participants receiving the first dose developed asymptomatic infection, assessed at the time of second dose with nasal swabs, compared with 0.27% of the unvaccinated group 21 days after the first dose, which is suggestive of 61.4% efficacy against asymptomatic carriage of the wild-type strain. Participants in this trial were negative for COVID-19 by RT-PCR or antibody testing at baseline.<sup>32</sup>

A phase 2/3 randomized-control trial by Ali et al. in 3732 seronegative adolescents (age 12-17 years) in the USA evaluating the vaccine efficacy of the Moderna vaccine is ongoing.<sup>51</sup> Participants were randomly assigned in a 2:1 ratio to receive two injections of either the Moderna vaccine (n=2139), each containing 100 µg or placebo (n=1042), 28 days apart. The study reported 0.97% of participants with an asymptomatic infection of those with a PCRpositive swab with no respiratory symptoms present 14 days after the first dose compared to 2.70% of unvaccinated adolescents with asymptomatic infection; this is suggestive of 59.5% (95% CI: 28.4-77.3) vaccine efficacy against asymptomatic carriage of the wild-type strain.<sup>51</sup> Fourteen days after the second dose, 1.17% of adolescents were had an asymptomatic infection as confirmed by the presence of a PCR-positive swab compared to 1.54% of unvaccinated adolescents with asymptomatic carriage of wild-type strain.<sup>51</sup> Full vaccination of Moderna is suggestive of 39.2% (95% CI: -24.7-69.7) vaccine efficacy against asymptomatic carriage of the wild-type strain.<sup>51</sup>

# Moderna Vaccine Effectiveness in the General Population

First Dose Moderna Vaccine

Wild Type



A test-negative case-control study by Chemaitelly et al. in Qatar suggested a vaccine efficacy of 47.3% (95% CI: 36.7-55.5) against asymptomatic infection, defined as a PCR-positive test conducted with no reported symptoms compatible with a respiratory tract infection, at least 14 days after a first dose.<sup>11</sup>

#### Variants of Concern

Another Qatari study by Tang et al. found that use of at least one dose of the Moderna vaccine past the 14<sup>th</sup> day from the date of vaccination reduced asymptomatic infection by 57.4% (95% CI: 0-92.9) against the B.1.617.2 (Delta) variant.<sup>12</sup>

Full Dose Moderna Vaccine

#### Wild type

Chemaitelly et al. found that a full dose of Moderna was 92.5% effective (95% CI: 84.8-96.9) against asymptomatic carriage 14 days after full vaccination.<sup>11</sup>

#### Variants of Concern

Tang et al. found 80.2% vaccine effectiveness (95% CI: 54.2-92.6) against asymptomatic carriage of the Delta strain 14 days after full vaccination.<sup>12</sup>

#### Janssen Vaccine Efficacy in the General Population

No new additional studies investigating vaccine efficacy or effectiveness of the J&J vaccine against asymptomatic infections were included in this update.

#### Full Dose Janssen vaccine

This is a single dose vaccine. The J&J vaccine did not show statistically significant efficacy against asymptomatic infection in the first 29 days of follow-up. However, after 29 days post-vaccination, asymptomatic infection, assessed via surveillance swabs at unspecified intervals among baseline seronegative participants, was significantly lower among vaccinated participants (74%, 95% CI: 46.8-88.4%).<sup>33</sup> Asymptomatic infection in this trial was assessed by lack of symptoms on the day preceding, the day of, or any time after a positive PCR test. Furthermore, efficacy as demonstrated by seroconversion in previously asymptomatic participants was 74.2% compared with placebo (95% CI: 47.1; 88.6).<sup>33</sup>



Table 6: Observational Studies of Vaccine Effectiveness Against Transmission to Household Contacts

			_	Follow-up		
Vaccine	Author	Country	Dose	days*	Outcomes*	Vaccine Effectiveness (95%CI)*
	1.1			44.40	Transmission to household	
	Harris et al.57	England	1	14-16	contact	OR: 0.73 (0.62-0.83) [Estimated VE: 25%]
	1.1			2.04	Transmission to household	
	Harris et al.57	England	1	≥21	contact	aOR: 0.51 (0.44, 0.59) [Estimated VE: 46%]
					Transmission to household	
	Harris et al.57	England	1	28-34	contact	OR: 0.62 (0.52-0.74) [Estimated VE:35%]
					Transmission to household	
	de Gier et al. <sup>2</sup>	Netherlands	1	>14	contact	Adjusted VE 26% (95%CI: 12-37)
				-	Transmission to household	
	de Gier et al. <sup>2</sup>	Netherlands	2	>7	contact	Adjusted VE 70% (95% CI: 61-77)
	0 1 - 1 - 1 4	la na al		0.7	Transmission to vaccinated	
	Gazit et al.4	Israel	1	0-7	household contact	94% (95% Cl: 90.8-95.7)
	0 1 - 1 - 1 4	lawa al	0	7	Transmission to vaccinated	
	Gazit et al.4	Israel	2	>7	household contact	70.1% (95% CI: 61.3-76.9)
Pfizer, BioNTech	1	la na al	0	7	HCW transmission to	700( (050( 01 00 04)
(BNT162b2)	Layan et al.⁵	Israel	2	>7	household contact	78% (95% CI: 30-94)
		No the other state			Transmission to household	
	de Gier et al. <sup>2</sup>	Netherlands	1	>14	contact	Adjusted VE 51% (95%CI: 8-74)
	de Cienet el 2	N a the and a reade	~	. 7	Transmission to household	
Moderna (mRNA-1273)	de Gier et al. <sup>2</sup>	Netherlands	2	>7	contact	Adjusted VE 88% (95% CI: 50-97)
		No the other state			Transmission to household	
Janssen (Ad26.COV2.S)	de Gier et al. <sup>2</sup>	Netherlands	1	>14	contact	Adjusted VE 77% (95% CI: 6-94)
	Llarria at al 57	E a al a a d	1	44.40	Transmission to household	OD: 0.70 (0.00 0.00) [Estimated)/[E: 040/]
	Harris et al.57	England	1	14-16	contact	OR: 0.78 (0.66-0.92) [Estimated VE: 21%]
	Llarria at al 57	E a al a a d	1	≥21	Transmission to household	- OD: 0.52 (05% OL 0.42 0.62) [Estimated a)/[5: 44%]
	Harris et al.57	England		221	contact Transmission to household	aOR: 0.53 (95% CI 0.43-0.63) [Estimated aVE: 44%]
	Harris et al.57	England	1	28-34	contact	OP: 0.44 (0.34.0.58) [Estimated \/E: 549/1
	nams et al."	Englanu	1	20-34	Transmission to household	OR: 0.44 (0.34-0.58) [Estimated VE: 54%]
	de Gier et al.2	Netherlands	1	>14	contact	Adjusted V/E 159/ (059/ CI: 4.20)
AstraZeneca		inemenanus		>14	Transmission to household	Adjusted VE 15% (95%CI: 4-26)
(ChAdOx1 nCoV-19)	de Gier et al.2	Netherlands	2	>7	contact	Adjusted VE 58% (95% CI: -12-84)
	ue Giel el al.	inetheriarius	۷	21	HCW Transmission to	Aujusieu VE 30 /0 (93 /0 01 12-04)
	Shah et al.53	Scotland	1	7-13	household	-8% (95% Cl: -25 - 6)
BNT162b2 or ChAdOx1	Shall et al.	Scollariu		7-15	HCW Transmission to	-070 (9570 0125 - 0)
nCoV-19	Shah et al.53	Scotland	1	14-20	household	15% (95%Cl:1-27)
11004-13	Shan et al.	Scollanu	I	14-20	nousenoiu	1070 (907001.1-27)



					HCW Transmission to	
	Shah et al.53	Scotland	1	>28	household	36% (95% CI: 27-44)
					HCW Transmission to	
	Shah et al.53	UK	2	>14	household	54%(95% CI: 30-70)
					Transmission to household	
					contact (unvaccinated	
	Salo et al. <sup>3‡</sup>	Finland	1	14	spouse)	8.7% (95% CI: -28.9-35.4)
					Transmission to household	
					contact (unvaccinated	
	Salo et al. <sup>3‡</sup>	Finland	1	70	spouse)	42.9% (95% CI: 22.3-58.1)
					Transmission to household	
					contact (unvaccinated child	
	Salo et al. <sup>3‡</sup>	Finland	1	14	3-18 years)	-1% (95% CI: -53.9-33.7)
					Transmission to household	
					contact (unvaccinated child	
	Salo et al. <sup>3‡</sup>	Finland	1	70	3-18 years)	32.9% (95% CI: 4.1-53.0)
					Transmission to household	
					contact (unvaccinated child	
	Salo et al. <sup>3‡</sup>	Finland	1	42	3-12 years)	12.3% (95% CI: -37.4-44.0)
					Transmission to household	
					contact (unvaccinated child	
	Salo et al. <sup>3‡</sup>	Finland	1	70	3-12 years)	22.3% (95%Cl: -34.4-55.2)
					Transmission to household	
					contact (unvaccinated child	
	Salo et al. <sup>3‡</sup>	Finland	1	42	13-18 years)	16.7% (95% CI: -17.7-41.0)
					Transmission to household	
Pfizer, BioNTech or					contact (unvaccinated child	
Moderna, mRNA-1273	Salo et al. <sup>3‡</sup>	Finland	1	70	13-18 years)	38% (95% CI: 1.2-61.1)
					Transmission to household	
	de Gier et al.2	Netherlands	1	>14	contact	Adjusted VE 21% (95% CI: 12-28)
					Transmission to	
					unvaccinated household	
	de Gier et al.2	Netherlands	1	>14	contact	Adjusted VE 23% (95% CI: 14-32)
					Transmission to any other	
	de Gier et al.2	Netherlands	1	>14	close contact	Adjusted VE 22% (95% CI: 9-33)
					Transmission to any	
	de Gier et al.2	Netherlands	1	>14	unvaccinated close contact	Adjusted VE 22% (95% CI: 8-34)
					Transmission to household	
	de Gier et al.2	Netherlands	2	>7	contact	Adjusted VE 71% (95% CI: 63-77)
Pfizer, BioNTech or					Transmission to	
Moderna, mRNA-1273 or					unvaccinated household	
AstraZeneca, ChAdOx1	de Gier et al.2	Netherlands	2	>7	contact	Adjusted VE 73% (95% CI: 65-79)
nCoV-19 or					Transmission to any other	
Janssen, Ad26.COV2.S	de Gier et al.2	Netherlands	2	>7	close contact	Adjusted VE 22% (95% CI: -5-43)



					Transmission to any	
d	de Gier et al. <sup>2</sup>	Netherlands	2	>7	unvaccinated close contact	Adjusted VE 24% (95% CI: -5-43)

\*None of the studies excluded other sources of exposure. VE: Vaccine Effectiveness, OR: Odds Ratio, + VE =1-RR (or HR) x100%, where RR is the reported relative risk or Hazard ratio; or derived from reported baseline prevalence in unvaccinated group and OR<sup>64</sup>.

<sup>‡</sup>All studies included participants with unknown baseline serology except for Salo et al. whom included participants who were seronegative



#### Table 7: First-dose Vaccine Efficacy or Effectiveness Against Asymptomatic Infection

Vaccine	Author	Country	Strain targeted by PCR	Baseline Serology	Dosing Schedule	Follow-up days*	Outcomes	Vaccine Efficacy or Effectiveness (95%Cl) <sup>+</sup>
	Voysey et al. <sup>7</sup> (RCT)	UK	Milet to me	Manativa		. 04	Asymptomatic	7.00/ ( 40.7.40.4)
	Voysey et al. <sup>7</sup> (RCT)	UK	Wild type	Negative	LD or SD	>21	or unknown	7.8% (-46.7-42.1)
	voysey et al. (RCT)	UK/Brazil/South.	Wild type	Negative	LD or SD	>21	Any PCR+	46.3% (31.8-57.8)
	Voysey et al. <sup>6</sup> (RCT)	Afrca	Wild type	Negative	LD or SD	22-90	Any PCR+	67% (49-78)
	Voysey et al. <sup>6</sup> (RCT)	UK/Brazil/South. Afrca	Wild type	Negative	SD	22-30	Asymptomatic or Unknown	0.2 (-209-68)
	Voysey et al. <sup>6</sup> (RCT)	UK/Brazil/South. Afrca	Wild type	Negative	SD	31-60	Asymptomatic or Unknown	17% (-172-75)
	Voysey et al. <sup>6</sup> (RCT)	UK/Brazil/South. Afrca	Wild type	Negative	SD	22-90	Asymptomatic or unknown	16% (-88-62)
	Pritchard et al. <sup>17</sup>	UK	Wild type and B.1.1.7	Both	NA	0-7	Asymptomatic	OR: 0.45(0.35 to 0.57)
AstraZenec	Pritchard et al. <sup>17</sup>	UK	Wild type and B.1.1.7	Both	NA	8-20	Asymptomatic	OR: 0.47(0.37 to 0.6)
<b>a</b> (ChAdOx1	Pritchard et al. <sup>17</sup>	UK	Wild type and B.117	Both	NA	≥ 29	Asymptomatic	OR: 0.39(0.3 to 0.51)
nCoV-19)	Souza et al.54	Brazil	B.1.1.7 (UK)	NR	NA	>23 days	asymptomatic	NR
Janssen Biotech	Janssen Biotech <sup>33</sup> (RCT)	Multiple	Wild type	Negative	NA	1-29	Asymptomatic	20% (-7-40.4)
(Ad26.COV2 .S)	Janssen Biotech <sup>33</sup> (RCT)	Multiple	Wild type	Negative	NA	≥ 29	Asymptomatic	74% (46.8-88.4)
	Amit et al. <sup>52</sup>	Israel	Wild type	Unknown	NA	1-14	Asymptomatic or unknown	NR##
	Amit et al. <sup>52</sup>	Israel	Wild type	Unknown	NA	15-28	Asymptomatic or unknown	NR###
	Dagan et al. <sup>13</sup>	Israel	Wild type and B.1.1.7	Unknown	NA	14-20	Asymptomatic	29% (17-39)
	Dagan et al. <sup>13</sup>	Israel	Wild type and B.1.1.7	Unknown	NA	21-27	Asymptomatic	52% (41-60)
	Hall et al. <sup>56</sup>	UK	Wild type	Unknown	NA	21 days after 1 <sup>st</sup> dose and 7 days after 2 <sup>nd</sup>	Asymptomatic or unknown	97.2%#
	Haas et al. <sup>14</sup>	Israel	Wild type and B.1.1.7	Unknown	NA	14-21 days	Asymptomatic	52% (48·9–55·0)
	Pritchard et al. <sup>17</sup>	UK	Wild type and B.117	Both	NA	0-7	Asymptomatic	OR: 0.48(0.39 to 0.6)
Pfizer, BioNTech	Pritchard et al. <sup>17</sup>	UK	Wild type and B.117	Both	NA	8-20	Asymptomatic	OR:0.54(0.45 to 0.65)
(BNT162b2)	Pritchard et al. <sup>17</sup>	UK	Wild type and B.1.1.7	Both	NA	≥ 29	Asymptomatic	OR: 0.44(0.36 to 0.55)



Vaccine	Author	Country	Strain targeted by PCR	Baseline Serology	Dosing Schedule	Follow-up days*	Outcomes	Vaccine Efficacy or Effectiveness (95%Cl) <sup>+</sup>
	Regev-Yochay et al. <sup>25</sup>	Israel	Wild type	Both	NA	4-10	Asymptomatic at first test	28(-18 to 57)
	Regev-Yochay et al. <sup>25</sup>	Israel	Wild type	Both	NA	4-10	Asymptomatic (who never became symptomatic)	27(-38 to 61)
	McEllistrem et al.55	USA	Wild Type	Unknown	NA	12-15 days	Asymptomatic	NR
	Jones et al. <sup>23</sup>	UK	Wild type and B.1.1.7	Unknown	NA	<12 and >12	Asymptomatic	NR
	Tang et al.12	Qatar	VOC delta	NR	NA	≥ 14	Asymptomatic	25.2 (0.0-78.7)
	Angel et al. <sup>9</sup>	Israel	Wild type	Seronegativ e	NA	7-28	Asymptomatic	adjusted IRR (95%CI): 0.48(0.19- 1.26); p=0.12
	Tang et al.8	USA	Wild type	Seronegativ e	NA	0-11	Asymptomatic	IRR(95%CI): 0.58 (0.3-1.12)
	Tang et al.8	USA	Wild type	Seronegativ e	NA	>12	Asymptomatic	IRR(95%CI): 0.58 (0.3-1.13)
	McEllistrem el al.24	USA	Wild type	Unclear	NA	1-21	Asymptomatic	NR
	McEllistrem el al.24	USA	Wild type	Unclear	NA	14-21	Asymptomatic	NR
	Baden et al. <sup>32</sup> (RCT)	USA	Wild type	Negative	NA	From day 1	Asymptomatic	61.4% <sup>#</sup>
	Ali et al. <sup>51</sup> (RCT)	USA	Wild type	Negative	NA	>14	Asymptomatic	59.5%(28.4 to 77.3)
Moderna	Tang et al.12	Qatar	VOC delta	NR	NA	≥ 14	Asymptomatic	57.4 (0.0-92.9)
<b>(</b> mRNA- 1273)	Chemaitelly et al. <sup>11</sup>	Qatar	Wild type and VOC B.1.1.7; B.1.351; B.1.617	Unknown	NA	14	Asymptomatic	47.3%(37.6-55.5%)
mRNA vaccines (BNT162b2	Tande et al. <sup>15</sup>	USA		Unknown	NA	From day 1, at least one dose	PCR+ in asymptomatic	56% (40-67)
or Moderna mRNA-	Bouton et al. <sup>60</sup>	USA	Wild type	Unknown	NA	From day 1, 1 dose	Asymptomatic	RR calculated: -1% (-57-35)
1273)	Tang et al. <sup>12</sup>	Qatar	VOC delta	NR	NA	≥ 14	Asymptomatic	44.3 (0.0-78.4)

\* Time PCR was conducted after first or second dose; LD: Low dose, SD: Standard dose, #Calculated from raw values. ## 2.7 cases per 10000 person-days in vaccine vs 2.4 cases per 10000 person-days in control. ###1.7 cases per 10000 person-days in vaccine vs 2.4 cases per 10000 person-days, Calculated from raw values).\* Efficacy reported for RCTs and Effectiveness for observational studies. All Pfizer BioNtech's studies except Dagan et al. involved healthcare workers. 0.44 (95% CI: 0.33-0.60). Studies are observational except otherwise stated. , \* VE =1-RR (or HR) x100%



Table 8: Full-dose Vaccine Efficacy or Effectiveness Against Asymptomatic Infection

Vaccine	Author	Country	Strain targeted by PCR	Baseline Serology	Dosing Schedule	Follow-up days*	Outcomes	Vaccine Efficacy or Effectiveness (95%CI)*
					LD or SD		Asymptomatic or	
	Voysey et al. <sup>7</sup> (RCT)	UK	Wild type	Negative	and SD	>14	unknown	27.3% (-17-54.9)
					LD and		Asymptomatic or	
	Voysey et al.7 (RCT)	UK	Wild type	Negative	SD	>14	unknown	58.9% (1-82.9)
	$\lambda$ (average et al. <sup>7</sup> (DCT)	UK	Wild type	Negotivo	SD and SD	- 14	Asymptomatic or	2.90/(72.4.46.2)
	Voysey et al. <sup>7</sup> (RCT)	UK	Wild type	Negative	LD or SD	>14	unknown	3.8% (-72.4-46.3)
	Voysey et al.7 (RCT)	UK	Wild type	Negative	and SD	>14	Any PCR+	55.7% (41.1-66.7)
					LD or SD		Í	
	Voysey et al.6 (RCT)	UK	Wild type	Negative	and SD	>14	Any PCR+	54.1% (44.7%, 61.9%)
					LD or SD		Asymptomatic	
	Voysey et al. <sup>6</sup> (RCT)	UK	Wild type	Negative	and SD	>14	or unknown	22.2% (-9.9-45)
					SD and		Asymptomatic or	
	Voysey et al.6 (RCT)	UK	Wild type	Negative	SD	>14	unknown	2.0% (-50.7-36.2)
	Voysey et al. <sup>6</sup> (RCT)	UK	Wild type	Negotivo	LD and SD	>14	Asymptomatic or unknown	40,29/ (7,4,72,2)
		UK		Negative	LD or SD	>14	UNKNOWN	49.3% (7.4-72.2)
	Emary et al. <sup>16</sup> (RCT)	UK	Wild type, B.1.1.7, Other	Negative	and SD	>14	Asymptomatic	15.7% (-10.7-35.8)
				0	LD or SD			
AstraZeneca	Emary et al. <sup>16</sup>	UK	B.1.1.7	Negative	and SD	>14	Asymptomatic	26.5% (-112-74.5)
(ChAdOx1					LD or SD			
nCoV-19)	Emary et al. <sup>16</sup>	UK	Variants not B.1.1.7	Negative	and SD	>14	Asymptomatic	75.4% (39.9-89.9)
Janssen Biotech	Janssen Biotech <sup>33</sup> (RCT)	Multiple	Wild type	Negative	NA	1-29	Asymptomatic	20% (-7-40.4)
(Ad26.COV2.S)	Janssen Biotech <sup>33</sup> (RCT)	Multiple	Wild type	Negative	NA	≥ 29	Asymptomatic	74% (46.8-88.4)
	Dagan et al. <sup>13</sup>	Israel	Wild type and B.1.1.7	Unknown	NA	>7	Asymptomatic	90% (83-94)
						21 days after 1 <sup>st</sup> dose		
						and 7 days after	Asymptomatic or	
	Hall et al. <sup>56</sup>	UK	Wild type	Unknown	NA	2 <sup>nd</sup>	unknown	NR
	Haas et al.14	Israel	Wild type and B.1.1.7	Unknown	NA	≥7	Asymptomatic	91.5% (90.7–92.2)
	Haas et al.14	Israel	Wild type and B.1.1.7	Unknown	NA	≥14	Asymptomatic	93.8% (93.3–94.2)
Pfizer	Regev-Yochay et al. <sup>25</sup>	Israel	Wild	Both	NA	>10	Asymptomatic at first testing	65% (45 to 79)
BioNTech	Denou Veebou et al 25	lana al		Deth	NIA	. 10	Asymptomatic	700// 40.000/)
(BNT162b2)	Regev-Yochay et al. <sup>25</sup>	Israel	Wild	Both	NA	>10	(who never	72%( 48-86%).



Vaccine	Author	Country	Strain targeted by PCR	Baseline Serology	Dosing Schedule	Follow-up days*	Outcomes	Vaccine Efficacy or Effectiveness (95%CI) <sup>+</sup>
							became symptomatic)	
	Pritchard et al.17	UK	Wild type and B.117	Both	NA	NR	Asymptomatic	OR:0.48(0.36 to 0.66)
	Pfizer [Press Release] <sup>50</sup>	Israel	Wild type and B.1.1.7	Unknown	NA	>14	Asymptomatic	94%
	Tang et al.12	Qatar	VOC Delta	NR	NA	≥14	Asymptomatic	35.9 (11.1-53.9)
	Sansone et al. <sup>62</sup>	Italy	Wild type and VOC B.1.1.7 (Alpha/UK), B1.525	NR	NA	>14	Asymptomatic	OR: 0.38 (0.16-0.88)
	McEllistrem et al.24	USA	Wild type	Unclear	NA	7	Asymptomatic	NR
	Bailly et al.27	France	VOC 501Y.V2	Both	NA	NR	Asymptomatic	NR
	Andrejko et al. <sup>10</sup>	USA	Wild type	Seronegative	NA	>15	Asymptomatic	68.3% (27.9-85.7%)
	Tang et al.8	USA	Wild type	Seronegative	NA	0-6	Asymptomatic	IRR: 0.35 (0.11-1.09)
	Tang et al.8	USA	Wild type	Seronegative	NA	>7	Asymptomatic	IRR: 0.10 (0.04-0.22)
	Angel et al.9	Israel	Wild type	Seronegative	NA	>7	Asymptomatic	adjusted IRR: Vax 0.09(0.03-0.25); p<0.001
	Angel et al.9	Israel	Wild type	Seronegative	NA	>21	Asymptomatic	adjusted IRR: Vax 0.09(0.01-0.35); p=0.002
	Ali et al. <sup>51</sup> (RCT)	USA	Wild type	Seronegative	NA	>14	Asymptomatic	39.2% (-24.7 to 69.7)
Moderna	Chemaitelly et al.11	Qatar	Wild type and VOC B.1.1.7; B.1.351; B.1.617	NR	NA	>14	Asymptomatic	92.5%(84.8-96.9%)
(mRNA-1273)	Tang et al.12	Qatar	VOC Delta	NR	NA	≥14	Asymptomatic	80.2 (54.2-92.6)
BNT162b2 and ChAdOx1	Shah et al. <sup>53</sup>	UK	Wild type	Unknown	NA	>14	HCW Transmission to household	HR=0.46 (95% CI: 0.30-0.70)
nCoV-19	Tang et al.12	Qatar	VOC Delta	NR	NA	≥14	Asymptomatic	40.0 (18.2-56.1)

\* Time PCR was conducted after first or second dose; LD: Low dose, SD: Standard dose, #Calculated from raw values.\* Efficacy reported for RCTs and Effectiveness for observational studies. Hall et al. was the only full-dose study healthcare workers' data. Studies are observational, except otherwise state



Table 9: Ct Values

Vaccine	Author	Country	Virus type	Baseline Serology	1st or 2nd dose	Follow- up days	Sub-population	Ct Values (Vaccinated), Median (IQR), unless otherwise specified	Ct Values (Unvaccinated), Median (IQR), unless otherwise specified	Effect size/p-values
							Breakthrough	27.8 (21.1-32.7)	25.8 (19.5-31.4)	p <0.001
	Abu-	Qatar	Wild type and B.1.1.7, B.1.351,	Unknown	2	<u>≥</u> 14	infection vs primary infection in unvaccinated	Mean=26.8 (95% Cl: 26.5- 27.2)	Mean=25.5 (95% Cl: 25.2- 25.8)	Mean difference (95% CI): 1.3 (0.9- 1.8), p<0.001
	Raddad <sup>26</sup>	Qatar	B.1.617.2	OTIKHOWIT			Breakthrough	28.2 (21.1-33.1)	31.2 (24.3-33.9)	p <0.001
					2	<u>≥</u> 14	infection vs reinfection in unvaccinated	Mean=27 (95% CI: 26.3- 27.6)	Mean=28.9 (95% Cl: 28.3- 29.5)	Mean difference (95% Cl): 2.0 (1.1- 2.8), p <0.001
	Bailly <sup>27</sup>	France	501Y.V2	Both	2	NR		21 (13-32)	15 (12-17)	p=0.05
	loannou <sup>28</sup>	Greece	B.1.1.7		2	>14		18 (15.5-25.5)	18.5 (13.5-24)	ns
	Jones <sup>23</sup>	UK	Wild-type and B.1.1.7	Unknown	1	<u>&gt;</u> 12		30.3 (25.5-35.1)	23.3 (13.5-33)	ns
BNT162b2	Levine- Tiefenbrun <sup>22</sup>				1	<12		NR	NR	no significant differences in the Ct values for any of the 3 genes (RdRp, N and E)
			Wild type	Unknown	1	12-28		NR	NR	the Ct values for the 3 genes were significantly higher among infected vaccinated persons than controls (p<10 <sup>-8</sup> )
	McEllistrem <sup>24</sup>	USA	Wild type	Unknown	1	NR	Asymptomatic COVID-19	19.4 (18.9-22.5)	12.8 (12.4-14.9)	p=0.009
	Muhsen58	Israel	Wild type	Seronegative	2	>14		32 (14.5)	26.7 (8.8)	p=0.008
	Regev- Yochay <sup>25</sup>	Israel	Wild type	Both	2	<u>&gt;</u> 11		Mean=27.3 (SD=2.2)	Mean=22.2 (SD=1.0)	Mean difference (95% CI): 5.09 (2.8- 7.4), p<0.001



Vaccine	Author	Country	Virus type	Baseline Serology	1st or 2nd dose	Follow- up days	Sub-population	Ct Values (Vaccinated), Median (IQR), unless otherwise specified	Ct Values (Unvaccinated), Median (IQR), unless otherwise specified	Effect size/p-values
	Baltas <sup>20</sup>	UK	Wild type and B.1.1.7	Unknown	1	9-24		30.8 (25.9-35.4)	28.8 (25.3-33.7)	P=0.053
	Lumley <sup>21</sup>	England	Wild type and B.1.1.7 (35% of unvaccinated seronegative; 65% vaccinated)	Seronegative	NR	NR		Change in median: 2.7 (-0.5 to 6.7)		NR
		England	Wild type and VOC	Seronegative	NR	NR		Mean=19.66 (95% Cl: 15.01- 27.53)	Mean=18.39 (95% Cl: 14.00- 25.57)	p=0.19
BNT162b2 or ChAdOx1	Mostafa <sup>19</sup>	USA	Wild type, P.1, B.1.17 (61%), B.1.351, B.1.526 (9%), and B.1.526.1 (4.5%)	Unknown	2	2 to 100		19.26 (Q1, Q3: 16.56 to 21.96)	19.6 (Q1, Q3: 16.28 to 22.66)	ns
		UK	Wild type	Both	1	0-7		31.2 (20.6-33.7)	28.4 (20.1-32.9)	p<0.001
	Duitak and <sup>17</sup>				1	8-20		31 (23.5 to 33.8)	28.4 (20.1 to 32.9)	p<0.001
	Pritchard <sup>17</sup>	UK			1	≥21		31.7 (26.9 to 33.7)	28.4 (20.1 to 32.9)	p<0.001
					2	NR		33.1 (30.5 to 34.2)	28.4 (20.1 to 32.9)	p<0.001
	Shrotri <sup>18</sup>	UK	Wild type	Determined (adjusted for in analysis)	1	0-27		26.9 (25.19- 26.62)	26.6 (26-27.1)	0.158
					1	0 to 20		30.93 (Q1, Q3: 22.93 to 33.71)		ns from unvaccinated but previously
BNT162b2, ChAdOx1, or Moderna	Pouwels <sup>59</sup>	UK	Alpha, Delta	Both	1 or 2	>21 for dose 1 or 0-13 for dose 2	Alpha (1 Dec 2020 to 16 May 2021)	31.71(Q1, Q3: 26.64 to 33.57)	28.7 (Q1, Q3: 20.4, 32.9) for not previously PCR positive; 32.8 (Q1, Q3: 30.9-	PCR/antibody- positive (age/sex- adjusted p=0.72), but significantly higher than in those
mRNA- 1273						>14		33.3 (Q1, Q3: 31.6 to 34.0)	34.2) for previously PCR positive	unvaccinated and not previously PCR/antibody- positive, age/sex- adjusted p=0.02).



Vaccine	Author	Country	Virus type	Baseline Serology	1st or 2nd dose	Follow- up days	Sub-population	Ct Values (Vaccinated), Median (IQR), unless otherwise specified	Ct Values (Unvaccinated), Median (IQR), unless otherwise specified	Effect size/p-values
					1	0 to 20		29.93 (Q1, Q3: 22 to 34.21)	21.5 (Q1, Q3: 16.5 to 31.64) for	
					1 or 2	>21 for dose 1 or 0-13 for dose 2	Delta (17 May 2021 to 13 Jun 2021)	30.07 (Q1, Q3: 18.64 to 33.64)	not previously PCR positive; 30.86 (Q1, Q3: 29.5 to 34.28) for	NR
				2	>14		32.29 (Q1, Q3: 26.07 to 33.93)	previously PCR positive		
					1	0 to 20		25.64 (Q1, Q3: 21.64 to 30.79)	25.71 (Q1, Q3:	
					1 or 2	>21 for dose 1 or 0-13 for dose 2	Delta (14 Jun 2021 to 1 Aug 2021)	24.64 (Q1, Q3: 18.86 to 31.29	19.07 to 30.71) for not previously PCR positive; 22.29 (Q1, Q3: 16.57 to 30.29)	NR
					2	>14		25.29 (Q1, Q3: 19.21 to 31.29)	for previously PCR positive	
BNT162b2					1	≤ 14	Early post vaccination vs. unvaccinated	Mean=22.6 (SD=7)	Mean=23 (SD=7.4)	NR
or Moderna mRNA- 1273	Jacobson <sup>29</sup>	USA	Wild type and B.1.427/B.1.429 (34.3%)	Both	1 or 2	up to 14 after 2nd dose	Partially vaccinated vs unvaccinated	Mean=27.7 (SD=8.7)	Mean=23 (SD=7.4)	NR
					2	>14	Fully vaccinated vs unvaccinated	Mean=28.5 (SD=7.4)	Mean=23 (SD=7.4)	NR
BNT162b2.			Wild type, B.1.1.7,		1 or 2	<u>≥</u> 14	Vaccination breakthrough infections	27 (13-42)	<u>&lt;</u> 30	NR
Moderna, or Janssen	Duerr <sup>30</sup>	USA	others	Unknown	1 or 2	<u>≥</u> 14	Vaccination breakthrough infections that passed quality control	24 (13-36)	<u>≤</u> 30	NR
ChAdOx1	Emary <sup>16</sup>	UK	Wild type and B.1.1.7	Unknown	2	<u>&gt;</u> 14	Asymptomatic	30.25 (24.81- 34.20)	28.15 (19.51- 32.35)	p=0.0040



Vaccine	Author	Country	Virus type	Baseline Serology	1st or 2nd dose	Follow- up days	Sub-population	Ct Values (Vaccinated), Median (IQR), unless otherwise specified	Ct Values (Unvaccinated), Median (IQR), unless otherwise specified	Effect size/p-values
					2	<u>&gt;</u> 14	Symptomatic	20.49 (15.43- 24.44)	17.9 (15.06- 25.06)	p=0.1534
					2	<u>≥</u> 14	Symptomatic and asymptomatic B1.1.7	19.34 (15.39- 21.62)	15.03 (12.51- 16.59)	p=0.0113
					2	<u>≥</u> 14	Symptomatic and asymptomatic not sequenced	29.52 (23.29- 33.59)	25.57 (19.22- 31.44)	p=0.0164
					2	<u>&gt;</u> 14	Symptomatic and asymptomatic non- B.1.1.7 only	22.93 (17.54- 29.4)	18.26 (15.15- 25.57)	p=0.0201
Moderna (mRNA- 1273)	Abu- Raddad <sup>26</sup>	Qatar	Wild type and B.1.1.7, B.1.351, B.1.617.2	Unknown	2	<u>≥</u> 14	Breakthrough infection vs primary infection in unvaccinated	33.3 (29.6-34.8)	30.5 (23.5-33.7)	p<0.001
								Mean=31.2 (95% CI: 30.4- 32.1)	Mean=28 (95% Cl: 27-29.1)	Mean difference (95% Cl): 3.2 (1.8- 4.5), p<0.001
					2	<u>&gt;</u> 14	Breakthrough infection vs reinfection in unvaccinated	33.1 (26.5-34.8)	33.1 (31.1-34.6)	p=0.104
								Mean=30 (95% Cl: 28.3- 31.7)	Mean=31.7 (95% Cl: 30.5- 32.9)	Mean difference: (95% CI): 1.7 (-0.4- 3.8), p=0.104
NR	Riemersma <sup>36</sup>	USA	Wild type and B.1.617.2	NR	2	<u>&gt;</u> 14		NR	NR	p=0.84
					2	<u>&gt;</u> 14		NR	NR	p=0.99
					2	<u>&gt;</u> 14		NR	NR	p=0.85
					2	<u>&gt;</u> 14		NR	NR	p=0.61

Newly identified observational studies shaded in blue.



# Cycle Threshold (Ct) Values

Twenty studies reported on Ct values, which is an inverse proxy for viral load. Eleven of these are new to the updated version of this report (Table 9).

Results from Phase 2/3 vaccine efficacy studies of AZ vaccine compared with a comparator meningococcal vaccine in the United Kingdom, showed that the Ct values in infected vaccinated participants were statistically significantly higher than the comparator (p<0.0001), after 14 days of the second dose in baseline seronegative efficacy cohorts.<sup>16</sup> Furthermore, the vaccine recipients were PCR-positive for a significantly shorter period of time (p<0.0001). The Ct values in asymptomatic cases were also significantly higher among vaccine recipients than control (p=0.0040); however, this difference was not significant for primary symptomatic cases (p=0.1534). Vaccine recipients infected with the B.1.1.7 variant also showed significantly higher Ct values than control (p=0.0113).<sup>16</sup>

A longitudinal UK household survey by Pritchard el al. found statistically significant increase in the median Ct values of PfBnT or AZ single or full dose vaccinated individuals compared with unvaccinated individuals at any time point before or after 21 days post-vaccination (p<0.001).<sup>17</sup> Similarly, in another UK study by Shrotri et al., the mean Ct value of unvaccinated individuals within 27 days of vaccination was 26.6 (95% CI: 26-27.1) compared with 26.6 (95% CI: 25.19-26.62) with one dose of PfBnT or AZ, which was not significantly different (p=0.158).<sup>18</sup> However, after 28 days, there was a statistically significant decrease in the mean Ct between vaccinated and unvaccinated persons (mean Ct 26.6 (95% CI: 26-27.1) vs 31.3 (95% CI: 29.6-32.9), p<0.001).<sup>18</sup> Monthly routine PCR testing was conducted in these patients; however, the baseline serology was not reported.<sup>18</sup> In a longitudinal cohort study of HCWs who were offered voluntary nasal and oropharyngeal swab PCR testing every two weeks as well as serological testing, a small study of 49 people vaccinated with either PfBNT or AZ and 96 unvaccinated people in the USA by Mostafa et al. demonstrated non-significant differences in median Ct values (19.26 [Q1, Q3: 16.56-21.96] vs 19.6 [Q1, Q3: 16.28-22.66], respectively).<sup>19</sup> Similar non-significance in the median Ct values of PfBNT or AerZ-vaccinated people vs unvaccinated was found in a UK study by Baltas et al. (Median=30.8 [IQR: 25.9-35.4] vs. Median=28.8 [IQR: 25.3-33.7], p=0.053).<sup>20</sup> Lumley et al., found vaccination with either PfBnT or AZ to non-significantly increase Ct value by a mean of 2.7.21

A retrospective study of PfBnT mRNA vaccine recipients compared with demographically matched control group of unvaccinated individuals in Israel, found no significant differences in the Ct values for any of the three genes (RdRp, N and E) measured less than 12 days after the first dose in infected persons. However, between 12 and 28 days after the first dose, the Ct values for the three genes were significantly higher among infected vaccinated persons than controls (p<10<sup>-8</sup>).<sup>22</sup> In another UK study of one dose of BNT162b2 vaccine, the median Ct values of infected HCWs were reported to have shown a non-significant trend towards increase between unvaccinated (median=20.3) and vaccinated HCWs after 12 days post-vaccination (median=30.3), suggesting that samples from infected vaccinated individuals had lower viral loads.<sup>23</sup> A study by McEllistrem et al. among community living centre residents reported five cases of asymptomatic infections (determined by surveillance nasal swabs every 2-5 days) among baseline PCR negative PfBnT vaccinated and unvaccinated residents. The

Transmissibility of COVID-19 among vaccinated individuals



median Ct values among unvaccinated residents (12.8, IQR: 12.4-14.9) were significantly lower (p=0.009) than vaccinated residents (19.4, IQR: 18.9-25.5).<sup>24</sup> Furthermore, viral load was -2.4 mean log10 lower among the vaccinated cohort (p=0.004).<sup>24</sup> In another large cohort study of HCWs at a large medical centre in Israel by Regev-Yochay et al, the mean Ct values among PfBNT fully vaccinated HCWs (27.3±2.2) was significantly higher (mean difference 5.09, 95% CI: 2.8-7.4, p<0.001) than unvaccinated HCWs (22.2±1.0).<sup>25</sup> A matched casecontrol study by Abu-Raddad et al. from Qatar, evaluating the Ct values of people with two doses of PfBNT with breakthrough infections compared to Ct values of infections in unvaccinated individuals, found statistically significant higher median Ct values in vaccinated individuals (27.8; IQR: 21.1-32.7) than the median Ct value of unvaccinated individuals (25.8 (IQR: 19.5-31.4; p<0.001).<sup>26</sup> However, studies in France<sup>27</sup> and Greece<sup>28</sup> found no statistically significant differences between PfBNT vaccinated individuals' Ct values and the Ct values of those who were unvaccinated. Bailly et al. found that the Ct values of PfBNT fully vaccinated long-term care residents did not differ from the Ct values of unvaccinated residents (Median=21 [IQR:13-32] vs 15 [IQR: 12-17]; p=0.05).<sup>27</sup> Similarly, Ioannou et al.'s study of fully vaccinated healthcare workers in a Greek hospital amidst an outbreak found no significant differences between the median Ct values of those vaccinated and unvaccinated (18 [15.5-25.5] vs 18.5 [13.5-24]).<sup>28</sup>

A USA study investigating the Ct values of mRNA-based vaccinated healthcare workers (PfBNT or Moderna) compared to unvaccinated healthcare workers found that there was no statistically significant difference in mean Ct values in the early post-vaccination period defined as less than 14 days post vaccination  $(22.6\pm7 \text{ vs. } 23\pm7.4)$  for partially-vaccinated healthcare workers more than 14 days past first dose but before the second dose  $(27.7\pm8.7 \text{ vs. } 23\pm7.4)$  or for fully vaccinated healthcare workers at least 14 days past vaccination  $(28.5\pm7.4 \text{ vs. } 23\pm7.4)$ .<sup>29</sup> Another similar study by Duerr et al. reported Ct values for vaccinated individuals in the community but lumped all unvaccinated comparators as under a Ct value equal or less than 30 therefore, no effect size was presented<sup>30</sup> (Table 9).

A longitudinal UK study from December 2020 to August 2021 found significant differences in Ct values in individuals vaccinated with either one or two doses of PfBNT, AZ, or Moderna<sup>59</sup>. From 1<sup>st</sup> December 2020 to 16<sup>th</sup> May 2021, Pouwels et al. assessed Ct values of people vaccinated with either PfBNT, Moderna, or AZ against the B.1.1.7 (Alpha) variant compared to their unvaccinated counterparts. Individuals with at least one dose of a vaccine had significantly higher Ct values (median=31.71) compared to the seronegative unvaccinated (median=28.7; p=0.02) or seropositive unvaccinated individuals (median=32.8, p=0.72) (Table 9).<sup>59</sup> From the 17<sup>th</sup> of May 2021 to 1<sup>st</sup> August 2021, Ct values of vaccinated individuals were assessed for the B.1.617.2 (Delta) variant compared to unvaccinated individuals and on average found that there appeared to be a higher cycle threshold among vaccinated people. However, there were no reported effect sizes or p-values for these periods.<sup>59</sup>

The study by Abu-Raddad et al. from Qatar also evaluated the median Ct values of people with breakthrough infections those with two doses of Moderna and found that there was statistically significantly higher median Ct values in the vaccinated (33.3; IQR: 29.6-34.8) than the median Ct value of unvaccinated individuals (30.5 (IQR: 23.5-33.7; p<0.001).<sup>26</sup>



In a pre-print by Riemersma et al., the authors included the Ct values of RT-PCR SARS-CoV-2 positive people from a single Wisconsin commercial laboratory with self-reported vaccination status between June 28 through July 24, 2021.<sup>36</sup> There were 291 specimens positive for SARS-CoV-2, with 79 people indicating that they were fully vaccinated and 212 unvaccinated individuals. The authors reported no significant differences in Ct values of vaccinated and unvaccinated people (p=0.85); the mean or median values or the previous serology of the sample were not reported. Of the 42 people identified with a Delta infection it was reported that there was no difference in Ct values (p=0.61).<sup>36</sup>

# **Viral Load**

Two USA studies reported on viral load; one study is new to this updated version of the report. A prospective cohort study of baseline seronegative vaccinated and unvaccinated healthcare workers across Arizona had their mid-turbinate nasal swabs assessed for viral load.<sup>31</sup> Thompson et al. found that the mean viral RNA load for partially and fully vaccinated healthcare workers, with a mRNA-based vaccine, who were at least 14 days past the date of vaccination had lower presence of virus compared to their unvaccinated counterparts  $(2.3\pm1.7 \text{ Log}_{10} \text{ copies/mL vs. } 3.8\pm1.7 \text{ Log}_{10} \text{ copies/mL}).^{31}$  This represented at least 40.2% lower viral RNA load after at least partial vaccination.<sup>31</sup>

A second retrospective cohort study of five vaccinated and five unvaccinated asymptomatic nursing home residents in a single nursing home evaluated the effectiveness of at least one dose of the PfBNT vaccine on attenuating viral load.<sup>24</sup> Viral load was -2.4 mean log10 lower among the vaccinated cohort (p=0.004).<sup>24</sup>

# **Discussion**

In this update, 25 additional studies were included. Therefore, this review has a total of 45 included studies. Four new studies from the Netherlands, Finland, and Israel evaluating household transmission following vaccination found that PfBNT, Moderna, AZ, and J&J vaccines significantly reduce the risk of household transmission.<sup>2-5</sup> The majority of the vaccines included in this review demonstrated efficacy and effectiveness against asymptomatic wild-type COVID-19 infections. There is some limited evidence that there is moderate efficacy and effectiveness of vaccines against the B.1.617.2 (Delta) strain of the virus.

The AZ, PfBnT, and Moderna vaccines were found to be significantly associated with higher Ct values than their respective comparators, suggesting that these vaccines may potentially reduce viral load and consequently lower the risk of transmission. It is however noteworthy that the relationship between viral load, viral shedding, infectivity, and the duration of infectivity are not well understood. Ct values are also subject to error.<sup>65</sup> Furthermore, although there were statistically significant differences in median Ct values between vaccinated and unvaccinated individuals, most of the Ct values for both the vaccinated and unvaccinated individuals were  $\leq$ 29, which clinically both indicate that the cycle thresholds were strongly positive indicative of an abundance of target viral nucleic acid in the samples.<sup>66,67</sup> A couple of studies found that vaccination with an mRNA-based vaccine reduced the viral load.<sup>31,55</sup>

Transmissibility of COVID-19 among vaccinated individuals



There were significant limitations to many of the included studies. It was not possible to directly compare findings across studies owing to variations in the assessment of asymptomatic status, the testing used, and timing of these assessments. Also, the possibility of persistent PCR positivity after COVID-19 infection<sup>68</sup> could not be excluded in some of the studies without baseline PCR assessment. Few studies included surveillance nasal swabs for PCR positivity. Most of the current data were around viral detection, rather than evidence of cultivatable virus. Therefore, there was limited data to evaluate the efficacy or effectiveness of COVID-19 vaccines in decreasing viral loads. In addition, there are only a limited number of epidemiologic data addressing evidence of forward transmission after vaccination.

# **Emerging Evidence**

Variants of SARS-CoV-2 continue to surface, and the B.617.2 (Delta) variant is currently the one of most concern.<sup>34,35</sup> There has been emerging evidence that indicates that although a full vaccination series might reduce an individual's overall risk of becoming infected, there seems to be a limited difference in the Ct values between those vaccinated and unvaccinated.<sup>34,36</sup> Furthermore, certain outbreaks amongst vaccinated individuals in the USA have led to expanded prevention strategies, such as universal masking in indoor spaces.<sup>35</sup>

Comparisons of the proxy viral load measurement, Ct value, have found that, regardless of vaccination status, there was no difference in Ct values once an individual was infected with the B.1.617.2 (Delta) variant.<sup>34,36</sup> Public Health England (PHE) released a technical report on the NHS Test and Trace case data of the median and mean Ct values for all cases in the country. They reported that since May 2021 to July 2021, the median Ct value for unvaccinated individuals was 17.8 compared to the median of 18.0 in those who were vaccinated.<sup>34</sup> PHE suggested that there was limited difference in infectiousness due to the similarity in Ct values; the case data was not age stratified.<sup>34</sup>

The CDC reported in the Morbidity and Mortality Weekly Report on an outbreak of SARS-CoV-2 infections in vaccinated individuals in Barnstable County, Massachusetts.<sup>35</sup> During July 2021, 469 cases were associated with multiple summer and large public gathering events, with 74% (n=346) of cases occurring in fully vaccinated persons who had completed a 2-dose course of an mRNA vaccine (PfBNT or Moderna) or received the single-dose J&J vaccine. Genomic sequencing of specimens from 133 patients identified the B.1.617.2 (Delta) variant in 90% of cases. Of the 346 identified breakthrough infections, 274 (79%) had symptoms, and among the five who were hospitalized, four were fully vaccinated.<sup>35</sup> Ct values in specimens from 127 vaccinated persons with breakthrough cases were similar to those from 84 persons who were unvaccinated, not fully vaccinated, or whose vaccination status was unknown (median = 22.77 vs median=21.54, respectively).<sup>35</sup> Persons with COVID-19 reported attending densely packed indoor and outdoor events at venues that included bars, restaurants, guest houses, and rental homes. After the events of Barnstable County, the CDC recommended that all persons, including fully vaccinated individuals wear masks indoors in public settings especially in attendance of large public gatherings.

This emerging evidence suggests that the B.1.617.2 (Delta) variant is highly transmissible and that there may not be a difference between the viral load of those vaccinated compared Transmissibility of COVID-19 among vaccinated individuals 59



to those unvaccinated. However, more studies need to be conducted to fully understand the protection gleaned from vaccination. Until then, expanded prevention strategies, such as universal indoor masking, may help prevent the spread of the Delta variant.

# **Recommendations**

Based on the current evidence, we suggest the following:

1) All vaccinees should self-isolate and seek testing after the development of COVID-19 compatible symptoms.

2) Following exposure, the risk of contracting COVID-19 and subsequent forward transmission from asymptomatic or pauci-symptomatic viral carriage should be considered in light of whether the exposed individual was vaccinated, the time elapsed since immunization, and the consequent expected degree of protection on a case-by-case basis for those in vulnerable setting. When possible, a case-by-case consideration for whether exposed persons are immunized, is necessary. Low-moderate risk exposures could potentially be managed with careful use of personal protective equipment (PPE), and self-monitoring.

3) If a vaccinated HCW is assessed as having a significant exposure before the period of expected robust immunity, high risk exposures may be managed as for unvaccinated persons.

4) All vaccinated persons should continue to use recommended PPE when in close contact with unvaccinated persons.

5) Population and public health data being collected on positive COVID-19 tests occurring after vaccination should be combined with laboratory data on Ct values, identification of variant strain infections, and epidemiologic contact tracing data to prospectively monitor for evidence of forward transmission of infection from vaccinated persons.

# Conclusion

Four months since the publication of the previous version of this report, 24 additional relevant studies have been conducted. Four of these were large household surveillance studies from the Netherlands, Finland, and Israel suggesting that a full dose of PfBNT, Moderna, AZ, or J&J vaccines may prevent household transmission after 14 days of vaccination. Twelve additional studies found that vaccines significantly reduce the risk of asymptomatic infection, with multiple studies finding that vaccines decreased the viral RNA load or increased the cycle threshold, suggestive of reduced viral load. Some studies, such as the AZ vaccine RCTs, included data on cross sectional prevalence of positive SARS-CoV-2 RT-PCR from routine swabbing, which suggested efficacy against asymptomatic infection, although this was not routinely assessed in a comparable way across studies. Evidence regarding the Ct values for the AZ, PfBnT, and Moderna vaccine suggest their potential to reduce viral load and possibly transmission. Further research is needed to evaluate post-vaccination infectivity and transmission of variants of concern especially the B.1.617.2 (Delta) strain from other jurisdictions.

Transmissibility of COVID-19 among vaccinated individuals



# References

- 1. Egunsola O, Mastikhina L, Dowsett L, et al. Transmissibility of COVID-19 among vaccinated individuals: Update #1. 2021.
- 2. de Gier B, Andeweg S, Joosten R, et al. Vaccine effectiveness against SARS-CoV-2 transmission and infections among household and other close contacts of confirmed cases, the Netherlands, February to May 2021. *Euro Surveill.* 2021;26(31).
- 3. Salo J, Hägg M, Kortelainen M, et al. The indirect effect of mRNA-based Covid-19 vaccination on unvaccinated household members. *medRxiv*. 2021:2021.2005.2027.21257896.
- 4. Gazit S, Mizrahi B, Kalkstein N, et al. BNT162b2 mRNA Vaccine Effectiveness Given Confirmed Exposure; Analysis of Household Members of COVID-19 Patients. *medRxiv*. 2021:2021.2006.2029.21259579.
- 5. Layan M, Gilboa M, Gonen T, et al. Impact of BNT162b2 vaccination and isolation on SARS-CoV-2 transmission in Israeli households: an observational study. *medRxiv*. 2021:2021.2007.2012.21260377.
- 6. Voysey M, Clemens SAC, Madhi SA, et al. Single-dose administration and the influence of the timing of the booster dose on immunogenicity and efficacy of ChAdOx1 nCoV-19 (AZD1222) vaccine: a pooled analysis of four randomised trials. *The Lancet.* 2021.
- Voysey M, Clemens SAC, Madhi SA, 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. *The Lancet.* 2021;397(10269):99-111.
- 8. Tang L, Hijano DR, Gaur AH, et al. Asymptomatic and Symptomatic SARS-CoV-2 Infections After BNT162b2 Vaccination in a Routinely Screened Workforce. *JAMA*. 2021;325(24):2500-2502.
- 9. Angel Y, Spitzer A, Henig O, et al. Association Between Vaccination With BNT162b2 and Incidence of Symptomatic and Asymptomatic SARS-CoV-2 Infections Among Health Care Workers. *Jama.* 2021;325(24):2457-2465.
- 10. Andrejko KL, Pry J, Myers JF, et al. Prevention of COVID-19 by mRNA-based vaccines within the general population of California. *medRxiv*. 2021:2021.2004.2008.21255135.
- 11. Chemaitelly H, Yassine HM, Benslimane FM, et al. mRNA-1273 COVID-19 vaccine effectiveness against the B.1.1.7 and B.1.351 variants and severe COVID-19 disease in Qatar. *Nat Med.* 2021;27(9):1614-1621.
- 12. Tang P, Hasan MR, Chemaitelly H, et al. BNT162b2 and mRNA-1273 COVID-19 vaccine effectiveness against the Delta (B.1.617.2) variant in Qatar. *medRxiv*. 2021:2021.2008.2011.21261885.
- 13. Dagan N, Barda N, Kepten E, et al. BNT162b2 mRNA Covid-19 Vaccine in a Nationwide Mass Vaccination Setting. *New England Journal of Medicine.* 2021;384:1412-1423.
- 14. Haas EJ, Angulo FJ, McLaughlin JM, et al. Impact and effectiveness of mRNA BNT162b2 vaccine against SARS-CoV-2 infections and COVID-19 cases, hospitalisations, and deaths following a nationwide vaccination campaign in Israel: an observational study using national surveillance data. *The Lancet.* 2021.
- 15. Tande AJ, Pollock BD, Shah ND, et al. Impact of the COVID-19 Vaccine on Asymptomatic Infection Among Patients Undergoing Pre-Procedural COVID-19 Molecular Screening. *Clinical Infectious Diseases.* 2021.



- 16. Emary KR, Golubchik T, Aley PK, et al. Efficacy of ChAdOx1 nCoV-19 (AZD1222) vaccine against SARS-CoV-2 variant of concern 202012/01 (B.1.1.7): an exploratory analysis of a randomised controlled trial. *Lancet.* 2021;397:1351–1362.
- 17. Pritchard E, Matthews PC, Stoesser N, et al. Impact of vaccination on SARS-CoV-2 cases in the community: a population-based study using the UK's COVID-19 Infection Survey. *medRxiv*. 2021.
- Shrotri M, Krutikov M, Palmer T, et al. Vaccine Effectiveness of the First Dose of ChAdox1 nCOV-19 and BNT162b2 Against SARS-CoV-2 Infection in Residents of Long Term Care Facilities (VIVALDI Study). 2021.
- 19. Mostafa HH, Luo CH, Morris CP, et al. SARS-CoV-2 Infections in mRNA Vaccinated Individuals are Biased for Viruses Encoding Spike E484K and Associated with Reduced Infectious Virus Loads that Correlate with Respiratory Antiviral IgG levels. *medRxiv.* 2021.
- 20. Baltas I, Boshier FAT, Williams CA, et al. Post-vaccination COVID-19: A case-control study and genomic analysis of 119 breakthrough infections in partially vaccinated individuals. *Clin Infect Dis.* 2021.
- 21. Lumley SF, Rodger G, Constantinides B, et al. An observational cohort study on the incidence of SARS-CoV-2 infection and B. 1.1. 7 variant infection in healthcare workers by antibody and vaccination status. *medRxiv.* 2021.
- 22. Levine-Tiefenbrun M, Yelin I, Katz R, et al. Initial report of decreased SARS-CoV-2 viral load after inoculation with the BNT162b2 vaccine. *Nature Medicine*. 2021;27:790-792.
- 23. Jones NK, Rivett L, Seaman S, et al. Single-dose BNT162b2 vaccine protects against asymptomatic SARS-CoV-2 infection. *eLife.* 2021;10:e68808.
- 24. McEllistrem MC, Clancy CJ, Buehrle DJ, Lucas A, Decker BK. Single dose of a mRNA SARS-CoV-2 vaccine is associated with lower nasopharyngeal viral load among nursing home residents with asymptomatic COVID-19. *Clinical Infectious Diseases*. 2021.
- 25. Regev-Yochay G, Amit S, Bergwerk M, et al. Decreased infectivity following BNT162b2 vaccination. 2021.
- 26. Abu-Raddad LJ, Chemaitelly H, Ayoub HH, et al. Effect of vaccination and of prior infection on infectiousness of vaccine breakthrough infections and reinfections. *medRxiv.* 2021:2021.2007.2028.21261086.
- 27. Bailly B, Guilpain L, Bouiller K, et al. BNT162b2 mRNA vaccination did not prevent an outbreak of SARS COV-2 variant 501Y.V2 in an elderly nursing home but reduced transmission and disease severity. *Clin Infect Dis.* 2021.
- 28. Ioannou P, Karakonstantis S, Astrinaki E, et al. Transmission of SARS-CoV-2 variant B.1.1.7 among vaccinated health care workers. *Infect Dis (Lond).* 2021;53(11):876-879.
- 29. Jacobson KB, Pinsky BA, Rath MEM, et al. Post-vaccination SARS-CoV-2 infections and incidence of the B.1.427/B.1.429 variant among healthcare personnel at a northern California academic medical center. *medRxiv.* 2021.
- 30. Duerr R, Dimartino D, Marier C, et al. Dominance of Alpha and lota variants in SARS-CoV-2 vaccine breakthrough infections in New York City. *J Clin Invest.* 2021;131(18).
- 31. Thompson MG, Burgess JL, Naleway AL, et al. Prevention and Attenuation of Covid-19 with the BNT162b2 and mRNA-1273 Vaccines. *New England Journal of Medicine*. 2021;385(4):320-329.
- 32. Baden LR, El Sahly HM, Essink B, et al. Efficacy and safety of the mRNA-1273 SARS-CoV-2 vaccine. *New England Journal of Medicine.* 2021;384(5):403-416.
- 33. Janssen Biotech I. Vaccines and Related Biological Products Advisory Committee Meeting February 26, 2021: FDA Briefing Document. 26 February 2021 2021.



- 34. England PH. SARS-CoV-2 variants of concern and variants under investigation in England. Public Health England; August 6 2021 2021.
- 35. Brown CM, Vostok J, Johnson H, et al. Outbreak of SARS-CoV-2 Infections, Including COVID-19 Vaccine Breakthrough Infections, Associated with Large Public Gatherings - Barnstable County, Massachusetts, July 2021. *MMWR Morb Mortal Wkly Rep.* 2021;70(31):1059-1062.
- 36. Riemersma KK, Grogan BE, Kita-Yarbro A, et al. Vaccinated and unvaccinated individuals have similar viral loads in communities with a high prevalence of the SARS-CoV-2 delta variant. *medRxiv.* 2021:2021.2007.2031.21261387.
- 37. Johns Hopkins University. COVID-19 Dashboard by the Center for Systems Science and Engineering (CSSE) at Johns Hopkins University (JHU). <u>https://www.covidtracker.com/</u>. Published 2021. Accessed 27 February 2021.
- 38. Shapiro J, Dean NE, Madewell ZJ, Yang Y, Halloran ME, Longini IM. Efficacy Estimates for Various COVID-19 Vaccines: What we Know from the Literature and Reports. *medRxiv*. 2021.
- 39. Lopez Bernal J, Andrews N, Gower C, et al. Effectiveness of Covid-19 Vaccines against the B.1.617.2 (Delta) Variant. *N Engl J Med.* 2021;385(7):585-594.
- 40. Weinreich DM, Sivapalasingam S, Norton T, et al. REGN-COV2, a neutralizing antibody cocktail, in outpatients with Covid-19. *N Engl J Med.* 2021:238-251.
- 41. Chen P, Nirula A, Heller B, et al. SARS-CoV-2 neutralizing antibody LY-CoV555 in outpatients with Covid-19. *New England Journal of Medicine*. 2021;384(3):229-237.
- 42. Lyngse FP, Mølbak K, Franck KT, Nielsen C, Skov RL, Kirkeby CT. Association between SARS-CoV-2 Transmission Risk, Viral Load, and Age: A Nationwide Study in Danish Households. *medRxiv.* 2021.
- 43. Marks M, Millat-Martinez P, Ouchi D, et al. Transmission of COVID-19 in 282 clusters in Catalonia, Spain: a cohort study. *The Lancet Infectious Diseases*. 2021.
- 44. Bjorkman KK, Saldi TK, Lasda E, et al. Higher viral load drives infrequent SARS-CoV-2 transmission between asymptomatic residence hall roommates. *medRxiv*. 2021.
- 45. Vibholm LK, Nielsen SS, Pahus MH, et al. SARS-CoV-2 persistence is associated with antigen-specific CD8 T-cell responses. *EBioMedicine*. 2021;64:103230.
- 46. Madewell ZJ, Yang Y, Longini IM, Halloran ME, Dean NE. Household Transmission of SARS-CoV-2: A Systematic Review and Meta-analysis. *JAMA network open.* 2020;3(12):e2031756-e2031756.
- 47. McGowan J, Sampson M, Salzwedel DM, Cogo E, Foerster V, Lefebvre C. PRESS peer review of electronic search strategies: 2015 guideline statement. *Journal of clinical epidemiology*. 2016;75:40-46.
- 48. Sterne JA, Hernán MA, Reeves BC, et al. ROBINS-I: a tool for assessing risk of bias in non-randomised studies of interventions. *BMJ.* 2016;355:i4919.
- 49. Higgins JP, Altman DG, Gøtzsche PC, et al. The Cochrane Collaboration's tool for assessing risk of bias in randomised trials. *Bmj.* 2011;343.
- 50. Real-World Evidence Confirms High Effectiveness of Pfizer-BioNTech COVID-19 Vaccine and Profound Public Health Impact of Vaccination One Year After Pandemic Declared [press release]. 2021.
- 51. Ali K, Berman G, Zhou H, et al. Evaluation of mRNA-1273 SARS-CoV-2 Vaccine in Adolescents. *New England Journal of Medicine.* 2021.
- 52. Amit S, Regev-Yochay G, Afek A, Kreiss Y, Leshem E. Early rate reductions of SARS-CoV-2 infection and COVID-19 in BNT162b2 vaccine recipients. *The Lancet*. 2021.
- 53. Shah AS, Gribben C, Bishop J, et al. Effect of vaccination on transmission of COVID-19: an observational study in healthcare workers and their households. *MedRxiv*. 2021.



- 54. Souza W, Muraro S, Souza G, et al. Clusters of SARS-CoV-2 Lineage B.1.1.7 Infection After Vaccination With Adenovirus-Vectored and Inactivated Vaccines: A Cohort Study. SSRN Electronic Journal. 2021.
- 55. McEllistrem MC, Clancy CJ, Buehrle DJ, et al. Introduction of the BNT162b2 vaccine during a COVID-19 nursing home outbreak. *Am J Infect Control.* 2021.
- 56. Hall VJ, Foulkes S, Saei A, et al. COVID-19 vaccine coverage in health-care workers in England and effectiveness of BNT162b2 mRNA vaccine against infection (SIREN): a prospective, multicentre, cohort study. *Lancet.* 2021;397:1725-1735.
- 57. Harris R, Hall J, Zaidi A, Andrews NJ, Dunbar JK, Dabrera G. Impact of vaccination on household transmission of SARS-COV-2 in England. *Preprint Disponible en:* <u>https://khub</u> net/documents/135939561/390853656/Impact+ of+ vaccination+ on+ household+ transmission+ of+ SARS-COV-2+ in+ England pdf/35bf4bb1-6ade-d3eb-a39e-9c9b25a8122a.
- 58. Muhsen K, et al. Effectiveness of BNT162b2 mRNA COVID-19 Vaccine Against Acquisitions of SARS-CoV-2 Among Health Care Workers in Long-Term Care Facilities: A Prospective Cohort Study. . 2021.
- 59. Pouwels KB, Pritchard E, Matthews PC, et al. Impact of Delta on viral burden and vaccine effectiveness against new SARS-CoV-2 infections in the UK. *medRxiv*. 2021:2021.2008.2018.21262237.
- 60. Bouton TC, Lodi S, Turcinovic J, et al. COVID-19 vaccine impact on rates of SARS-CoV-2 cases and post vaccination strain sequences among healthcare workers at an urban academic medical center: a prospective cohort study. *medRxiv*. 2021.
- 61. Antonelli M, Penfold RS, Merino J, et al. Risk factors and disease profile of postvaccination SARS-CoV-2 infection in UK users of the COVID Symptom Study app: a prospective, community-based, nested, case-control study. *The Lancet Infectious Diseases.*
- 62. Sansone E, Sala E, Tiraboschi M, et al. Effectiveness of BNT162b2 vaccine against SARS-CoV-2 among healthcare workers. *Med Lav.* 2021;112(3):250-255.
- 63. Dagan N, Barda N, Kepten E, et al. BNT162b2 mRNA Covid-19 Vaccine in a Nationwide Mass Vaccination Setting. *New England Journal of Medicine*. 2021.
- 64. Zhang J, Kai FY. What's the relative risk?: A method of correcting the odds ratio in cohort studies of common outcomes. *Jama.* 1998;280(19):1690-1691.
- 65. Pollock AM, Lancaster J. Asymptomatic transmission of covid-19. *BMJ.* 2020;371:m4851.
- 66. Laboratory WVD. *Real Time PCR Ct Values*. University of Wisconsin-Madison;2018.
- 67. Ontario) OAfHPaPPH. Focus on: an overview of cycle threshold values and their role in SARS-Cov-2 real-time PCR test interpretation.: Public Health Ontario;2020.
- 68. Kim S-m, Hwang YJ, Kwak Y. Prolonged SARS-CoV-2 detection and reversed RT-PCR results in mild or asymptomatic patients. *Infectious Diseases*. 2021;53(1):31-37.

# Appendix 1: Search Strategy

# Ovid Multifile

Database: EBM Reviews - Cochrane Central Register of Controlled Trials <July 2021>, Embase <1974 to 2021 August 20> , Ovid MEDLINE(R) and Epub Ahead of Print, In-



Process, In-Data-Review & Other Non-Indexed Citations and Daily <1946 to August 20, 2021>

Search Strategy:

\_\_\_\_\_

1 exp COVID-19 Vaccines/ (4322)

2 ((COVID-19 or COVID19) adj5 (immun\* or inoculat\* or vaccin\*)).tw,kf. (19693)

3 ((coronavirus\* or corona virus\*) adj5 (immun\* or inoculat\* or vaccin\*)).tw,kf. (4818)

4 ((2019-nCoV or nCoV or nCoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or

SARSCoV2 or SARS2) adj5 (immun\* or inoculat\* or vaccin\*)).tw,kf. (11949)

5 (((BNT162 or BNT162-01 or BNT162a1 or BNT162b1 or BNT162b2 or BNT162c2) and vaccin\*) or N38TVC63NU).tw,kf. (1144)

6 (((AZD1222 or ChAdOx1) and vaccin\*) or Covishield\$2 or B5S3K2V0G8).tw,kf. (766)

7 ((Moderna and vaccin\*) or EPK39PL4R4).tw,kf. (454)

8 ((mRNA adj3 vaccin\*) and (COVID-19 or COVID19 or coronavirus\* or corona virus\* or 2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).tw.kf. (1795)

9 ((messenger RNA adj3 vaccin\*) and (COVID-19 or COVID19 or coronavirus\* or corona virus\* or 2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV2 or SARSCoV2 or SARSC).tw,kf. (158)

10 (LV-SMENP-DC and vaccin\*).tw,kf. (5)

11 ((Ad5-nCoV and vaccin\*) or hAdOx1 nCoV-19).tw,kf. (37)

12 (("Ad26.COV2.S" or Ad26COVS1 or JNJ 78436735 or JNJ-78436735 or JT2NS6183B) and vaccin\*).tw,kf. (158)

13 Viral Vaccines/ and (Coronavirus/ or Betacoronavirus/ or Coronavirus Infections/) (1906)

14 or/1-13 [COVID-19 VACCINES] (33065)

15 COVID-19/pc [prevention & control] (8859)

16 Coronavirus Infections/pc [prevention & control] (10743)

17 Pandemics/pc [prevention & control] (14512)

18 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (COVID-19 or COVID19)).tw,kf. (27144)

19 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (coronavirus\* or corona virus\*)).tw,kf. (3792)

20 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).tw,kf. (6889)

21 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 spread\*).tw,kf. (52770)

22 COVID-19/ep [Epidemiology] (17717)

23 COVID-19/tm [Transmission] (3563)

- 24 COVID-19/vi [Virology] (6344)
- 25 Coronavirus Infections/ep [Epidemiology] (23509)
- 26 Coronavirus Infections/tm [Transmission] (4703)
- 27 Coronavirus Infections/vi [Virology] (7690)
- 28 exp Disease Transmission, Infectious/ (352747)

29 (transmit\* or transmissi\* or infectiousness\* or infectivit\*).tw,kf. (1235632)

30 ((COVID-19 or COVID19) adj5 (caus\* or pass or passed or passes or passing or spread\*)).tw,kf. (32763)

31 ((coronavirus\* or corona virus\*) adj5 (caus\* or pass or passed or passes or passing or spread\*)).tw,kf. (17810)

32 ((virus\* or infection\*) adj5 (caus\* or pass or passed or passes or passing or spread\*)).tw,kf. (386113)



33 ((2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2) adj5 spread\*).tw,kf. (5291)

- 34 (unvaccinat\* or nonvaccinat\* or non-vaccinat\* or "not vaccinat\*").tw,kf. (42643)
- 35 or/15-34 [TRANSMISSION] (1923307)
- 36 14 and 35 [COVID-19 VACCINES DISEASE TRANSMISSION] (16076)
- 37 (controlled clinical trial or randomized controlled trial or pragmatic clinical trial or equivalence trial).pt. (1248935)
- 38 "Clinical Trials as Topic"/ (314700)
- 39 exp "Controlled Clinical Trials as Topic"/ (382216)
- 40 (randomi#ed or randomi#ation? or randomly or RCT or placebo\*).tw,kf. (3721675)
- 41 ((singl\* or doubl\* or trebl\* or tripl\*) adj (mask\* or blind\* or dumm\*)).tw,kf. (731366)
- 42 trial.ti. (939819)
- 43 or/37-42 [RCT FILTER] (4670500)
- 44 36 and 43 [RCTs] (1647)
- 45 controlled clinical trial.pt. (186646)
- 46 Controlled Clinical Trial/ or Controlled Clinical Trials as Topic/ (573850)
- 47 (control\* adj2 trial).tw,kf. (691363)
- 48 Non-Randomized Controlled Trials as Topic/ (12839)
- 49 (nonrandom\* or non-random\* or quasi-random\* or quasi-experiment\*).tw,kf. (157674)
- 50 (nRCT or non-RCT).tw,kf. (1028)
- 51 Controlled Before-After Studies/ (210357)
- 52 (control\* adj3 ("before and after" or "before after")).tw,kf. (825048)
- 53 Interrupted Time Series Analysis/ (203342)
- 54 time series.tw,kf. (77728)
- 55 (pre- adj3 post-).tw,kf. (294878)
- 56 (pretest adj3 posttest).tw,kf. (16682)
- 57 Historically Controlled Study/ (220764)
- 58 (control\* adj2 study).tw,kf. (556350)
- 59 Control Groups/ (111904)
- 60 (control\* adj2 group?).tw,kf. (1573730)
- 61 trial.ti. (939819)
- 62 or/45-61 [nRCT FILTER] (4350878)
- 63 36 and 62 [nRCTs] (1075)
- 64 exp Cohort Studies/ (3093661)
- 65 cohort?.tw,kf. (1911901)
- 66 Retrospective Studies/ (1784445)
- 67 (longitudinal or prospective or retrospective).tw,kf. (3831913)
- 68 ((followup or follow-up) adj (study or studies)).tw,kf. (130616)
- 69 Observational study.pt. (107979)
- 70 (observation\$2 adj (study or studies)).tw,kf. (341258)
- 71 ((population or population-based) adj (study or studies or analys#s)).tw,kf. (47080)
- 72 ((multidimensional or multi-dimensional) adj (study or studies)).tw,kf. (276)
- 73 Comparative Study.pt. (2066781)
- 74 ((comparative or comparison) adj (study or studies)).tw,kf. (285012)
- 75 exp Case-Control Studies/ (1423752)
- 76 ((case-control\* or case-based or case-comparison) adj (study or studies)).tw,kf.

(272319)

- 77 Cross-Sectional Studies/ (690149)
- 78 (crosssection\* or cross-section\*).tw,kf. (1054262)
- 79 or/64-78 [OBSERVATIONAL STUDY FILTER] (9827573)
- 80 36 and 79 [OBSERVATIONAL STUDIES] (2568)
- 81 44 or 63 or 80 [ALL STUDY DESIGNS] (4117)



82 exp Animals/ not Humans/ (16123202)

83 81 not 82 [ANIMAL-ONLY REMOVED] (3982)

84 (202105\* or 202106\* or 202107\* or 202108\*).dt. (478611)

85 83 and 84 [UPDATE PERIOD] (576)

86 85 use ppez [MEDLINE RECORDS] (576)

87 SARS-CoV-2 vaccine/ (8936)

88 ((COVID-19 or COVID19) adj5 (immun\* or inoculat\* or vaccin\*)).tw,kw. (22548)

89 ((coronavirus\* or corona virus\*) adj5 (immun\* or inoculat\* or vaccin\*)).tw,kw. (5596)

90 ((2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or

SARSCoV2 or SARS2) adj5 (immun\* or inoculat\* or vaccin\*)).tw,kw. (13897)

91 (((BNT162 or BNT162-01 or BNT162a1 or BNT162b1 or BNT162b2 or BNT162c2) and vaccin\*) or N38TVC63NU).tw,kw. (1157)

92 (((AZD1222 or ChAdOx1) and vaccin\*) or Covishield\$2 or B5S3K2V0G8).tw,kw. (772)
 93 ((Moderna and vaccin\*) or EPK39PL4R4).tw,kw. (459)

94 ((mRNA adj3 vaccin\*) and (COVID-19 or COVID19 or coronavirus\* or corona virus\* or 2019-nCoV or nCoV or nCoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).tw,kw. (1814)

95 ((messenger RNA adj3 vaccin\*) and (COVID-19 or COVID19 or coronavirus\* or corona virus\* or 2019-nCoV or nCoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).tw,kw. (161)

96 (LV-SMENP-DC and vaccin\*).tw,kw. (5)

97 ((Ad5-nCoV and vaccin\*) or hAdOx1 nCoV-19).tw,kw. (37)

98 (("Ad26.COV2.S" or Ad26COVS1 or JNJ 78436735 or JNJ-78436735 or JT2NS6183B) and vaccin\*).tw,kw. (158)

99 (severe acute respiratory syndrome vaccine/ or virus vaccine/) and (coronavirinae/ or betacoronavirus/ or exp SARS-related coronavirus/ or coronavirus infection/) (1111)

100 or/87-99 [COVID-19 VACCINES] (36579)

101 coronavirus disease 2019/pc [prevention] (16383)

102 coronavirus infection/pc [prevention] (10779)

103 pandemic/pc [prevention] (14596)

104 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (COVID-19 or COVID19)).tw,kw. (27702)

105 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (coronavirus\* or corona virus\*)).tw,kw. (5416)

106 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).tw,kw. (7180)

107 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 spread\*).tw,kw. (52807)

108 coronavirus disease 2019/ep [epidemiology] (29877)

109 coronavirus infection/ep [epidemiology] (23549)

110 virus transmission/ (72848)

111 (transmit\* or transmissi\* or infectiousness\* or infectivit\*).tw,kw. (1241266)

112 ((COVID-19 or COVID19) adj5 (caus\* or pass or passed or passes or passing or spread\*)).tw,kw. (32798)

113 ((coronavirus\* or corona virus\*) adj5 (caus\* or pass or passed or passes or passing or spread\*)).tw,kw. (17833)

114 ((virus\* or infection\*) adj5 (caus\* or pass or passed or passes or passing or spread\*)).tw,kw. (386369)

115 ((2019-nCoV or nCoV or nCoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2) adj5 spread\*).tw,kw. (5298)

116 (unvaccinat\* or nonvaccinat\* or non-vaccinat\* or "not vaccinat\*").tw,kw. (42646)



- 117 or/101-116 [TRANSMISSION] (1771114)
- 118 100 and 117 [COVID-19 VACCINES DISEASE TRANSMISSION] (18078)
- 119 exp randomized controlled trial/ or controlled clinical trial/ (1492681)
- 120 clinical trial/ (1541965)
- 121 exp "controlled clinical trial (topic)"/ (217360)
- 122 (randomi#ed or randomi#ation? or randomly or RCT or placebo\*).tw,kw. (3785088)
- 123 ((singl\* or doubl\* or trebl\* or tripl\*) adj (mask\* or blind\* or dumm\*)).tw,kw. (760770)
- 124 trial.ti. (939819)
- 125 or/119-124 [RCT FILTER] (5250210)
- 126 118 and 125 [RCTs] (1545)
- 127 controlled clinical trial/ (558033)
- 128 "controlled clinical trial (topic)"/ (11831)
- 129 (control\* adj2 trial).tw,kw. (1058875)
- 130 (nonrandom\* or non-random\* or quasi-random\* or quasi-experiment\*).tw,kw.

#### (158708)

- 131 (nRCT or non-RCT).tw,kw. (1030)
- 132 (control\* adj3 ("before and after" or "before after")).tw,kw. (825052)
- 133 time series analysis/ (29877)
- 134 time series.tw,kw. (78764)
- 135 pretest posttest control group design/ (564)
- 136 (pre- adj3 post-).tw,kw. (294927)
- 137 (pretest adj3 posttest).tw,kw. (20506)
- 138 controlled study/ (8321938)
- 139 (control\* adj2 study).tw,kw. (985693)
- 140 control group/ (111799)
- 141 (control\* adj2 group?).tw,kw. (1574940)
- 142 trial.ti. (939819)
- 143 or/127-142 [nRCT FILTER] (11387363)
- 144 118 and 143 [nRCTs] (2998)
- 145 cohort analysis/ (1033210)
- 146 cohort?.tw,kw. (1917721)
- 147 retrospective study/ (2052657)
- 148 longitudinal study/ (308271)
- 149 prospective study/ (1296159)
- 150 (longitudinal or prospective or retrospective).tw,kw. (3853020)
- 151 follow up/ (1723966)
- 152 ((followup or follow-up) adj (study or studies)).tw,kw. (132434)
- 153 observational study/ (349424)
- 154 (observation\$2 adj (study or studies)).tw,kw. (343991)
- 155 population research/ (116549)
- 156 ((population or population-based) adj (study or studies or analys#s)).tw,kw. (55167)
- 157 ((multidimensional or multi-dimensional) adj (study or studies)).tw,kw. (277)
- 158 exp comparative study/ (3392659)
- 159 ((comparative or comparison) adj (study or studies)).tw,kw. (302872)
- 160 exp case control study/ (1423752)
- 161 ((case-control\* or case-based or case-comparison) adj (study or studies)).tw,kw.

(275780)

- 162 cross-sectional study/ (817603)
- 163 (crosssection\* or cross-section\*).tw,kw. (1058104)
- 164 major clinical study/ (4204970)
- 165 or/145-164 [OBSERVATIONAL STUDY FILTER] (13849196)
- 166 118 and 165 [OBSERVATIONAL STUDIES] (3519)



## 167 126 or 144 or 166 [ALL STUDY DESIGNS] (5730)

168 exp animal/ or exp animal experimentation/ or exp animal model/ or exp animal experiment/ or nonhuman/ or exp vertebrate/ (54468436)

- 169 exp human/ or exp human experimentation/ or exp human experiment/ (42855732)
- 170 168 not 169 (11614435)

171 167 not 170 [ANIMAL-ONLY REMOVED] (5365)

172 (202105\* or 202106\* or 202107\* or 202108\*).dc. (933210)

- 173 171 and 172 [UPDATE PERIOD] (1560)
- 174 173 use oemezd [EMBASE RECORDS] (1560)
- 175 exp COVID-19 Vaccines/ (4322)

176 ((COVID-19 or COVID19) adj5 (immun\* or inoculat\* or vaccin\*)).ti,ab,kw. (22546)

((coronavirus\* or corona virus\*) adj5 (immun\* or inoculat\* or vaccin\*)).ti,ab,kw. (5595)
 ((2019-nCoV or nCoV or nCoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or

SARSCoV2 or SARS2) adj5 (immun\* or inoculat\* or vaccin\*)).ti,ab,kw. (13896)

179 (((BNT162 or BNT162-01 or BNT162a1 or BNT162b1 or BNT162b2 or BNT162c2) and vaccin\*) or N38TVC63NU).ti,ab,kw. (1123)

180 (((AZD1222 or ChAdOx1) and vaccin\*) or Covishield\$2 or B5S3K2V0G8).ti,ab,kw. (741)

181 ((Moderna and vaccin\*) or EPK39PL4R4).ti,ab,kw. (392)

182 ((mRNA adj3 vaccin\*) and (COVID-19 or COVID19 or coronavirus\* or corona virus\* or 2019-nCoV or nCoV or nCoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).ti,ab,kw. (1813)

183 ((messenger RNA adj3 vaccin\*) and (COVID-19 or COVID19 or coronavirus\* or corona virus\* or 2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCOV-2

184 (LV-SMENP-DC and vaccin\*).ti,ab,kw. (1)

185 ((Ad5-nCoV and vaccin\*) or hAdOx1 nCoV-19).ti,ab,kw. (26)

186 (("Ad26.COV2.S" or Ad26COVS1 or JNJ 78436735 or JNJ-78436735 or

JT2NS6183B) and vaccin\*).ti,ab,kw. (145)

187 Viral Vaccines/ and (Coronavirus/ or Betacoronavirus/ or Coronavirus Infections/) (1906)

188 or/175-187 [COVID-19 VACCINES] (35531)

189 COVID-19/pc [prevention & control] (8859)

190 Coronavirus Infections/pc [prevention & control] (10743)

191 Pandemics/pc [prevention & control] (14512)

192 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (COVID-19 or COVID19)).ti,ab,kw. (27702)

193 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (coronavirus\* or corona virus\*)).ti,ab,kw. (5416)

194 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 (2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2)).ti,ab,kw. (7180)

195 ((control\* or decreas\* or halt\* or prevent\* or reduc\* or stop\*) adj5 spread\*).ti,ab,kw. (52807)

196 COVID-19/ep [Epidemiology] (17717)

197 COVID-19/tm [Transmission] (3563)

198 COVID-19/vi [Virology] (6344)

199 Coronavirus Infections/ep [Epidemiology] (23509)

- 200 Coronavirus Infections/tm [Transmission] (4703)
- 201 Coronavirus Infections/vi [Virology] (7690)
- 202 exp Disease Transmission, Infectious/ (352747)

203 (transmit\* or transmissi\* or infectiousness\* or infectivit\*).ti,ab,kw. (1241265)



204 ((COVID-19 or COVID19) adj5 (caus\* or pass or passed or passes or passing or spread\*)).ti,ab,kw. (32798)

205 ((coronavirus\* or corona virus\*) adj5 (caus\* or pass or passed or passes or passing or spread\*)).ti,ab,kw. (17833)

206 ((virus\* or infection\*) adj5 (caus\* or pass or passed or passes or passing or spread\*)).ti,ab,kw. (386369)

207 ((2019-nCoV or nCoV or n-CoV or SARS-CoV-2 or SARS-CoV2 or SARSCoV-2 or SARSCoV2 or SARS2) adj5 spread\*).ti,ab,kw. (5298)

208 (unvaccinat\* or nonvaccinat\* or non-vaccinat\* or "not vaccinat\*").ti,ab,kw. (42646) 209 or/189-208 [TRANSMISSION] (1929116)

- 210 188 and 209 [COVID-19 VACCINES DISEASE TRANSMISSION] (17254)
- 211 (202105\* or 202106\* or 202107\* or 202108\*).up. (1733210)
- 212 210 and 211 [UPDATE PERIOD] (4802)
- 213 212 use cctr [CENTRAL RECORDS] (233)
- 214 86 or 174 or 213 [ALL DATABASES] (2369)
- 215 remove duplicates from 214 (1983) [TOTAL UNIQUE RECORDS]
- 216 215 use ppez [MEDLINE UNIQUE RECORDS] (569)
- 217 215 use oemezd [EMBASE UNIQUE RECORDS] (1206)
- 218 215 use cctr [CENTRAL UNIQUE RECORDS] (208)
- 219 202104\*.up. (277685)
- 220 210 and 219 [APRIL UPDATE] (917)
- 221 220 use cctr [CENTRAL APRIL UPDATE] (25)
- 222 214 or 221 [ALL UPDATE PERIODS ALL DATABASES] (2394)
- remove duplicates from 222 (2007)
- 224 223 use ppez [MEDLINE UNIQUE RECORDS] (569)
- 225 223 use oemezd [EMBASE UNIQUE RECORDS] (1206)
- 226 223 use cctr [CENTRAL UNIQUE RECORDS] (232)
- 227 226 and 219 [CENTRAL UNIQUE RECORDS APRIL ONLY] (24)

\*\*\*\*\*

PICO - Prevention>SARS-CoV-2 Vaccines-Primary Studies 3819 records