

1 **Comparative effectiveness of BNT162b2 and mRNA-1273 booster dose after**
2 **BNT162b2 primary vaccination against the Omicron variants: A retrospective cohort**
3 **study using large-scale population-based registries in Japan**

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10 **Running title:** Heterologous booster against COVID-19

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12

1 **Abstract**

2 **Background:** Direct comparative effectiveness of booster doses of BNT162b2 and mRNA-
3 1273 after BNT162b2 primary vaccination is unknown.

4 **Methods:** We investigated comparative effectiveness of BNT162b2 and mRNA-1273
5 booster dose using data from registry systems for vaccination and COVID-19 infection in a
6 local city in Japan. We followed participants aged ≥ 16 years who completed the BNT162b2
7 primary vaccination between November 22, 2021, and April 15, 2022. We collected
8 information on age, sex, vaccination status, vaccine type, and infection status. Age was
9 categorized as 16–44, 45–64, 65–84, and ≥ 85 years. Vaccine effectiveness for mRNA-1273
10 and no booster vaccination against BNT162b2 was estimated using age-stratified Cox
11 regression adjusted for age, sex, and days since the second vaccination. The estimated
12 hazard ratios for mRNA-1273 and no booster vaccinations were integrated separately using
13 random effects meta-analyses.

14 **Results:** During the study period, we identified 62,586 (40.4%), 51,490 (33.2%), and
15 40,849 (26.4%) participants who received BNT162b2, mRNA-1273, and no booster dose,
16 respectively. The median age was 69, 71, and 47 years for BNT162b2, mRNA-1273, and
17 no booster dose, respectively. The integrated hazard ratio with reference to BNT162b2 was

1 1.72 for no booster vaccination and 0.62 for mRNA-1273. The comparative effectiveness
2 of mRNA-1273 was similar across age categories.

3 **Conclusions:** Both homologous and heterologous vaccinations are effective against
4 Omicron variants. In the head-to-head comparison, the effect was stronger in people who
5 received heterologous vaccination than in those who received homologous vaccination.
6 These findings may help improve logistics and decision making in future vaccination
7 programs.

8 **Keywords:** Omicron variants, BNT162b2, mRNA-1273, booster dose, population-based
9 registries

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1 **Introduction**

2 Previous studies reported acceptable safety profiles and sufficient levels of antibodies after
3 the booster dose of coronavirus disease (COVID-19) vaccines [1,2]. These studies showed
4 that the levels of IgG anti-SARS-CoV-2-binding antibody and neutralizing antibody
5 differed by the combination of booster vaccine and primary course. Noticeably, participants
6 who received heterologous mRNA-1273 (Moderna) booster dose after two primary doses
7 of BNT162b2 (Pfizer-BioNTech) showed higher antibody levels than those with three
8 homologous BNT162b2 doses.

9 Higher antibody levels are inversely associated with symptomatic COVID-19
10 [3,4]. However, the actual effectiveness of heterologous and homologous vaccination has
11 not been directly compared at the population level. A recent population level study in the
12 UK examined the effectiveness of several combinations of primary course and booster dose
13 [5]. The study found that BNT162b2 or mRNA-1273 booster vaccination after BNT162b2
14 primary course effectively protected the participants against COVID-19 when compared to
15 no booster dose. Although mRNA-1273 booster vaccination after the BNT162b2 primary
16 course showed higher vaccine effectiveness than the three doses of BNT162b2 vaccines,

1 the two vaccines were not directly compared because that study aimed to estimate the
2 vaccine effectiveness compared with no booster dose.

3 In Japan, BNT162b2 or half dose (50 μ g) of mRNA-1273 has been available as a
4 booster dose since December 1, 2021, for people aged ≥ 18 years, regardless of the type of
5 primary vaccine course. The effectiveness of vaccines should be fully notified, especially
6 when there is a choice of booster vaccine, as in many countries. In addition to individual
7 choice, this information is useful for future logistics in vaccination campaigns. We
8 therefore conducted a head-to-head comparison of vaccine effectiveness between
9 BNT162b2 and mRNA-1273 in people who completed the primary BNT162b2 vaccination
10 course in Japan using data from registry systems for vaccination and COVID-19.

11

12 **Methods**

13 *Data source*

14 We used vaccination records linked with the Health Center Real-time information-sharing
15 System on COVID-19 (HER-SYS) data [6], collected between February 19, 2021, and
16 April 15, 2022, in Shimonoseki City, a municipality located in the southwestern region of
17 Japan. Briefly, Shimonoseki has approximately 250,000 residents and 130,000 households

1 in an area of 716 sq. km. in 2022 [7]. There are 26 hospitals with 5,300 beds, and the
2 population per physician is approximately 350 [8].

3 The vaccination records contain information on all residents of Shimonoseki city,
4 including age, sex, number of doses (first, second, and booster) of COVID-19 vaccine, date
5 of vaccination, and type of vaccine (BNT162b2, mRNA-1273, and ChAdOx1-S
6 [Oxford/AstraZeneca] that were available in Japan during the study period). As of April 11,
7 2022, approximately 80% of the residents of Shimonoseki city completed two primary
8 doses of the COVID-19 immunization course, and approximately 50% received a booster
9 dose [9].

10 The HER-SYS was established to share information on patients with COVID-19 in
11 governments, community health centers, and medical facilities. Patients diagnosed with
12 COVID-19 and their close contacts were registered in the HER-SYS for monitoring and
13 health management. The HER-SYS data includes the date of report on COVID-19
14 diagnosis, date of hospital admission for COVID-19, and date of death from COVID-19.

15 The date of diagnosis in the HER-SYS was entered by a physician for patients with
16 definitive diagnosis of COVID-19, which was made by the physician based on the results
17 of polymerase chain reaction and/or antigen testing [10,11]. People who tested positive

1 outside the medical facilities still required physician's assessment to be registered as
2 infected cases during the study period. Although the HER-SYS was designed to collect
3 other clinically important information, such as symptoms of COVID-19, comorbidities,
4 smoking, and body mass index, we were unable to utilize the information because of
5 missing values, as input of details was not mandatory at the time of the study.

6 The vaccination records were linked to the HER-SYS data via the Basic Resident
7 Register of Shimonoseki City using anonymized ID created based on name, sex, and date of
8 birth. Almost all (i.e., 99.98%) vaccination records and 76.3% of HER-SYS data were
9 linked to the Basic Resident Register. All HER-SYS data linked to the Basic Resident
10 Register were linked with the vaccination records. The linkage rate of the HER-SYS data
11 with the Basic Resident Register was relatively low because the data included information
12 on some people who were diagnosed with COVID-19 at medical institutions in
13 Shimonoseki City but lived outside the city.

14 The study was approved by the Institutional Review Board of the University of Tokyo
15 (2021187NI-[3]) and followed the Strengthening the Reporting of Observational Studies in
16 Epidemiology (STROBE) reporting guideline. As our analyses involved a secondary use of
17 anonymized data that was routinely collected by the local government in Japan, the

1 requirement for individual informed consent was waived in accordance with the current
2 ethical guidelines for medical and health research involving human participants in Japan.

4 ***Participants***

5 Of the study population vaccinated with the primary course, >80% received two doses of
6 BNT162b2, while the others received two doses of mRNA-1273, heterologous vaccination,
7 or unknown type of vaccination as a primary course. We included Shimonoseki residents
8 aged ≥ 16 years and who had completed two primary doses of the BNT162b2 COVID-19
9 immunization course by November 22, 2021—the start of the sixth wave of COVID-19 in
10 Japan. We excluded participants who received ChAdOx1-S (AstraZeneca) as a booster
11 dose, and those who had already been diagnosed with COVID-19 by the start of the sixth
12 wave.

14 ***Variables***

15 The primary outcome was COVID-19 diagnosis, defined as availability of a diagnosis date
16 in the HER-SYS data between November 22, 2021 and April 15, 2022, when Omicron
17 variants predominantly circulated. The other outcomes were hospitalization and COVID-

1 19-related death, defined as availability of a date for these events in the HER-SYS data,
2 although we failed to conduct statistical adjustment for these outcomes due to their rarity.
3 The exposures were as follows: (i) receiving mRNA-1273 as a booster dose (heterologous
4 COVID-19 vaccination) and (ii) receiving no third dose. The effects of these exposures
5 were estimated with reference to receiving BNT162b2 as the booster dose (homologous
6 COVID-19 vaccination). In the mass vaccination program, those who completed the
7 primary COVID-19 vaccination course at least 6 months prior were eligible for the booster
8 dose [12]. For people aged ≥ 18 years, BNT162b2 or half dose (50 μg) of mRNA-1273 was
9 available as a booster dose regardless of the type of primary vaccine course, while only
10 BNT162b2 was available as a third dose for people aged ≤ 17 years.

11 The exposures were treated as time-dependent variables in the analyses, and the
12 vaccination status was changed from “no booster” to “vaccinated with BNT162b2 booster”
13 or “vaccinated with mRNA booster” seven days after the booster dose was administered.
14 For a sensitivity analysis, the time-dependent exposure was further divided based on the
15 time from the booster dose (post-booster week 2–6, 7–12, and 13–). The thresholds of post-
16 booster week category were set based on the half of maximum follow-up day, the
17 maximum follow-up day, and beyond in those with mRNA booster. Other variables used

1 for statistical adjustment were age, sex, and number of days from the second dose to the
2 start of the sixth wave. Age was categorized as 16–44, 45–64, 65–84, and ≥ 85 years for
3 stratified analyses.

4

5 *Statistical analyses*

6 We described the characteristics of the participants stratified according to vaccination status
7 at the end of the follow-up period. The effect of the booster COVID-19 vaccination dose on
8 infection was analyzed for each age category using Cox regression with time-dependent
9 exposures. We followed participants from November 22, 2021, the start of the sixth wave,
10 to the date of infection, the date of death from COVID-19, or April 15, 2022, whichever
11 occurred first. Death from other reasons and moving out of the city (i.e., out of the cohort)
12 were disregarded because such data were unavailable. The results of the regression analyses
13 for each category were integrated using a random effects meta-analysis. Integration was
14 performed for heterologous vaccination and for no vaccine separately, with homologous
15 vaccination as a reference category for both. Heterogeneity among age categories was
16 assessed using I^2 measure [13]. To account for time difference from the booster dose (i.e.,
17 waning) between the groups, we conducted a sensitivity analysis for the entire cohort using

1 Cox regression with time-dependent exposure further divided by post-booster week
2 category (post-booster week 2–6, 7–12, and 13–). The other variables included in the
3 model were age, sex, and number of days from the second dose to the start of the sixth
4 wave, as in the main analysis. With BNT162b2 post-booster week 2–6 as a reference, we
5 presented the hazard ratio (HR) for each exposure category. Statistical analyses were
6 conducted using R Statistical Software (version 4.2.0; R Foundation for Statistical
7 Computing, Vienna, Austria).

9 **Results**

10 Of the 259,361 residents of Shimonoseki City, we identified 258,234 participants aged ≥ 16
11 years and who had completed two primary doses of BNT162b2 by November 22, 2021
12 (**Supplementary Figure 1**). We excluded 3 participants who received ChAdOx1-S as a
13 booster dose and 1,124 participants who had already been diagnosed with COVID-19 by
14 the start of the sixth wave. Of the remaining 154,925 participants, 62,586 (40.4%) received
15 BNT162b2, 51,490 (33.2%) received mRNA-1273, and 40,849 (26.4%) received no
16 vaccine as a booster dose by the end of the follow-up period. The distribution of age
17 categories and days from the second dose to the start of the sixth wave were similar in the

1 BNT162b2 and mRNA-1273 groups, while participants without booster doses were much
2 younger and had fewer days from the second dose to the start of the sixth wave than those
3 in the other two groups (**Table 1**). The duration (mean [standard deviation] days) from the
4 start of the sixth wave to the administration of BNT162b2 booster was shorter than that of
5 mRNA-1273 (96 [31] vs. 103 [16] days) (**Table 1**). The number of COVID-19 cases and
6 the number of booster dose administration in the cohort are shown in **Figure 1**. The
7 maximum follow-up days after the booster dose were 133 for those with BNT162b booster
8 and 80 for those with mRNA-1273 booster. Comparison of no vaccination with
9 homologous vaccination showed that the proportions of infection (4.9% vs. 1.4%) and
10 hospitalization (0.3% vs. 0.1%) for COVID-19 were higher in participants with no
11 vaccination (Table 2). Further, participants who received mRNA-1273 were less likely to
12 be diagnosed with COVID-19 than those who received BNT162b2 (0.7% vs. 1.4%).

13 After adjusting for age, sex, and days from the second dose to the start of the sixth
14 wave, the risk of infection was higher in participants without a booster dose than in those
15 with a BNT162b2 booster dose across all age categories (**Figure 2A**). The integrated results
16 showed that participants without a booster COVID-19 vaccine dose were more likely to be
17 infected (HR: 1.72, 95% confidence interval [CI]: 1.22–2.22, $I^2 = 80\%$) than participants

1 with BNT162b2 booster dose. In contrast, participants with the mRNA-1273 booster (i.e.,
2 heterologous vaccination) dose had a lower risk of COVID-19 than those with BNT162b2
3 booster (i.e., homologous vaccination) dose (HR: 0.62, 95% CI: 0.50–0.74, $I^2 = 0\%$)
4 (**Figure 2B**). The effects of heterologous vaccination were consistent across all age
5 categories. In the sensitivity analysis that accounted for time from the booster dose, those
6 with heterologous vaccination had consistently lower risk of COVID-19 over time than
7 those with homologous vaccination (HR: 0.61, 95% CI: 0.49–0.7 for post-booster week
8 2–6) (**Table 3**). Although the effect of homologous vaccination waned over time, those
9 with BNT162b2 booster had lower risk of COVID-19 than those without the booster dose
10 throughout the follow-up period.

12 **Discussion**

13 To the best of our knowledge, this is the first study to compare the effectiveness of
14 heterologous vaccination (mRNA-1273 after two doses of BNT162b2) and homologous
15 vaccination (three doses of BNT162b2), directly. The registry systems for vaccination and
16 infection, a relatively homogenous primary vaccination strategy, and the availability of two
17 different mRNA booster doses in Japan allowed a large-scale direct comparison of

1 vaccination regimens. While homologous vaccination was associated with less risk of
2 COVID-19 than no booster dose, heterologous vaccination had an even more protective
3 effect against COVID-19 than homologous vaccination.

4 Previous studies indicated that heterologous vaccination may have a greater
5 protective effect than homologous vaccination based on neutralizing antibody response
6 [1,2]. The IgG serum binding antibody and pseudovirus neutralizing antibody were higher
7 in participants who received mRNA-1273 booster dose after two doses of BNT162b2 than
8 in those who received three doses of BNT162b2 [1]. Later, a study from the UK confirmed
9 that the effectiveness of heterologous vaccination was indeed higher than that of the
10 homologous vaccination at the population level. However, these studies used no booster
11 vaccination as a reference and did not conduct direct comparisons between heterologous
12 and homologous vaccinations. The present study directly compared two mRNA booster
13 vaccination regimens and suggested that heterologous vaccination was more effective
14 against infections by Omicron variants than homologous vaccination at the population
15 level. Informing the population about the better vaccine effectiveness of the mRNA-1273
16 booster dose than BNT162b2 may further help in individual decision-making for COVID-
17 19 vaccination.

1 Heterologous booster vaccination has also attracted the attention of the world to
2 optimize logistics under an ongoing global inequity in vaccine distribution [14]. Despite the
3 accumulating evidence of the safety and effectiveness for homologous vaccination [15–17],
4 comparative effectiveness of heterologous vaccination remains uncertain. Nonetheless,
5 many countries have adopted heterologous booster regimens to respond to the surge of new
6 variants [18–21]. Because of a safety profile similar to that of homologous vaccination[1,2]
7 and the higher effectiveness confirmed by the present study, heterologous vaccination could
8 be a viable option for many countries. The booster dose in general reinforces protection
9 against COVID-19[5]; thus, an indiscriminative promotion of homologous and
10 heterologous booster dose in vaccination campaigns may further contribute to protecting
11 people from COVID-19.

12 This study has several limitations. First, the generalizability of the results may be
13 limited because the current study was conducted at a local municipality in Japan, where
14 almost all residents were Japanese. The effects of heterologous vaccination may differ in
15 other races. In addition, most of the viruses circulating during the study period were
16 Omicron variants [22]. Thus, the effects of the third dose of COVID-19 vaccine may differ
17 for other or future variants. Second, information on the participants was limited. We were

1 unable to adjust for known risk factors for COVID-19, such as comorbidities, smoking, and
2 body mass index, due to the lack of data. Imbalance of these factors between the groups
3 may have led to biased results in either direction depending on their distributions.
4 Similarly, we were unable to control for healthcare seeking behavior and access to
5 healthcare. The comparative effectiveness of homologous vaccination against no booster
6 vaccination may have been underestimated if those without booster dose did not seek
7 healthcare for COVID-19. Moreover, we were unable to censor the participants who died of
8 causes other than COVID-19 and those who moved out of the municipality due to the lack
9 of data. According to the census of Shimonoseki City, the number of individuals aged ≥ 15
10 years decreased by 1,663 between the end of November 2021 and end of April 2022 [7].
11 The results of comparison, especially between homologous vaccination and no vaccination,
12 may have been affected by a lack of proper censoring. However, we hypothesize that the
13 results of the comparison between homologous and heterologous vaccinations are still
14 meaningful due to the absence of a reason to claim that censoring occurred
15 disproportionately between the two groups. To optimize vaccination program, future
16 studies are warranted to compare effectiveness of several existing regimens of COVID-19
17 vaccines while addressing these concerns.

1 In conclusion, two different types of mRNA booster doses after two primary doses
2 of the BNT162b2 COVID-19 immunization course were associated with a lower risk of
3 COVID-19. In the head-to-head comparison, the protective effect against infection was
4 stronger with heterologous vaccination than with homologous vaccination. This
5 information can be utilized for individual decision-making in future vaccination campaigns.
6 Moreover, policymakers should consider heterologous mRNA vaccination as a viable
7 option, along with homologous vaccination, to protect people from COVID-19.

8

9

1 **NOTES**

2 **Author contributions**

3 SO, NM, and H Yasunaga conceived and designed the study. SO, MN, and H Yasunaga
4 were responsible for data acquisition. SO and NM analyzed data. All authors interpreted
5 data. SO wrote the first draft of the manuscript. All authors reviewed and edited the
6 manuscript critically for important intellectual content. All authors approved the final draft
7 of the manuscript.

8 **Ethical considerations**

9 This study was approved by the Research Ethics Committee of the Graduate School of
10 Medicine and Faculty of Medicine, University of Tokyo (approval no. 2021187NI-[3]). As
11 our analyses involved a secondary use of anonymized data that was routinely collected by
12 the local government in Japan, the requirement for individual informed consent was waived
13 in accordance with the current ethical guidelines for medical and health research involving
14 human participants in Japan.

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1 **Data availability statement**

2 The database used in this study was maintained by Shimonoseki city, Japan. Restrictions
3 applied to the availability of data that were used with permission for this study. Therefore,
4 data are not publicly available.

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10 **Conflict of interest**

11 SO is a member of the Department of Eat-loss Medicine, a cooperative program between
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3 potential conflicts of interest.

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ACCEPTED MANUSCRIPT

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1 **Table 1.** Characteristics of participants in each regimen

Characteristics	Booster dose		
	BNT162b2	mRNA-1273	None
n	62,586	51,490	40,849
Female, n (%)	39,200 (62.6)	29,149 (56.6)	22,019 (53.9)
Age, years, median (IQR)	69 (55–78)	71 (62–79)	47 (34–61)
Age category, n (%)			
16–44	8,649 (13.8)	3,694 (7.2)	17,814 (43.6)
45–64	15,400 (24.6)	12,160 (23.6)	14,279 (35.0)
65–84	31,080 (49.7)	29,429 (57.2)	5,935 (14.5)
≥85	7,457 (11.9)	6,207 (12.1)	2,821 (6.9)
Days from the second dose to the start of wave 6, mean (SD)	135 (39)	121 (21)	70 (43)
Days from the start of wave 6 to booster dose, mean (SD)	96 (31)	103 (16)	—

2 IQR, interquartile range; SD, standard deviation

1 **Table 2.** Unadjusted outcomes in each regimen

Outcomes	Booster dose		
	BNT162b2	mRNA-1273	None
n	62,586	51,490	40,849
Infection, n (%)	863 (1.4)	376 (0.7)	1,990 (4.9)
Hospitalization, n (%)	84 (0.1)	33 (0.1)	122 (0.3)
Death, n (%)	0 (0.0)	1 (0.0)	9 (0.0)

ACCEPTED MANUSCRIPT

- 1 **Table 3.** The risk of COVID-19 for each post-booster week category with reference to
 2 2–6 week category in BNT162b2 group.

Booster	Post-booster week category	Hazard ratio [95% CI]
No booster		1.82 [1.57, 2.11]
BNT162b2	WEEK 2–6	Reference
BNT162b2	WEEK 7–12	1.52 [1.27, 1.83]
BNT162b2	WEEK 13–*	1.22 [0.93, 1.60]
mRNA-1273	WEEK 2–6	0.61 [0.49, 0.76]
mRNA-1273	WEEK 7–12**	0.69 [0.45, 1.04]
mRNA-1273	WEEK 13–	–

3 * Maximum follow-up days after BNT162b booster was 133

4 ** Maximum follow-up days after mRNA-1273 booster was 80

5 The hazard ratio for each category was estimated by Cox regression, adjusted for age,

6 sex, and number of days from the second dose to the start of the sixth wave of

7 COVID-19. Abbreviation: CI, confidence interval; COVID-19, coronavirus disease.

1 **Figure Legends**

2 **Figure 1.** Time from the start of sixth wave to COVID-19 diagnosis and booster dose.

3 Panel A shows the number of COVID-19 cases at each time point. Panel B shows the
4 number of booster dose administration at each time point.

5 Abbreviation: COVID-19, coronavirus disease

6

7 **Figure 2.** The risk of COVID-19 for no booster dose group and for mRNA-1273

8 booster group with reference to BNT162b2 booster group.

9

10 Panel A shows the risk of COVID-19 for no booster group with reference to BNT162b2
11 booster group. Panel B shows the risk of COVID-19 for mRNA-1273 group with
12 reference to BNT162b2 booster group.

13 The results for each age category were estimated by Cox regression, adjusted for age,
14 sex, and number of days from the second dose to the start of the sixth wave of COVID-

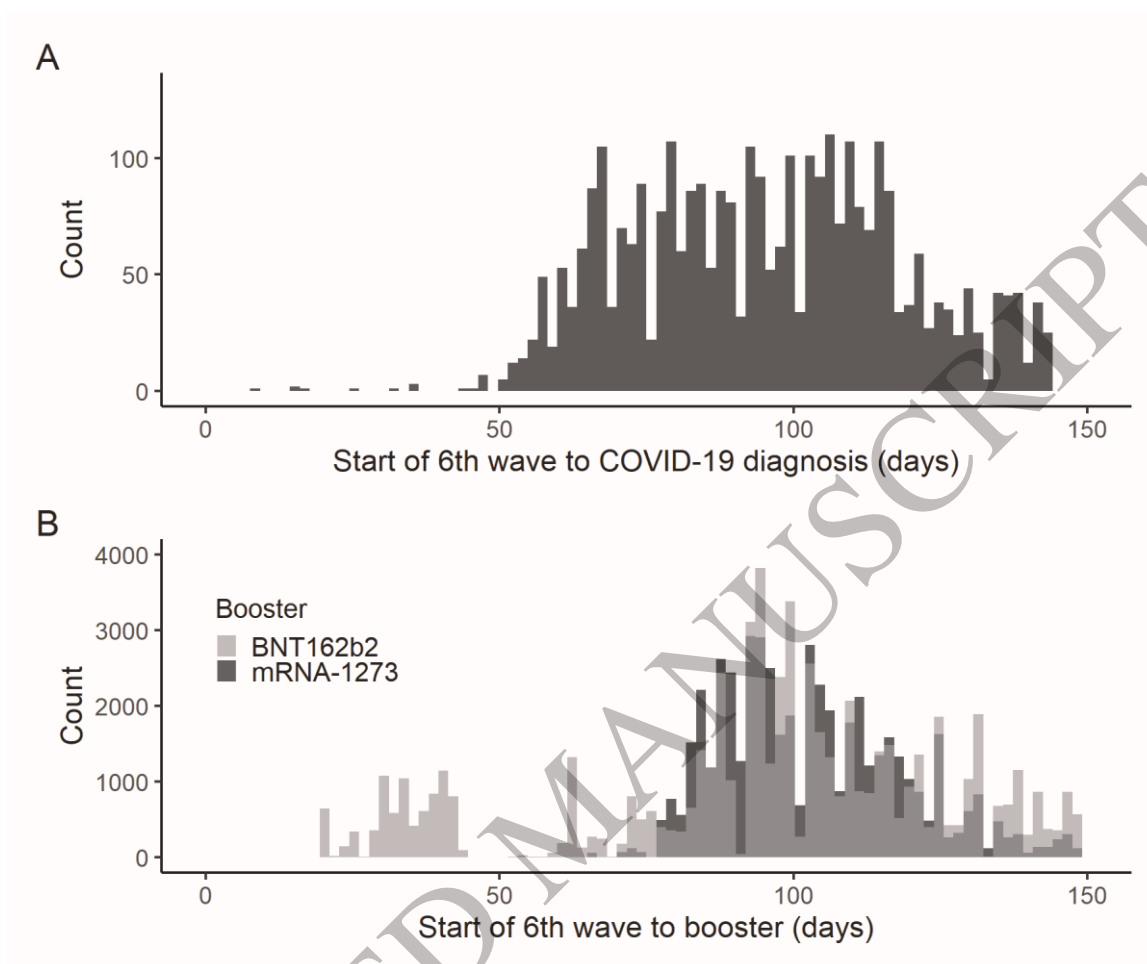
15 19. Hazard ratios from the age categories were integrated using the random-effects

16 meta-analysis method. Abbreviation: CI, confidence interval; COVID-19, coronavirus

17 disease.

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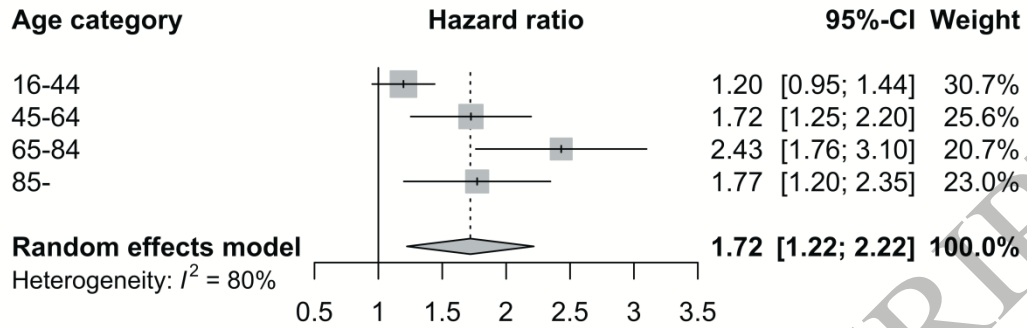
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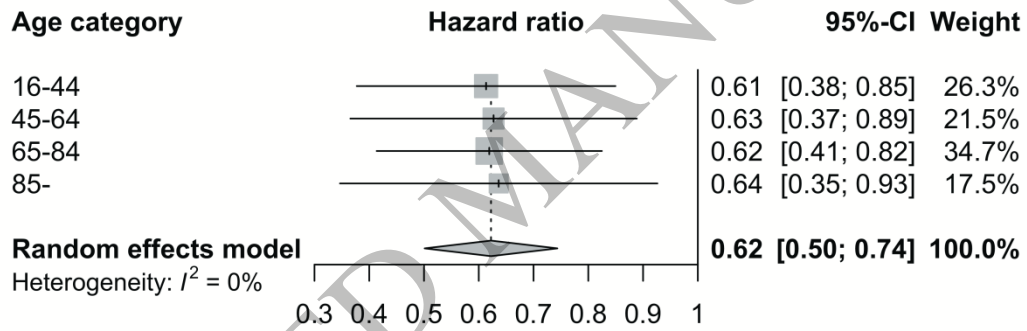
Figure 1

152x127 mm (x DPI)

A



B



1
2
3

Figure 2
356x356 mm (x DPI)