

# Immunogenicity, efficacy, and safety of SARS-CoV-2 vaccine dose fractionation: a systematic review and meta-analysis

Bingyi Yang (✉ [yangby@hku.hk](mailto:yangby@hku.hk))

The University of Hong Kong <https://orcid.org/0000-0002-0811-8332>

Xiaotong Huang

The University of Hong Kong

Huizhi Gao

The University of Hong Kong

Nancy Leung

HKU

Tim Tsang

The University of Hong Kong <https://orcid.org/0000-0001-5037-6776>

Benjamin Cowling

University of Hong Kong <https://orcid.org/0000-0002-6297-7154>

---

## Article

**Keywords:** SARS-CoV-2, vaccine, fractional dose

**Posted Date:** June 2nd, 2022

**DOI:** <https://doi.org/10.21203/rs.3.rs-1571821/v2>

**License:** © ⓘ This work is licensed under a Creative Commons Attribution 4.0 International License.

[Read Full License](#)

---

1 **Immunogenicity, efficacy, and safety of SARS-CoV-2 vaccine dose fractionation: a**  
2 **systematic review and meta-analysis**

3 Bingyi Yang<sup>1</sup>, Xiaotong Huang<sup>1</sup>, Huizhi Gao<sup>1</sup>, Nancy H. Leung<sup>1</sup>, Tim K. Tsang<sup>1,2</sup>, Benjamin J.  
4 Cowling<sup>1,2</sup>

5 **Affiliations:**

- 6 1. WHO Collaborating Centre for Infectious Disease Epidemiology and Control, School  
7 of Public Health, Li Ka Shing Faculty of Medicine, The University of Hong Kong, Hong  
8 Kong, China  
9 2. Laboratory of Data Discovery for Health Limited, Hong Kong Science and Technology  
10 Park, New Territories, Hong Kong, China

11 **Corresponding author:**

12 Benjamin J. Cowling, School of Public Health, Li Ka Shing Faculty of Medicine, The University  
13 of Hong Kong, 7 Sassoon Road, Pokfulam, Hong Kong

14 Tel: +852 3917 6711; Email: [bcowling@hku.hk](mailto:bcowling@hku.hk)

15 or

16 Bingyi Yang, School of Public Health, Li Ka Shing Faculty of Medicine, The University of Hong  
17 Kong, 7 Sassoon Road, Pokfulam, Hong Kong

18 Tel: +852 3917 6911; Email: [yangby@hku.hk](mailto:yangby@hku.hk)

19

20 **Word count:** (Abstract: 155)

21 (Main text: 2, 933)

22 **Abstract**

23 **Background:** Dose fractionation of Coronavirus Disease 2019 (COVID-19) vaccine could  
24 effectively accelerate global vaccine coverage, while supporting evidence of efficacy,  
25 immunogenicity, and safety are unavailable, especially with emerging variants.

26 **Methods:** We systematically reviewed clinical trials reported dose-finding results and  
27 estimated the dose-response relationship of neutralizing antibodies (nAbs) of COVID-19  
28 vaccines using generalized additive model. We predicted the vaccine efficacy against both  
29 ancestral and variants, using previously reported correlates of protection and cross-  
30 reactivity. We also reviewed and compared seroconversion to nAbs, T-cell responses and  
31 safety profiles between fractional and standard dose groups.

32 **Results:** We found that dose fractionation of mRNA and protein subunit vaccines could  
33 induce SARS-CoV-2 specific nAbs and T-cells that confer a reasonable level of protection  
34 (i.e., vaccine efficacy > 50%) against ancestral strains and variants up to Omicron. Safety  
35 profiles of fractional doses were non-inferior to the standard dose.

36 **Conclusion:** Dose fractionation of mRNA and protein subunit vaccines may be safe and  
37 effective.

## 38 Introduction

39 Three years into the pandemic, COVID-19 continues to threaten global health with emerging  
40 variants. While vaccinations are effective in preventing hospitalizations and deaths <sup>1,2</sup>, there  
41 has been unequal distribution of vaccinations across the globe. Despite that the current  
42 vaccine supply would cover most of the global population, a portion of the supply were  
43 prioritized for the fourth or fifth dose in high income countries, while only 15.7% of people  
44 in lower income countries had received at least one vaccine dose as of 16 May 2022 <sup>3</sup>. Dose  
45 fractionation of vaccines has been previously recommended to ease global supply shortage  
46 and accelerate vaccine coverage in low-income countries, where a larger proportion of the  
47 population could have access to vaccination while each individual would receive a lower  
48 vaccine dose <sup>4,5</sup>. However, uncertainties and concerns about the vaccine efficacy using  
49 fractional doses against SARS-CoV-2 ancestral strains and emerging variants of concern  
50 (VoCs) <sup>6-8</sup>, and the potential differences between vaccine platforms, hindered the  
51 endorsement for dose fractionation of COVID-19 vaccines <sup>9,10</sup>. Nevertheless, a half-dose of  
52 the original Moderna vaccine has been recommended for the booster dose for adults who  
53 are not moderately and severely immunocompromised <sup>11</sup>.

54

55 Here, we conducted a systematic review and meta-analysis of phase I/II trials that reported  
56 dose-finding results of immunogenicity and safety profiles for COVID-19 vaccines (detailed  
57 search terms in Table S1). We estimated the pooled dose-response relationship of  
58 neutralizing antibodies (nAbs) against the ancestral strain, which were then used to predict  
59 the potential VE of fractional doses against infections of the ancestral strain and VoCs using  
60 a hypothesized relation between nAbs and protection <sup>12,13</sup>. We also reviewed the  
61 differences in seroconversion of nAbs, T-cell mediated immune responses and safety profile

62 between fractional and standard dose (i.e., doses used for final products or phase III trials)  
63 groups, to further assess the differences in immunogenicity and safety after receiving  
64 fractional and standard doses.

65

## 66 **Methods**

### 67 ***Search strategy and study selection***

68 We searched peer-reviewed publications on clinical trials of SARS-CoV-2 vaccines in PubMed  
69 on 9 December 2021. We searched with the following terms: (SARS-CoV-2 OR COVID-19)  
70 AND (vaccine AND dose) AND (antibod\* OR immun\*) (Table S1). We included dose-  
71 escalation studies that reported safety, neutralizing antibodies (nAbs, which were measured  
72 by PRNT<sub>50</sub> and/or sVNT), and/or T-cell mediated immunity among healthy individuals  
73 received SRAS-CoV-2 vaccines (Tables S2-4). We excluded 1) studies did not report  
74 immunological response or only reported binding antibody; 2) studies without dose-  
75 escalation; 3) studies on non-human hosts; 4) studies on participants with specific health  
76 conditions (e.g., cancer, organ transplantation) or pregnancy; 5) studies specifically designed  
77 for hybrid immunity (i.e., natural infection or heterogenous vaccinations); and 6) reviews or  
78 commentaries (Figure S1). We assessed the quality of included studies using the Cochrane  
79 Risk of Bias tool 2.0 for randomized trials<sup>14</sup> (Figure S2).

80

### 81 ***Data extraction and processing***

82 Two reviewers (BY and XH) independently screened the titles and full texts of articles  
83 according to the inclusion and exclusion criteria. For each included study, we extracted  
84 relevant information of the vaccines and participants onto a standardized form, which

85 includes vaccine name, manufacture, platform, dose fraction, vaccination and sampling  
86 schedule, age group, and sizes of vaccinated subjects. Dose fraction ( $F_{i,j}$ ) was defined as the  
87 ratio of each examined dose group ( $d_{i,j}$ ) and the standard dose ( $d_{ref,j}$ , defined as the dose  
88 selected for the approved vaccine product or phase III trials) for each study ( $j$ ):

$$F_{i,j} = \frac{d_{i,j}}{d_{ref,j}} \quad (1)$$

89

### 90 ***Differences in seroconversion of neutralizing antibodies after fractional and standard dose***

91 We compared the seroconversion proportion to nAbs against ancestral strains after  
92 receiving fractional doses compared with the standard dose group, where seroconversion  
93 was predefined by each study as at least four-fold increase in nAbs and/or changing from  
94 negative to positive after vaccinations (details about definitions for positive threshold and  
95 seroconversion were shown in Table S3). We chose to estimate the pooled risks ratio of  
96 seroconversion over the pooled proportion of seroconversion, to minimize the impacts of  
97 measurement variations between laboratories. Sample size and the number of  
98 seroconverted participants were extracted for each dose group, which were then used to  
99 estimate the pooled log risk ratio of seroconversion between fractional and standard dose  
100 group using random effects (RE) model, stratified by vaccine type. We fitted mixed effects  
101 meta-regressions to assess the effects of vaccine platform and dose fractions on  
102 seroconversion, after accounting for age group and assay methods. We also repeated the  
103 above analysis for higher dose group, which results can be found in our data repository.

104

105 ***Dose-response relationship of neutralizing antibodies***

106 For each study  $j$ , we extracted the mean ( $\mu_{i,j}$ ) and standard deviations ( $\sigma_{i,j}$ ) of nAbs titers in  
107 different dose groups  $i$ ; if not reported, we estimated  $\mu_{i,j}$  and  $\sigma_{i,j}$  from 1) individual data  
108 points, or 2) median, interquartile (IQR) and sample sizes<sup>15</sup>. We then standardized the  
109 vaccine-induced nAbs level ( $z_{i,j}$ ) using the nAbs measured in convalescent sera ( $\mu_{c,j}$ ) for  
110 each study:

$$z_{i,j} = \frac{\mu_{i,j}}{\mu_{c,j}} \quad (2)$$

111 We summarized standardized nAbs among different dose groups (i.e., fractional, standard,  
112 and higher dose groups) at different time points (i.e., days after 1 or 2 doses). To quantify  
113 the non-linear dose-response relationship of vaccination and the standardized nAbs ( $z_{i,j}$ ),  
114 we fitted a generalized additive model (GAM; Table S5) that accounted for the vaccine  
115 platform ( $V$ ), vaccine schedule (i.e., total dosages  $D$  and days after full vaccination  $T_{i,j}$ ), age  
116 group ( $A$  = children, adult, or elderly) and antigen used for neutralizing assay ( $M$  = live or  
117 pseudo virus):

$$\log_2 z_{i,j} = \beta_0 + \beta_{TS}(T_{i,j}) + \beta_{FS}(\log_2 F_{i,j}) + \beta_V V_j + \beta_D D_{i,j} + \beta_A A_{i,j} + \beta_M M_{i,j} \quad (3)$$

118 With estimates from equation 3, we predicted the standardized nAbs (assuming measured  
119 by live virus and in adults; same for the following) against SARS-CoV-2 ancestral strain 14  
120 days after fully vaccinated (i.e., 1 dose for vector and 2 for the rest) with fractional doses  
121 ( $F_{i,j}$ ) for different vaccine platforms. We validated our model predictions and raw data and  
122 performed ten-fold cross-validation (Figure S3 and Table S6).

123

124 ***Vaccine efficacy predicted from neutralizing antibodies***

125 We applied the established CoP protection of standardized nAbs<sup>12,13</sup> to predict the dose-  
126 fractioning vaccine efficacy ( $\Phi_i$ ) against symptomatic infections of SARS-CoV-2 ancestral  
127 strain for different vaccine platform ( $V$ ):

$$\Phi_{i,V,E} = \frac{1}{1 + e^{-k_E \log_{10} \frac{z_{i,V}}{z_{50,E}}}} \quad (4)$$

128 We obtained the log-transformed 50% protective efficacy ( $\log_{10} z_{50,E}$ ) and steepness  
129 parameter  $k_E$  for both symptomatic and severe infection from the previous study<sup>12</sup>.  $z_{i,V}$  is  
130 the standardized nAbs at 14 days after fully vaccinated of fractional doses ( $F_{i,j}$ ) for each  
131 vaccine platform ( $V$ ), which was estimated from equation 3 with coefficients shown in Table  
132 S5.

133

134 We used previously reported<sup>6,13</sup> fold of reduction ( $\delta_S$ ; Table S7) in nAbs to estimate the  
135 level of standardized nAbs ( $\delta_S z_{i,V}$ ) against the variant  $S$ , which was then applied to equation  
136 4 to predict the vaccine efficacy of dose fractioning against infections of SARS-CoV-2 variant  
137 of concerns (VoCs). To validate our predicted vaccine efficacy against VoCs (Supplementary  
138 Figure 9), we compared the predicted vaccine efficacy against symptomatic infections after  
139 standard dose and observations (Table S8) reported previously by Pearson correlation.  
140 Standard doses were used for comparison since there were no empirical data regarding half-  
141 dose.

142

143 ***T-cell responses***

144 Since assays and measurements used for T-cell mediated responses vary across studies, we  
145 reviewed if T-cell responses elicited by dose-fractioning vaccines 1) would be higher than

146 that at pre-vaccination level and 2) would be lower than that elicited by the standard dose  
 147 vaccine within the same study. Briefly, we extracted mean ( $\bar{x}_{i,j,k}$ ; log-transformed if  
 148 originally measured in log-scale; same for SE), standard error (SE,  $\hat{\sigma}_{\bar{x}_{i,j,k}}$ ) and sample size ( $n_i$ )  
 149 of specific measurement  $k$  for T-cell responses for each dose group or reference group (i.e.,  
 150 pre-vaccination or post standard dose vaccination)  $i$  in study  $j$ . Specific measurement ( $k$ )  
 151 includes T-cell types (i.e., CD4+ or CD8+) and/or cytokines for T helper type 1 (Th1, including  
 152 interferon- $\gamma$  (IFN-  $\gamma$ ), tumor necrosis factor (TNF-  $\alpha$ ) and Interleukin-2 (IL-2)) and T helper  
 153 type 2 (Th2, including IL-4, IL-5, IL-13). If mean and SE were not reported, we estimated  
 154 these metrics from individual original data points or median, IQR and sample sizes<sup>15</sup>. We  
 155 determined the statistical significance of difference in (log-)means ( $\Delta_{i,j,k}$ ) assuming it  
 156 follows a normal distribution.

$$\bar{\Delta}_{i,j,k} = \bar{x}_{i,j,k} - \bar{x}_{ref,j,k} \quad (5)$$

$$\hat{\sigma}_{\bar{\Delta}_{i,j,k}} = \sqrt{\frac{\hat{\sigma}_{\bar{x}_{i,j,k}}^2}{n_i} + \frac{\hat{\sigma}_{\bar{x}_{ref,j,k}}^2}{n_{ref}}} \quad (6)$$

157

## 158 **Safety**

159 We compared the safety profiles after receiving fractional dose compared with the standard  
 160 dose group. We extracted the sample size and the number of adverse events (AEs, i.e.,  
 161 solicited local and/or systemic events, unsolicited events, and any AEs) for each dose group.  
 162 Individual manifestations within each AE category were extracted and assessed. We  
 163 estimated the pooled log risk ratio of experiencing AEs between fractional and standard  
 164 dose group using RE model and stratifying by specific AE and vaccine platform. We  
 165 calculated the  $I^2$  to measure the heterogeneity of the included estimates. We also repeated  
 166 the above analysis for higher dose group, which results can be found in our data repository.

167

## 168 **Results**

169 In total 1,733 records were returned from PubMed search with 44 duplicates. After titer and  
170 abstract screening, 136 records were eligible for full-text screen (Figure S1). Thirty-eight  
171 studies were included in the analyses <sup>16-53</sup>, among which inactivated vaccines (29%, n = 11)  
172 were studied the most, followed by protein subunit (“subunit” hereafter; 26%, n = 10),  
173 mRNA (24%, n = 9), non-replicating viral vector (“vector” hereafter; 13%, n = 5) and others  
174 (Figure S1 and Table S2). We found overall low risks of bias of the included studies, expect  
175 that seven adopted the non-randomized, and non-double-blinded design (Figure S2)

176 <sup>16,26,27,32,37,39,53</sup>.

177

### 178 ***Seroconversion of neutralizing antibodies after fractional doses***

179 We estimated the pooled risk ratio (RR) of the seroconversion against ancestral strains  
180 among individuals who completed fractional and standard dose from 14 studies of 9  
181 vaccines (Figure 1). The probability of seroconversion to ancestral strains was 2.1% (95%  
182 confidence interval (CI) 0.4% to 3.6%;  $I^2 = 52.0\%$ , P-value < 0.01) lower among individuals  
183 with fractional doses compared to standard doses within the same trial. However, we found  
184 no association between dose fractionation (1.4%, 95% CI, -20.4% to 29.3% per fold increase  
185 in dose) and seroconversion proportions between lower and standard dose groups after  
186 accounting for vaccine platform, age group and assay methods (i.e., live or pseudo virus)  
187 (Table S9).

188

189 ***Dose-relationship of neutralizing antibodies and predicted vaccine efficacy***

190 Twenty-four studies reported nAbs against live (n = 20) and/or pseudo (n = 7) ancestral  
191 viruses from both post-vaccination and convalescent sera (Figures S4-6 and Table S3). We  
192 estimated that prime with one half-dose would elicit less than 10% of nAbs in convalescent  
193 sera, while prime-boost with two half-doses elicited higher nAbs than a single standard dose  
194 across all platforms (Figure 2A and Figures S3-7).

195

196 We estimated that two half-dose mRNA vaccines would elicit 2.6 (95% CI, 2.1 to 3.3,  
197 measured on day 14) fold of the nAbs against the ancestral strain in convalescent sera  
198 (Figure 2A), which is expected to prevent 97% (95% CI, 95% to 97%) of symptomatic  
199 infections of the ancestral strains, respectively (Figure 2B). Whereas two half-dose  
200 inactivated vaccines would elicit 0.28 (95% IC 0.20 to 0.37) -fold of nAbs against the  
201 ancestral strains in convalescent sera, corresponding to 61% (95% CI, 51% to 70%) and 95%  
202 (95% CI, 92% to 96%) efficacy against symptomatic and severe infections of the ancestral  
203 strains, respectively. Overall, our predictions suggested that the reduction in vaccine  
204 efficacy was smaller than dose fractionation across all vaccine platforms (Figure 2C); half-  
205 doses may provide more than half of protection efficacy of standard doses.

206

207 Further incorporating the reported fold reduction in of vaccine-induced nAbs against VoCs  
208 (Table S7)<sup>6,13</sup>, we projected that two half-dose mRNA vaccines would confer the highest  
209 efficacy against symptomatic infections of VoCs (94%, 95% CI, 92% to 95% against Alpha,  
210 63%, 54% to 70% against Beta, 85%, 79% to 89% against Gamma, 83%, 78% to 87% against  
211 Delta and 32%, 26% to 40% against Omicron), followed by subunit, vector and inactivated  
212 vaccines (Figure 3 and Figure S8). Our predicted efficacy against symptomatic infections of

213 VoCs for standard dose highly correlated (Pearson correlation 0.705, p-value < 0.01; Figure  
214 S9) with empirical data <sup>54-64</sup>, while we were not able to validate predictions for fractional  
215 doses due to lack of data. Results from ten-fold validations further supported our model  
216 fitting (Table S6).

217

### 218 ***T-cell responses after fractional doses***

219 We first reviewed whether T-cell mediated immune responses elicited by dose fractioning  
220 vaccines would be higher than pre-vaccination level. All 7 studies of 5 vaccines reported  
221 significant increase in SARS-CoV-2 specific CD4+/CD8+ or CD4+ T helper type 1 (Th1)  
222 responses after vaccinated with fractional doses compared to pre-vaccination (Figure 4A),  
223 which were all biased to Th1 cells.

224

225 We then reviewed whether T-cell responses would be lower than that elicited by the  
226 standard dose vaccine. Three vaccines (BNT162b1<sup>30,37</sup>, MVC-COV1901<sup>26</sup> and Sf9 cells<sup>31</sup>)  
227 reported that dose fractionation elicited similar level of CD4+ and/or CD8+ T-cells compared  
228 to standard dose (Figure 4B). Quarter-dose of mRNA-1273 <sup>16,27</sup> was reported to induce  
229 significantly lower CD4+ Th1 cells compared to standard dose, while half-dose of BBV152  
230 were reported to induce significantly higher Th1 cytokines in one of two trials. We also  
231 compared the cellular responses between standard and higher dose groups and found no  
232 evidence for dose-dependent relationship for 7 out of 9 vaccines (Figure S10).

233

234 ***Safety profiles after fractional doses***

235 We reviewed the safety profile for 34 studies and found that, compared to standard dose  
236 group, people in the fractional dose groups tended to experience adverse events at similar  
237 or lower frequency (Figures 11-16). Particularly, the risk of experiencing solicited, and  
238 unsolicited adverse events were 9.5% (95% CI, 3.9% to 14.8 %) and 24.4% (3.9% to 41.1%)  
239 lower in individuals who received fractional dose of mRNA vaccines compared to standard  
240 doses (Figures 14-15). One inactivated (BBIBP-CorV in children <sup>46</sup>) and two subunit (NVX-  
241 CoV2373 and Livzon in adults <sup>22,38</sup>) vaccines reported higher risk of solicited systemic  
242 reactions in groups that received lower dose than the standard dose.

243

244 **Discussion**

245 Our findings suggested that vaccine-induced nAbs varied substantially across dose fractions,  
246 number of dosages, and vaccine platforms. For vaccine platforms (e.g., mRNA and subunit)  
247 which standard doses could elicit higher nAbs levels than convalescent sera, fractionation of  
248 prime-boost doses could induce robust nAbs against the ancestral strains and similar  
249 seroconversion proportion with standard doses. nAbs induced by fractional vaccines of  
250 mRNA and subunit were predicted to confer  $\geq 65\%$  efficacy against symptomatic infections  
251 of SARS-CoV-2 variants, except for Beta and Omicron. Fractionation of vaccine doses  
252 seemed to be safe and induce robust Th1 biased T-cell responses that were similar to  
253 standard doses except for mRNA-1273.

254

255 We found that dose fractionation of COVID-19 vaccines would induce, though lower than  
256 standard doses, detectable nAbs against ancestral strains. Based on previously established  
257 CoP <sup>12,13,65</sup>, these nAbs may confer reasonable protection (i.e., > 50%) against symptomatic

258 infections of ancestral strains, but not the subsequent VoCs, especially Omicron. Previous  
259 modelling study suggested that dose fractionation could be a cost-effective strategy in low-  
260 income countries, if vaccination could confer at least 50% of protection against symptomatic  
261 infections of variants with low or moderate transmissibility (i.e., basic reproduction number  
262  $R_0 < 5$ )<sup>66</sup>. Given these findings, our results of nAbs and vaccine efficacy predictions  
263 suggested that dose fractionation could have been cost-effective strategy to control the  
264 emergence of some early VoCs (e.g., Alpha and Delta), but not for the currently circulating  
265 Omicron given the significant immunity breakthrough<sup>6,7</sup> and higher transmissibility<sup>67</sup>.

266

267 While there is no established correlate of protection against severe COVID, we found that  
268 fractional doses of most studied vaccines could induce detective and likely robust T-cell  
269 responses, which may contribute to protection against severe outcomes given that SARS-  
270 CoV-2 specific T-cells could broadly cross-react to a range of VoCs (including Omicron) and  
271 were associated with better outcomes<sup>68,69</sup>. Therefore, dose fractionation of COVID-19  
272 vaccines might still be able to avert a considerable number of hospitalizations and deaths,  
273 even with the emergence of new variants with higher rates of breakthrough infections.

274

275 We were not able to assess the durability of the immune responses elicited by fractional  
276 doses of COVID-19 vaccines, as most trials reported limited follow up that was typically just  
277 one month after vaccination. Therefore, our efficacy estimates may only be indicative for  
278 short-term protection. Waning SARS-CoV-2 specific aAbs, T-cells and vaccine efficacy  
279 (against both ancestral and VoCs) may be expected, as suggested by evidence from  
280 individuals receiving standard doses after 6 months<sup>7,70-72</sup>. For standard doses, both

281 homogeneous and heterogenous boosters could substantially increase nAbs and vaccine  
282 efficacy against VoCs<sup>8</sup>, while such data were lacking for fractional doses.

283

284 To minimize the impacts of measurement variations between laboratories, we compared  
285 the differences in seroconversion of nAbs and T-cell responses within each trial and  
286 quantified the dose-relationship using nAbs that were standardized to convalescent sera.

287 Calibration to recommended international standard may further reduce the between  
288 laboratory variations, which was, however, not reported by the included trials.

289

290 We did not look at the nAbs induced by individual vaccine manufacturers due to limited  
291 data, while we found consistent seroconversion proportion and dose-relationship within  
292 platform (Figures 3-7). Nevertheless, disparities in nAbs levels and durability were reported  
293 for individual vaccines from the same platform (e.g., mRNA-1273 vs. BNT162b1 vaccines<sup>9</sup>).

294

295 Our study only focused on the immunogenicity and safety and the projected efficacy of dose  
296 fractionation of COVID-19 vaccines, and therefore findings should be interpreted within this  
297 scope. Endorsement of dose fractionation of vaccines by regulatory agencies would likely  
298 need stronger efficacy data, and other considerations would include the evolving supply  
299 situation, logistics restrictions, and vaccine communications.

300

301 To summarise, fractionation of vaccine doses, especially mRNA and protein subunit  
302 vaccines, are safe and would induce antibody and T-cell responses that likely confer a  
303 reasonable level of protection against severe infections of SARS-CoV-2 ancestral and VoCs  
304 up to Omicron. The use of vaccines with lower antigen content earlier in the pandemic

305 might have been an efficient approach to save even more lives, while further clinical  
306 investigation of fractional booster doses would certainly be worthwhile.

307

308

### 309 **Data availability**

310 All data were collected from publicly available literatures, with detailed description in the  
311 Methods and Supplementary Tables. Data used for the analysis can be assessed at:  
312 [https://github.com/byyangyby/fractional\\_dose\\_review](https://github.com/byyangyby/fractional_dose_review).

313

### 314 **Code availability**

315 The authors declare that all codes for analyzing the data are made available at  
316 [https://github.com/byyangyby/fractional\\_dose\\_review](https://github.com/byyangyby/fractional_dose_review).

317

### 318 **Acknowledgements**

319 This project was supported by the Theme-based Research Scheme (Project No. T11-712/19-  
320 N) of the Research Grants Council of the Hong Kong SAR Government. BJC also  
321 acknowledges the support AIR@InnoHK administered by Innovation and Technology  
322 Commission of the Government of the Hong Kong Special Administrative Region.

323

### 324 **Author contributions**

325 All authors meet the ICMJE criteria for authorship. BY and BJC conceived the study. BY and  
326 XH performed the literature review and screening. BY, XH and HG extracted data. BY  
327 analyzed the data and wrote the first draft of the manuscript. All authors provided critical  
328 review and revision of the text and approved the final version.

329

330 **Completing interest statement**

331 B.J.C. consults for AstraZeneca, GSK, Moderna, Roche, Sanofi Pasteur, and Pfizer. The

332 remaining authors declare no competing interests.

333 **References**

- 334 1. Haas EJ, McLaughlin JM, Khan F, et al. Infections, hospitalisations, and deaths  
335 averted via a nationwide vaccination campaign using the Pfizer-BioNTech BNT162b2 mRNA  
336 COVID-19 vaccine in Israel: a retrospective surveillance study. *Lancet Infect Dis* 2022; **22**(3):  
337 357-66.
- 338 2. Mesle MM, Brown J, Mook P, et al. Estimated number of deaths directly averted in  
339 people 60 years and older as a result of COVID-19 vaccination in the WHO European Region,  
340 December 2020 to November 2021. *Euro Surveill* 2021; **26**(47).
- 341 3. Mathieu E, Ritchie H, Ortiz-Ospina E, et al. A global database of COVID-19  
342 vaccinations. *Nat Hum Behav* 2021; **5**(7): 947-53.
- 343 4. Cowling BJ, Lim WW, Cobey S. Fractionation of COVID-19 vaccine doses could extend  
344 limited supplies and reduce mortality. *Nat Med* 2021; **27**(8): 1321-3.
- 345 5. World Health Organization. Fractional dose yellow fever vaccine as a dose-sparing  
346 option for outbreak response. 2016.
- 347 6. Carreno JM, Alshammary H, Tcheou J, et al. Activity of convalescent and vaccine  
348 serum against SARS-CoV-2 Omicron. *Nature* 2021.
- 349 7. Andrews N, Stowe J, Kirsebom F, et al. Covid-19 Vaccine Effectiveness against the  
350 Omicron (B.1.1.529) Variant. *New England Journal of Medicine* 2022.
- 351 8. Cheng SMS, Mok CKP, Leung YWY, et al. Neutralizing antibodies against the SARS-  
352 CoV-2 Omicron variant BA.1 following homologous and heterologous CoronaVac or  
353 BNT162b2 vaccination. *Nat Med* 2022.
- 354 9. Wilder-Smith A, Desai S, Cravioto A, Nohynek H, Hombach J. Caution before  
355 fractionating COVID-19 vaccines. *Nature Medicine* 2021; **27**(11): 1856-7.
- 356 10. World Health Organization. Interim statement on dose-sparing strategies for COVID-  
357 19 vaccines (fractionated vaccine doses). 2021. [https://www.who.int/news/item/10-08-  
358 2021-interim-statement-on-dose-sparing-strategies-for-covid-19-vaccines-\(fractionated-  
359 vaccine-doses\)](https://www.who.int/news/item/10-08-2021-interim-statement-on-dose-sparing-strategies-for-covid-19-vaccines-(fractionated-vaccine-doses)) (accessed 6 March 2022).
- 360 11. Centers for Diseases Control and Prevention. Interim Clinical Considerations for Use  
361 of COVID-19 Vaccines | CDC. May 23, 2022. [https://www.cdc.gov/vaccines/covid-19/clinical-  
362 considerations/covid-19-vaccines-us.html#considerations-covid19-vax-booster](https://www.cdc.gov/vaccines/covid-19/clinical-considerations/covid-19-vaccines-us.html#considerations-covid19-vax-booster) (accessed  
363 June 1 2022).
- 364 12. Khoury DS, Cromer D, Reynaldi A, et al. Neutralizing antibody levels are highly  
365 predictive of immune protection from symptomatic SARS-CoV-2 infection. *Nat Med* 2021;  
366 **27**(7): 1205-11.
- 367 13. Cromer D, Steain M, Reynaldi A, et al. Neutralising antibody titres as predictors of  
368 protection against SARS-CoV-2 variants and the impact of boosting: a meta-analysis. *Lancet*  
369 *Microbe* 2022; **3**(1): e52-e61.
- 370 14. Sterne JAC, Savovic J, Page MJ, et al. RoB 2: a revised tool for assessing risk of bias in  
371 randomised trials. *BMJ* 2019; **366**: l4898.
- 372 15. Wan X, Wang W, Liu J, Tong T. Estimating the sample mean and standard deviation  
373 from the sample size, median, range and/or interquartile range. *BMC Med Res Methodol*  
374 2014; **14**: 135.
- 375 16. Anderson EJ, Roupheal NG, Widge AT, et al. Safety and Immunogenicity of SARS-CoV-  
376 2 mRNA-1273 Vaccine in Older Adults. *N Engl J Med* 2020; **383**(25): 2427-38.

- 377 17. Chappell KJ, Mordant FL, Li Z, et al. Safety and immunogenicity of an MF59-  
378 adjuvanted spike glycoprotein-clamp vaccine for SARS-CoV-2: a randomised, double-blind,  
379 placebo-controlled, phase 1 trial. *Lancet Infect Dis* 2021.
- 380 18. Che Y, Liu X, Pu Y, et al. Randomized, double-blinded and placebo-controlled phase II  
381 trial of an inactivated SARS-CoV-2 vaccine in healthy adults. *Clin Infect Dis* 2020.
- 382 19. Chu L, McPhee R, Huang W, et al. A preliminary report of a randomized controlled  
383 phase 2 trial of the safety and immunogenicity of mRNA-1273 SARS-CoV-2 vaccine. *Vaccine*  
384 2021; **39**(20): 2791-9.
- 385 20. Ella R, Reddy S, Jogdand H, et al. Safety and immunogenicity of an inactivated SARS-  
386 CoV-2 vaccine, BBV152: interim results from a double-blind, randomised, multicentre, phase  
387 2 trial, and 3-month follow-up of a double-blind, randomised phase 1 trial. *Lancet Infect Dis*  
388 2021; **21**(7): 950-61.
- 389 21. Ella R, Vadrevu KM, Jogdand H, et al. Safety and immunogenicity of an inactivated  
390 SARS-CoV-2 vaccine, BBV152: a double-blind, randomised, phase 1 trial. *Lancet Infect Dis*  
391 2021; **21**(5): 637-46.
- 392 22. Formica N, Mallory R, Albert G, et al. Different dose regimens of a SARS-CoV-2  
393 recombinant spike protein vaccine (NVX-CoV2373) in younger and older adults: A phase 2  
394 randomized placebo-controlled trial. *PLoS Med* 2021; **18**(10): e1003769.
- 395 23. Goepfert PA, Fu B, Chabanon AL, et al. Safety and immunogenicity of SARS-CoV-2  
396 recombinant protein vaccine formulations in healthy adults: interim results of a randomised,  
397 placebo-controlled, phase 1-2, dose-ranging study. *Lancet Infect Dis* 2021; **21**(9): 1257-70.
- 398 24. Guo W, Duan K, Zhang Y, et al. Safety and immunogenicity of an inactivated SARS-  
399 CoV-2 vaccine in healthy adults aged 18 years or older: A randomized, double-blind,  
400 placebo-controlled, phase 1/2 trial. *EClinicalMedicine* 2021; **38**: 101010.
- 401 25. Han B, Song Y, Li C, et al. Safety, tolerability, and immunogenicity of an inactivated  
402 SARS-CoV-2 vaccine (CoronaVac) in healthy children and adolescents: a double-blind,  
403 randomised, controlled, phase 1/2 clinical trial. *Lancet Infect Dis* 2021.
- 404 26. Hsieh SM, Liu WD, Huang YS, et al. Safety and immunogenicity of a Recombinant  
405 Stabilized Prefusion SARS-CoV-2 Spike Protein Vaccine (MVC-COV1901) Adjuvanted with  
406 CpG 1018 and Aluminum Hydroxide in healthy adults: A Phase 1, dose-escalation study.  
407 *EClinicalMedicine* 2021; **38**: 100989.
- 408 27. Jackson LA, Anderson EJ, Roupheal NG, et al. An mRNA Vaccine against SARS-CoV-2 -  
409 Preliminary Report. *N Engl J Med* 2020; **383**(20): 1920-31.
- 410 28. Keech C, Albert G, Cho I, et al. Phase 1-2 Trial of a SARS-CoV-2 Recombinant Spike  
411 Protein Nanoparticle Vaccine. *N Engl J Med* 2020; **383**(24): 2320-32.
- 412 29. Kremsner PG, Mann P, Kroidl A, et al. Safety and immunogenicity of an mRNA-lipid  
413 nanoparticle vaccine candidate against SARS-CoV-2 : A phase 1 randomized clinical trial.  
414 *Wien Klin Wochenschr* 2021: 1-11.
- 415 30. Li J, Hui A, Zhang X, et al. Safety and immunogenicity of the SARS-CoV-2 BNT162b1  
416 mRNA vaccine in younger and older Chinese adults: a randomized, placebo-controlled,  
417 double-blind phase 1 study. *Nat Med* 2021; **27**(6): 1062-70.
- 418 31. Meng FY, Gao F, Jia SY, et al. Safety and immunogenicity of a recombinant COVID-19  
419 vaccine (Sf9 cells) in healthy population aged 18 years or older: two single-center,  
420 randomised, double-blind, placebo-controlled, phase 1 and phase 2 trials. *Signal Transduct*  
421 *Target Ther* 2021; **6**(1): 271.
- 422 32. Momin T, Kansagra K, Patel H, et al. Safety and Immunogenicity of a DNA SARS-CoV-  
423 2 vaccine (ZyCoV-D): Results of an open-label, non-randomized phase I part of phase I/II

424 clinical study by intradermal route in healthy subjects in India. *EClinicalMedicine* 2021; **38**:  
425 101020.

426 33. Mulligan MJ, Lyke KE, Kitchin N, et al. Phase I/II study of COVID-19 RNA vaccine  
427 BNT162b1 in adults. *Nature* 2020; **586**(7830): 589-93.

428 34. Pan HX, Liu JK, Huang BY, et al. Immunogenicity and safety of a severe acute  
429 respiratory syndrome coronavirus 2 inactivated vaccine in healthy adults: randomized,  
430 double-blind, and placebo-controlled phase 1 and phase 2 clinical trials. *Chin Med J (Engl)*  
431 2021; **134**(11): 1289-98.

432 35. Richmond P, Hatchuel L, Dong M, et al. Safety and immunogenicity of S-Trimer (SCB-  
433 2019), a protein subunit vaccine candidate for COVID-19 in healthy adults: a phase 1,  
434 randomised, double-blind, placebo-controlled trial. *Lancet* 2021; **397**(10275): 682-94.

435 36. Sadoff J, Le Gars M, Shukarev G, et al. Interim Results of a Phase 1-2a Trial of  
436 Ad26.COVS.2 Covid-19 Vaccine. *N Engl J Med* 2021; **384**(19): 1824-35.

437 37. Sahin U, Muik A, Derhovanessian E, et al. COVID-19 vaccine BNT162b1 elicits human  
438 antibody and T(H)1 T cell responses. *Nature* 2020; **586**(7830): 594-9.

439 38. Shu YJ, He JF, Pei RJ, et al. Immunogenicity and safety of a recombinant fusion  
440 protein vaccine (V-01) against coronavirus disease 2019 in healthy adults: a randomized,  
441 double-blind, placebo-controlled, phase II trial. *Chin Med J (Engl)* 2021; **134**(16): 1967-76.

442 39. Tebas P, Yang S, Boyer JD, et al. Safety and immunogenicity of INO-4800 DNA vaccine  
443 against SARS-CoV-2: A preliminary report of an open-label, Phase 1 clinical trial.  
444 *EClinicalMedicine* 2021; **31**: 100689.

445 40. Walsh EE, Frenck RW, Jr., Falsey AR, et al. Safety and Immunogenicity of Two RNA-  
446 Based Covid-19 Vaccine Candidates. *N Engl J Med* 2020; **383**(25): 2439-50.

447 41. Walter EB, Talaat KR, Sabharwal C, et al. Evaluation of the BNT162b2 Covid-19  
448 Vaccine in Children 5 to 11 Years of Age. *N Engl J Med* 2021.

449 42. Ward BJ, Gobeil P, Séguin A, et al. Phase 1 randomized trial of a plant-derived virus-  
450 like particle vaccine for COVID-19. *Nat Med* 2021; **27**(6): 1071-8.

451 43. Wu S, Huang J, Zhang Z, et al. Safety, tolerability, and immunogenicity of an  
452 aerosolised adenovirus type-5 vector-based COVID-19 vaccine (Ad5-nCoV) in adults:  
453 preliminary report of an open-label and randomised phase 1 clinical trial. *Lancet Infect Dis*  
454 2021.

455 44. Wu Z, Hu Y, Xu M, et al. Safety, tolerability, and immunogenicity of an inactivated  
456 SARS-CoV-2 vaccine (CoronaVac) in healthy adults aged 60 years and older: a randomised,  
457 double-blind, placebo-controlled, phase 1/2 clinical trial. *Lancet Infect Dis* 2021; **21**(6): 803-  
458 12.

459 45. Xia S, Duan K, Zhang Y, et al. Effect of an Inactivated Vaccine Against SARS-CoV-2 on  
460 Safety and Immunogenicity Outcomes: Interim Analysis of 2 Randomized Clinical Trials.  
461 *Jama* 2020; **324**(10): 951-60.

462 46. Xia S, Zhang Y, Wang Y, et al. Safety and immunogenicity of an inactivated SARS-CoV-  
463 2 vaccine, BBIBP-CorV: a randomised, double-blind, placebo-controlled, phase 1/2 trial.  
464 *Lancet Infect Dis* 2021; **21**(1): 39-51.

465 47. Xia S, Zhang Y, Wang Y, et al. Safety and immunogenicity of an inactivated COVID-19  
466 vaccine, BBIBP-CorV, in people younger than 18 years: a randomised, double-blind,  
467 controlled, phase 1/2 trial. *Lancet Infect Dis* 2021.

468 48. Yang S, Li Y, Dai L, et al. Safety and immunogenicity of a recombinant tandem-repeat  
469 dimeric RBD-based protein subunit vaccine (ZF2001) against COVID-19 in adults: two

470 randomised, double-blind, placebo-controlled, phase 1 and 2 trials. *Lancet Infect Dis* 2021;  
471 **21**(8): 1107-19.

472 49. Zhang J, Hu Z, He J, et al. Safety and immunogenicity of a recombinant interferon-  
473 armed RBD dimer vaccine (V-01) for COVID-19 in healthy adults: a randomized, double-  
474 blind, placebo-controlled, Phase I trial. *Emerg Microbes Infect* 2021; **10**(1): 1589-97.

475 50. Zhang Y, Zeng G, Pan H, et al. Safety, tolerability, and immunogenicity of an  
476 inactivated SARS-CoV-2 vaccine in healthy adults aged 18-59 years: a randomised, double-  
477 blind, placebo-controlled, phase 1/2 clinical trial. *Lancet Infect Dis* 2021; **21**(2): 181-92.

478 51. Zhu F, Jin P, Zhu T, et al. Safety and immunogenicity of a recombinant adenovirus  
479 type-5-vectored COVID-19 vaccine with a homologous prime-boost regimen in healthy  
480 participants aged 6 years and above: a randomised, double-blind, placebo-controlled, phase  
481 2b trial. *Clin Infect Dis* 2021.

482 52. Zhu FC, Guan XH, Li YH, et al. Immunogenicity and safety of a recombinant  
483 adenovirus type-5-vectored COVID-19 vaccine in healthy adults aged 18 years or older: a  
484 randomised, double-blind, placebo-controlled, phase 2 trial. *Lancet* 2020; **396**(10249): 479-  
485 88.

486 53. Zhu FC, Li YH, Guan XH, et al. Safety, tolerability, and immunogenicity of a  
487 recombinant adenovirus type-5 vectored COVID-19 vaccine: a dose-escalation, open-label,  
488 non-randomised, first-in-human trial. *Lancet* 2020; **395**(10240): 1845-54.

489 54. Abu-Raddad LJ, Chemaitelly H, Butt AA, National Study Group for C-V. Effectiveness  
490 of the BNT162b2 Covid-19 Vaccine against the B.1.1.7 and B.1.351 Variants. *N Engl J Med*  
491 2021; **385**(2): 187-9.

492 55. Chemaitelly H, Yassine HM, Benslimane FM, et al. mRNA-1273 COVID-19 vaccine  
493 effectiveness against the B.1.1.7 and B.1.351 variants and severe COVID-19 disease in Qatar.  
494 *Nat Med* 2021; **27**(9): 1614-21.

495 56. Li XN, Huang Y, Wang W, et al. Effectiveness of inactivated SARS-CoV-2 vaccines  
496 against the Delta variant infection in Guangzhou: a test-negative case-control real-world  
497 study. *Emerg Microbes Infect* 2021; **10**(1): 1751-9.

498 57. Lopez Bernal J, Andrews N, Gower C, et al. Effectiveness of Covid-19 Vaccines against  
499 the B.1.617.2 (Delta) Variant. *N Engl J Med* 2021; **385**(7): 585-94.

500 58. Madhi SA, Baillie V, Cutland CL, et al. Efficacy of the ChAdOx1 nCoV-19 Covid-19  
501 Vaccine against the B.1.351 Variant. *N Engl J Med* 2021; **384**(20): 1885-98.

502 59. Nasreen S, Chung H, He S, et al. Effectiveness of COVID-19 vaccines against  
503 symptomatic SARS-CoV-2 infection and severe outcomes with variants of concern in  
504 Ontario. *Nat Microbiol* 2022; **7**(3): 379-85.

505 60. Sheikh A, McMenamin J, Taylor B, Robertson C, Public Health S, the EIIIC. SARS-CoV-2  
506 Delta VOC in Scotland: demographics, risk of hospital admission, and vaccine effectiveness.  
507 *Lancet* 2021; **397**(10293): 2461-2.

508 61. Tang P, Hasan MR, Chemaitelly H, et al. BNT162b2 and mRNA-1273 COVID-19  
509 vaccine effectiveness against the SARS-CoV-2 Delta variant in Qatar. *Nat Med* 2021; **27**(12):  
510 2136-43.

511 62. Shinde V, Bhikha S, Hoosain Z, et al. Efficacy of NVX-CoV2373 Covid-19 Vaccine  
512 against the B.1.351 Variant. *N Engl J Med* 2021; **384**(20): 1899-909.

513 63. Tseng HF, Ackerson BK, Luo Y, et al. Effectiveness of mRNA-1273 against SARS-CoV-2  
514 Omicron and Delta variants. *Nat Med* 2022.

515 64. Heath PT, Galiza EP, Baxter DN, et al. Safety and Efficacy of NVX-CoV2373 Covid-19  
516 Vaccine. *N Engl J Med* 2021.

517 65. Padmanabhan P, Desikan R, Dixit NM. Modeling how antibody responses may  
518 determine the efficacy of COVID-19 vaccines. *Nature Computational Science* 2022; **2**(2): 123-  
519 31.

520 66. Du Z, Wang L, Pandey A, et al. Modeling comparative cost-effectiveness of SARS-  
521 CoV-2 vaccine dose fractionation in India. *Nat Med* 2022.

522 67. Nishiura H, Ito K, Anzai A, Kobayashi T, Piantham C, Rodriguez-Morales AJ. Relative  
523 Reproduction Number of SARS-CoV-2 Omicron (B.1.1.529) Compared with Delta Variant in  
524 South Africa. *J Clin Med* 2021; **11**(1).

525 68. Sette A, Crotty S. Adaptive immunity to SARS-CoV-2 and COVID-19. *Cell* 2021; **184**(4):  
526 861-80.

527 69. Rydzynski Moderbacher C, Ramirez SI, Dan JM, et al. Antigen-Specific Adaptive  
528 Immunity to SARS-CoV-2 in Acute COVID-19 and Associations with Age and Disease Severity.  
529 *Cell* 2020; **183**(4): 996-1012 e19.

530 70. Ward H, Whitaker M, Flower B, et al. Population antibody responses following  
531 COVID-19 vaccination in 212,102 individuals. *Nat Commun* 2022; **13**(1): 907.

532 71. Abu-Raddad LJ, Chemaitelly H, Bertollini R, National Study Group for C-V. Waning  
533 mRNA-1273 Vaccine Effectiveness against SARS-CoV-2 Infection in Qatar. *N Engl J Med* 2022.

534 72. Peng Q, Zhou R, Wang Y, et al. Waning immune responses against SARS-CoV-2  
535 variants of concern among vaccinees in Hong Kong. *eBioMedicine* 2022; **77**.

536

537

538 **Figure legends**

539 **Figure 1. Pooled risk ratio (in log scale) of seroconversion between fractional and standard**  
540 **dose group of COVID-19 vaccines.** Number of seroconversion individuals and sample sizes  
541 were shown for the standard and nonstandard group, respectively.

542

543 **Figure 2. Dose-response relationship of neutralizing antibodies (nAbs) and vaccine efficacy**  
544 **(VE) against ancestral strains induced by COVID-19 vaccines.** A 2-dose schedule was  
545 assumed for RNA, protein subunit and inactivated vaccines, while 1-dose schedule was  
546 assumed for non-replicating viral vector (as suggested by the included trials). (A) Dose-  
547 response relationship of nAbs against ancestral strains. nAbs were standardized as the ratio  
548 to the convalescent sera. Dashed horizontal line indicates the average level of nAbs against  
549 ancestral strains in convalescent sera. (B) Dose-response relationship of predicted vaccine  
550 efficacy against symptomatic infections of ancestral strains. (C) Association between  
551 reduction in vaccine efficacy and dose fractionation. Reductions in vaccine efficacy were  
552 measured as the ratio between vaccine efficacy against symptomatic infections of ancestral  
553 strains between fractional and standard dose groups.

554

555 **Figure 3. Predicted vaccine efficacy against SARS-CoV-2 variants of concern after fully**  
556 **vaccinated with half-dose vaccines.** Vaccine efficacy against symptomatic infections after  
557 full vaccinations (i.e., one dose non-replicating viral vector and two doses for the rest) of  
558 half-dose are shown, with the complete dose-dependent effectiveness are shown in  
559 Supplementary Figure 8. A-E for Alpha, Beta, Gamma, Delta, and Omicron.

560

561 **Figure 4. Comparison of T-cell responses against the ancestral strains elicited by dose**  
562 **fractioning of COVID-19 vaccines.** Size of dots represent the total sample sizes of the  
563 standard and non-standard dose groups. (A) Compared to pre-vaccination. If the mean and  
564 95% CI of the difference in mean T-cell levels before and after the fractional doses were all  
565 greater than 0, we determined T-cell responses were significantly higher between the  
566 groups. (B) Compared to people who received standard doses. If the mean and 95% CI of the  
567 difference in mean T-cell levels between the fractional and standard dose groups were all  
568 greater or less than 0, we determined T-cell responses were significantly higher or lower  
569 than that elicited by the standard dose.

# Figures

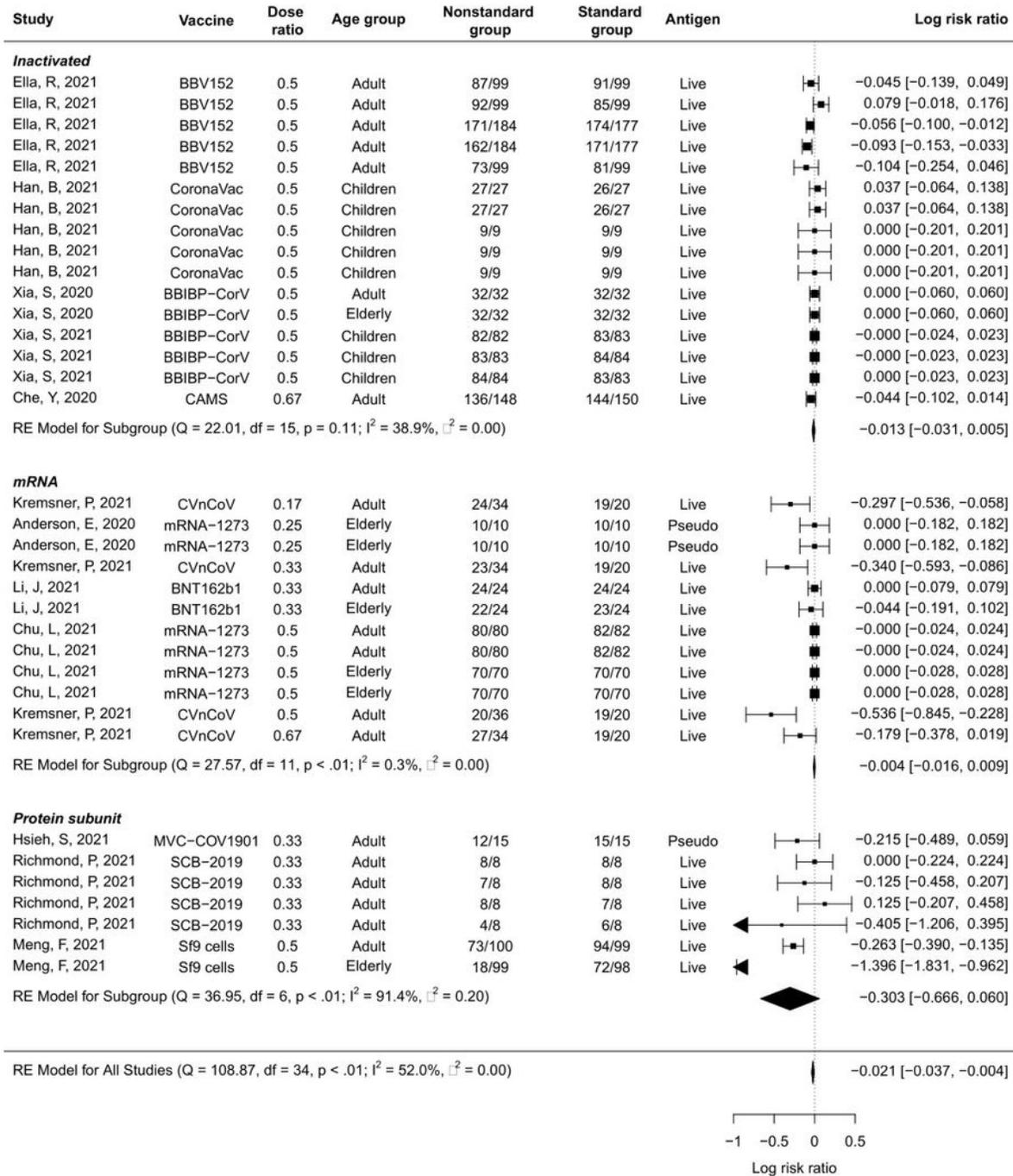


Figure 1

Pooled risk ratio (in log scale) of seroconversion between fractional and standard dose group of COVID-19 vaccines. Number of seroconversion individuals and sample sizes were shown for the standard and nonstandard group, respectively.

Figure 2

**Figure 2. Dose-response relationship of neutralizing antibodies (nAbs) and vaccine efficacy (VE) against ancestral strains induced by COVID-19 vaccines.** A 2-dose schedule was assumed for RNA, protein subunit and inactivated vaccines, while 1-dose schedule was assumed for non-replicating viral vector (as suggested by the included trials). (A) Dose-response relationship of nAbs against ancestral strains. nAbs were standardized as the ratio to the convalescent sera. Dashed horizontal line indicates the average level of nAbs against ancestral strains in convalescent sera. (B) Dose-response relationship of predicted vaccine efficacy against symptomatic infections of ancestral strains. (C) Association between reduction in vaccine efficacy and dose fractionation. Reductions in vaccine efficacy were measured as the ratio between vaccine efficacy against symptomatic infections of ancestral strains between fractional and standard dose groups.

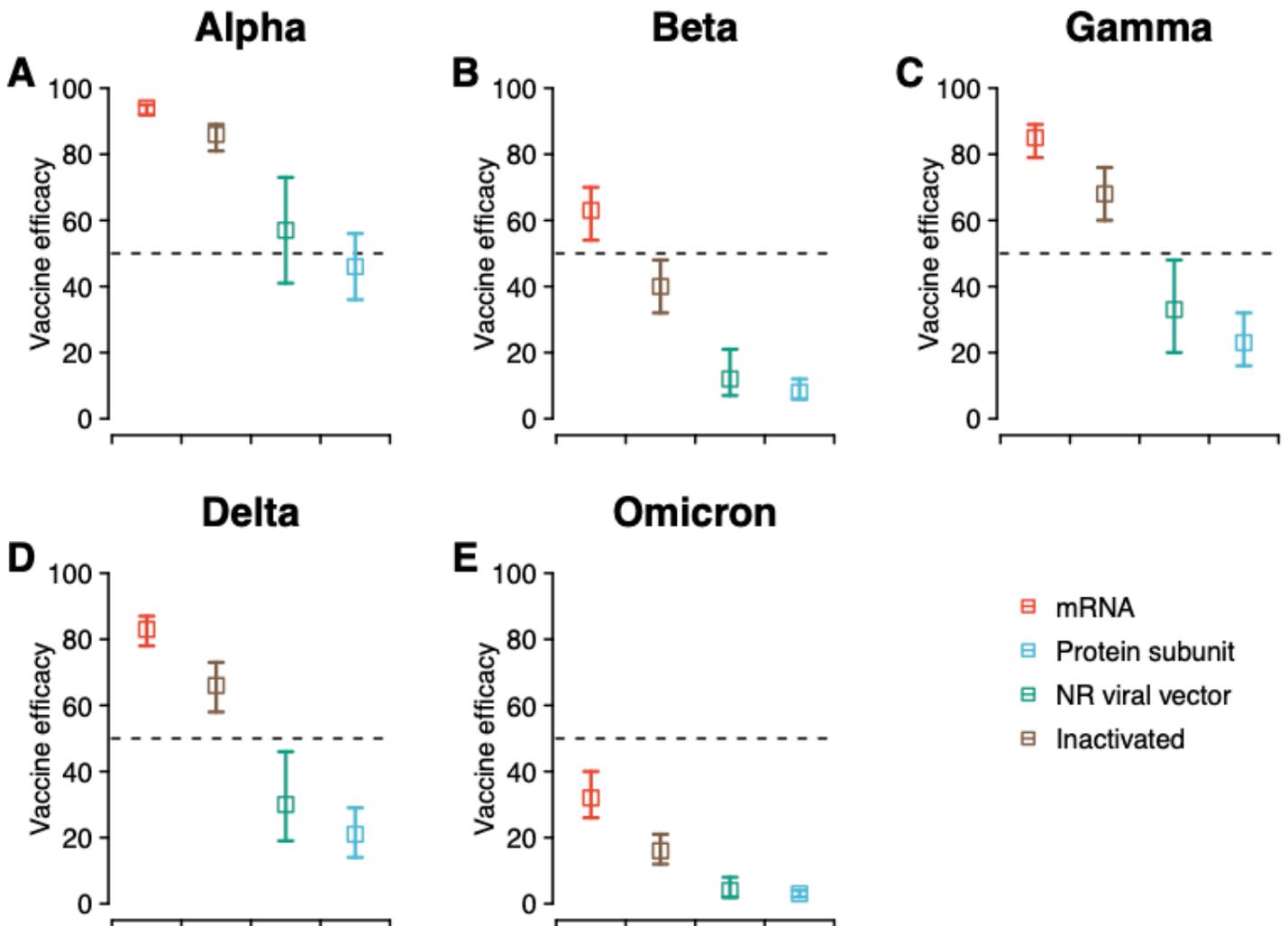


Figure 3

**Figure 3. Predicted vaccine efficacy against SARS-CoV-2 variants of concern after fully vaccinated with half-dose vaccines.** Vaccine efficacy against symptomatic infections after full vaccinations (i.e., one dose non-replicating viral vector and two doses for the rest) of half-dose are shown, with the complete dose-dependent effectiveness are shown in Supplementary Figure 8. A-E for Alpha, Beta, Gamma, Delta, and Omicron.

#### Figure 4

**Figure 4. Comparison of T-cell responses against the ancestral strains elicited by dose fractioning of COVID-19 vaccines.** Size of dots represent the total sample sizes of the standard and non-standard dose groups. (A) Compared to pre-vaccination. If the mean and 95% CI of the difference in mean T-cell levels before and after the fractional doses were all greater than 0, we determined T-cell responses were significantly higher between the groups. (B) Compared to people who received standard doses. If the mean and 95% CI of the difference in mean T-cell levels between the fractional and standard dose groups were all greater or less than 0, we determined T-cell responses were significantly higher or lower than that elicited by the standard dose.

## Supplementary Files

This is a list of supplementary files associated with this preprint. Click to download.

- [Supplementary.docx](#)