

Correlates of Neutralizing/SARS-CoV-2-S1-binding Antibody Response with Adverse Effects and Immune Kinetics in BNT162b2-Vaccinated Individuals

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Abstract

While mRNA vaccines against SARS-CoV-2 are exceedingly effective in preventing symptomatic infection, their immune response features remain to be clarified. In the present prospective study, 225 healthy individuals in Japan, who received two BNT162b2 doses, were enrolled. Correlates of BNT162b2-elicited SARS-CoV-2-neutralizing activity (50% neutralization titer: NT₅₀; assessed using infectious virions) with various determinants were examined and the potency of serums against variants of concerns was determined. Significant rise in NT₅₀s was seen in serums on day 28 post-1st dose. A moderate inverse correlation was seen between NT₅₀s and ages, but no correlation seen between NT₅₀s and adverse effects. NT₅₀s and SARS-CoV-2-S1-binding-IgG levels on day 28 post-1st dose and pain scores following the 2nd shot were greater in women than in men. The average half-life of NT₅₀s was ~ 68 days and the estimated average time length till the total disappearance of neutralizing activity was ~ 198 days. While serums from elite-responders (NT₅₀s > 1,500-fold: the top 4% among the participants) potentially to moderately blocked all variants of concerns examined, some serums with low NT₅₀s failed to block the B.1.351-beta strain. Since BNT162b2-elicited immunity against SARS-CoV-2 is short, an additional vaccine or other protective measures are needed.

Introduction

Since the emergence of coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) in Wuhan, China, the disease quickly spread to the world. As of June 29, 2021, more than 180 million SARS-CoV-2-infected individuals and almost 4 million death cases have been reported in over 200 countries¹⁻⁴. Since the beginning of the pandemic, researchers and pharmaceutical companies around the world have been working on developing vaccines⁵. Currently, more than 10 vaccines have been authorized for public use worldwide. The development of vaccines against SARS-CoV-2 was achieved time- and efficacy-wise beyond our expectations within a single calendar year from the availability of the viral sequence to the initiation of immunization of many people in several countries^{6,7}.

Among various vaccines, two RNA vaccines (BNT162b2 and mRNA-1273/TAK-919) have been shown to be as much as 94–95% effective and safe⁸⁻¹⁰. In addition, inactivated vaccines or viral vector vaccines have also been available in certain countries and areas^{5,7,9,10}. For example, the adenovirus-vector-based vaccine (ChAdOx1 nCoV-19/AZD1222) has reportedly achieved 62% efficacy in initial trials¹¹. The phase 3 reports of another adenovirus-based vaccine (Ad26.COVS.2.S) has indicated 85% efficacy against severe disease or death^{12,13}.

However, the recent emergence of various SARS-CoV-2 variants with mutations in the spike region is raising concerns about the efficacy of vaccines. The D614G and B.1.1.7 (alpha/N501Y) variants appear to be without antigenic escape^{14,15}. However, the B.1.351 (beta) variant is reportedly represents a neutralization escape variant to convalescent sera¹⁶. The phase 3 results of NVX-CoV2373 (a

nanoparticle, protein-based vaccine) from the United Kingdom indicated 89% efficacy with over 50% of cases attributable to the more transmissible alpha variant¹⁷. However, a phase 2b trial in South Africa showed 60% efficacy, in which approximately 90% of the endpoints occurred in subjects infected with the beta variant^{18,19}, suggesting that the beta variant is less susceptible to antibodies elicited with the original Wuhan strain antigens, which is in the composition of all the vaccines currently being evaluated⁷. Another recent concern is the emergence of a B.1.617.2 (delta) variant, which was first detected in India, is now spreading around the world. This variant of concern (VOC) seems to have less susceptibility to vaccine-elicited protection and increased transmissibility beyond alpha strains²⁰.

In the present study, we examined neutralizing activity and S1-binding-antibody response in BNT162b2-vaccinated health care workers (n = 225) in Japan. We also investigated the correlation among neutralizing activity levels, S1-binding-IgG and -IgM levels, genders, and adverse events. Decline of BNT162b2-elicited immune response and activity of the elite and moderate responders against VOCs were also investigated.

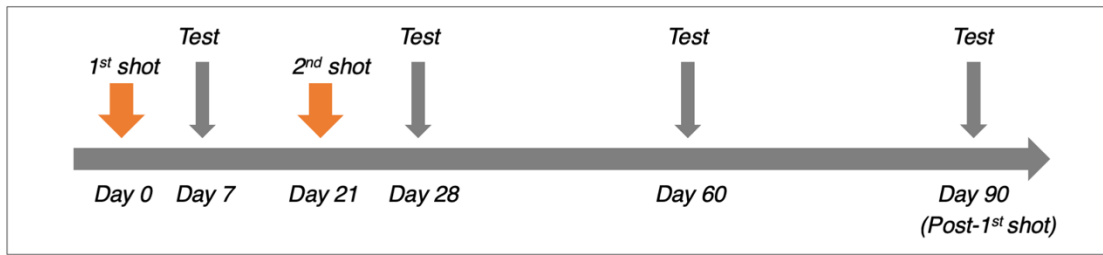
Results

Demographic characteristics and immune response in the participants.

We obtained blood samples for antibody testing from a total of 225, 220, 211, and 210 vaccine recipients on days 7, 28, 60, and 90 post-1st shot, respectively (Table 1). Demographic characteristics of the participants are shown in Table 1. As of the time of enrollment, the average age of the participants was 41.8 years (range: 21 to 72 years), and 69.8% of the participants were female serving as a physician, nurse, paramedical staff, or administrative staff. None of the participants was in the immunodeficient state or was receiving immunosuppressants or steroids.

Table 1: Study protocol and demographic characteristics of the participants.

We first determined the neutralizing activity against SARS-CoV-2 in serum samples taken on day 7 post-1st shot from 225 participants; however, none of the samples showed detectable neutralization activity ($NT_{50} < 20$ -fold). We then determined the neutralizing activity in samples taken on day 28 post-1st shot from 220 vaccinated participants. As shown in Fig. 1A, NT_{50} levels were substantially diverse among the participants: the geometric mean of NT_{50} values was 375.2 (range 25.6–2680.0). Very low or no correlation of the NT_{50} values with ages was identified (Fig. 1A: Spearman's $\rho = -0.22$; 95% CI -0.34 to -0.09). We also examined the levels of S1-binding-IgG and -IgM levels using the HISCL system that enables quantitative and highly sensitive determination of S1-binding-IgG and -IgM levels²¹. The geometric mean of S1-binding-IgG values was 527.0 (range 44.6-3212.2), while that of S1-binding-IgM was 85.1 (range 10.3-1406.5). There was a high positive correlation of the NT_{50} values with S1-binding-IgG levels (Spearman's $\rho = 0.71$; 95% CI 0.63 to 0.77) as examined on day 28 post-1st shot, while there was only a low positive correlation of the NT_{50} values with S1-binding-IgM levels (Spearman's $\rho = 0.43$; 95% CI 0.31 to 0.53), suggesting that the neutralizing activity largely resides in IgG fraction of the serum



		Day7 (Post 1 st dose)	(%)	Day28	Day60	Day90
All		225		220	211	210
Age	20-39	97	43.1	92	84	84
	40-59	110	48.9	110	109	108
	≥60	18	8.0	18	18	18
	(Average)	(41.8 y.o.)				
Gender	Men	68	30.2	68	63	61
	Women	157	69.8	52	148	149
Job	Physicians	36	16.0			
	Nurses	125	55.6			
	Others	64	28.4			

of vaccinated participants around on day 28 post-1st shot (Figs. 1B and 1C). However, when examined on day 60 post-1st shot, the correlations of NT₅₀ levels with both IgG and IgM levels became moderate or low (Spearman's $\rho = 0.56$ and 0.32 , respectively) (Supplementary Fig. 1).

The occurrence of adverse effects has no association with the BNT162b2-elicited neutralizing activity levels.

Commonly observed adverse events reported following BNT162b2 vaccination include injection-site pain, systemic fever, headache, and fatigue¹⁰. In the present study, the events were observed largely more often following the 2nd shot (Supplementary Fig. 2) as previously reported by Polack *et al.*¹⁰ Pains in the inject-site were reported by 67.6 and 61.6% of the participants and systemic fever ($\geq 37.1^\circ\text{C}$) was reported by 3.6 and 46.4% of the participants following the 1st and 2nd shots, respectively. Since the severity of pains can be relatively more quantitatively rated than that of other adverse effects such as headache and fatigue, the possible correlate of the NT₅₀ values with the severity of pains rated with the short form McGill pain questionnaire²² was first examined. No correlation was seen between the NT₅₀ values and the pain grades assessed following the 2nd shot (Spearman's $\rho = 0.14$; 95% CI 0.00 to 0.26). The correlation was also negligible between the NT₅₀ values and the incidence of systemic fever (Spearman's $\rho = 0.26$; 95% CI 0.13 to 0.38)(Figs. 2A and 2B).

The average half-life of neutralizing activity in the vaccinees is approximately 67.8 days and the average time-length for their serums to lose the detectable activity is 198.3 days.

Considering that recent multiple clinical studies strongly suggest that the presence of high-level neutralizing antibodies is generally sufficient to confer protection against SARS-CoV-2 infection and that the protection against COVID-19 development is largely explained by robust SARS-CoV-2-neutralizing antibody response⁸⁻¹⁰. If so, the once-established neutralizing antibody levels will decrease in time. We thus examined at what rate the levels of NT₅₀ and S1-binding-IgG and -IgM levels change by determining those levels from the data on day 28 (n = 220), day 60 (n = 211), and day 90 (n = 210) post-1st shot (Figs. 3A-C). The reduction of all NT₅₀, IgG, and IgM levels from day 28 through day 90 post-1st shot was found to occur virtually linearly. By computation, the predicted average half-life of all the NT₅₀ values turned out to be 67.8 days and those of S1-binding-IgG and IgM levels were 53.5 days and 43.6 days, respectively (Fig. 3D). The half-life of the NT₅₀ values and that of S1-binding-IgG were reasonably comparable, corroborating that the neutralizing activity largely resides in the S1-binding IgG fraction. Based on the chronologically linear nature of the reduction identified, we attempted to extrapolate from such half-life values and tried to predict the average time-length for the serums of the participants having significant NT₅₀ values to lose the activity down to under the undetectable level (UDL) (<20-fold) (Fig. 3D). The predicted average time-length for the serums to lose the activity was computed to be 198.3 days, while that of the top 10% participants to lose the activity was 204.3 days. The time-length of the middle 10% (between the top 45% and 55%) participants to lose the activity was 187.6 days. For all participants, it was predicted that day 160 after the 1st shot was when the 80% lower limit of predicted NT₅₀ levels drops under the detection level (UDL), while day 237 after the 1st shot was when the 80% upper limit of predicted NT₅₀ levels drops below UDL. Similarly, for the top 10% participants, the estimated days dropping below UDL were 171 and 243 for lower and upper limits, respectively, while for the middle 10%, the estimated days were 155 and 224, respectively. As for IgG, the predicted time-length for the serums to get undetected was 346.8 days, and days reaching below UDL were 333 and 362 for lower and upper limits, respectively. Likewise, the estimated day reaching below UDL for IgM was 183.6, and the lower and upper limits were 170 and 198 days, respectively (Fig. 3D). We also asked whether the chronological decay rate of neutralization titers and S1-binding-IgG and -IgM differs among three age subgroups: (i) 20–39 yo, (ii) 40–59 yo, and (iii) 60's and beyond. No significant difference was identified among the three age subgroups in the levels of neutralizing titers, IgG, or IgM levels ($p = 0.60, 0.16, \text{ and } 0.11$, respectively: Supplementary Figs. 3A-C). The present data suggest that vaccinated individuals with good neutralization response would lose BNT162b2's protection in 6 to 7 months without regard to age subgroups unless such people achieve robust immune boost response upon the future exposure to SARS-CoV-2. Otherwise, they should be protected by another booster vaccine shot or by other protective means.

Neutralization titers, S1-binding-IgG levels, and pain scores in the injection site were greater in women than in men.

We then asked whether there were differences between genders in neutralization activity levels, S1-binding-IgG and -IgM levels, injection-site pain scores, and systemic fever grades. Statistically significant differences were identified in the levels of neutralization determined on 60 and 90 days post-1st shot ($p = 0.002$ and 0.002 , respectively), S1-binding-IgG levels determined on 28, 60, and 90 days ($p < 0.001$, $p =$

0.001, and $p = < 0.001$, respectively) post-1st -shot, and S1-binding-IgM levels on 60 and 90 days post-1st shot ($p = 0.025$ and 0.044 , respectively)(Supplementary Figs. 4A-C). The injection-site pain score was greater in women ($p < 0.001$) (Supplementary Fig. 4D). However, there was no difference in systemic fever grades between genders (Supplementary Fig. 4E). However, no difference was seen in the decline rates of neutralization activity, S1-binding-IgG and -IgM levels between men and women (Supplementary Fig. 5).

Some serums retain potent activity against various VOCs, but others showed substantially less potent or undetectable activity.

We finally asked whether the neutralizing antibodies elicited by BNT162b2 vaccination blocked the infectivity and replication of various variants of concerns (VOCs). To this end, we employed serum samples from 6 elite responders (NT_{50} values $> 1,500$ -fold: the top 4% of all participants' NT_{50} values as determined on 28 days post-1st dose) and serum samples from twelve randomly-selected moderate responders (NT_{50} values = $200 \sim 1,500$ -fold) and tested them for their inhibition of the infectivity and cytopathic effect of each variant in the VeroE6_{TMPRSS2} cell-based assay²³. As shown in Fig. 4, serums from the elite responders ($n = 6$) showed potent inhibition against SARS-CoV-2_{05-2N} (Wuhan strain, PANGO lineage B), while they showed less activity against SARS-CoV-2_{QHN001} and SARS-CoV-2_{QK002} (alpha), SARS-CoV-2₅₃₅₆ (kappa), SARS-CoV-2₁₇₃₄ (delta), and SARS-CoV-2_{TY8-612} (beta). Serums from moderate responders ($n = 12$) exerted less activity against the Wuhan strain than those from the elite responders. Some serums from the moderate responders also showed substantially low potency to all the VOCs tested. Notably, three serums from the moderate responders showed only marginal activity against SARS-CoV-2_{TY8-612} (beta). Two of those three samples had no detectable inhibitory activity against SARS-CoV-2_{TY8-612} (Fig. 4).

Discussion

In this prospective observational study, 225 healthy individuals [physicians ($n = 36$), nurses ($n = 125$), and other healthcare professionals ($n = 64$)], who received two doses of $30 \mu\text{g}$ BNT162b2 (Pfizer–BioNTech) vaccine in February 2021, were enrolled, and the correlates of neutralization activity represented by 50% neutralization titers (NT_{50}) determined by employing the target living VeroE6_{TMPRSS2} cells and live SARS-CoV-2 with ages, adverse effects (AEs) that occur often such as injection-site pain and systemic fever were examined. The kinetics of NT_{50} values and S1-binding antibody levels were also examined. There was a significant rise in the NT_{50} values as determined on day 28 post-1st shot (a week after post 2nd shot) compared to those on day 7 post-1st shot. Correlation was negligible between NT_{50} values and ages or systemic fever grades. In this regard, most adverse effects that occur within 1–3 days following vaccine shots are thought to be caused by the release of certain pyrogenic and inflammatory cytokines (*e.g.*, interleukin-1, interleukin-6, and tumor-necrosis factor) from antigen-presenting cells (APCs) such as macrophages and dendritic cells when they ingest and process the exogenous antigens (*i.e.*, SARS-CoV-2 spike antigens) and transmit the antigenic information to relevant immune cells. Such early-phase defense events include response to antigenic determinants irrelevant to neutralizing activity but those

eliciting S1-binding antibodies. The release of the pyrogenic and inflammatory cytokines mostly subsides within days following the vaccine shot. The released cytokines activate the antigen-specific-antibody-producing B-cells, which respond to the processed antigenic determinants presented by the APCs and start to produce antigen-specific antibodies such as neutralizing antibodies as well as non-neutralizing but S1-binding antibodies. Such antigen-specifically-activated and antibody-producing B-cells continue to produce antibodies. In the case of BNT162b2 vaccination, it appears that it takes 10 to 12 days from the 1st vaccine shot for the vaccinated individuals to achieve the amounts of neutralizing antibodies that are enough to block the infection of substantial numbers of the virally-targeted cells and to inhibit further spread of the infection^{9,10}. It is thought that the release of pyrogenic and inflammatory cytokines and the build-up of the protective antibody levels are different events, occurring chronologically ~ 10–12 days apart. These two different events appear to have resulted in the absence of significant correlates between NT₅₀ levels and AEs examined in the present study.

In the present study, the NT₅₀ values had a substantial correlation with S1-binding-IgG levels but had only moderate correlation with S1-binding IgM levels, suggesting that the major neutralizing activity resides within the S1-binding IgG fraction. Interestingly, the approximate half-life of NT₅₀ values (67.8 days) and that of S1-binding-IgG levels (53.5 days) were reasonably close to each other, corroborating the assumption of the presence of the major fraction of neutralizing antibodies within IgG fraction. In human body, IgG has concentration-dependent half-life of approximately 21 days and IgM around 5–6 days²⁴. By contrast, the half-lives of NT₅₀, S1-binding-IgG, and -IgM levels determined in the present study were much longer with 43.6–67.8 days. This discrepancy is perhaps attributed to the persistence of continuously-antibody-producing B cells over weeks or months in the body of the participants following the vaccination^{25,26}, thereby the half-lives of neutralizing activity and S1-binding-IgG and -IgM levels have been extended as compared to the physiological half-lives of IgG and IgM. Of note, the assumption of the half-lives in the present work was based on the chronologically linear nature of the reduction observed during the present study period (Fig. 3D). Recently, Doria-Rose et al. reported that anti-SARS-CoV-2 antibodies persist through 6 months after the second dose of mRNA-1273 administration²⁷. However, the study was of a relatively small scale (n = 33), and more definite data are needed for constructing more protective measures.

There is a growing body of evidence that COVID-19 results in more severe symptoms and greater mortality among men than among women^{28,29}. A cohort study of 17 million adults in England has revealed a strong association between male sex and the risk of death from COVID-19 (hazard ratio 1.59, 95% confidence interval 1.53–1.65)³⁰. In the present data set, significantly greater levels of NT₅₀, S1-binding-IgG and -IgM were documented in women than in men when examined on 28, 60, and 90 days post-1st shot, while there was no difference in either of NT₅₀, S1-binding-IgG or -IgM levels on day 7 post-1st dose (Supplementary Figs. 4A-C). These results apparently relate to the findings by others reporting that women, in general, have more robust ability to control infectious pathogens (*i.e.*, SARS-CoV-2) than men^{30,31}. Indeed, there is increasing evidence indicating strong correlation between SARS-CoV-2-

neutralizing antibody titers and clinical efficacy, suggesting that a neutralizing antibody response provides the primary contribution to protection against COVID-19³² and that the presence of high levels of neutralizing antibody is largely sufficient for protection against SARS-CoV-2 infection and clinical onset upon exposure to the virus^{33,34}. In fact, Imai *et al.* have reported that the administration of convalescent plasma from previously-SARS-CoV-2-infected hamsters completely protected newly SARS-CoV-2-exposed hamsters from contracting viral pneumonitis³⁵. Thus, the greater neutralizing activity in women than in men observed in the present study can contribute at least in part to the gender differences in COVID-19 disease outcomes.

We also examined how the BNT162b2-elicited neutralizing antibodies blocked the infectivity and cytopathic effect of five different variants of concerns in the cell-based assays using various infectious variants (one Wuhan strain, two alpha strains, one strain each of beta, delta and kappa strains). Six selected serums from elite responders showed quite potent activity to the alpha, kappa, and delta variants, while they exerted relatively moderate activity against the beta strain (Fig. 4). On the other hand, one of the randomly-selected 12 serums from moderate responders showed a marginal activity (NT₅₀ value of 40-fold) and two of them failed to show detectable activity (NT₅₀ values < 20-fold) against the beta strain (Fig. 4). These data suggest that BNT162b2-receiving vaccinees who develop high magnitudes of neutralizing antibody should probably be well protected against the infection by most variants; however, those who develop only low levels of neutralizing antibody may be vulnerable to the infection by certain variants such as beta strains. If so, to such low-responders to BNT162b2 even after the 2nd shot, an additional 3rd shot may be needed. If the 3rd dose of the same vaccine fails to elicit good levels of neutralizing antibodies, new types of vaccines with different platform have to be stratified.

Methods

Participants and serum specimens.

Serum samples were collected from 225 vaccinated health care workers at JCHO Kumamoto General Hospital (Kumamoto, Japan). All the 225 participants were of Japanese citizen. Serum samples were analyzed at the National Center for Global Health and Medicine (NCGM) in Tokyo. The Ethics Committees from the Kumamoto General Hospital and NCGM approved this study (Kumamoto General Hospital No. 180, and NCGM-G-004176-00, respectively). Each participant provided a written informed consent, and this study abided by the Declaration of Helsinki principles. The vaccination (on days 0 and 21) and serum collection (from day 7 through day 90 post-1st shot) were conducted as shown in Table 1.

Cells and viruses.

VeroE6_{TM_{PRSS2}} cells²³ were obtained from Japanese Collection of Research Bioresources (JCRB) Cell Bank (Osaka, Japan). VeroE6_{TM_{PRSS2}} cells were maintained in DMEM supplemented with 10% FCS, 100 µg/ml of penicillin, 100 µg/ml of streptomycin, and 1 mg/mL of G418. SARS-CoV-2 NCGM-05-2N strain (SARS-CoV-2_{05-2N}) was isolated from nasopharyngeal swabs of a patient with COVID-19, who was

admitted to the NCGM hospital^{36,37}. Five clinically isolated SARS-CoV-2 mutant strains were used in the current study: two B.1.1.7 (alpha) strains [hCoV-19/Japan/QHN001/2020 (SARS-CoV-2_{QHN001}, GISAID Accession ID; EPI_ISL_804007) and hCoV-19/Japan/QK002/2020 (SARS-CoV-2_{QK002}, GISAID Accession ID; EPI_ISL_768526)] and a B.1.351 (beta) strain [hCoV-19/Japan/TY8-612-P0/2021 (SARS-CoV-2_{TY8-612})] were obtained from National Institute of Infectious Diseases, Tokyo, Japan. A B.1.617.1 (kappa) strain [TKYTK5356_2021 (SARS-CoV-2₅₃₅₆, DDBJ Accession ID; LC633761)] and a B.1.617.2 (delta) strain [hCoV-19/Japan/TKYK01734/2021 (SARS-CoV-2₁₇₃₄, GISAID Accession ID; EPI_ISL_2080609)] were provided from Tokyo Metropolitan Institute of public Health, Tokyo, Japan. Each variant was confirmed to contain each VOC-specific amino acid substitutions before the assays conducted in the present study (*vide infra*).

Neutralization assay.

The neutralizing activity of serums from vaccinated individuals was determined by quantifying the serum-mediated suppression of the cytopathic effect (CPE) of each SARS-CoV-2 strain in VeroE6_{TMPRSS2} cells as previously described with minor modifications³⁶. In brief, each serum was 4-fold serially diluted in culture medium. The diluted sera were incubated with 100 50% tissue culture infectious dose (TCID₅₀) of viruses at 37°C for 20 min (final serum dilution range of 1:20 to 1:4000), after which the serum-virus mixtures were inoculated to VeroE6_{TMPRSS2} cells (1.0 x 10⁴/well) in 96-well plates. For SARS-CoV-2 strains used in this assay are as follows: a wild-type strain, SARS-CoV-2_{05-2N} (PANGO lineage B)^{36,37}, two alpha variants (SARS-CoV-2_{QHN001} and SARS-CoV-2_{QK002}), a beta variant SARS-CoV-2_{TY8-612}, a delta variant SARS-CoV-2₁₇₃₄, and a kappa variant SARS-CoV-2₅₃₃₆. After culturing the cells for 3 days, the levels of CPE observed in SARS-CoV-2-exposed cells were determined using the WST-8 assay employing Cell Counting Kit-8 (Dojindo, Kumamoto, Japan). The serum dilution that gave 50% inhibition of CPE was defined as the 50% neutralization titer (NT₅₀). Each serum was tested in duplicate.

Measurement of anti-SARS-CoV-2 antibody titers.

Measurement of 3 anti-SARS-CoV-2 antibody levels (anti-S1-IgG, anti-S1-IgM, and anti-N-IgG) in each participant was performed using the chemiluminescence enzyme immunoassay (CLEIA) platform (HISCL) manufactured by Sysmex Co. (Kobe, Japan) as previously reported²¹.

Statistical analyses.

Out of the 225 participants, one participant, who had been infected by SARS-CoV-2 with PCR positivity documented was primarily excluded. Demographic characteristics of the participants are described in Table 1. Correlates of neutralizing activity levels with S1-binding-IgG and -IgM levels, ages, genders, pain scores in the injection-site, and systemic fever up to 40°C were examined by Spearman rank correlation coefficient. Also, neutralizing activity levels, S1-binding-IgG, and -IgM, pain scores, systemic fever were compared between genders using the Wilcoxon rank sum test. As for participants with normal fever, their

temperature was treated as 36.89 degree, a normal body temperature in Japanese³⁸. Percentage of the adverse events reported in writing following the 1st and 2nd dose administration were determined and assessed in regard to gender. The differences in neutralization activity between each measurement time point were tested by the Wilcoxon rank sum test, and were assessed among categorized age subgroups. Similarly, difference of S1-binding-IgG and -IgM levels among time points were tested. Decline of neutralizing activity over 90 days post-1st shot was assessed using the mixed-effects model including time as a main effect and intercept as a random effect. Also, the prediction slope and its 80% prediction interval were generated by drawing a sampling distribution for the fixed effects and then estimating the fitted value across that distribution. The merTools package (version 0.5.2) in R software was used for prediction. The decline for S1-binding-IgG and -IgM levels was assessed similarly. All the analyses were conducted with the use of R, version 4.1.0 (R Foundation for Statistical Computing).

Declarations

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Author Contributions

KM and HM had access to all data in this study and took and hold all responsibility for the integrity of the data and the accuracy of the data analysis. KM and HM: Concept and design. KM, MA, KT, TM, KN, YT, HG, KY, SO, TS, and TY: Acquisition, analysis, or interpretation of data. YU, YS, AF, and HM: Statistical analysis. AF, YI, HN, MK, SM, AM, WS, and SS: Administrative, technical, or material support. KM and HM: Original draft writing. All authors: Reading and writing manuscript.

Completing Interests

Matsushima, Noda, Sato, and Yoshida are employees of Sysmex Corporation. Other authors declare that there is no conflict of interest.

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Figures

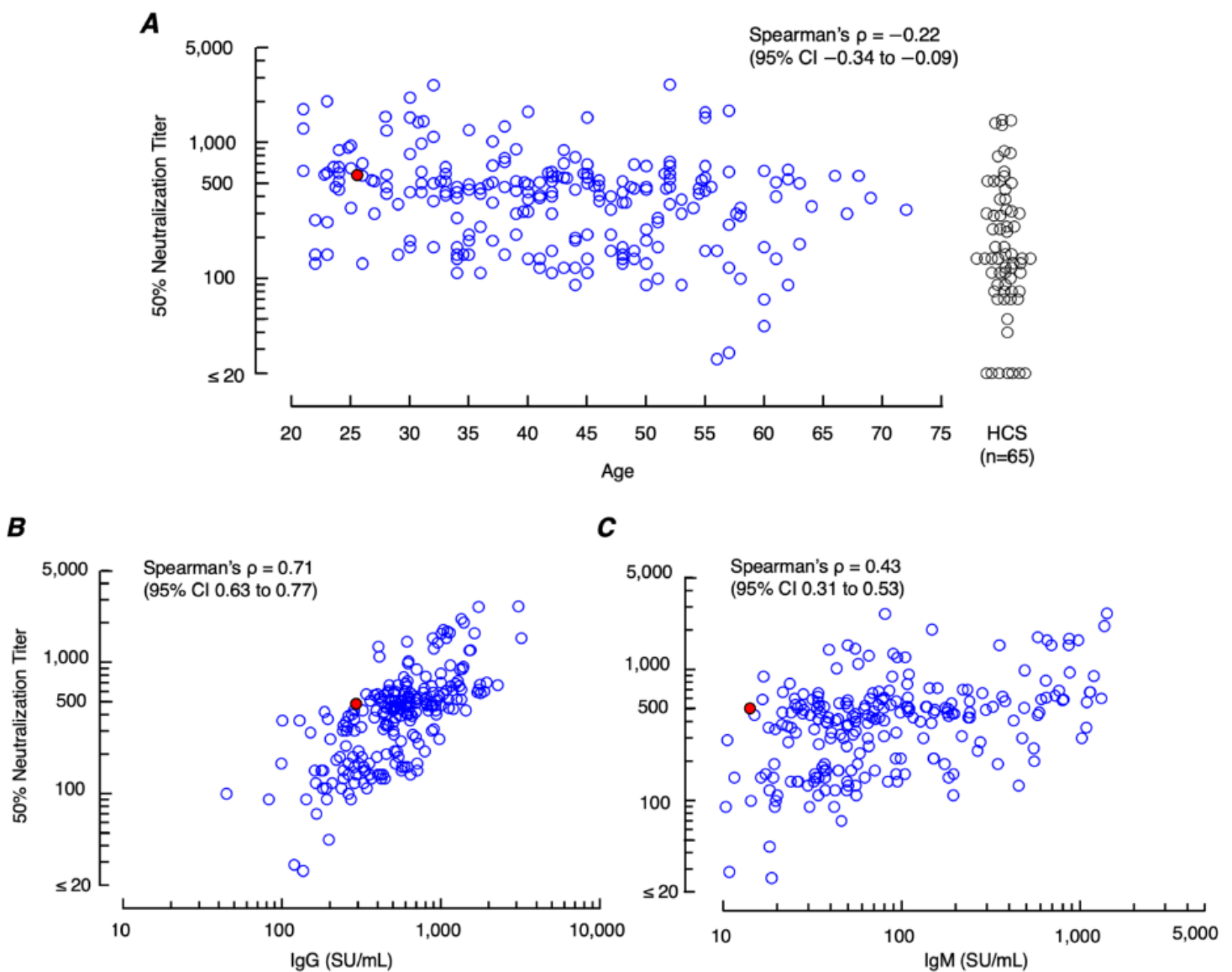


Figure 1

Correlations of neutralizing titers with ages and S1-binding-IgG and -IgM levels. A. Correlation between neutralizing titers (NT50s) and ages (on day 28 post-1st shot). The age range of the study participants was 21 to 72 (average 41.8 y.o.). A correlation is negligible between NT50 values and ages (Spearman's $\rho = -0.22$; 95% CI -0.34 to -0.09). The geometric mean NT50 of the values from all participants ($n = 225$) was 375.2-fold (range 25.6-2,680-fold), greater by a factor of 2.3 than the geometric mean NT50 from 65 COVID-19-convalescent patients (geometric mean = 163.0-fold; range 20.0-1470-fold) shown as references on the far right (human COVID-19-convalescent serum: HCS). B. A high correlation is identified (Spearman's $\rho = 0.71$; 95% CI 0.63 to 0.77) between NT50 values and S1-binding-IgG levels in samples obtained on day 28 post-1st dose. C. Moderate correlation is seen between neutralizing titers and S1-binding-IgM levels (Spearman's $\rho = 0.43$; 95% CI 0.31 to 0.53). One participant, who had been infected with SARS-CoV-2 with PCR-positivity documented, is indicated as a solid-red solitary circle. This participant was excluded from all analyses at later timepoints.

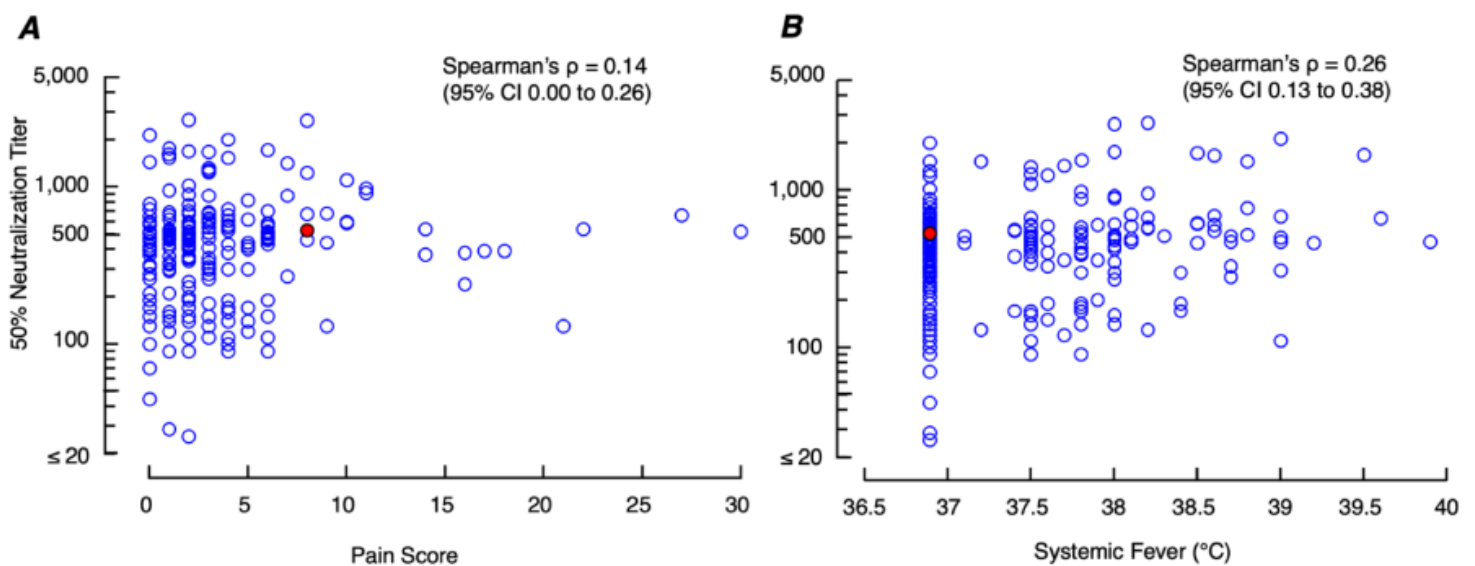


Figure 2

Correlations of neutralizing titers with injection-site pain scores and systemic fever grades. A. No correlation was seen between NT50 values and injection-site pain (Spearman's $\rho = 0.14$; 95% CI 0.00 to 0.26). The injection-site pain following the 2nd BNT162b2 dose was scored by using the short-form McGill Pain Questionnaire22. B. Correlation was negligible between NT50 values and systemic fever grades (Spearman's $\rho = 0.26$; 95% CI 0.13 to 0.38). A solid-red circle indicates a person with previous SARS-CoV-2 infection documented.

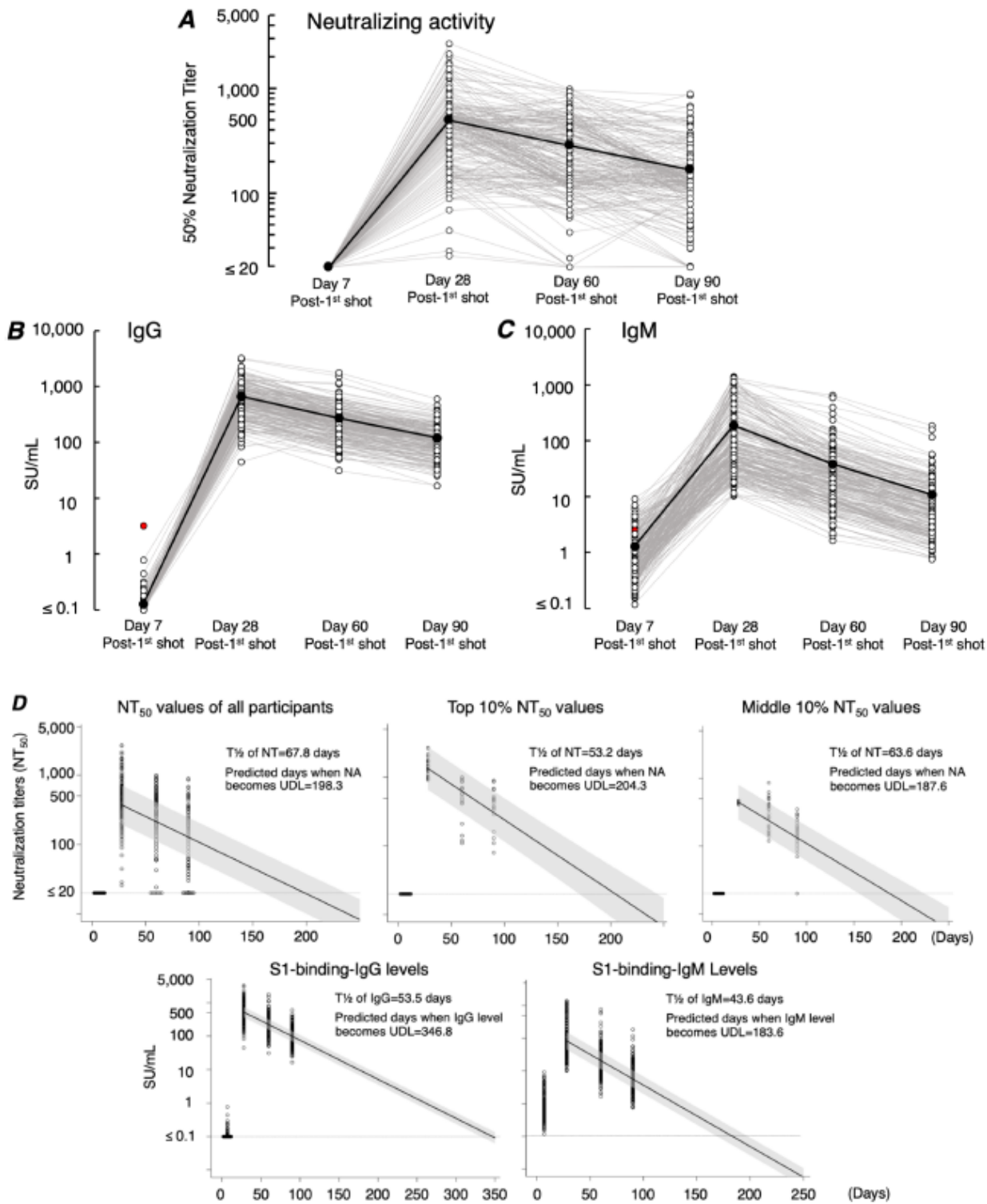


Figure 3

Kinetics of neutralizing activity and S1-binding-IgG and -IgM levels. Time-course analyses of neutralizing activity for 90 days were conducted. The 1st vaccine was administered on day 0, and the 2nd vaccine on day 21. Blood samples from vaccinated individuals were obtained on days 7, 28, 60, and 90 post 1st shot as illustrated in Table 1. A. Neutralizing activity is shown as NT50 (50% neutralizing titer). The NT50 value of 20-fold is the detection limit and values determined to be less than 20-fold were treated as 20-

fold. B and C. Kinetics of S1-binding-IgG and -IgM levels are shown. The average values of each data point are shown in black solid circles, which are connected with solid black lines. One participant, who had been infected with SARS-CoV-2 with PCR-positivity documented, is indicated as a solid-red solitary circle in B and C. This participant was excluded from all analyses at later timepoints. D. Decline of neutralizing activity, S1-binding-IgG and -IgM over 90 days post-1st shot. The solid-black lines consist of predicted values estimated by mixed effects model, and the shaded areas denote corresponding 80% prediction intervals. The dashed horizontal lines in the upper three panels denote the NT50 detection limit of 20-fold. NT50 values determined to be less than 20-fold were treated as 20-fold. The lowest detection limit for S1-binding-IgG and -IgM quantification shown as dashed horizontal lines in the two lower panels was 0.1 SU/ml and the values lower than 0.1 SU/ml were calculated as 0.1 SU/ml.

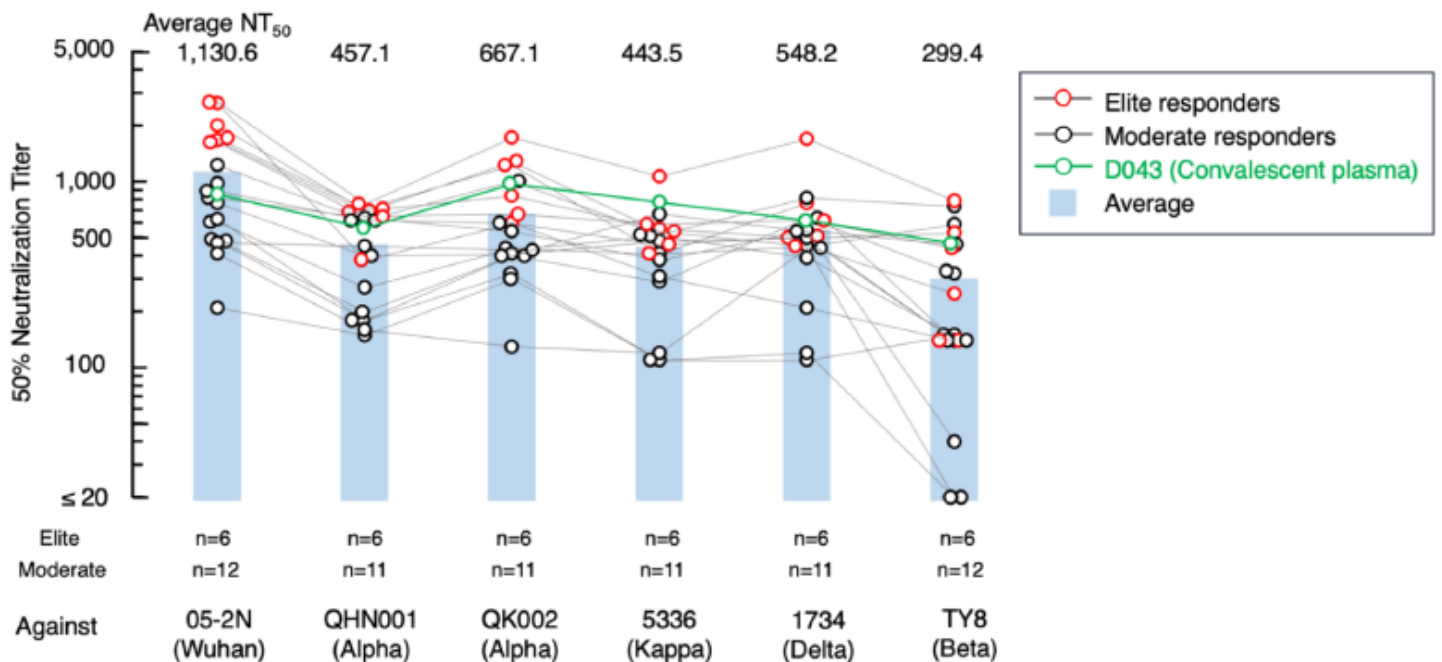


Figure 4

Blockade of the infectivity and replication of SARS-CoV-2 variants by vaccinees' serums. The activity of vaccinees' serums to block the infectivity and replication of 5 SARS-CoV-2 variants (alpha variants: SARS-CoV-2QHN001 and SARS-CoV-2QK002; a beta strain: SARS-CoV-2TY8-612; a delta strain: SARS-CoV-21734; and a kappa strain: SARS-CoV-25356) was evaluated. A Wuhan strain SARS-CoV-205-2N 37 was employed as a reference SARS-CoV-2. Six serums were from elite responders (NT50 >1,500-fold) and 12 serums were from randomly-selected moderate responders (NT50=200~1,500-fold). The NT50 titers of each serum against 6 SARS-CoV-2 strains are shown in red circles (for 6 elite responders) and in black circles (for 12 moderate responders). D043 is a serum from a COVID-19-convalescent patient 39 and served as an internal control in the assays.

Supplementary Files

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- [MaedaetalSupplementaryfigures.pdf](#)