

COVID-19 Vaccination Response in Participants Receiving Efgartigimod IV or Efgartigimod PH20 SC in ADAPT+ or ADAPT-SC+

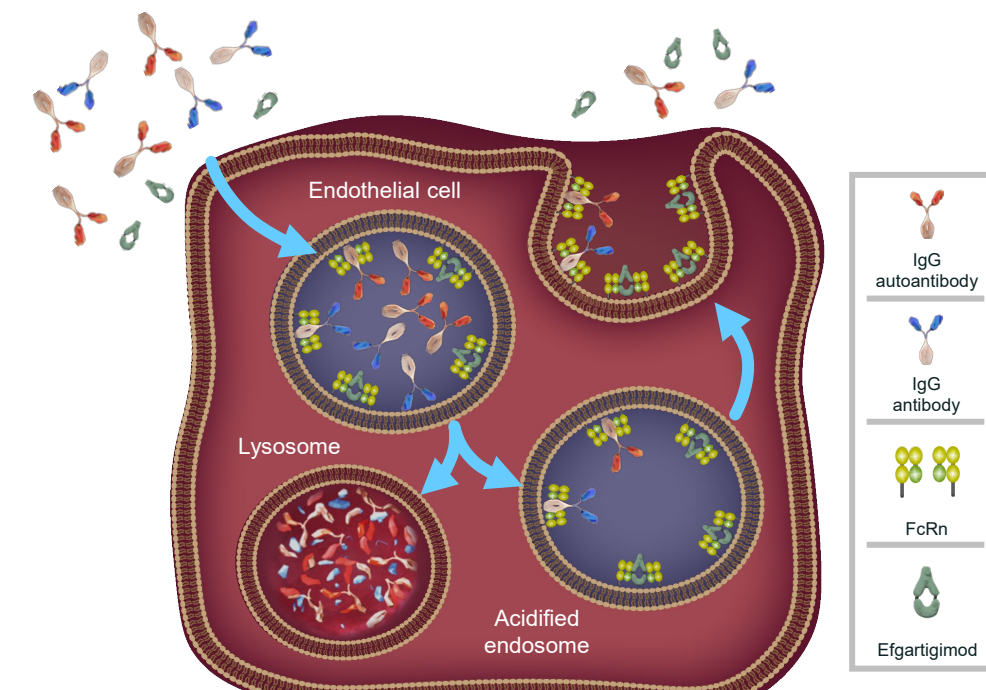
Tuan Vu,¹ Francesco Saccà,² James F. Howard Jr,³ John W. Sleasman,⁴ Fien Gistelincx,⁵ Paul Duncombe,⁵ Benjamin Van Hoorick,⁵ Sophie Steeland,⁵ Renato Mantegazza,⁶ Jan L. De Bleeker,⁷ Antoine Azar,⁸ Kevin Winthrop⁹

¹Department of Neurology, University of South Florida Morsani College of Medicine, Tampa, Florida, USA; ²GENESIS Department, Federico II University of Naples, Naples, Italy; ³Department of Neurology, The University of North Carolina, Chapel Hill, North Carolina, USA; ⁴Duke University School of Medicine, Division of Allergy, Immunology, and Pulmonary Medicine, Durham, North Carolina, USA; ⁵argenx, Ghent, Belgium; ⁶Department of Neuroimmunology and Neuromuscular Diseases, Fondazione IRCCS Istituto Neurologico Carlo Besta, Milan, Italy; ⁷Department of Neurology, Ghent University Hospital, Ghent, Belgium; ⁸Johns Hopkins University School of Medicine, Division of Allergy and Clinical Immunology, Baltimore, Maryland, USA; ⁹Oregon Health and Science University, Division of Infectious Disease, Portland, Oregon, USA



INTRODUCTION

Efgartigimod Mechanism of Action: Blocking FcRn



- FcRn recycles IgG antibodies and albumin. This recycling and salvage from lysosomal degradation results in IgG antibodies having the longest half-life and being the most abundant of all immunoglobulins in the plasma¹⁻³
- Efgartigimod is an IgG1 antibody Fc fragment that has been engineered for increased affinity to FcRn, and is uniquely composed of the only part of the IgG antibody that normally binds FcRn¹
- Efgartigimod selectively reduces IgG by blocking FcRn-mediated IgG recycling without impacting antibody production, albumin levels, or other parts of the immune system^{1,4,5}
- Efgartigimod PH20 SC is a coformulation of efgartigimod and recombinant human hyaluronidase PH20 (rHuPH20), which allows for rapid SC administration of larger volumes^{6,7}
 - PK/PD modeling and phase 3 data (ADAPT-SC) have demonstrated 4 once-weekly administrations of 1000 mg efgartigimod PH20 SC and 10 mg/kg efgartigimod IV result in comparable decreases in IgG levels^{6,8}

Some immunosuppressive therapies used in the treatment of gMG may increase susceptibility to infections and impair immune response to vaccines⁹

- Glucocorticoids, mycophenolate mofetil, and B-cell-depleting therapies can substantially reduce immunogenicity of mRNA vaccines to SARS-CoV-2^{10,11}
- In previous studies of participants with IgG-mediated autoimmune diseases efgartigimod did not impair generation of IgG responses to antigenic challenges, and levels of both naturally- and vaccine-induced protective antibody titers closely followed total IgG reduction kinetics⁵

RESULTS

Table 1. Baseline Demographics of Participants in ADAPT+/ADAPT-SC+ Receiving COVID-19 Vaccines^a

Characteristic	Participants (N=68)
Age, y, mean (SD)	49.0 (14.2)
Age category, n (%)	
18-64 y	55 (80.9)
65-74 y	11 (16.2)
≥75 y	2 (2.9)
Sex at birth, n (%)	
Female	44 (64.7)
Male	24 (35.3)
BMI ^b (kg/m), mean (SD)	28.6 (8.0)
Race, n (%)	
Asian	9 (13.2)
White	57 (83.8)
Multiple	2 (2.9)

Table 2. COVID-19 Vaccine Received by Participants in ADAPT+/ADAPT-SC+^a

COVID-19 Vaccine, n (%)	Participants (N=68)
Oxford–AstraZeneca	1 (1.5)
Janssen	1 (1.5)
Spikevax (Moderna)	12 (17.6)
Sputnik V	3 (4.4)
Pfizer-BioNTech	48 (70.6)
Unknown	7 (10.3)

Table 3. Concomitant MG Therapies^a

Therapy, n (%)	Participants (N=68)
AChEi	56 (82.4)
NSISTS	42 (61.8)
Steroids	45 (66.2)

^aParticipant data are included only for those who had a prevaccination titer sample and ≥1 postvaccination titer sample available. ^bBMI data were unavailable for 2 individuals.

ABBREVIATIONS

AChEi, acetylcholinesterase inhibitor; BMI, body mass index; COVID-19, coronavirus disease 2019; ELISA, enzyme-linked immunosorbent assay; EFG, efgartigimod; Fc, fragment crystallizable region; FcRn, neonatal Fc receptor; gMG, generalized myasthenia gravis; Ig, immunoglobulin; IV, intravenous; LLOQ, lower limit of quantification; MG, myasthenia gravis; mRNA, messenger RNA; -N, nucleocapsid protein; NSISTS, nonsteroidal immunosuppressive therapy; PD, pharmacodynamic; PK, pharmacokinetic; P-V, prevaccination; -RBD, receptor-binding domain of S protein; rHuPH20, recombinant human hyaluronidase PH20; SARS-CoV-2, severe acute respiratory syndrome coronavirus 2; -S, spike protein; SARS-CoV-2-IgG, severe acute respiratory syndrome coronavirus 2 specific IgG; SC, subcutaneous; V, vaccination; V1, first vaccination; V2, second vaccination; V3, third vaccination.

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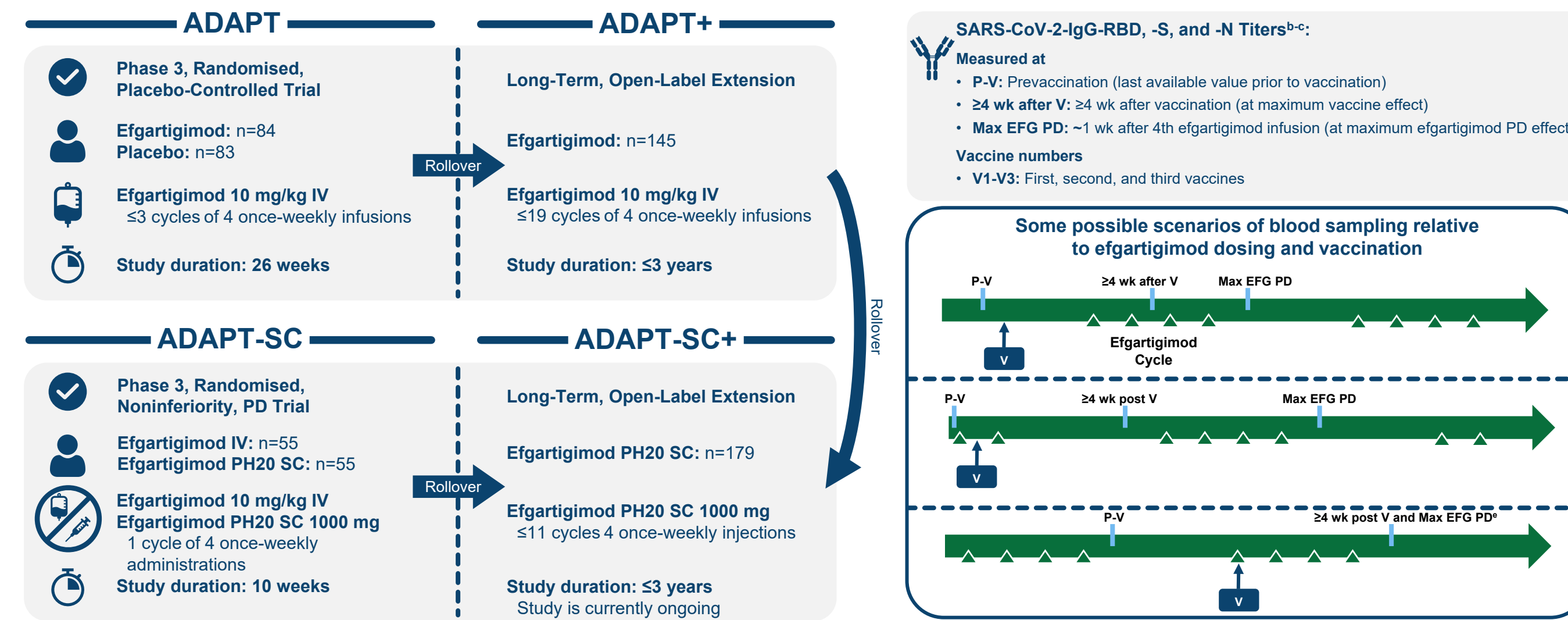
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ACKNOWLEDGMENTS AND DISCLOSURES: The authors gratefully acknowledge the ADAPT+/ADAPT-SC+/ADAPT-SC+ trial participants and investigators. **TV:** Alexion, argenx, CSL Behring, Novartis/AbbVie, AstraZeneca, Ra/UCB, Horizon, Regeneron, Janssen/Momenta, Immunovant, Cartesian Therapeutics, and Sanofi. **FS:** Alexion, Horizon, Mylan, Novartis, Roche, Sanofi, Teva, Almirall, argenx, Avexis, Forward, Leveo, Merck, Novartis, Pomona, Takeda, and Prilexia. **JFH:** Ad Scintiam, Alexion AstraZeneca Rare Disease, argenx, Cartesian, Centers for Disease Control and Prevention, MGFA, Muscular Dystrophy Association, NIH, PCORI, UCB Pharma Academic/CME, Alexion AstraZeneca Rare Disease, argenx, Biologic, CheckRare CME, Hoffmann-La Roche, Horizon Therapeutics plc (now Amgen), Medscape CME, Merck EMD Serono, NMD, Novartis, PeerView CME, Physicians' Education Resource (PER) CME, Platform CME, Regeneron, Sanofi, Zai Labs, and Tolerancia AB. **JWS:** National Institutes of Health, Collective, Enzyvant, Jeffrey Modell Foundation, and argenx. **SS, FG, PD, and BVH:** argenx. **RM:** Alexion, argenx, Ra, Biomarin, Catalyst, UCB, TEVA, Merck, Roche, and Biogen. **JLDB:** argenx, Alexion, CSL, UCB, Alnylam, Janssen, and Sanofi. **AA:** X4, Grifols, Takeda, Pfizer, Janssen, and argenx. **KW:** Bristol Myers Squibb, Pfizer, AbbVie, UCB, Eli Lilly, Galapagos, GSK, Roche, Gilead, Regeneron, Sanofi, AstraZeneca, and Novartis. The ADAPT+ and ADAPT-SC+ trials were funded by argenx. Medical writing and editorial support for this presentation were provided by Precision AQ and funded by argenx.

METHODS

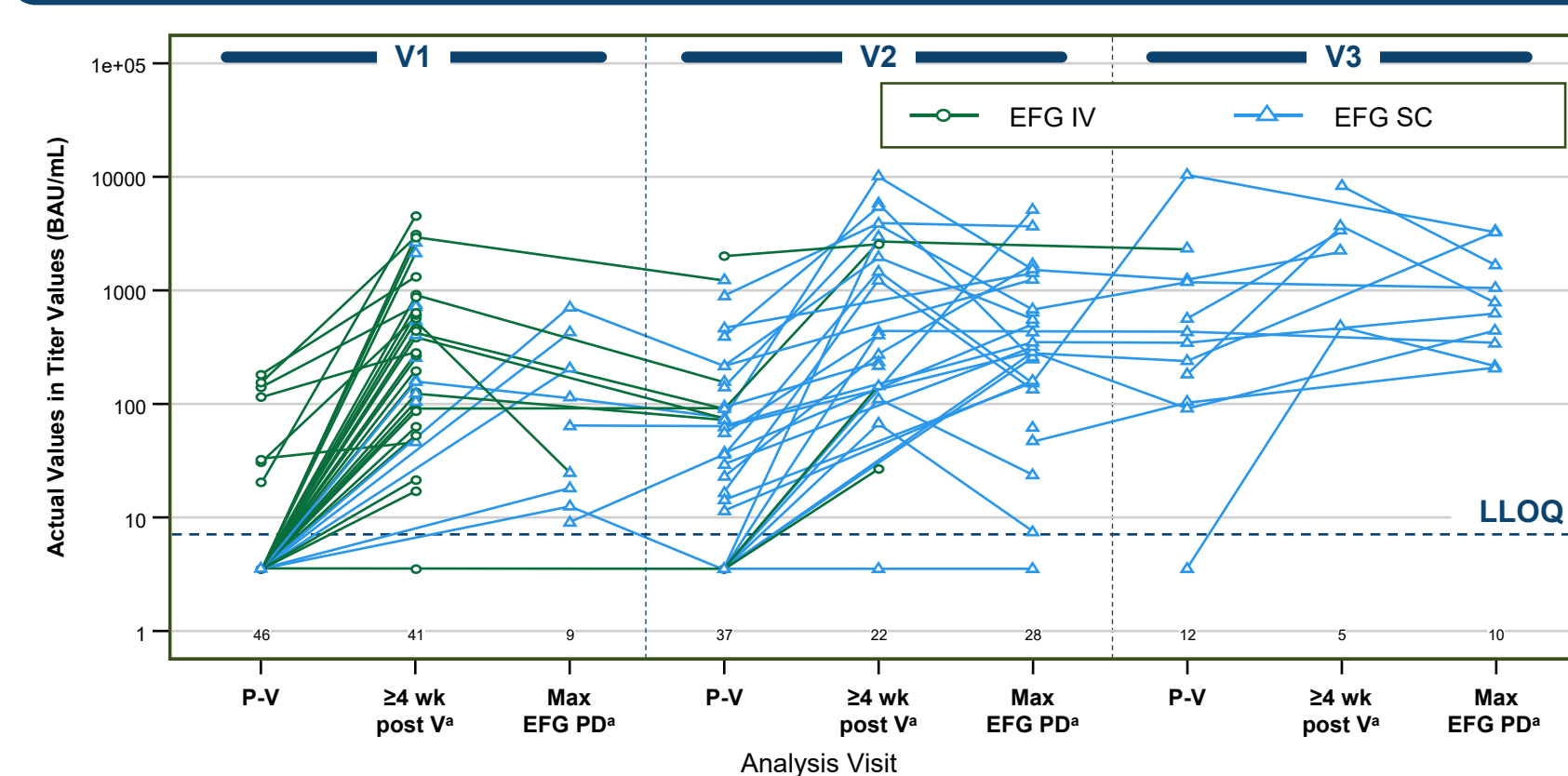
OBJECTIVE

Assess the effects of efgartigimod IV and efgartigimod PH20 SC treatment on humoral immune responses to COVID-19 vaccination in participants with gMG during ADAPT+ and ADAPT-SC+^a



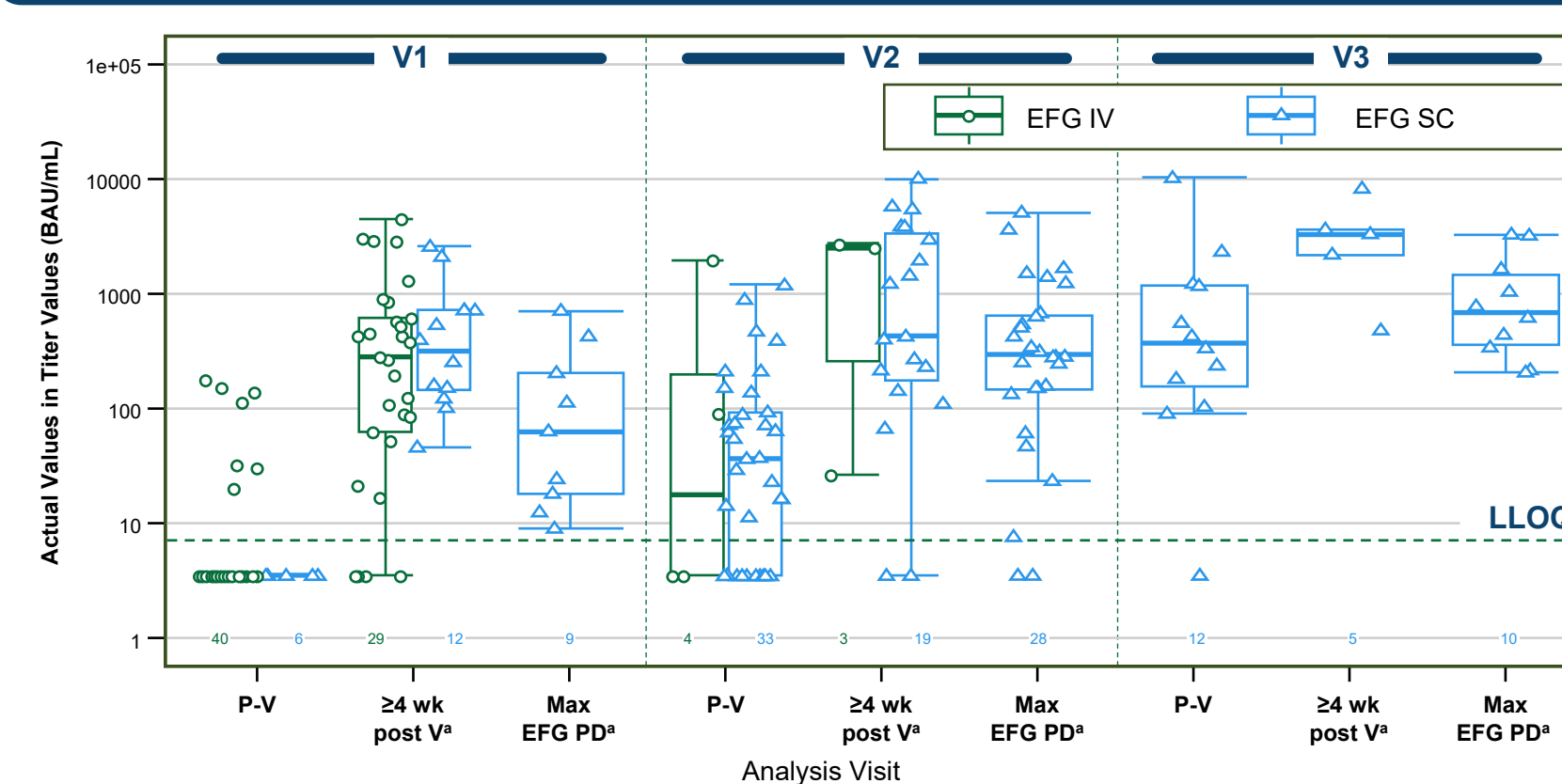
^aNot all participants enrolled in the ADAPT+ or ADAPT-SC+ received vaccines targeting COVID-19. ^bSARS-CoV-2-IgG-RBD, -S, and -N antibodies were assessed using an ELISA assay. ^cFor COVID-19 vaccinations consisting of multiple injections (ie, injections within 45 days of each other), only blood samples before the first injection and after the last injection were selected. ^dSamples were collected from the first cycle that occurred after the vaccination. ^eOne sample was collected if postvaccination time points (24 weeks after V at maximum vaccine effect and Max EFG PD) coincided with each other.

Figure 1. Individual Trend of Actual SARS-CoV-2-IgG-RBD Titer Values Over Time (Combined ADAPT+/ADAPT-SC+)



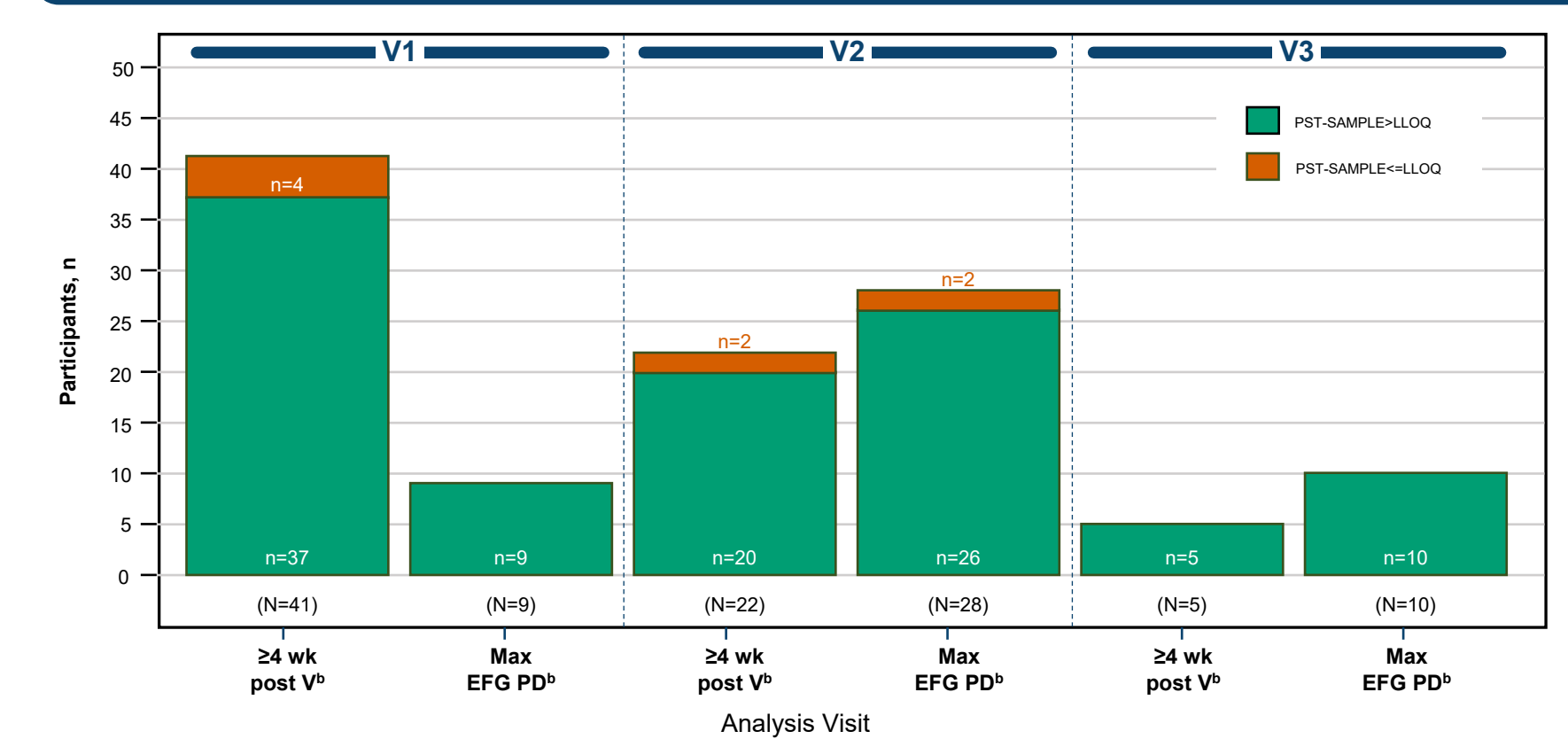
[†]If postvaccination timepoints (24 weeks after V and Max EFG PD) coincided with each other, the sample is presented at Max EFG PD timepoint.

Figure 2. Individual SARS-CoV-2-IgG-RBD Titer Values With Boxplots Over Time (Combined ADAPT+/ADAPT-SC+)



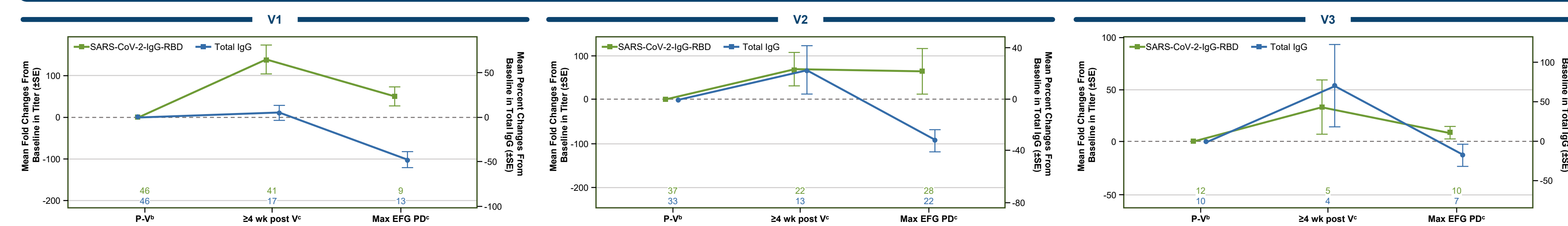
[†]If postvaccination timepoints (24 weeks after V and Max EFG PD) coincided with each other, the sample is presented at Max EFG PD timepoint.

Figure 3. Seroconversion^a of SARS-CoV-2-IgG-RBD Over Time (Combined ADAPT+/ADAPT-SC+)



^aSeroconversion was defined as the percentage of participants with SARS-CoV-2-IgG-RBD levels above the LLOQ. [†]If postvaccination timepoints (24 weeks after V and Max EFG PD) coincided with each other, the sample is presented at Max EFG PD timepoint.

Figure 4. Mean Fold Changes in SARS-CoV-2-IgG-RBD Titer Values and Mean Percent Changes in Total IgG Values Over Time (Combined ADAPT+/ADAPT-SC+)^a



^aMean fold change from baseline was imputed with a "1" and the mean percent changes from baseline with a "0" at P-V. [†]Includes all participants with baseline values available. ^{††}If postvaccination timepoints (24 wk post V and Max EFG PD) coincided with each other, the sample is presented at Max EFG PD timepoint.

SARS-CoV-2-IgG-S and SARS-CoV-2-IgG-N Titer Values

- SARS-CoV-2-IgG-S titer values showed a similar trend in response to vaccination as SARS-CoV-2-IgG-RBD titer values
- There was no effect of vaccination on SARS-CoV-2-N titer values (positive SARS-CoV-2-N titer values indicate a previous/resolving COVID-19 infection)¹²

