

Assessing Relationship Between Immunoglobulin G Level and Nipocalimab Efficacy Using Myasthenia Gravis-Activities of Daily Living Scale



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Introduction

- Generalized myasthenia gravis (gMG), a rare chronic condition, is characterized by pathogenic immunoglobulin G (IgG) autoantibody-mediated impairment of neurotransmission leading to fatigable muscle weakness¹²
- Nipocalimab, a fully human monoclonal antibody, binds the neonatal Fc receptor (FcRn) with high affinity, inhibiting IgG recycling and thereby reducing circulating total IgG, including pathogenic IgG autoantibodies^{3,4}
- Nipocalimab is approved for the treatment of gMG in anti-acetylcholine receptor (AChR) and anti-muscle-specific tyrosine kinase (MuSK)-antibody-positive adult and adolescent patients in the United States and the EU^{5,6}
 - In the phase 3 Vivacity-MG3 study (NCT04951622), nipocalimab added to standard-of-care (SOC) demonstrated substantial reduction in circulating IgG with rapid and sustained disease control over 24-weeks in a population of seropositive patients with gMG⁷
- Total serum IgG serves as a direct pharmacodynamic marker and an indirect marker of clinical improvement for anti-FcRn treatments in gMG and development of pharmacometrics model quantifying this relationship could help assess between-patient variability and evaluate whether dose adjustments are needed in specific patient subgroups⁸

Objectives

 To characterize the longitudinal relationship between reduction in total serum IgG level and clinical efficacy, measured by change from baseline in Myasthenia Gravis Activities of Daily Living (MG-ADL) score, using semi-mechanistic pharmacometrics modeling

Methods

Data

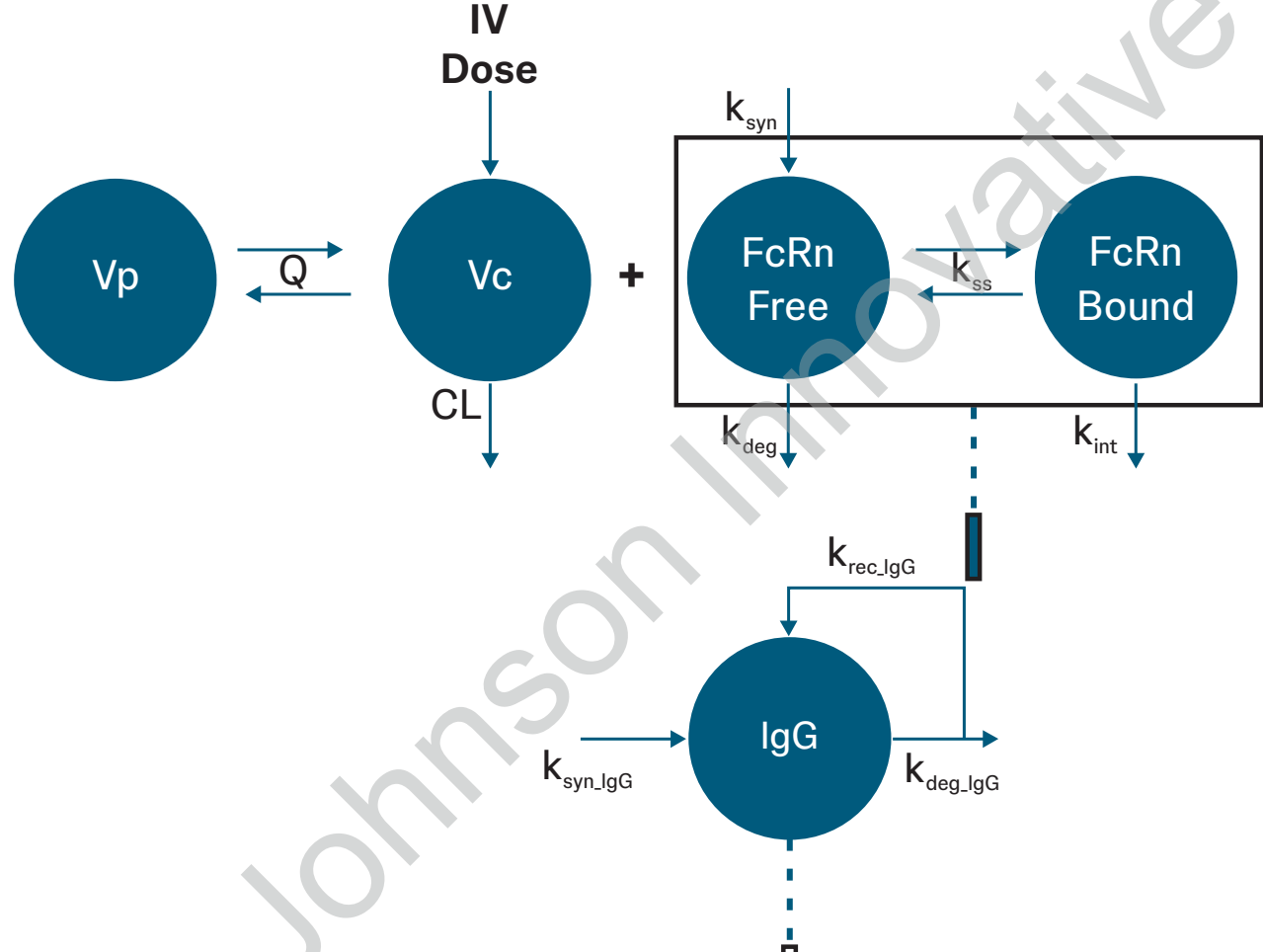
- Data were pooled from 7 clinical studies of nipocalimab
 - 5 phase 1 studies in healthy volunteers
 - 1 phase 2 study (Vivacity-MG) in patients with gMG⁹
 - 1 phase 3 study (Vivacity-MG3)⁷ in patients with gMG: total of 196 patients, of which 153 were seropositive (anti-AChR+, anti-MuSK+ and/or anti-LRP4+)

Dataset	Sample size
Serum nipocalimab concentrations	3,429 samples (n=277)
FcRn receptor occupancy	1,247 samples (n=78)
Total serum IgG concentrations	4,441 samples (n=421)
MG-ADL change from baseline	2,317 observations (n=220)

Pharmacokinetic/Pharmacodynamic Modeling

- A previously developed nonlinear mixed-effects model was used (NONMEM v7.4.3)¹⁰
- Investigated nipocalimab intravenous (IV) dose regimens:
 - 0.3, 3, 10, 15, 30, 45, 60 mg/kg, 300 mg, and 1200 mg single dose; 15 mg/kg once weekly (QW) and 30 mg/kg QW in healthy volunteers; 5 mg/kg every 4 weeks (Q4W), 30 mg/kg Q4W, 60 mg/kg single dose, and 60 mg/kg every 2 weeks (Q2W) in the phase 2 study
 - 30 mg/kg loading dose followed by 15 mg/kg Q2W in the phase 3 study

Figure 1. Schematic of the nipocalimab PK/RO/IgG/MG-ADL model¹⁰



$MG\text{-}ADL = MG\text{-}ADL_p + \Delta MG\text{-}ADL_p + \Delta MG\text{-}ADL_{\text{Dose}}$

the model parameters was evaluated

Figure reproduced with permission from Valezuela B, et al. CPT Pharmacometrics Syst Pharmacol. 2025;14(12):2074–2085. **CL** = linear clearance, **FcRn bound**= FcRn receptor bound to nicipolnib, **FcRn free**= FcRn receptor available for nicipolnib binding, **IgG**= total serum immunoglobulin G, **MG-ADL**= MG-ADL at baseline, **K_{deg}**= first-order degradation rate constant of the free FcRn, **K_{deg} IgG**= first-order IgG degradation rate constant, **K_{int}**= internalization rate constant for FcRn–nicipolnib complex, **K_{rec} IgG**= first-order IgG recycling rate constant, **K_{ss}**= quasi-steady-state FcRn–nicipolnib dissociation equilibrium constant, **K_{syn}**= zero-order FcRn free production rate, **K_{syn} IgG**= zero-order IgG production rate, **Q**= intercompartmental clearance, **RO**= FcRn receptor occupancy, **V_p**= central volume of distribution, **V_p**= peripheral volume of distribution, **ΔMG-ADL**= MG-ADL change from baseline in absence of nicipolnib, **ΔMG-ADL_{Dose}**= placebo-corrected MG-ADL change from baseline in presence of nicipolnib.

- After dosing, nipocalimab concentration was distributed into a central compartment (V_c) and peripheral compartment (V_p). It was subject to linear catabolism (CL), as well as binding to FcRn and subsequent internalization of the bound FcRn-nipocalimab complex
- IgG dynamics were described by a turnover model with synthesis rate (K_{syn}) and a natural degradation composed of catabolism (K_{deg}) and an FcRn-mediated recycling rate (K_{rec}), which was dependent on the available free FcRn compared to baseline FcRn
- MG-ADL was described by baseline ($MG-ADL_0$), a placebo signal ($MG-ADL_p$) and a drug effect ($MG-ADL_{Drug}$) that linearly scaled with IgG reduction from baseline
- Impact of age, body weight, sex, race/ethnicity, renal and hepatic impairments, autoantibody status and concomitant gMG therapy on the model parameters was evaluated

Results

Pharmacokinetics

- Pharmacokinetics (PK) of nipocalimab were well described using a two-compartment target-mediated drug disposition model (**Figure 2A**)
- Elimination occurred through both: linear catabolic clearance and nonlinear FcRn-mediated clearance
- Estimated central volume of distribution was 2.41 L and total volume of distribution 3.17 L in patients with gMG from Vivacity-MG3
- No systemic accumulation of nipocalimab was observed with repeated Q2W IV dosing of nipocalimab
- Inter-individual variability in model-predicted PK parameters was low to moderate (<20%)

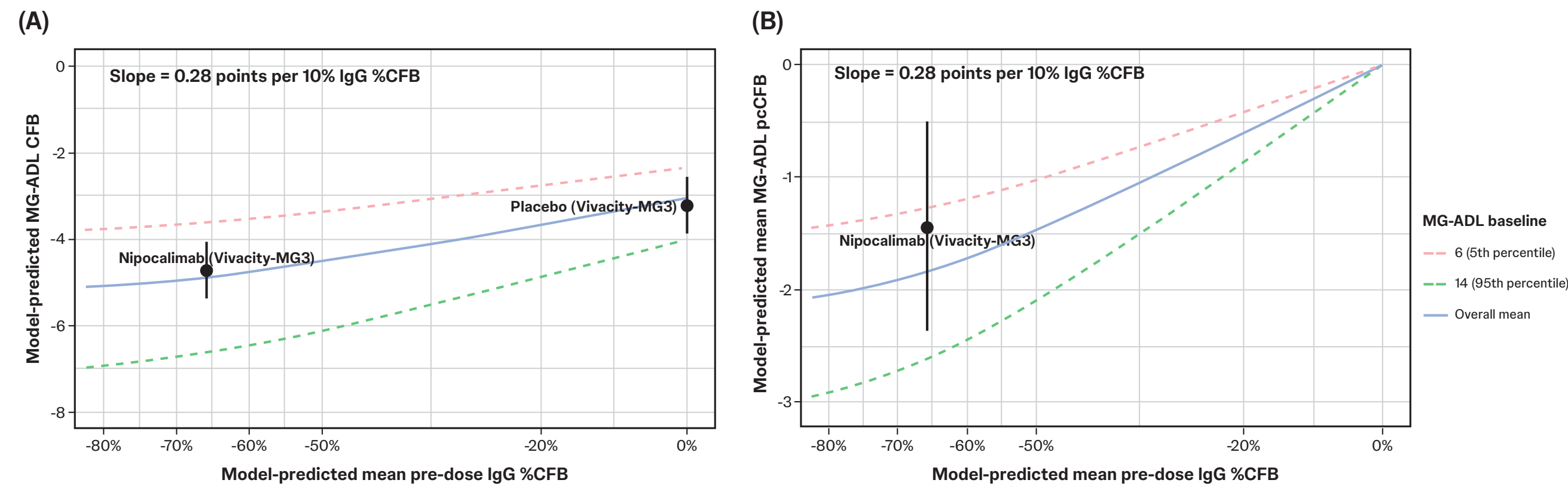
Pharmacokinetics of nipocalimab and relationship with total serum IgG reduction

- Nipocalimab Q2W dosing demonstrated rapid, substantial and sustained reductions in total serum IgG, well described by the model (**Figure 2C**)
- Model-predicted mean (SD) reductions in total serum IgG:
 - Week 2 reduction after loading dose: -71.2% (7.08)
 - Pre-dose steady-state: -65.9% (8.22)
 - Average steady-state : -74.7% (6.21)
 - Maximum predicted IgG reduction, steady state: -80.4% (4.95)
- FcRn blockade with nipocalimab reduced IgG half-life from 17.6 days to 2.75 days
- Total serum IgG concentrations were predicted to return to >90% of baseline approximately 9 weeks after the last dose following treatment discontinuation
- Percent change from baseline (CFB) in anti-AChR linearly correlated with %CFB in total serum IgG, with a slope not significantly different from 1 (**Figure 3**)

IgG-MG-ADL relationship

- Placebo-corrected CFB in MG-ADL linearly correlated with an IgG-reduction effect compartment ($T_{1/2}$ = 5.87 days)
- The estimated slope was 0.28 points MG-ADL improvement per 10% reduction in total serum IgG
- When adjusted for placebo effect, model simulations predicted that a 70% reduction in total serum IgG would result in an additive ~2-point MG-ADL reduction in patients with gMG
- The magnitude of predicted clinical improvement was dependent on the baseline MG-ADL score
- Differences in predicted MG-ADL improvement across tertiles of the nipocalimab-induced reduction in total serum IgG in the phase 3 study were minimal

Figure 4. Model-predicted relationship between mean %CFB in total serum IgG and (A) MG-ADL CFB (B) MG-ADL placebo-corrected CFB



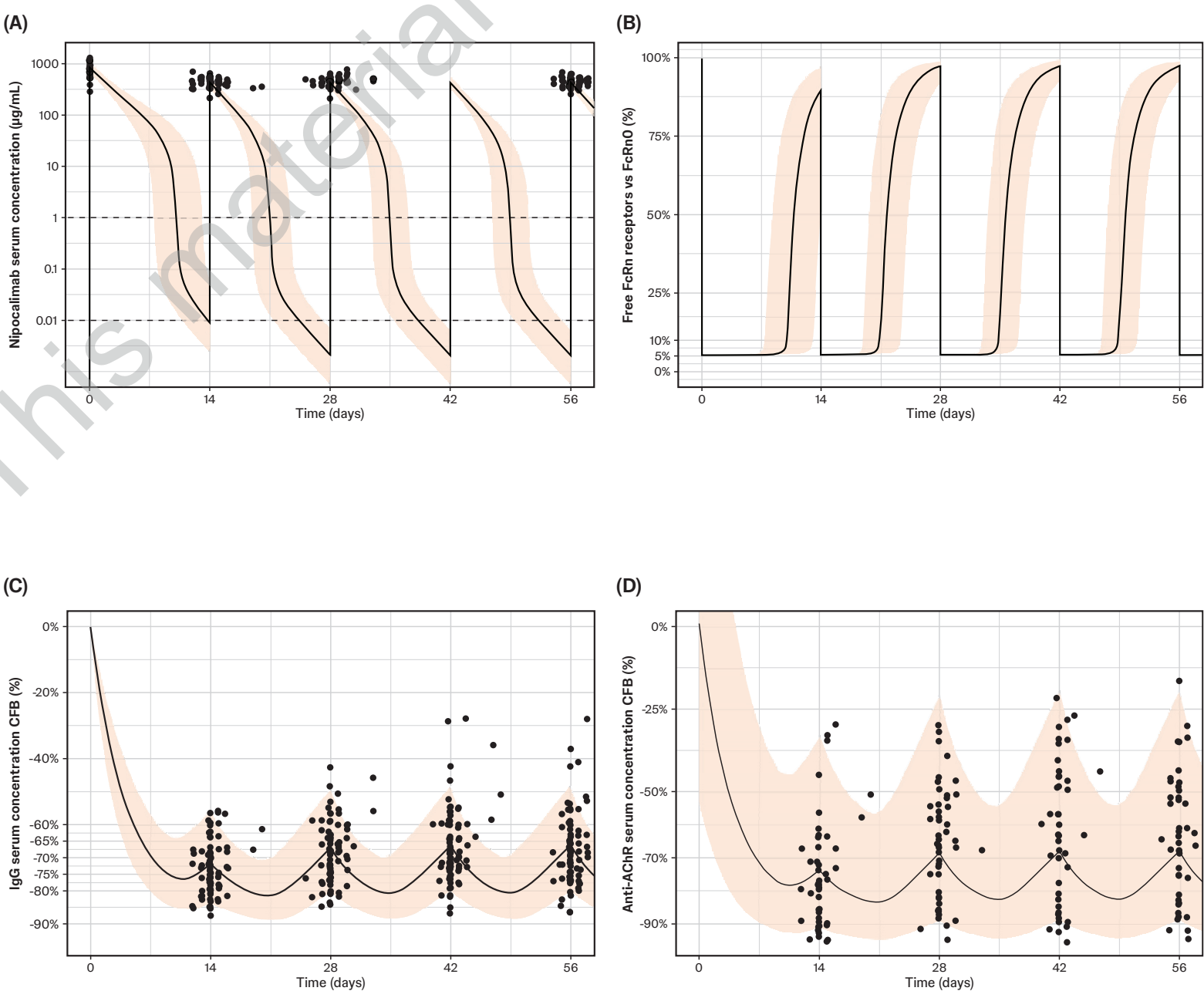
CFB=change from baseline, **pcCFB**=placebo-corrected change from baseline.

- The longitudinal MG-ADL model adequately reproduced placebo-corrected outcomes (i.e., between-group differences vs placebo) for multiple clinical endpoints in the gMG population from the Vivacity-MG3 study⁷
- Predicted improvement was observed when adjusted for age, body weight, sex, race/ethnicity, renal and hepatic impairments, autoantibody status and concomitant gMG therapy

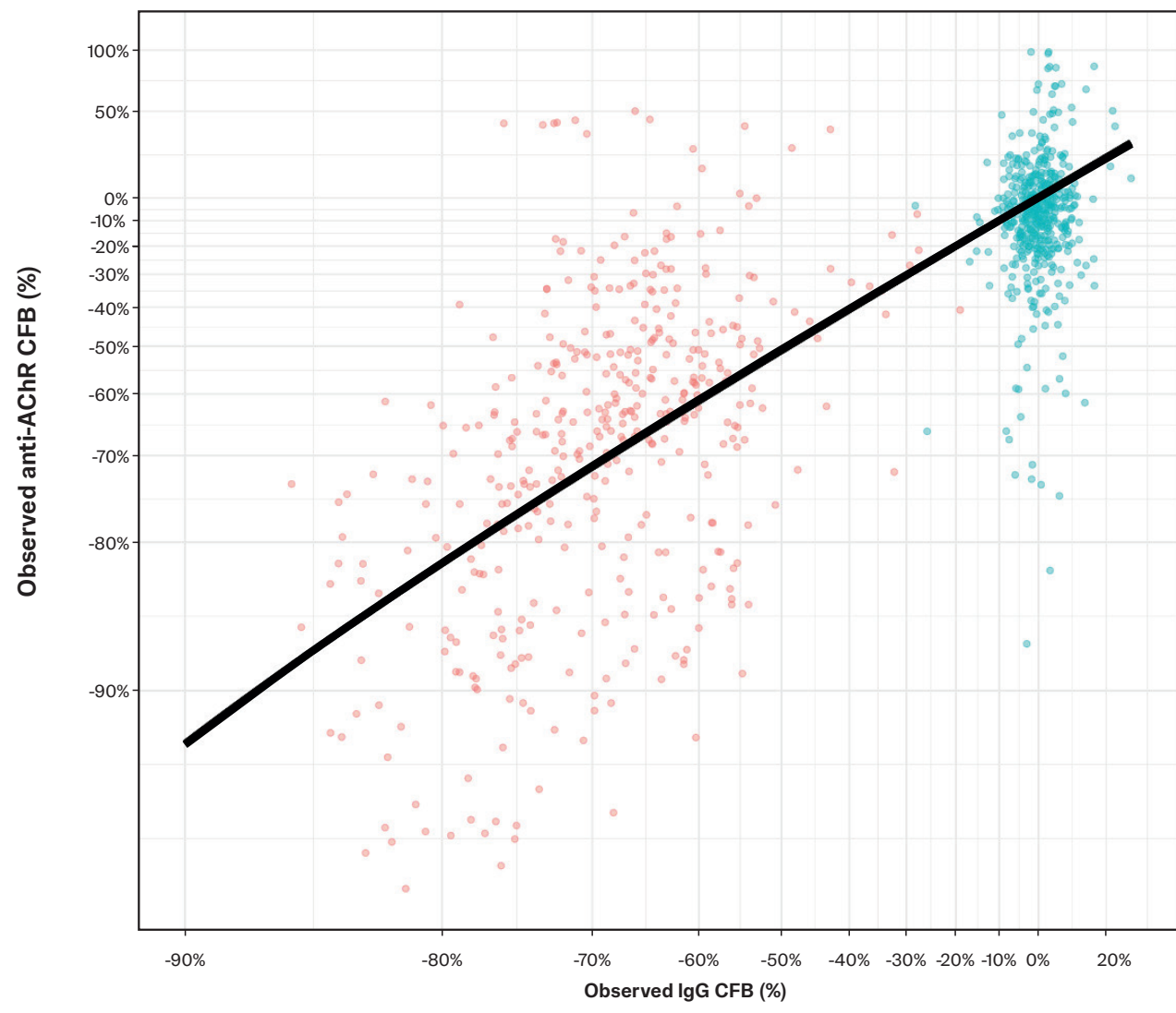
Table 1. Numerical predictive check of clinical endpoints⁷

Clinical endpoint	Nipocalimab		Placebo		Between-group difference	
	Observed	Model prediction (95% CI)	Observed	Model prediction (95% CI)	Observed	Model prediction (95% CI)
MG-ADL CFB (Weeks 22 to 24)	-4.98	-4.77 (-3.81; -5.67)	-3.22	-2.84 (-2.08; -3.56)	-1.75	-1.93 (-3.07; -0.67)
Responders (Weeks 22 to 24), %	82.6	75.4 (63.8; 85.5)	66.1	59.7 (48.4; 72.6)	16.5	15.3 (-1.8; 31.2)
Early responders (≤2 weeks), %	62.3	72.7 (63.6; 81.8)	56.2	60.3 (47.9; 71.2)	6.2	12.7 (-1.2; 28.4)
Sustained response (Weeks 4 to 24), %	60.5	52.6 (42.1; 63.2)	34.2	30.1 (20.5; 39.7)	26.3	22.6 (9.0; 37.1)
≥50% improvement in MG-ADL, %	56.5	50.7 (39.1; 62.3)	32.3	29.0 (19.3; 41.9)	24.3	21.7 (4.6; 38.1)

Observed data were derived through t-test (for continuous endpoints) and chi-squared test (for dichotomous endpoints) based on the Vivicity-MG3 PK/PD analysis dataset. Simulations were performed for 2,000 study replicates of same sample size as the phase 3 Vivicity-MG3 study. Responders were patients with ≥ 2 -point improvement (vs baseline) in average improvement in MG-ADL total score over Weeks 22, 23, and 24 of the double-blind placebo-controlled phase. CFB=change from baseline. CF=confidence interval.



Black line represents model-predicted median and shaded areas represent 95% prediction intervals. Dots represent observed post-dose PK, pre-dose IgG %CFB and pre-dose anti-AChR %CFB data. Pre-dose PK not shown because of fraction below limit of quantification (BLOQ) data. **CFB**=change from baseline, **FcRn RO**=FcRn receptor occupancy



Dots are observed IgG and anti-AChR CFB (%). Black line shows typical model prediction at %anti-AChR=1.03*%IgG-0.0250. CFB=change from baseline, LD=loading dose.