Xencor Q3 2022 Financial Results

XmAb[®]564 Data Presentation

November 7, 2022



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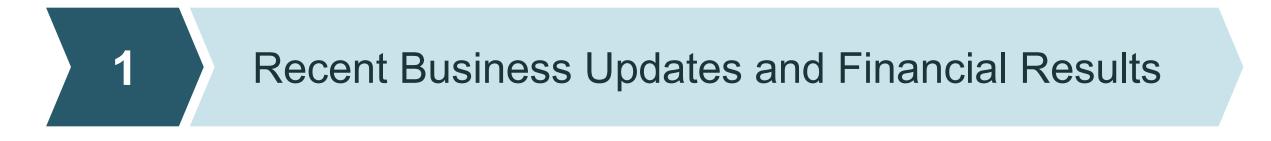


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Agenda for Today's Call







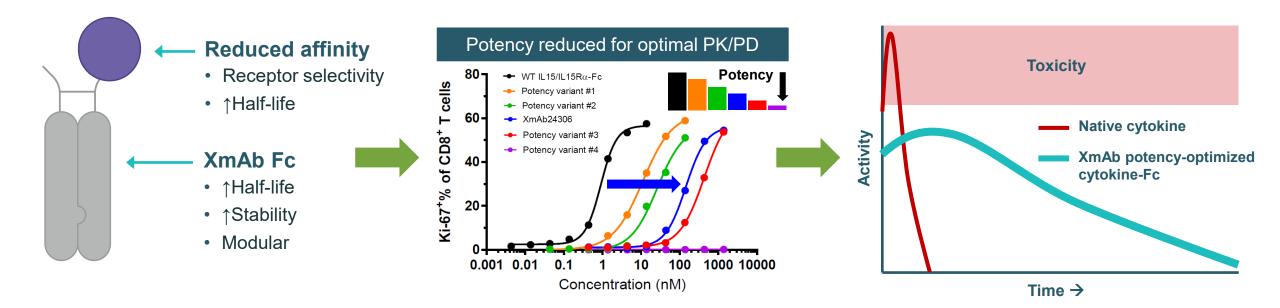


XmAb[®]564 Single Dose Phase 1a Data Presentation

Reduced potency IL2-Fc For selective expansion of regulatory T cells In development for patients with autoimmune diseases

Xencor®

XmAb[®] Cytokines: Potency-tuned to Enhance Half-life and Tolerability



Xencor's general approach for creating cytokine therapies

- Overcomes native cytokine short half-life and high toxicity
- Systematically engineer a broad portfolio of cytokines



XmAb®564 Phase 1a Clinical Trial Top-line Data Summary

- Single ascending dose of subcutaneously administered XmAb564 in healthy volunteers
- Well tolerated; no serious adverse events or dose limiting toxicities observed
- Selective expansion of CD25^{bright} regulatory T cells (Tregs) of 10x and higher beginning at 3rd dose level and reaching 117x increase over baseline at highest dose
- Exceptional durability of Treg expansion relative to reported third-party data
 - Provides opportunity to explore differentiated multi-week dosing schedules
- Minimal increases in natural killer (NK) cells and conventional T cells (Tcons)

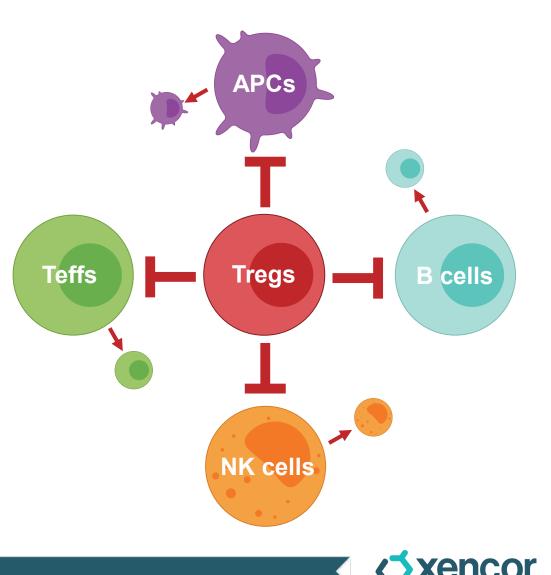
Second potency-tuned XmAb Cytokine program showing marked target cell expansion and good tolerability in human clinical trials

 XmAb306, an IL15-IL15Rα-Fc fusion in oncology, showed consistent and robust dose-dependent NK cell expansion and accumulation upon repeat dosing, reaching 40-100x higher than baseline in higher dose cohorts (Nov. 2021)

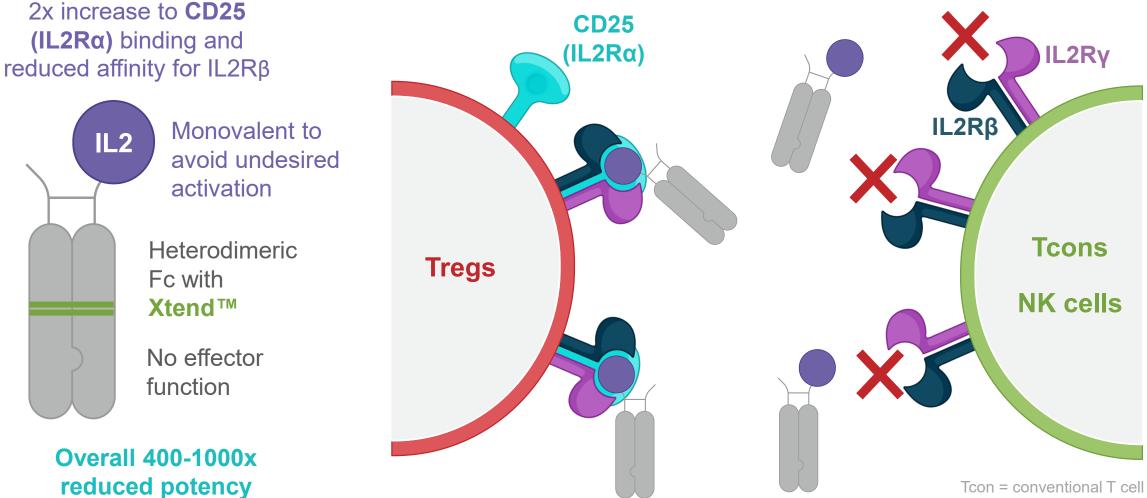


Tregs Important Role in Homeostasis and in Autoimmune Disease

- Regulatory T cells (Tregs) are CD4⁺FoxP3⁺ cells expressing CD25 (IL-2Rα) that maintain immune tolerance in tissues by suppressing the function of both CD4 and CD8 effector T cells
- Tregs are dysfunctional in most autoimmune diseases
- A therapeutic approach has been to restore Treg numbers and function via a low-dose IL-2 regimen
 - Treg homeostasis depends on IL-2
 - IL-2 as a drug suffers from fast *in vivo* clearance and a narrow therapeutic index



IL-2 Activates Multiple T Cell Types XmAb564 is Engineered to Improve for Treg Selectivity

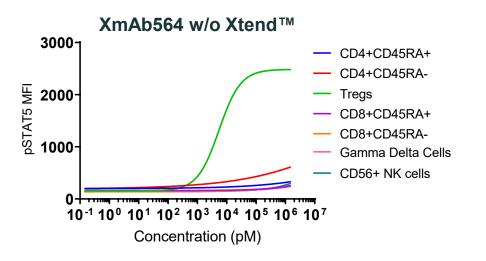


WT = wild-type

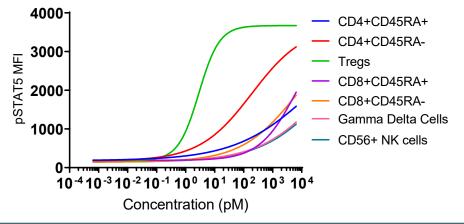


XmAb564 Design Reduces Potency and Improves Treg Selectivity

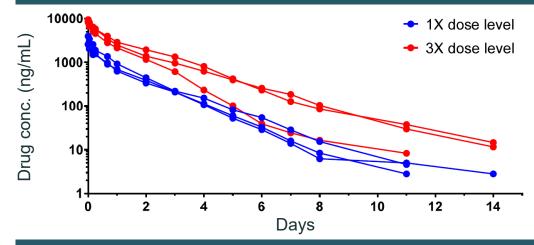
XmAb564 Selectively Promotes Treg Signaling in Human T Cells



Recombinant human WT IL-2



Sustained PK up to Several Days in Non-Human Primates



XmAb564 exhibits extended half-life and good tolerability in NHPs (not shown)

XmAb564 selectively promotes Tregs and has extended half-life due to low potency and Xtend[™] Fc domain



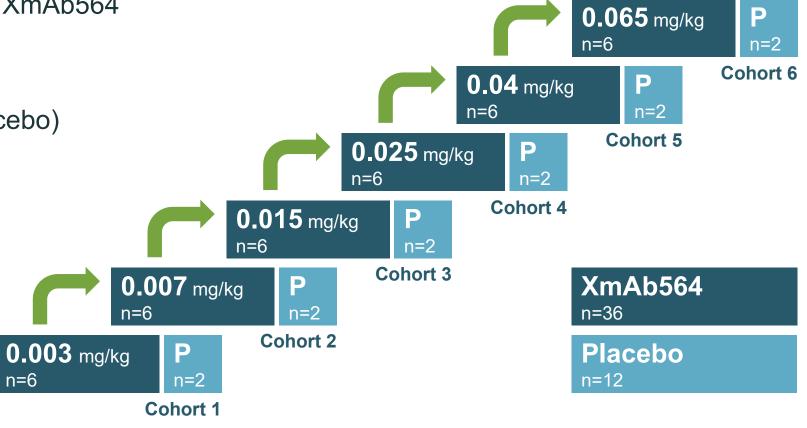
XmAb564 Phase 1a Healthy Volunteer Study Design

Phase 1a single-ascending dose (SAD) study

- Randomized and double-blinded
- Subcutaneous administration of XmAb564
- Heathy volunteers (n=48)
- 6 dose level cohorts
- Randomized 6:2 (XmAb564:placebo)

Outcome measures

- Safety and tolerability
- Pharmacokinetics and activity biomarkers (e.g., T-cell populations)





XmAb564 Was Well Tolerated

Well tolerated including at the highest dose evaluated (0.065 mg/kg)

- All adverse events (AEs) were grade 1 or 2 (mild-to-moderate) and self-limited
- Injection site reaction was the most reported AE
- No serious AEs, dose-limiting toxicities or early discontinuations due to AEs
- No clinically significant abnormalities in laboratory values, vital signs and ECGs

Laboratory findings and pharmacokinetics

- Some subjects had transient, reversible elevations in blood eosinophils
 - No eosinophil-related AEs were observed
 - Possibly related to mechanism-of-action, reported in other third-party CD25-targeting IL-2 programs
- No other clinically significant abnormalities in safety laboratory studies were observed
- Terminal half-life is estimated to be 9-10 days at lower doses and 6-7 days at the highest dose, consistent with an increase in CD25 target-mediated clearance on the expanding Treg population

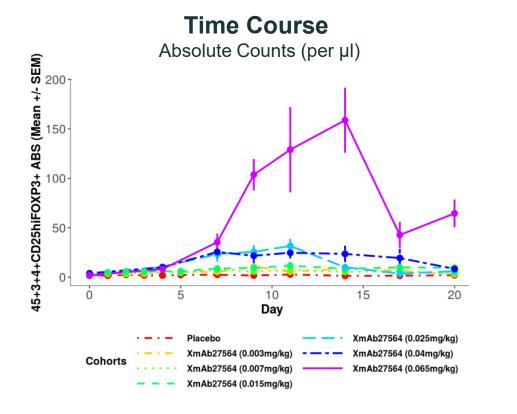




XmAb564 Promotes Robust & Durable Expansion of CD25^{bright} Tregs

300-

ABS

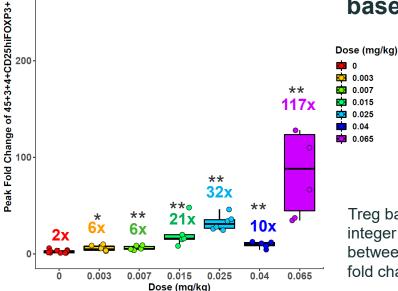


Durable, dose-dependent increases in CD25^{bright} Tregs

Peak Fold Change¹

0

117-fold mean peak expansion over baseline in Cohort 6



Treg baseline counts are low integer values, leading to between group variability in fold change calculations

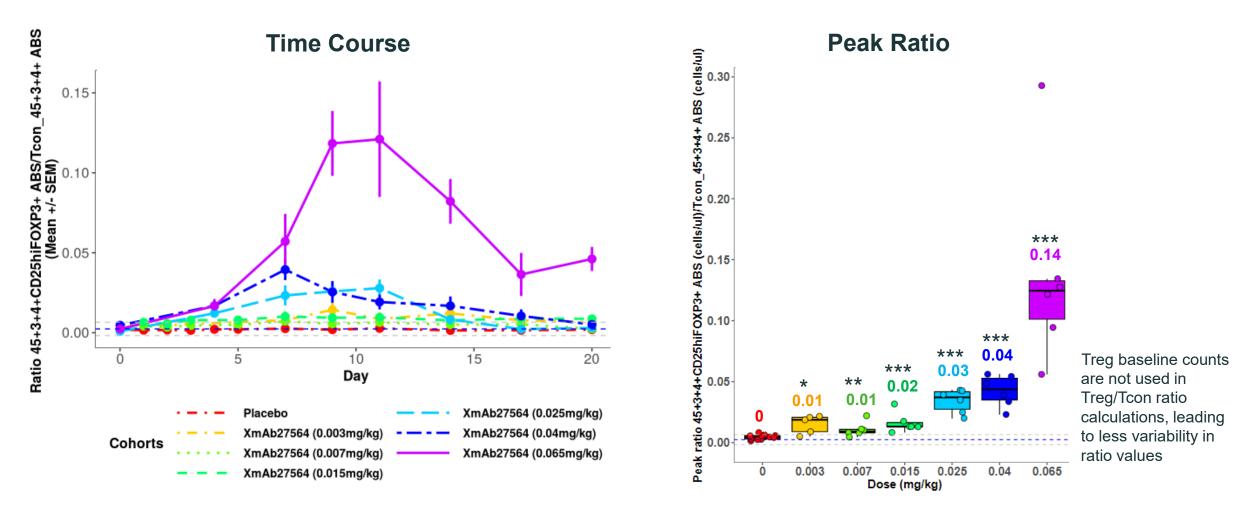
Highest reported CD25^{bright} Treg expansion

Numeric values in "Peak Fold Change" plot are Mean

¹ Peak fold change: Peak CD25^{bright}FoxP3⁺ CD4 Treg cell absolute count at a post-treatment time point divided by absolute count at baseline NS: p>0.05, *: $p\leq0.05$, **: $p\leq0.01$, ***: $p\leq0.001$ compared with placebo treated cohort, Wilcoxon test



XmAb564 Promotes Robust & Durable Increases in CD25^{bright} Treg/Tcon ratio

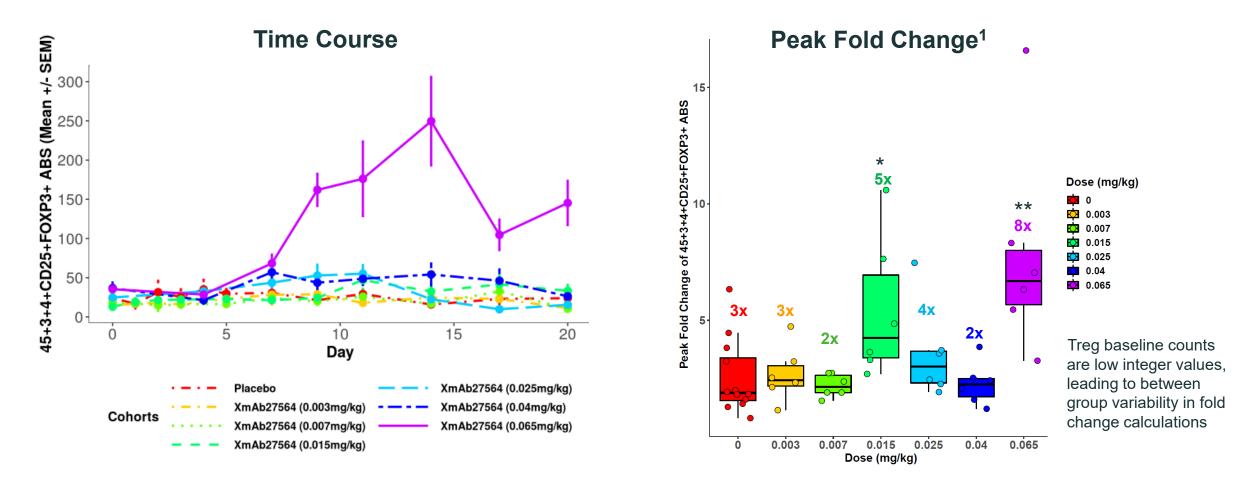


Numeric values in "Peak Fold Change" plot are Mean

Dashed horizontal blue line represents the average of pre-dose values from all 48 subjects with ± 2 SD shown in grey lines NS: p>0.05, *: p<0.05, *: p<0.01, ***: p<0.01 compared with placebo treated cohort, Wilcoxon test



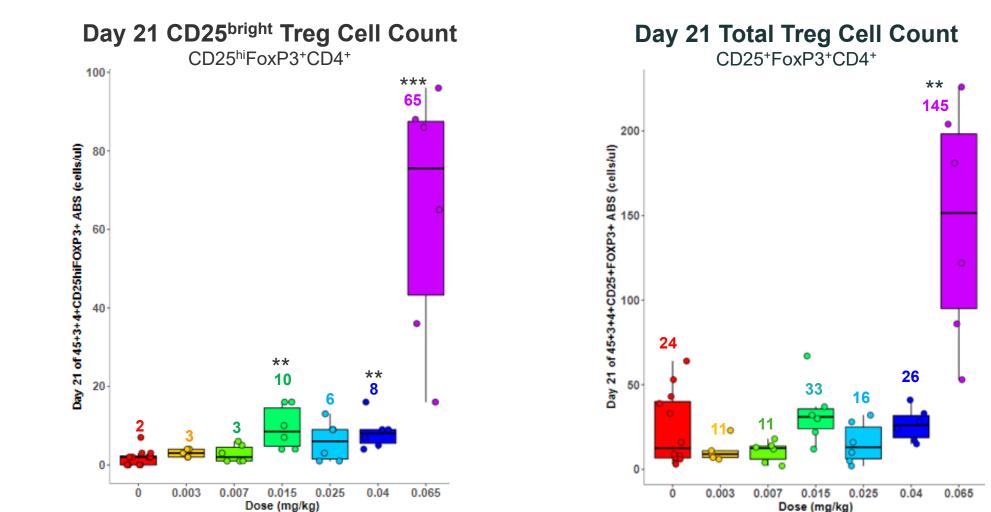
XmAb564 Promotes Robust & Durable Expansion of Total Tregs



Numeric values in "Peak Fold Change" plot are Mean

¹ Peak fold change: Peak CD25⁺FoxP3⁺ CD4 Treg cell absolute count at a post-treatment time point divided by absolute count at baseline NS: p>0.05, *: $p\leq0.05$, *: $p\leq0.01$, **: $p\leq0.01$, compared with placebo treated cohort, Wilcoxon test



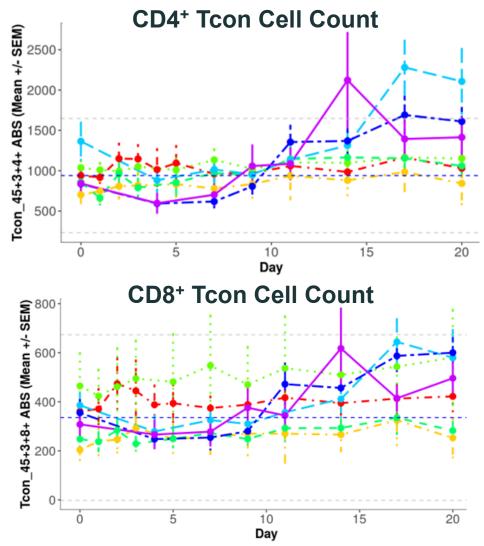


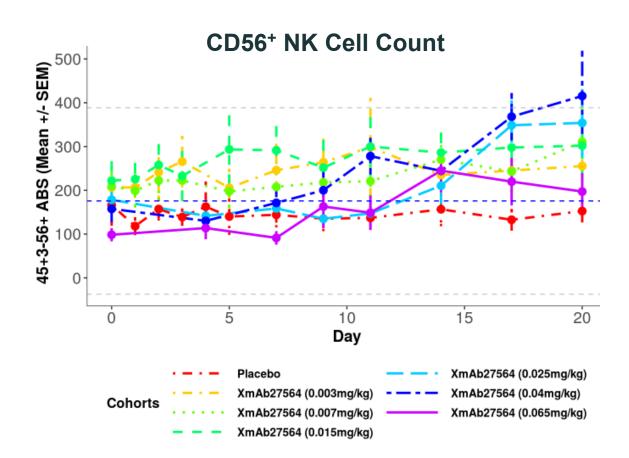
CD25^{bright} and Total Treg Remain Elevated for at Least 3 Weeks

Numeric values in plots are Mean NS: p>0.05, *: p≤0.05, **: p≤0.01, ***:p≤0.001 compared with placebo treated cohort, Wilcoxon test



XmAb564 Induces Minimal Increases in Conventional T cells and NK Cells





Dashed horizontal blue line represent the average of pre-dose values from all 48 subjects with ±2SD shown in grey lines



XmAb564 Phase 1a Topline Summary

XmAb564 is well tolerated and generates a durable, dose-dependent and selective expansion of Tregs with a single dose

- 117-fold mean peak expansion over baseline in CD25^{bright} Tregs and 8-fold mean peak expansion in total Tregs at highest dose
- Marked elevation of Tregs sustained through at least day 21: CD25^{bright} and total Tregs increased 44-fold and 4.5-fold at highest dose, respectively
- Treg/Tcon ratio increased significantly in a dose-dependent manner
- All AEs Grade 1/2 and resolved without intervention

Phase 1b study in patients

- First patient dosed in a newly initiated Phase 1b, multiple-ascending dose (MAD) study of XmAb564 in patients with atopic dermatitis and psoriasis
- Multiple dosing schedules to be explored based on pharmacodynamic data



Xencor's Growing Portfolio of Potency-Optimized XmAb® Cytokines

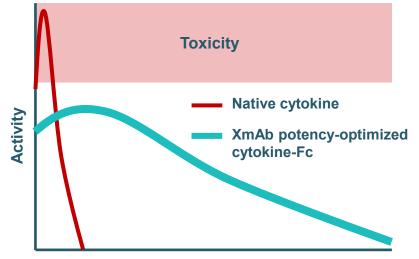


Second validation of Xencor's approach to cytokine therapeutics

- XmAb306 (IL-15) Phase 1 demonstrated 40- to 100-fold increases in activated NK cells and accumulation upon multiple doses
- In two first-in-human studies, reduced potency and long half-life has translated to improved tolerability compared to the high toxicities generated by native cytokines

XmAb Cytokines

- Engineered to expand select immune cell populations
- Designed to be tolerable, active and easy to use



Time \rightarrow



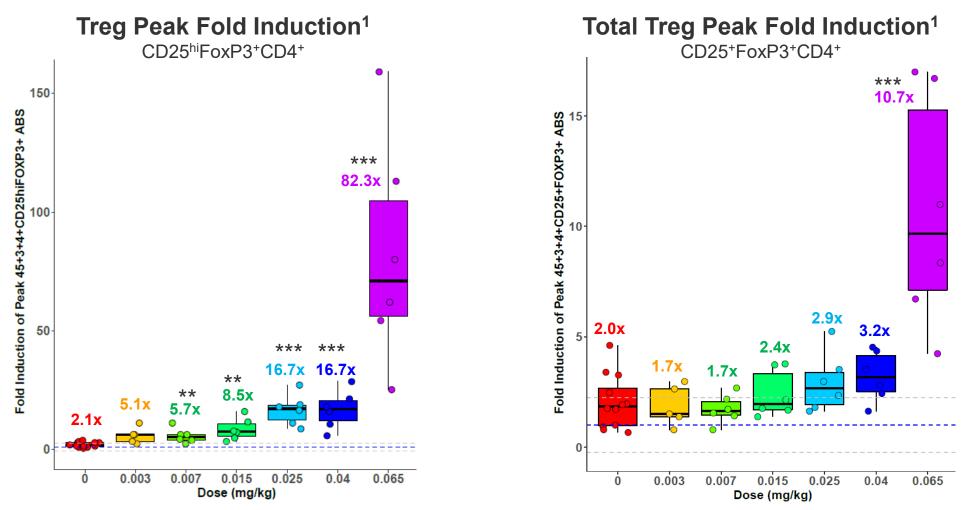




Data Appendix



Peak Fold Induction of CD25^{bright} and Total Treg Cells Shows Consistent Dose Response of Treg Populations to XmAb564

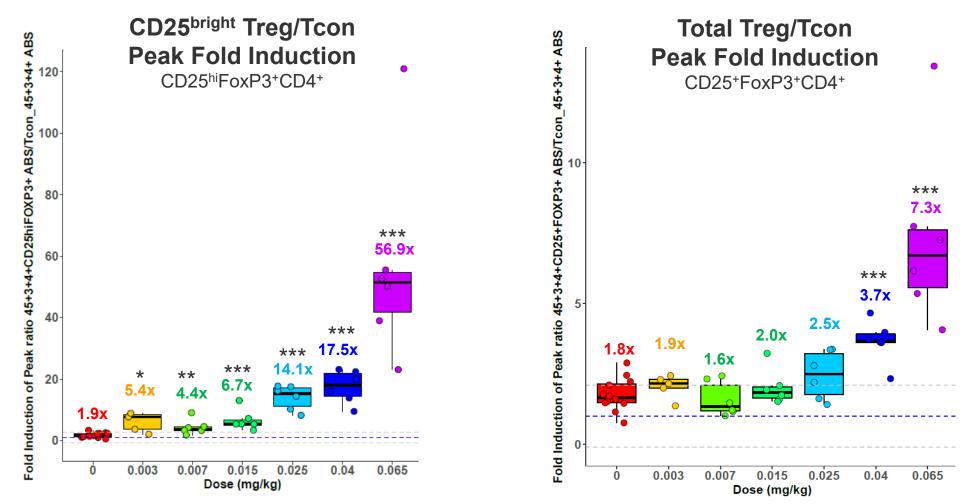


Numeric value in "Peak Fold Change" plot is Mean

¹ Peak fold induction: Peak CD25^{bright} Treg or total Treg cell absolute counts of each subject divided by the average of all pre-dose values from all subjects (n=48) NS: p>0.05, *: $p\leq0.05$, **: $p\leq0.01$, ***: $p\leq0.001$ compared with placebo treated cohort, Wilcoxon test



Ratio of CD25^{bright}/Tcon and Total Treg/Tcon Peak Fold Induction Shows Consistent Dose Response of Treg Populations to XmAb564



Numeric value in "Peak Fold Change" plot is Mean

Peak fold induction: Peak CD25^{bright} Treg/Tcon or total Treg/Tcon ratios of each subject divided by the average of all pre-dose values from all subjects (n=48) NS: p>0.05, *: p≤0.05, *: p≤0.01, ***:p≤0.001 compared with placebo treated cohort, Wilcoxon test

