Supplementary figures



Sup figure 1. Serum cytokine level at 2 hr and 10 day after bone marrow transplantation Recipient BALB/c mice were irradiated 8.8-Gy and transferred C57BL/6 bone marrow cells plus Tcons with or without 1.0×10^5 sorted Foxp3^{GFP+} Tregs followed by administration of ctrl-lipo or α -GalCer-lipo. Serum samples were collected at 2 hr (A) and 10 day (B) after BMT. Mean ± SD from pooled data of two independent experiments. P-value *<0.05, **<0.01, ***<0.001, *****<0.0001 calculated by ordinary one way ANOVA with Holm-Sidak correction. ns: not significant.



Sup figure 2. Murine survival curve from day 14 after BMT.

The mice that survived beyond day 14 were implemented into the analysis. P-value **<0.01, ****<0.001, ****<0.0001 calculated by Log-rank test between indicated two groups. Pooled data of four independent experiments is shown. ns: not significant.



Sup figure 3. Sorting strategy for CD4⁺Foxp3^{GFP+} Tregs.

CD25⁺ cells were magnetic bead-enriched from single cell suspension obtained from secondary lymphoid tissues of B6-Foxp3^{GFP+} mice, and further stained with CD4, CD8, and LIVE/DEAD Aquamine. CD4⁺CD⁻CD25^{hi}Foxp3^{GFP+} cells were gated and acquired by flow cell sorter.



Sup figure 4. Administration of α -GalCer-lipo significantly improved murine survival when combined with 5 \times 10⁴ Treg transfer

Flow sorted Foxp3^{GFP+} Tregs were injected into recipient mice at 1:20 Treg:Tcon ratio at the time of bone marrow transplantation. P-value ***<0.001, ****<0.0001 calculated by Log-rank test between indicated two groups. Pooled data of two independent experiments was shown.





в



Е









 Ctrl-lipo
 o-GalCer-lipo Treg + ctrl-lipo
Treg + α-GalCer-lipo

FA





F

G



αGalCer-lipo alone





29.6

5.45E-3

0.51 👄

#13

#17

#3





47.1

6.23 👄

#15

6.32









4.36

4.24 🗢

#16



4.48 🗢 #18







57.2 0.62

4

Sup figure 5. Gating strategy for UMAP plot.

(A) All data from one representative experiment were concatenated. Singlet-living lymphocytes were gated and TCR β^+ CD19⁻ gated cells were implemented for downstream analysis. UMAP was calculated according to expression level of CD4, CD8a, CD45.1, CD45.2, Foxp3 (intracellular stained), Foxp3^{GFP}, CD25, CD44, CD62L, ICOS, and Ki67. (B) CD4⁺Foxp3⁻ cells (orange box: CD4⁺ Tcons), CD8⁺ cells (green box: CD8⁺ Tcons), CD4⁺Foxp3⁺Foxp^{GFP-} cells (blue box: Tcon-derived Tregs), and CD4⁺Foxp3^{GFP+} cells (red box: transferred Tregs) were gated, and overlaid on UMAP plot (main figure 2E). (C) CD25⁺, CD44⁺, CD62L⁺, ICOS⁺, and Ki67⁺ cells (brown dots) that were overlaid on representative UMAP plot of each group. (D) Four major clusters (1 – 4) identified and gated for the following analysis. (E) The proportion of cluster 1 – 4 in each treatment group. P-value **<0.01, ***<0.001, ****<0.0001 calculated by student t-test between indicated two groups. (F) Concatenated data was divided into an individual mouse by gating on each sample number; #1-5: Ctrl-lipo, #6-10: α -GalCer-lipo, #11-14: Ctrl-lipo + Treg, #15-18: α -GalCer-lipo + Treg. (G) UMAP plots gated on individual mouse. The data from one representative of two individual experiment. ns: not significant.



Sup figure 6. T cell phenotype on day 3, day 6, and day 10.

Splenocytes from ctrl- and α -GalCer-lipo treated recipients were obtained on day 3, day 6, day 10 after BMT and analyzed by Flow cytometry. The proportion (above) and Ki67 expression level (bottom) of CD4+Foxp3^{GFP+} Tregs, CD45.1+CD4+ Tcons, and CD45.1+CD8+ Tcons at the indicated time points.



Sup figure 7. Immunohistochemistry of small intestine

Images of immunofluorescent staining for CD4 (red), Foxp3GFP (Green), and DAPI (Blue). Left; low magnification (10x, white bar indicates 100 μ m), right; high magnification of inset region (40x, white bar indicates 20 μ m). One representative image of four independent mice.