| 2 | Jing Zhao ^{1,2†} , Liwei Jiang ^{1,2†} , Mayuko Uehara ^{1,2} , Naima Banouni ^{1,2} , Basmah S Al Dulaijan ^{1,2} , |
|----------------|---|
| 3 1 | Jing Zhao ^{1,2†} , Liwei Jiang ^{1,2†} , Mayuko Uehara ^{1,2} , Naima Banouni ^{1,2} , Basmah S Al Dulaijan ^{1,2} , |
| 5 5 | |
| 4 J | Jamil Azzi ^{1,2} , Takaharu Ichimura ² , Xiaofei Li ^{1,2} , Petr Jarolim ³ , Paolo Fiorina ⁴ , Stefan G.Tullius ⁵ , |
| 5 J | Ioren C. Madsen ^{6,7} , Vivek Kasinath ^{1,2‡} , Reza Abdi ^{1,2‡} |
| 6 | |
| 7 ¹ | Transplantation Research Center, Brigham and Women's Hospital, Harvard Medical School, |
| 8 E | Boston, MA 02115, USA |
| 9 2 | Renal Division, Brigham and Women's Hospital, Harvard Medical School, Boston, MA 02115, |
| 10 U | USA |
| 11 3 | Department of Pathology, Brigham and Women's Hospital, Harvard Medical School, Boston, |
| 12 N | MA 02115, USA |
| 13 4 | Department of Nephrology, Boston Children's Hospital, Harvard Medical School, Boston, MA |
| 14 0 | 02115, USA |
| 15 5 | Division of Transplant Surgery, Department of Surgery, Brigham and Women's Hospital, |
| 16 H | Harvard Medical School, Boston, MA, 02115, USA |
| 17 6 | Center for Transplantation Sciences, Department of Surgery, Massachusetts General Hospital |
| 18 a | and Harvard Medical School, Boston, MA, 02114USA |
| 19 7 | Division of Cardiac Surgery, Department of Surgery, Massachusetts General Hospital and |
| 20 H | Harvard Medical School, Boston, MA, 02114, USA |
| 21 | |
| 22 | Correspondence: |

23 Reza Abdi, MD

- 24 Transplantation Research Center, Brigham and Women's Hospital, 221 Longwood Ave, Boston
- 25 MA 02115, USA
- 26 Tel: 617-732-5259, Fax: 617-732-5254, Email: rabdi@rics.bwh.harvard.edu
- 27 † Co-first authors; ‡ co-senior authors
- 28
- 29 **Conflict-of-interest:** The authors have declared that no conflict of interest exists.









33 populations from Treg assay in untreated control and ACTH-treated groups, using lymphocytes 34 from Wild-type (WT) and CD28 knock-out (CD28KO) C57BL/6 background mice. (B) qPCR of 35 Tregs shows significantly increased *Foxp3* gene expression in ACTH-treated group compared to 36 untreated control group from CD28KO mice, but no significant difference is seen with WT mice. 37 No significant difference in gene expression levels of Nfatc2, Ap1, E2a, and Cre1 between 38 untreated control and ACTH-treated group from WT and CD28KO mice was seen. (C) Flow 39 cytometric analysis reveals that ACTH induced a higher CD25 expression (by mean fluorescence 40 intensity (MFI)) than no treatment (control) in a Treg induction assay, using splenocytes from WT 41 and CD28KO mice. Experiments were performed independently in triplicate. (D) Luminex assay 42 revealed no significant difference in IL-2 concentration between WT and CD28KO mice following 43 ACTH treatment. (E) Flow cytometric analysis of T cells from WT spleen shows no difference of 44 percentage in Teff cells following either treatment with ACTH or no treatment (Control). (F) Flow 45 cytometric analysis of T cells collected from WT spleen shows no significant difference in the 46 percentages of nature Tregs (nTregs) between the ACTH-treated group and the untreated control 47 group. (G) Flow cytometric analysis of T cells collected from WT spleen shows significantly 48 higher percentage of induced Tregs (iTregs) in ACTH-treated group than untreated control group. 49 (H) Flow cytometric analysis of Tregs collected from WT spleen shows no significant difference 50 in the percentages of Ki-67, Caspase 3, PD-1, TIM-3, LAG-3 and CTLA-4 between the ACTH-51 treated group and untreated control group. Data presented as the mean \pm SEM, *p<0.05, ** p<0.01, *** p<0.001. 52



54

55 Supplemental Figure 2. (A) Survival curve demonstrates no significant difference in heart allograft 56 survival between C57BL/6 recipients of BALB/c hearts that received ACTH (n=4 mice/group, 57 MST=7.5 days) versus untreated allograft recipients (control; n= 6 mice/group, MST=7 days). (B) 58 Flow cytometric analysis of LNs collected from CD28KO recipients at Day 7 shows no difference 59 in percentages of Tregs and Treg/Teff ratio (CD4⁺CD25⁺Foxp3⁺/ CD44⁺CD62L⁻) between the 60 ACTH-treated group and untreated control group. (C) Flow cytometric analysis of spleens 61 collected from CD28KO recipients at day 7 shows no significant difference in Tregs but 62 significantly higher Treg/Teff ratio in the ACTH-treated group than the untreated control group. 63 (n=3 mice/group) Data presented as the mean \pm SEM, ** p<0.01.





Supplemental Figure 3. (A) Semi-quantitative analysis by MFI of florescence micrographs of heart
allograft sections revealed fewer CD3⁺ T cells and CD11b⁺ cells, but more Foxp3⁺ Tregs in group
treated with CTLA-4-Ig + ACTH in comparison to group treated with CTLA-4-Ig alone (n=3)

69 allograft/group) (B) qPCR of heart allografts showed significantly higher expression of *Foxp3* and Ccr2 genes in CTLA-4-Ig + ACTH group, as compared to group treated with CTLA-4-Ig alone. 70 No significant difference in gene expression of Il10, Ifng, Tnfa and Il17 was observed (n=4 71 72 mice/group). (C) Flow cytometric analysis of LNs collected from C57BL/6 recipients at Day 28 73 showed no difference in the percentages of Tregs and the Treg/Teff ratio between the CTLA-4-Ig 74 group and CTLA-4-Ig + ACTH group. (D) Flow cytometric analysis of spleens collected from 75 C57BL/6 recipients at Day 28 showed no difference in the percentages of Tregs, but significantly higher Treg/Teff ratio was observed in the CTLA-4-Ig + ACTH group in comparison to CTLA-4-76 Ig group (n=3 mice/group). Data presented as mean \pm SEM, * p<0.05, ** P<0.01. 77





Supplemental Figure 4. (A) Flow cytometric analysis of LNs collected from C57BL/6 recipients
at 4 weeks shows no significant difference in the percentages of Tregs between ACTH-treated
group and untreated control group. (B) Flow cytometric analysis of spleens collected from

- 83 C57BL/6 recipients at 4 weeks shows no significant difference in the percentages of Tregs and
- 84 CD4⁺IL-17⁺ cells between the ACTH-treated group and the untreated control group. (n=4
- 85 mice/group) Data presented as the mean \pm SEM.