

**Tuberous sclerosis complex 1 (Tsc1) enforces quiescence of naive T cells to  
promote immune homeostasis and function**

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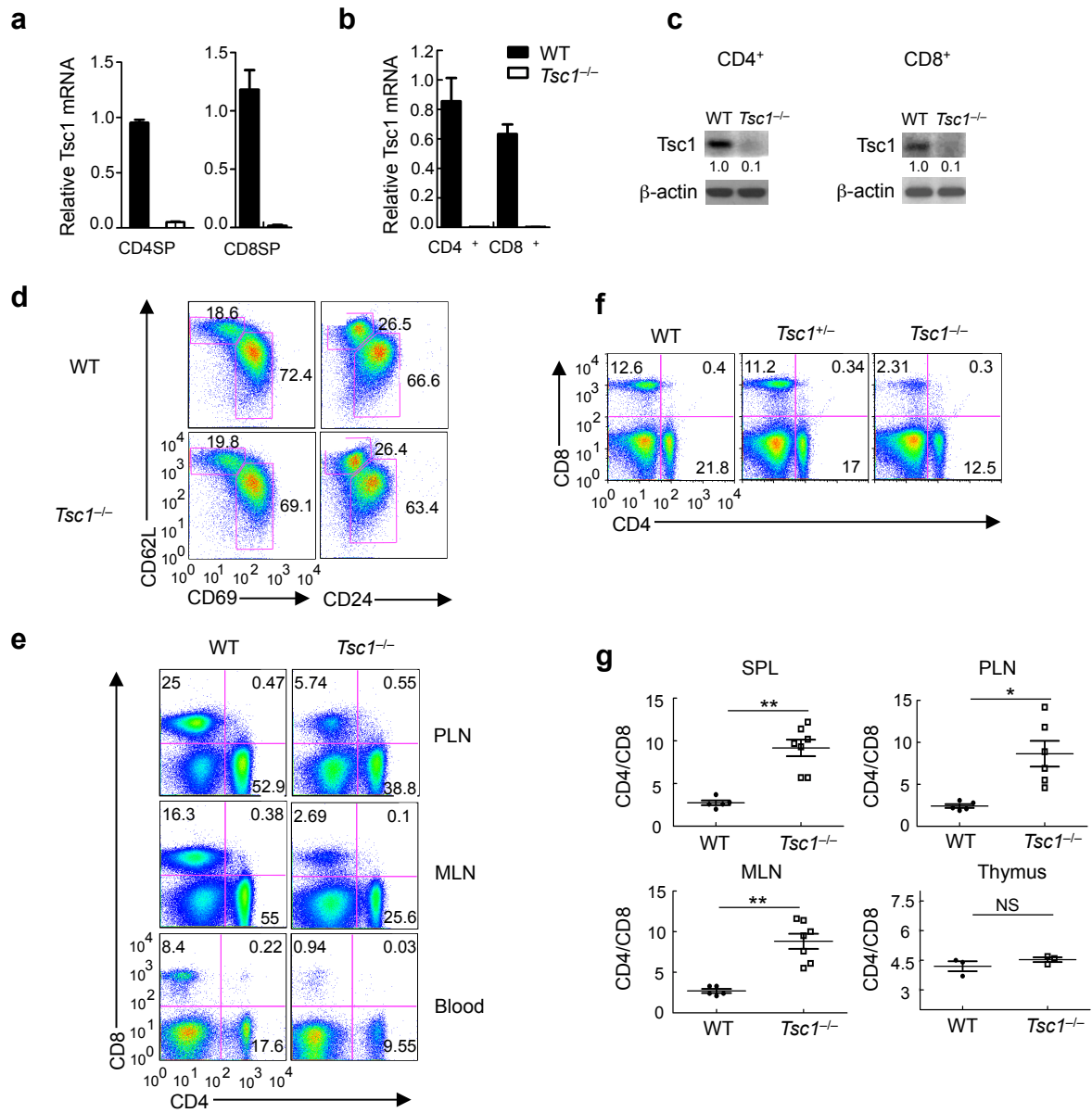
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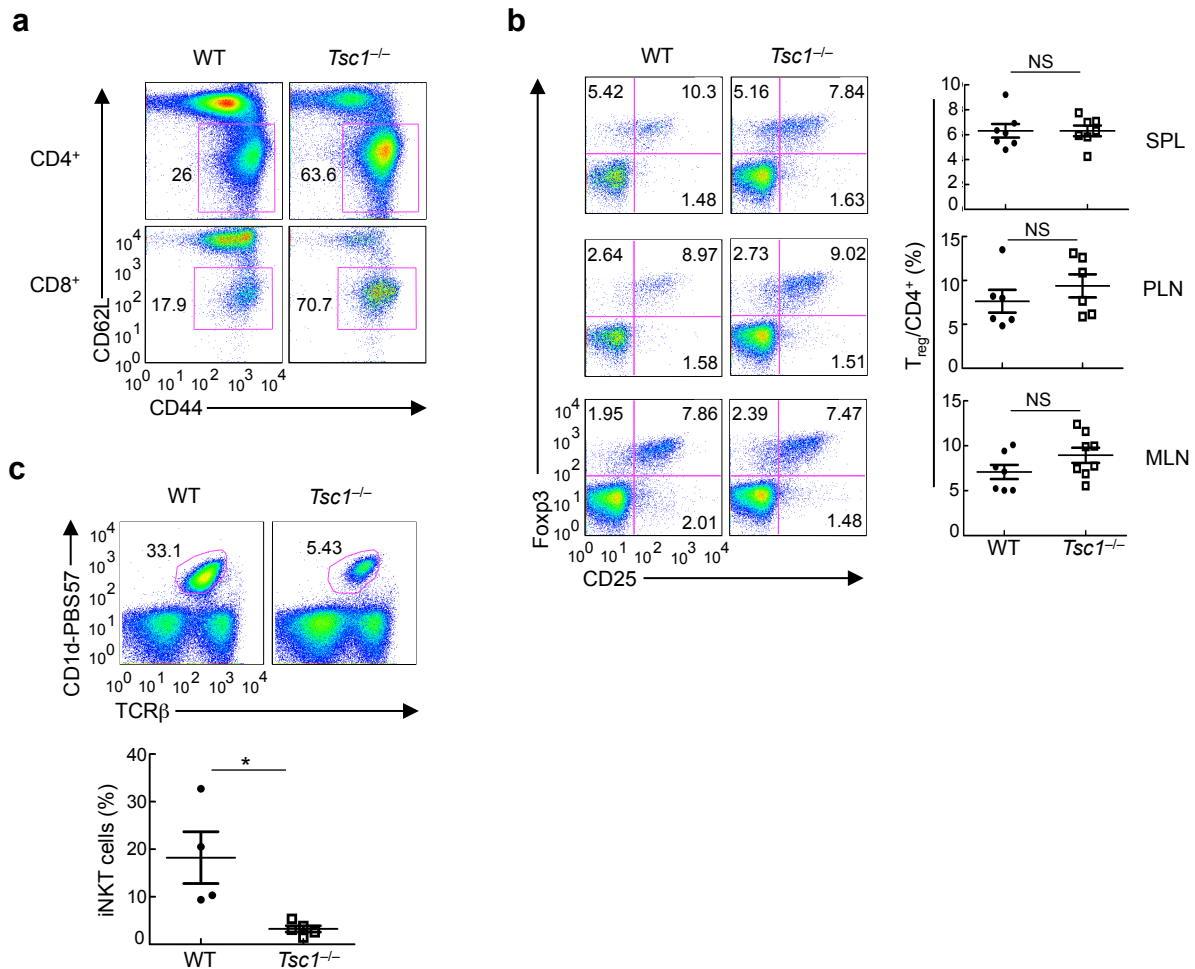
[hongbo.chi@stjude.org](mailto:hongbo.chi@stjude.org).

## Supplementary Figure 1



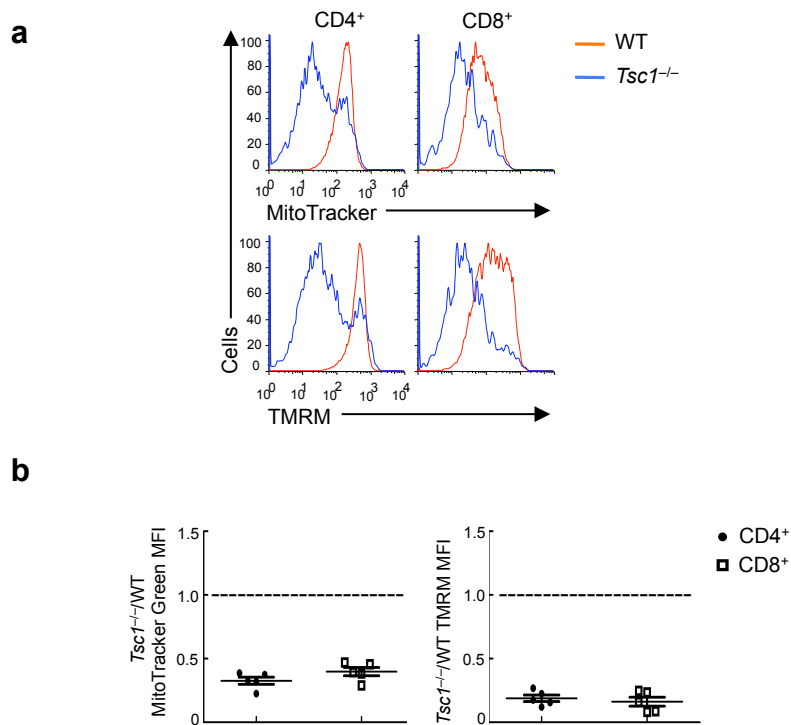
**Supplementary Figure 1. T cell-specific deletion of *Tsc1* results in altered homeostasis of peripheral but not thymic T cells.** (a,b) Analysis of *Tsc1* mRNA expression in CD4SP and CD8SP thymocytes (a) and in peripheral CD4 and CD8 T cells (b). (c) Western blot of *Tsc1* expression in peripheral CD4 and CD8 T cells. Numbers below lanes indicate band intensity relative to that of the loading control  $\beta$ -actin. (d) Normal expression of maturation markers on *Tsc1*<sup>-/-</sup> CD4SP thymocytes. (e) Reduced CD4 and CD8 populations in peripheral lymph nodes (PLN), mesenteric lymph nodes (MLN) and blood of *Tsc1*<sup>-/-</sup> mice. (f) Normal distribution of CD4 and CD8 populations in the spleen of *Tsc1*<sup>+/-</sup> mice. (g) Increased ratios of CD4 vs CD8 T cells in the peripheral but not thymic T cells from *Tsc1*<sup>-/-</sup> mice (n=3-7). SPL, spleen. \*  $P < 0.01$ ; \*\*  $P < 0.001$ ; NS, not significant. Data are representative of 3-4 independent experiments.

## Supplementary Figure 2



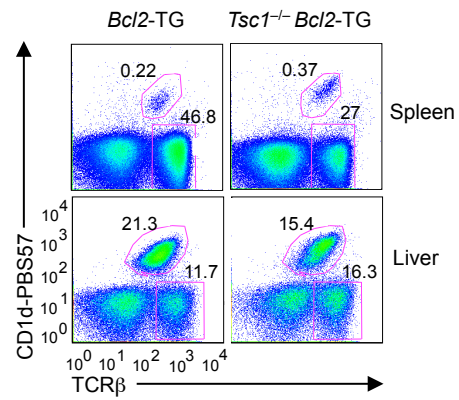
**Supplementary Figure 2. Analysis of T cell subsets in the periphery of *Tsc1*<sup>-/-</sup> mice. (a)** Flow cytometry of CD62L and CD44 expression in mice at 8 months of age showed expansion of the CD62L<sup>lo</sup>CD44<sup>hi</sup> population in *Tsc1*<sup>-/-</sup> cells. **(b)** Flow cytometry of Foxp3 and CD25 expression revealed normal frequency of T<sub>reg</sub> cells in the periphery of *Tsc1*<sup>-/-</sup> mice (n=6-8). **(c)** Flow cytometry analysis of iNKT cells (TCR $\beta$ <sup>+</sup>CD1d-PBS57<sup>+</sup>) in the livers of WT and *Tsc1*<sup>-/-</sup> mice (n=4-5). \* *P*<0.05; NS, not significant. Data are representative of 4 independent experiments.

### Supplementary Figure 3



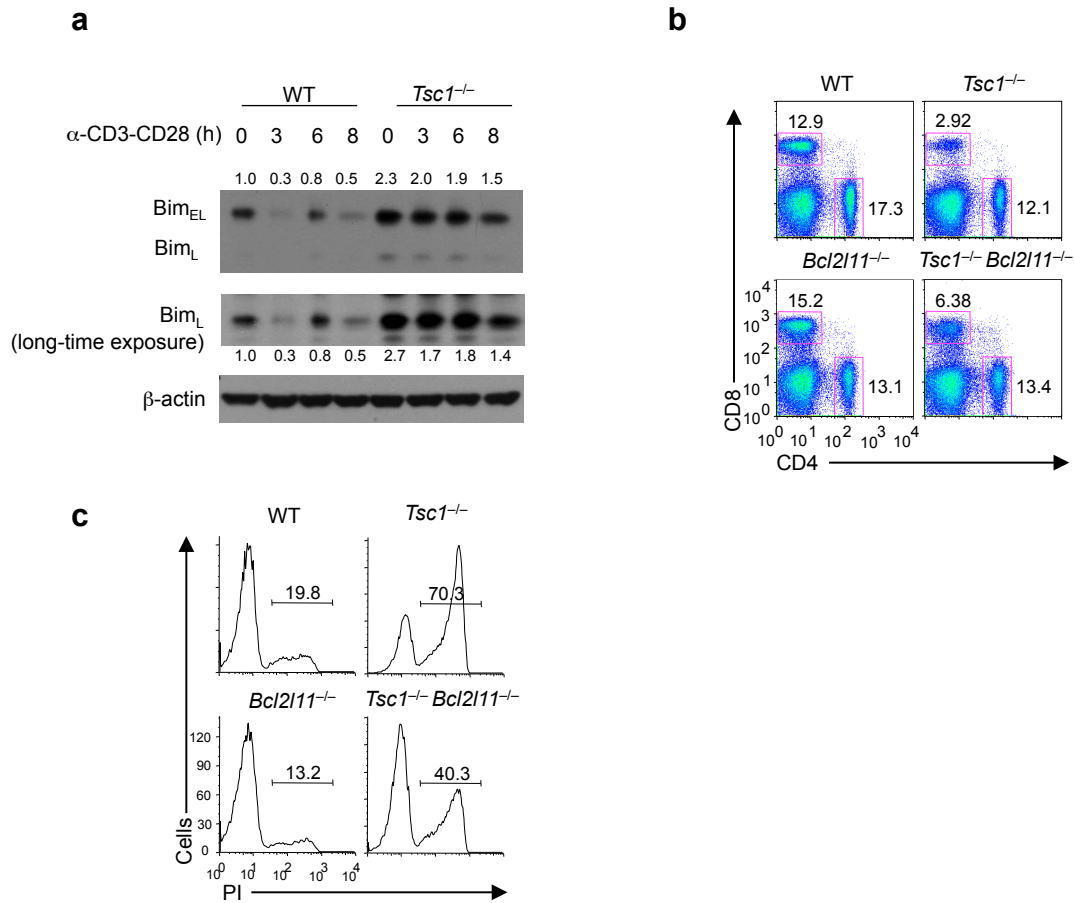
**Supplementary Figure 3. Altered mitochondrial homeostasis in *Tsc1*<sup>-/-</sup> T cells.** (a) CD4 and CD8 T cells from WT and *Tsc1*<sup>-/-</sup> mice were stained with MitoTracker Green and TMRM and analyzed by flow cytometry. (b) Ratios of *Tsc1*<sup>-/-</sup>/WT for the mean fluorescent intensity (MFI) of MitoTracker Green and TMRM staining (n=5). Data are representative of 4 independent experiments.

## Supplementary Figure 4



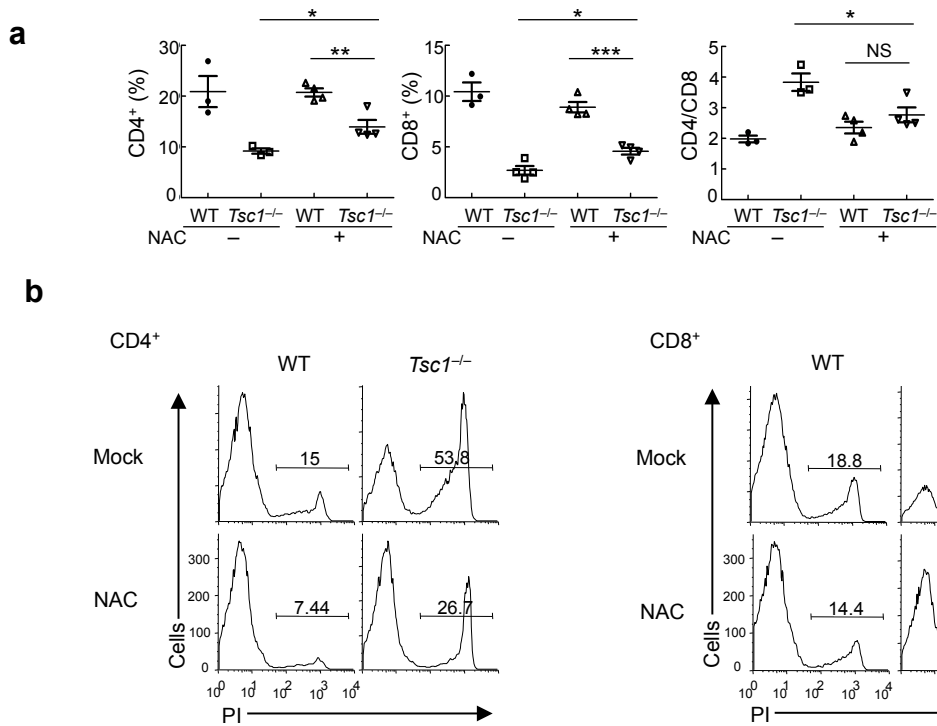
**Supplementary Figure 4. The *Bcl2* transgene rescues the diminished iNKT cells in *Tsc1*<sup>-/-</sup> mice.** Flow cytometry of iNKT cells (TCRβ<sup>+</sup>CD1d-PBS57<sup>+</sup>) in the spleens and livers of *Bcl2*-TG and *Tsc1*<sup>-/-</sup> *Bcl2*-TG mice. Data are representative of 3 independent experiments.

## Supplementary Figure 5



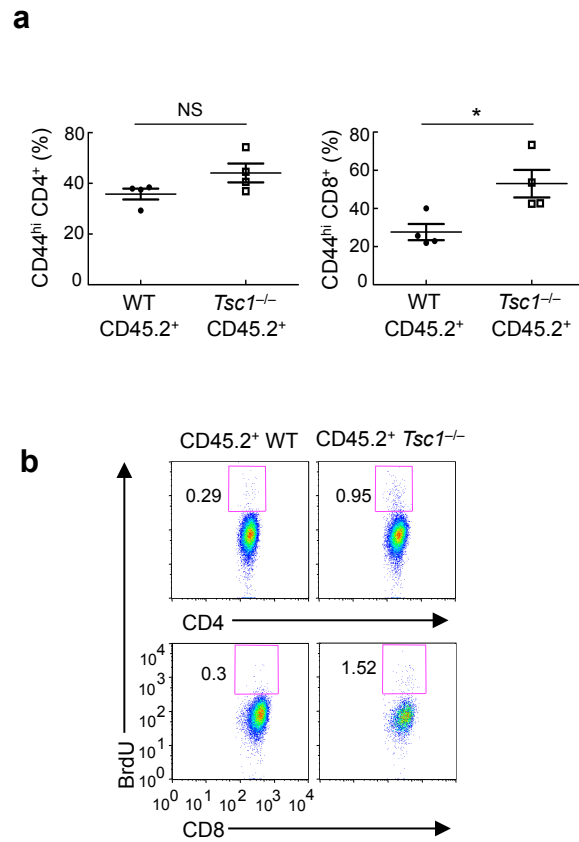
**Supplementary Figure 5. Role of Bim in apoptosis of *Tsc1*-deficient T cells.** (a) Western blot of Bim<sub>EL</sub> and Bim<sub>L</sub> expression in WT and *Tsc1*<sup>-/-</sup> CD8 T cells. A long-time exposure was shown in the middle panel to highlight Bim<sub>L</sub> levels. Numbers above (Bim<sub>EL</sub>) and below (Bim<sub>L</sub>) lanes indicate band intensity relative to that of the loading control  $\beta$ -actin. (b) Flow cytometry of CD4 and CD8 T cells in the spleens of WT, *Tsc1*<sup>-/-</sup>, *Bcl2/11*<sup>-/-</sup>, and *Tsc1*<sup>-/-</sup> *Bcl2/11*<sup>-/-</sup> mice. (c) Flow cytometry detection of PI<sup>+</sup> apoptotic cells after 16 hours of stimulation of CD4 T cells with anti-CD3-CD28. Data are representative of 2 independent experiments.

## Supplementary Figure 6



**Supplementary Figure 6. NAC treatment partially rescues the increased apoptosis of *Tsc1*<sup>-/-</sup> T cells.** (a) WT and *Tsc1*<sup>-/-</sup> mice were treated with NAC or mock for 5 weeks, followed by flow cytometry of CD4 and CD8 T cells in the spleens (n=3-4). \*  $P < 0.05$ ; \*\*  $P < 0.005$ ; \*\*\*  $P < 0.0005$ ; NS, not significant. (b) Naïve CD4 (left) and CD8 (right) T cells were purified from these mice and stimulated with anti-CD3-CD28 for 16 hours, followed by flow cytometry of PI<sup>+</sup> apoptotic cells. Data are representative of 2 independent experiments.

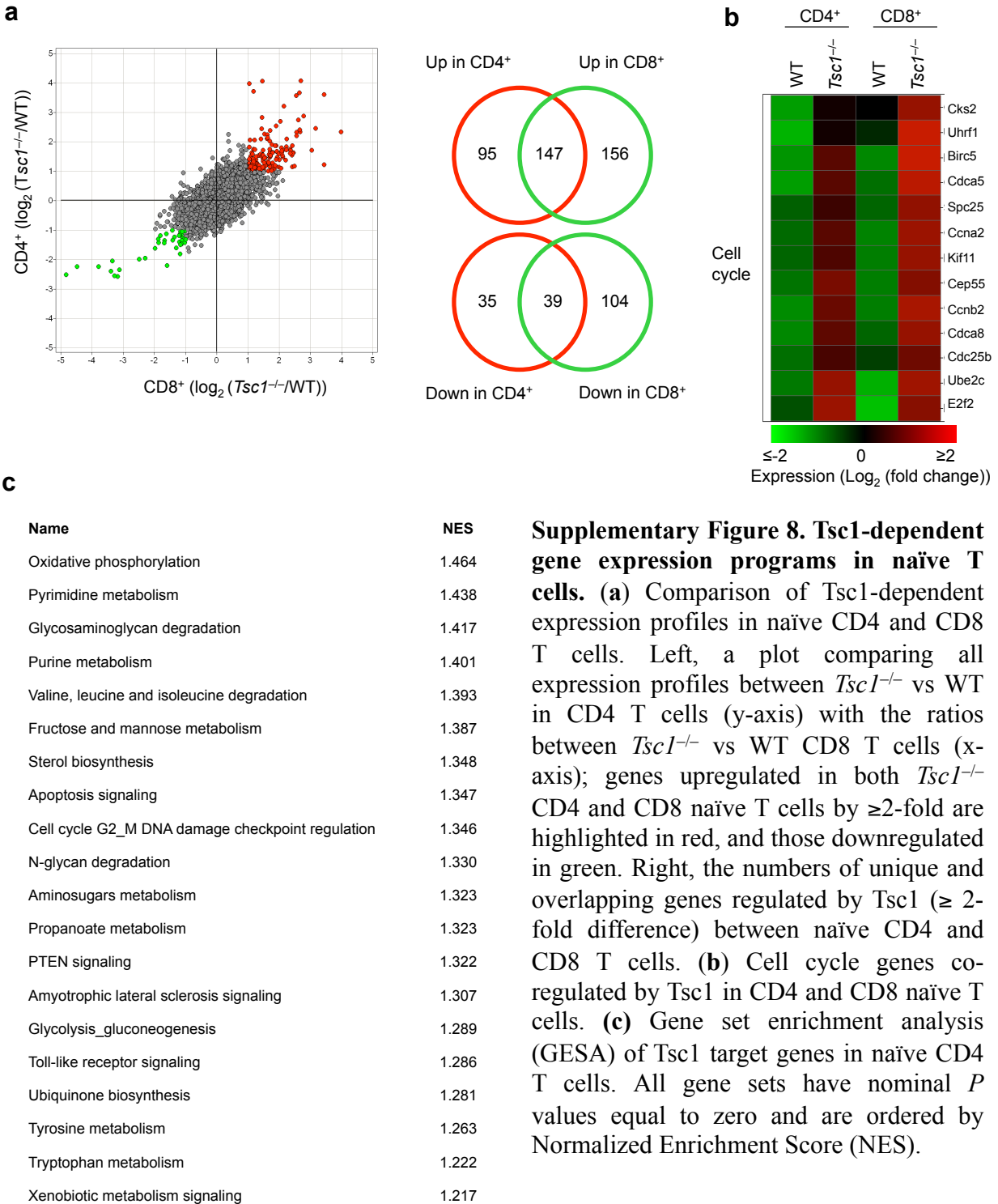
## Supplementary Figure 7



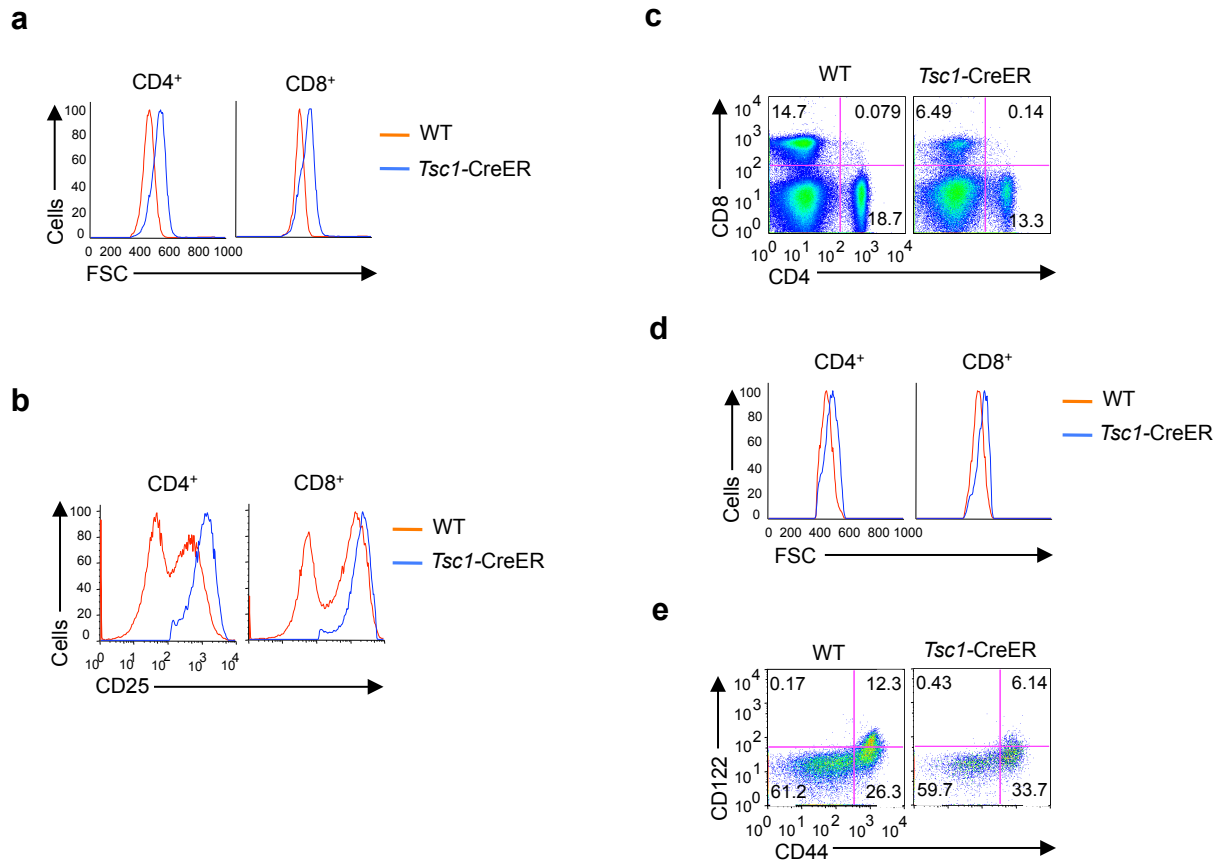
**Supplementary Figure 7. Analysis of T cell homeostasis and *in vivo* cycling in mixed bone marrow chimeras. (a)** Proportions of CD44<sup>hi</sup> populations in CD45.2<sup>+</sup> donor-derived CD4 (left) and CD8 (right) cells in the spleens of mixed bone marrow chimeras. \*  $P < 0.05$ ; NS, not significant. **(b)** BrdU staining in CD45.2<sup>+</sup> donor-derived naïve cells in the mixed bone marrow chimeras 16 hours after injection with BrdU. Data are representative of 2 independent experiments.



## Supplementary Figure 8



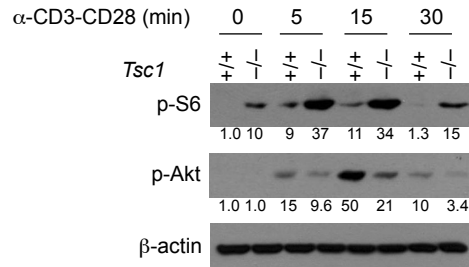
## Supplementary Figure 9



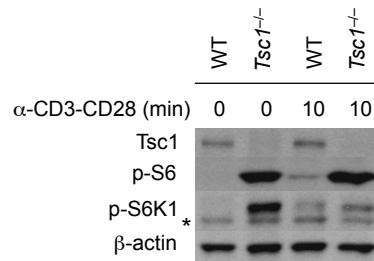
**Supplementary Figure 9. Effects of inducible deletion of *Tsc1* on T cell quiescence and homeostasis.** (a,b) WT and *Tsc1*-CreER mice were treated with tamoxifen, and 2 weeks later, naïve T cells were purified and stimulated with anti-CD3-CD28 for 16 hours. Cell size (a) and expression of activation marker CD25 (b) in stimulated cells were examined by flow cytometry. (c-e) WT and *Tsc1*-CreER mice were treated with tamoxifen, and analyzed 5 weeks later for the peripheral T cell populations (c), cell size (d), and distribution of CD44<sup>hi</sup>CD122<sup>+</sup> and CD44<sup>hi</sup>CD122<sup>-</sup> populations (e). Data are representative of 2 independent experiments.

## Supplementary Figure 10

**a**

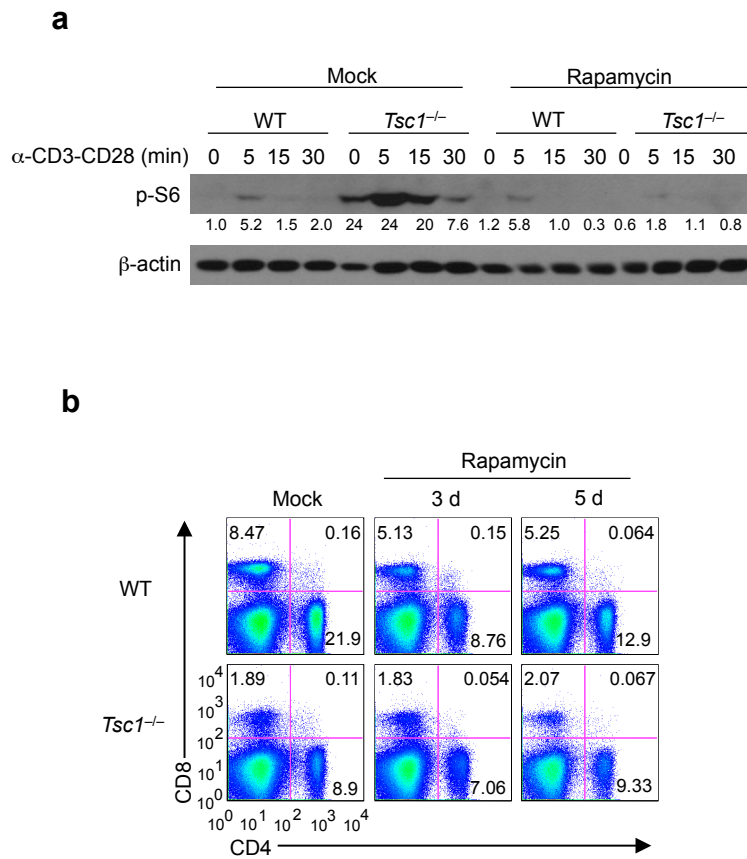


**b**



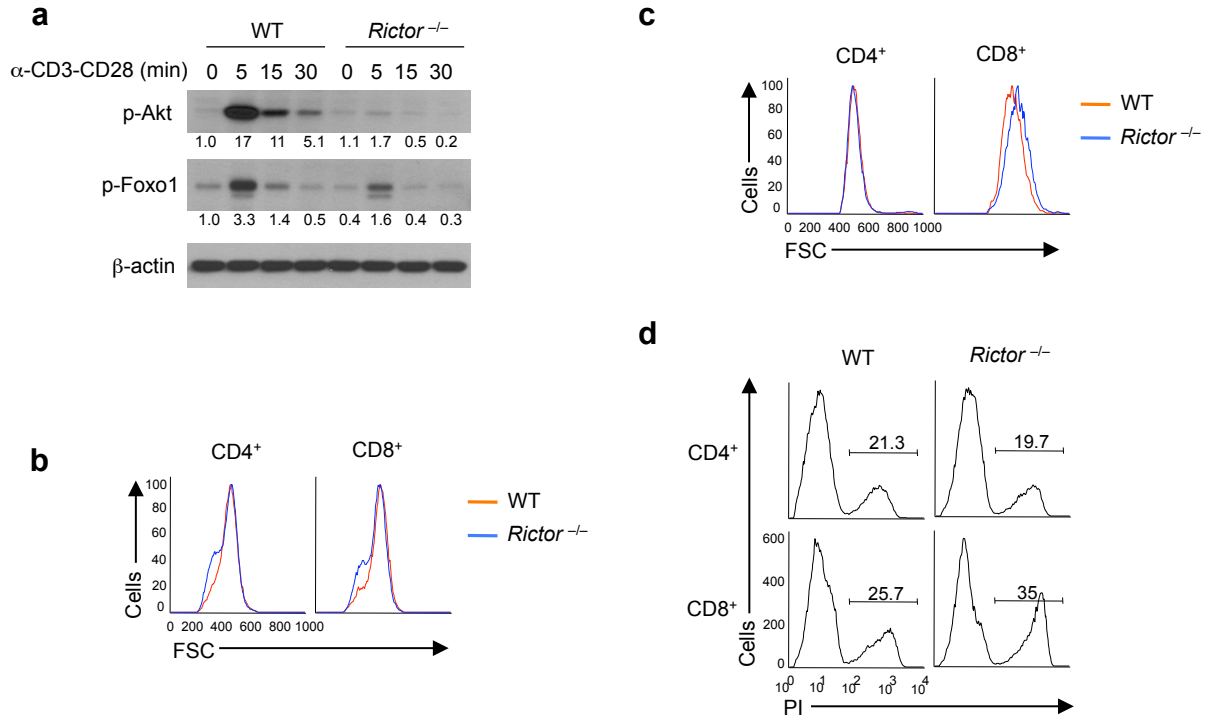
**Supplementary Figure 10. *Tsc1* regulates mTOR signaling in CD8 T cells and mature thymocytes. (a)** Western blot of S6 and Akt (Ser473) phosphorylation in WT and *Tsc1*<sup>-/-</sup> naïve CD8 T cells before and after anti-CD3-CD28 stimulation. Numbers below lanes indicate band intensity relative to that of the loading control  $\beta$ -actin. **(b)** Western blot of S6 and S6K1 phosphorylation in WT and *Tsc1*<sup>-/-</sup> CD4SP thymocytes before and after anti-CD3-CD28 stimulation. \*, non-specific bands. Data are representative of 2-3 independent experiments.

## Supplementary Figure 11



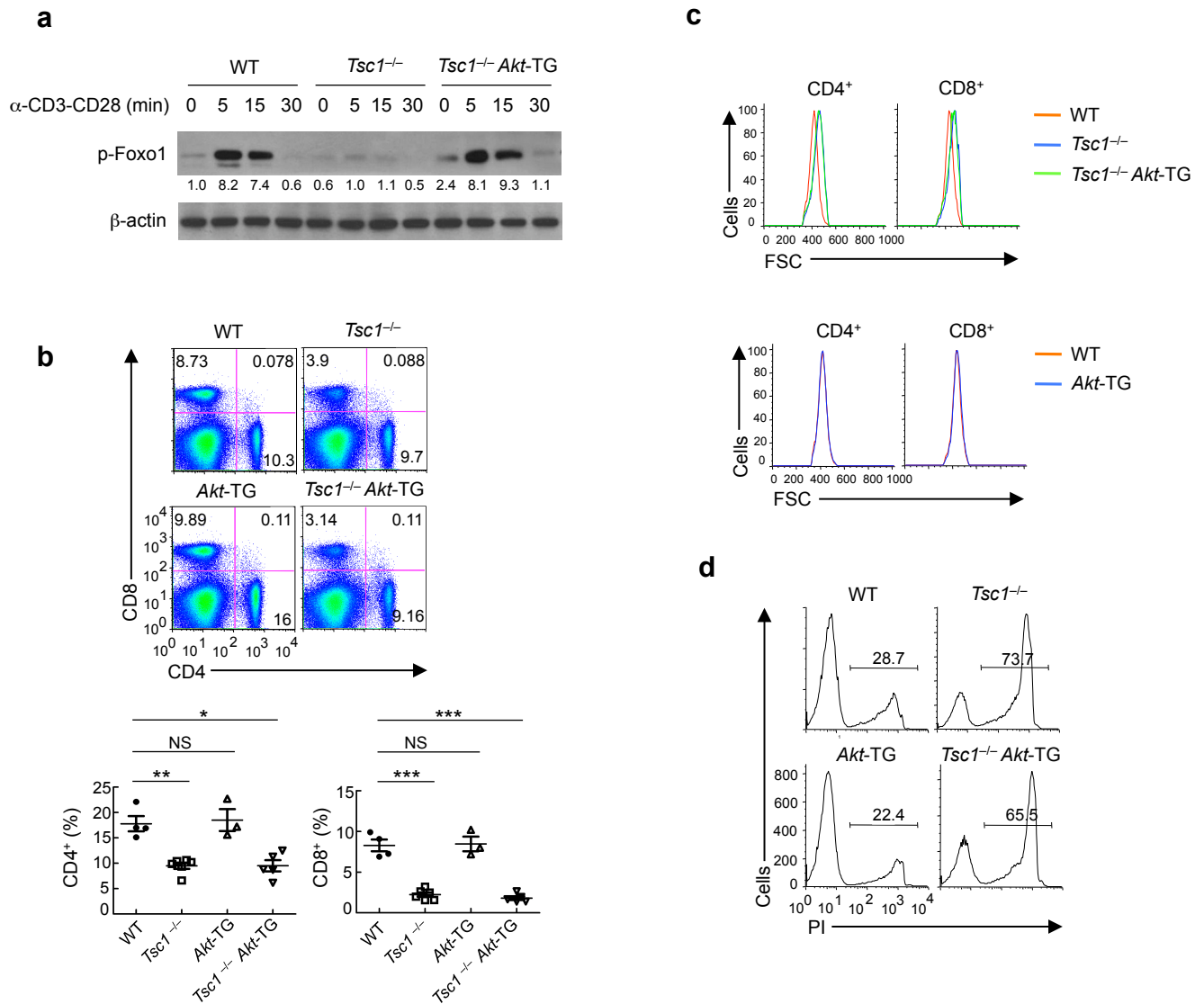
**Supplementary Figure 11. Effects of short-term rapamycin treatment on *Tsc1<sup>-/-</sup>* T cells.** (a) Sorted naïve T cells from WT and *Tsc1<sup>-/-</sup>* mice following rapamycin or mock treatment were activated with anti-CD3-CD28 for various times, and phosphorylation of S6 was examined by Western blots. Numbers below lanes indicate band intensity relative to that of the loading control  $\beta$ -actin. (b) Flow cytometry of CD4 and CD8 T cells in the spleens of rapamycin or mock treated WT and *Tsc1<sup>-/-</sup>* mice. Data are representative of 2-3 independent experiments.

## Supplementary Figure 12



**Supplementary Figure 12. Analysis of homeostasis and activation of *Rictor*<sup>-/-</sup> T cells.** (a) Western blot of Akt (Ser473) and Foxo1 phosphorylation in WT and *Rictor*<sup>-/-</sup> naïve T cells before and after anti-CD3-CD28 stimulation. Numbers below lanes indicate band intensity relative to that of the loading control  $\beta$ -actin. (b) Flow cytometry of cell size for WT and *Rictor*<sup>-/-</sup> splenocytes. (c,d) Sorted naïve T cells from WT and *Rictor*<sup>-/-</sup> mice were activated with anti-CD3-CD28 for 16 hours, followed by measurements of cell size (c) and PI<sup>+</sup> apoptotic cells (d). Data are representative of 2-3 independent experiments.

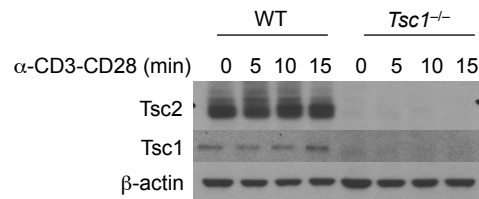
## Supplementary Figure 13



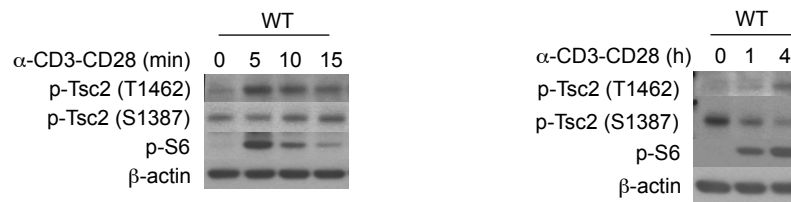
**Supplementary Figure 13. The *Akt*-transgene fails to rectify the defective homeostasis and activation of *Tsc1*<sup>-/-</sup> T cells.** (a) Western blot analysis of Foxo1 phosphorylation in WT, *Tsc1*<sup>-/-</sup>, *Akt*-TG, and *Tsc1*<sup>-/-</sup> *Akt*-TG naïve T cells before and after anti-CD3-CD28 stimulation. Numbers below lanes indicate band intensity relative to that of the loading control  $\beta$ -actin. (b) Flow cytometry of CD4 and CD8 T cells in the spleens of WT, *Tsc1*<sup>-/-</sup>, *Akt*-TG, and *Tsc1*<sup>-/-</sup> *Akt*-TG mice. Lower panels, proportions and absolute numbers of CD4<sup>+</sup> and CD8<sup>+</sup> T cells in the spleens of WT and *Tsc1*<sup>-/-</sup> mice (n=3-6). \*  $P < 0.01$ ; \*\*  $P < 0.001$ ; \*\*\*  $P < 0.001$ ; NS, not significant. (c) Flow cytometry of cell size of WT, *Tsc1*<sup>-/-</sup> and *Tsc1*<sup>-/-</sup> *Akt*-TG T cells (upper panels). For clarity, comparison between WT and *Akt*-TG T cells was shown in the lower panels. (d) Sorted naïve T cells from WT, *Tsc1*<sup>-/-</sup>, *Akt*-TG, and *Tsc1*<sup>-/-</sup> *Akt*-TG mice were activated with anti-CD3-CD28 for 16 hours, followed by flow cytometry detection of PI<sup>+</sup> apoptotic cells. Data are representative of 3 independent experiments.

## Supplementary Figure 14

**a**

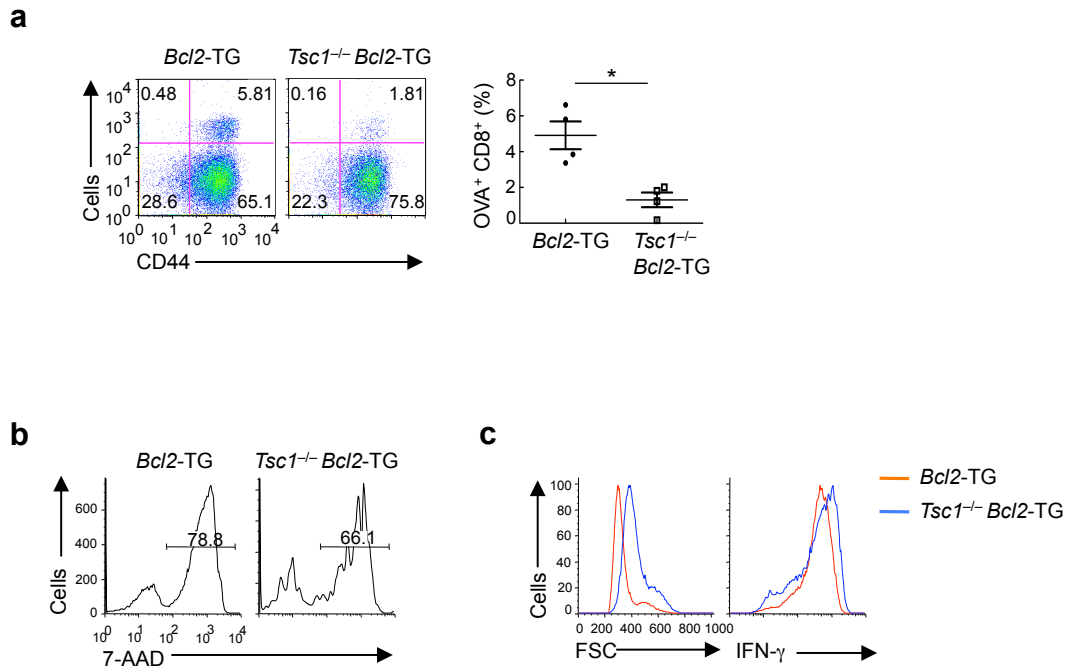


**b**



**Supplementary Figure 14. Regulation of Tsc1 and Tsc2 in T cells.** (a) Western blot of Tsc1 and Tsc2 expression in WT and *Tsc1*<sup>-/-</sup> naïve T cells before and after anti-CD3-CD28 stimulation. (b) Western blot of Tsc2 phosphorylation in WT naïve T cells after they were stimulated with anti-CD3-CD28 for short-term (left) or long-term (right) time points. Data are representative of 2-3 independent experiments.

## Supplementary Figure 15



**Supplementary Figure 15. Effects of *Tsc1* deficiency on CD8 T cell response and survival *in vivo* and differentiation *in vitro*.** (a,b) *Bcl2*-TG and *Tsc1*<sup>-/-</sup> *Bcl2*-TG mice were infected with LM-OVA. After 6-7 days, OVA-reactive tetramer-positive CD8 T cells were analyzed by flow cytometry (a). \*  $P < 0.01$ . Tetramer-positive CD8 T cells were purified, cultured in the medium overnight, and stained with 7-AAD to measure the extent of cell death (b). (c) Naïve CD8 T cells from *Bcl2*-TG and *Tsc1*<sup>-/-</sup> *Bcl2*-TG mice were stimulated with anti-CD3, anti-CD28 and IL-2 for 5 days, followed by brief PMA-ionomycin stimulation and analysis of cell size (left) and intracellular staining of IFN- $\gamma$  (right). Data are representative of 2 (a,b) and 4 (c) independent experiments.



Table S1. 186 probe sets with  $\geq 2$ -fold difference in both CD4 *Tsc1*<sup>-/-</sup> vs WT and CD8 *Tsc1*<sup>-/-</sup> vs WT naïve T cells

HT_HG430_PM	Gene Symbol	CD4 WT 0h	CD4 KO 0h	CD8 WT 0h	CD8 KO 0h
		Average	Average	Average	Average
1419758_PM_at	Abcb1a	4.41	8.49	6.30	8.99
1421920_PM_a_at	Ccr9	5.19	9.25	9.59	11.05
1427419_PM_x_at	Ccr9	6.66	10.65	10.77	11.82
1421919_PM_a_at	Ccr9	2.65	6.37	6.77	7.95
1419759_PM_at	Abcb1a	4.66	8.33	6.27	8.91
1434428_PM_at	D330028D13Rik	4.86	8.47	3.99	7.43
1457976_PM_at	2010002M12Rik	4.66	8.23	4.64	7.18
1421492_PM_at	Hpgds	2.84	6.05	2.78	4.93
1434909_PM_at	Rragd	3.97	6.85	3.87	6.66
1453748_PM_a_at	Kif23	3.40	6.23	5.50	6.95
1421396_PM_at	Pcsk1	2.71	5.46	3.16	5.77
1418722_PM_at	Ngp	3.46	6.16	4.30	6.34
1419147_PM_at	Rec8	3.11	5.66	2.59	4.49
1437445_PM_at	Trpm1	3.67	6.13	6.12	9.29
1449310_PM_at	Ptger2	3.53	5.97	3.53	4.97
1430735_PM_at	Patl2	3.66	6.07	3.90	6.56
1435313_PM_at	Cd200r4	4.92	7.31	5.37	7.08
1420805_PM_at	Myl10	4.50	6.84	3.69	7.67
1425385_PM_a_at	Igh	2.67	5.01	3.04	5.71
1417094_PM_at	Acot7	5.19	7.52	5.44	7.99
1419598_PM_at	Ms4a6d	3.64	5.91	4.62	6.15
1419599_PM_s_at	Ms4a6d	4.99	7.24	6.18	7.80
1430210_PM_at	C330024D12Rik	4.93	7.14	5.05	7.61
1440666_PM_at	4933430I17Rik	2.17	4.32	2.20	3.83
1438311_PM_at	2010002M12Rik	2.79	4.90	2.66	4.27
1418025_PM_at	Bhlhe40	4.47	6.57	4.67	6.91
1439831_PM_at	---	3.44	5.53	3.19	4.60
1451335_PM_at	Plac8	6.96	9.03	10.54	11.60
1455990_PM_at	Kif23	6.69	8.75	8.27	9.48
1420375_PM_at	Kif3a	4.54	6.57	3.68	5.38
1424305_PM_at	Igj	6.13	8.04	6.07	7.83
1434313_PM_at	Ccdc126	6.44	8.32	6.11	7.98
1419691_PM_at	Camp	3.42	5.29	3.30	5.20
1452954_PM_at	Ube2c	3.58	5.45	3.11	5.53
1448944_PM_at	Nrp1	4.62	6.47	3.96	6.19
1429524_PM_at	Myo1f	3.36	5.18	6.95	8.00
1424400_PM_a_at	Aldh1l1 /// LOC100047937	3.19	4.99	2.72	4.79
1434437_PM_x_at	Rrm2	4.86	6.61	4.87	7.22
1448756_PM_at	S100a9	5.95	7.70	6.28	7.98
1420505_PM_a_at	Stxbp1	3.66	5.41	3.91	5.08
1452202_PM_at	Pde2a	7.68	9.42	8.89	10.11
1444040_PM_at	Lair1	4.49	6.22	4.27	6.50
1416802_PM_a_at	Cdca5	3.39	5.12	3.75	5.88
1426817_PM_at	Mki67	3.99	5.71	3.74	6.78
1422734_PM_a_at	Myb	5.92	7.64	5.68	7.16
1448226_PM_at	Rrm2	3.62	5.33	3.63	5.99
1450920_PM_at	Ccnb2	3.15	4.85	3.26	5.34

1430447_PM_a_at	Lair1	4.81	6.49	4.76	6.68
1419394_PM_s_at	S100a8	5.56	7.23	5.76	7.68
1421317_PM_x_at	Myb	7.87	9.54	7.61	9.32
1450194_PM_a_at	Myb	7.38	9.03	7.11	8.89
1424278_PM_a_at	Birc5	3.34	4.98	3.43	5.88
1452242_PM_at	Cep55	2.86	4.50	2.79	4.58
1449841_PM_at	Kif3a	4.67	6.31	3.39	5.47
1437602_PM_at	Fam71b	3.30	4.93	5.22	6.34
1455790_PM_at	E2f2	5.93	7.55	4.91	7.26
1440627_PM_at	Atp8a2	3.37	4.97	3.39	5.82
1448943_PM_at	Nrp1	5.36	6.96	4.28	6.27
1455442_PM_at	Slc6a19	2.78	4.36	7.68	8.83
1449084_PM_s_at	Sh3d19	4.47	6.05	3.61	5.01
1416343_PM_a_at	Lamp2	5.12	6.69	5.15	6.82
1436574_PM_at	1700029I01Rik	5.47	7.03	6.05	7.52
1436434_PM_at	E2f2	6.05	7.60	5.15	7.57
1455049_PM_at	Igsf3	3.46	5.00	3.99	5.47
1436847_PM_s_at	Cdca8	4.36	5.90	4.63	6.35
1426565_PM_at	Igf1r	5.06	6.60	4.87	6.98
1425702_PM_a_at	Enpp5	3.87	5.41	3.54	4.88
1435680_PM_a_at	Dpp7	3.42	4.94	3.63	5.07
1421375_PM_a_at	S100a6	5.08	6.59	4.84	6.56
1424354_PM_at	Tmem140	4.09	5.60	4.62	5.71
1451461_PM_a_at	Aldoc	4.17	5.67	3.77	5.03
1418084_PM_at	Nrp1	4.66	6.15	3.71	5.83
1421014_PM_a_at	Clybl	6.41	7.89	6.59	7.68
1415810_PM_at	Uhrf1	3.89	5.36	5.00	6.70
1454694_PM_a_at	Top2a	4.61	6.08	4.47	6.41
1441623_PM_at	---	2.73	4.19	2.91	4.46
1457117_PM_at	Nfe2l2	5.91	7.36	5.76	7.29
1425763_PM_x_at	Igh /// Igh-2 /// Igh-VJ558 /// LOC544903	7.23	8.66	7.41	8.53
1450646_PM_at	Cyp51	5.77	7.20	5.22	6.71
1456328_PM_at	Bank1	5.57	6.99	5.66	6.71
1429381_PM_x_at	Igh /// Igh-2 /// Igh-VJ558 /// LOC544903	6.71	8.13	6.85	8.02
1421653_PM_a_at	Igh /// Igh-2 /// Igh-VJ558 /// LOC544903	8.15	9.56	8.26	9.39
1416957_PM_at	Pou2af1	6.03	7.44	6.37	7.63
1459643_PM_at	---	2.81	4.21	2.96	4.86
1436545_PM_at	Dtx4	5.18	6.57	5.78	7.27
1439255_PM_s_at	Gpr137b /// Gpr137b-ps /// LOC100044979	4.95	6.34	4.08	6.08
1417910_PM_at	Ccna2	3.71	5.09	3.59	5.79
1422567_PM_at	Fam129a	4.34	5.69	4.80	6.00
1443884_PM_at	Thada	7.12	8.47	7.11	8.59
1430985_PM_at	1810027O10Rik	7.30	8.64	7.72	9.36
1416441_PM_at	Pgcp	3.06	4.40	3.24	4.43
1428562_PM_at	2210403K04Rik	3.74	5.07	3.65	4.83
1417300_PM_at	Smpdl3b	2.96	4.29	3.17	4.81
1455048_PM_at	Igsf3	3.83	5.14	4.08	5.74
1457120_PM_at	Itk	5.63	6.93	5.72	7.24
1427455_PM_x_at	Gm10883 /// Gm1420 /// Gm7202 /// Igk /// Igk-C /// Igk-V28 /// LOC100047628	8.73	10.02	8.36	10.15
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1417457_PM_at	Cks2	6.64	7.92	7.76	8.81

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1449925_PM_at	Cxcr3	4.80	6.07	6.82	8.48
1448613_PM_at	Ecm1	6.73	8.00	5.07	6.29
1439259_PM_x_at	Abhd4	4.94	6.20	5.07	6.21
1455692_PM_x_at	1700097N02Rik	6.86	8.11	7.03	8.21
1424784_PM_at	OTTMUSG00000010657	3.93	5.18	4.29	5.57
1425738_PM_at	LOC100047222 /// LOC676193	3.53	4.78	3.30	4.99
1424631_PM_a_at	Ighg	2.96	4.19	2.19	5.62
1421963_PM_a_at	Cdc25b	7.02	8.24	7.40	8.54
1436479_PM_a_at	Dpp7	4.93	6.14	4.48	6.34
1460423_PM_x_at	Igk-V1	4.58	5.79	3.94	5.30
1417911_PM_at	Ccna2	3.85	5.04	3.57	5.33
1416344_PM_at	Lamp2	6.83	8.02	6.83	8.39
1439067_PM_at	Lair1	4.49	5.66	3.82	6.31
1448314_PM_at	Cdc2a	3.49	4.67	3.36	5.06
1457915_PM_at	4833442J19Rik	4.04	5.21	3.80	4.89
1435306_PM_a_at	Kif11	2.83	4.00	2.63	4.53
1433942_PM_at	Myo6	7.06	8.23	3.59	4.67
1417898_PM_a_at	Gzma	5.01	6.18	4.85	6.15
1428094_PM_at	Lamp2	7.23	8.40	6.98	8.49
1446303_PM_at	Igf1r	3.70	4.86	3.81	4.98
1452583_PM_s_at	Galm	5.74	6.90	5.67	6.70
1428283_PM_at	Cyp2s1	6.11	7.26	5.12	7.55
1428967_PM_at	Igf1r	6.92	8.07	6.78	8.25
1417185_PM_at	Ly6a	10.86	12.00	9.27	11.26
1419402_PM_at	Mns1	5.51	6.65	6.15	7.20
1429525_PM_s_at	Myo1f	4.05	5.19	6.65	7.88
1455711_PM_at	Dtx4	3.21	4.35	3.82	5.03
1417976_PM_at	Ada	5.91	7.04	6.26	7.31
1434503_PM_s_at	Lamp2	8.21	9.34	7.98	9.37
1436419_PM_a_at	1700097N02Rik	7.18	8.31	7.40	8.67
1419153_PM_at	2810417H13Rik	6.14	7.24	5.84	7.90
1457825_PM_x_at	Tcn2	4.95	6.05	4.90	6.26
1448143_PM_at	Aldh2	6.79	7.88	4.33	5.66
1452463_PM_x_at	Gm10883	8.42	9.50	8.08	9.51
1442183_PM_at	Gm13238	5.43	6.50	5.57	6.72
1417783_PM_at	Als2	5.37	6.44	5.37	7.20
1420394_PM_s_at	Gp49a /// Lilrb4	4.80	5.87	5.09	6.86
1427660_PM_x_at	Gm10883 /// Gm1420 /// Gm7202 /// Igk /// Igk-C /// Igk-V28 /// LOC100047628	9.86	10.93	9.52	11.03
1439256_PM_x_at	Gpr137b-ps	4.76	5.81	4.60	5.82
1448646_PM_at	Wdr12	7.59	8.63	7.29	8.31
1455065_PM_x_at	Gnpda1	7.73	8.77	7.44	8.64
1456225_PM_x_at	Trib3	5.43	6.45	4.64	5.96
1452982_PM_at	Igf1r	6.73	7.75	6.45	8.20
1424118_PM_a_at	Spc25	3.47	4.48	3.46	5.11
1449048_PM_s_at	Rab4a	3.29	4.30	3.52	4.76
1439293_PM_at	BC031353	6.78	7.78	6.77	7.96
1450881_PM_s_at	Gpr137b	4.61	5.61	4.03	5.33
1452417_PM_x_at	2010205A11Rik /// Gm10883 /// Gm1420 /// Gm7202 /// Igk /// Igk-C /// Igk-V28 /// LOC100047628	9.71	10.71	9.36	10.86
1435545_PM_at	BC032203	7.59	6.59	6.91	5.49
1418989_PM_at	Ctse	7.93	6.92	6.93	5.90

1427442_PM_a_at	App	8.13	7.10	7.62	6.59
1448297_PM_a_at	Tnk2	7.97	6.93	7.93	6.77
1444005_PM_at	---	4.64	3.58	4.51	3.41
1459595_PM_at	---	5.84	4.72	5.53	3.92
1448298_PM_at	Tnk2	5.25	4.12	5.11	3.98
1433977_PM_at	Hs3st3b1	7.07	5.92	6.00	4.82
1423059_PM_at	Ptk2	4.86	3.69	4.10	2.92
1434473_PM_at	Slc16a5	7.80	6.62	7.91	6.70
1437303_PM_at	Il6st	9.06	7.88	9.04	7.24
1425245_PM_a_at	Rgs11	5.30	4.08	5.54	4.18
1418210_PM_at	Pfn2	5.10	3.84	4.25	3.16
1423415_PM_at	Gpr83	5.52	4.24	4.28	3.25
1428122_PM_s_at	Fam125b	6.44	5.14	6.34	4.46
1437868_PM_at	Fam46a	7.86	6.54	8.52	7.52
1437064_PM_at	Ar	5.42	4.09	5.57	4.16
1424455_PM_at	Gprasp1	7.80	6.46	7.18	6.05
1448485_PM_at	Ggt1	5.16	3.81	8.03	6.36
1418216_PM_at	Ggt5	6.60	5.21	5.59	4.39
1434601_PM_at	Amigo2	6.28	4.86	5.29	4.26
1421991_PM_a_at	Igfbp4	5.59	4.17	5.16	3.29
1456751_PM_x_at	A530021J07Rik	5.68	4.25	3.78	2.09
1436585_PM_at	BB182297	5.88	4.42	5.45	4.32
1455647_PM_at	Ar	5.59	4.10	5.70	4.53
1452714_PM_at	Tanc1	6.60	5.00	7.09	5.79
1421840_PM_at	Abca1	7.35	5.74	7.87	5.89
1455747_PM_at	Ggt5	6.12	4.32	4.87	3.69
1445450_PM_x_at	A530021J07Rik	6.04	4.10	4.45	2.15
1456878_PM_at	AI646023	4.21	2.23	5.81	3.32
1437406_PM_x_at	Igfbp4	9.82	7.78	9.05	5.71
1458977_PM_at	A530021J07Rik	6.15	3.95	3.80	2.21
1423757_PM_x_at	Igfbp4	10.28	8.05	9.45	5.67
1437405_PM_a_at	Igfbp4	11.60	9.37	10.72	6.23
1421992_PM_a_at	Igfbp4	8.88	6.53	7.94	4.82
1427127_PM_x_at	Hspa1b	8.42	6.03	7.90	4.51
1423756_PM_s_at	Igfbp4	10.70	8.19	9.74	4.91
1452318_PM_a_at	Hspa1b	9.11	6.57	8.71	5.43
1427126_PM_at	Hspa1b	8.25	5.68	7.58	4.40

Table S2. 342 probe sets with  $\geq 2$ -fold difference in CD4 *Tsc1*<sup>-/-</sup> vs WT cells after 4 h TCR stimulation (FDR<0.1)

HT_HG430_PM	Gene Symbol	CD4 WT 0h Average	CD4 KO 0h Average	CD4 WT 4h Average	CD4 KO 4h Average
1419758_PM_at	Abcb1a	4.41	8.49	4.36	8.32
1419759_PM_at	Abcb1a	4.66	8.33	4.39	8.08
1439831_PM_at	---	3.44	5.53	5.93	9.15
1419394_PM_s_at	S100a8	5.56	7.23	4.55	7.67
1418722_PM_at	Ngp	3.46	6.16	3.34	6.35
1434909_PM_at	Rragd	3.97	6.85	3.76	6.77
1448756_PM_at	S100a9	5.95	7.70	5.06	7.98
1447285_PM_at	---	4.01	5.09	6.94	9.80
1419691_PM_at	Camp	3.42	5.29	3.02	5.52
1420505_PM_a_at	Stxbp1	3.66	5.41	4.12	6.46
1417094_PM_at	Acot7	5.19	7.52	4.86	7.16
1451335_PM_at	Plac8	6.96	9.03	5.47	7.75
1421396_PM_at	Pcsk1	2.71	5.46	2.78	5.02
1426817_PM_at	Mki67	3.99	5.71	3.36	5.58
1444588_PM_at	---	2.49	3.07	3.18	5.33
1449310_PM_at	Ptger2	3.53	5.97	4.66	6.80
1435172_PM_at	Eomes	5.64	6.89	4.88	6.95
1427419_PM_x_at	Ccr9	6.66	10.65	3.83	5.87
1450070_PM_s_at	Pak1	5.25	5.25	5.15	7.17
1422804_PM_at	Serpnb6b	5.58	6.67	5.25	7.26
1452954_PM_at	Ube2c	3.58	5.45	2.80	4.80
1444103_PM_at	Epsti1	5.35	6.07	3.64	5.58
1457976_PM_at	2010002M12Rik	4.66	8.23	4.52	6.44
1427455_PM_x_at	Gm10883 /// Gm1420 /// Gm7202 /// Igh /// Igh-C /// Igh-V28 /// LOC100047628	8.73	10.02	8.07	9.99
1425763_PM_x_at	Igh /// Igh-2 /// Igh-VJ558 /// LOC544903	7.23	8.66	6.60	8.51
1435313_PM_at	Cd200r4	4.92	7.31	3.08	4.95
1429381_PM_x_at	Igh /// Igh-2 /// Igh-VJ558 /// LOC544903	6.71	8.13	6.09	7.96
1424305_PM_at	Igj	6.13	8.04	6.17	8.04
1416592_PM_at	Glrx	7.23	8.33	5.39	7.25
1437502_PM_x_at	Cd24a	4.23	4.87	4.90	6.75
1426001_PM_at	Eomes	5.67	6.84	5.21	7.06
1421653_PM_a_at	Igh /// Igh-2 /// Igh-VJ558 /// LOC544903	8.15	9.56	7.38	9.21
1418248_PM_at	Gla	3.90	4.72	3.31	5.13
1425947_PM_at	Ifng	2.77	3.28	4.68	6.50
1432282_PM_a_at	Tlcd2	3.15	3.46	3.40	5.21
1440387_PM_at	6820431F20Rik	3.41	4.09	4.13	5.92
1452417_PM_x_at	2010205A11Rik /// Gm10883 /// Gm1420 /// Gm7202 /// Igh /// Igh-C /// Igh-V28 /// LOC100047628	9.71	10.71	9.04	10.80
1435529_PM_at	Gm14446	10.52	11.70	9.64	11.39
1427660_PM_x_at	Gm10883 /// Gm1420 /// Gm7202 /// Igh /// Igh-C /// Igh-V28 /// LOC100047628	9.86	10.93	9.26	11.01
1424354_PM_at	Tmem140	4.09	5.60	5.23	6.96
1436996_PM_x_at	Lyz1	7.14	7.83	4.81	6.53
1453851_PM_a_at	Gadd45g	5.27	5.76	5.00	6.71

1435245_PM_at	Gls2	2.49	2.55	2.92	4.63
1452087_PM_at	Epsti1	10.00	11.03	7.18	8.87
1418191_PM_at	LOC100048346 /// Usp18	8.26	7.99	7.24	8.92
1421375_PM_a_at	S100a6	5.08	6.59	3.61	5.29
1416034_PM_at	Cd24a	4.07	4.24	4.26	5.94
1417898_PM_a_at	Gzma	5.01	6.18	3.25	4.92
1423547_PM_at	Lyz2	6.46	6.91	3.60	5.26
1455692_PM_x_at	1700097N02Rik	6.86	8.11	3.67	5.32
1425546_PM_a_at	Trf	4.11	4.16	3.14	4.79
1449025_PM_at	Ifit3	8.38	8.13	7.58	9.21
1420979_PM_at	Pak1	4.21	4.04	3.64	5.27
1457780_PM_at	Stx11	2.80	3.50	5.28	6.90
1421571_PM_a_at	LOC100045833 /// Ly6c1 /// Ly6c2	10.91	12.16	9.62	11.24
1422734_PM_a_at	Myb	5.92	7.64	7.15	8.77
1426208_PM_x_at	Plagl1	3.08	2.98	6.87	8.48
1418025_PM_at	Bhlhe40	4.47	6.57	6.19	7.80
1419647_PM_a_at	Ier3	3.14	3.10	4.30	5.90
1452463_PM_x_at	Gm10883	8.42	9.50	7.90	9.49
1439255_PM_s_at	Gpr137b /// Gpr137b-ps /// LOC100044979	4.95	6.34	4.28	5.87
1438311_PM_at	2010002M12Rik	2.79	4.90	2.72	4.30
1434437_PM_x_at	Rrm2	4.86	6.61	4.81	6.38
1454942_PM_at	Fam129a	5.18	5.99	4.86	6.43
1416593_PM_at	Glrx	6.28	7.45	4.26	5.82
1449864_PM_at	Il4	2.28	2.30	3.08	4.64
1434830_PM_at	Mxd1	5.72	5.92	7.15	8.68
1448226_PM_at	Rrm2	3.62	5.33	3.48	5.01
1422567_PM_at	Fam129a	4.34	5.69	4.51	6.03
1428660_PM_s_at	LOC100047963 /// Tor3a	5.00	5.44	4.46	5.98
1416046_PM_a_at	Fuca2	5.94	7.22	3.65	5.16
1449773_PM_s_at	Gadd45b	6.04	6.17	8.79	10.31
1452426_PM_x_at	---	6.07	6.95	5.17	6.66
1415810_PM_at	Uhrf1	3.89	5.36	4.13	5.63
1424923_PM_at	Serpina3g	3.83	4.92	3.98	5.47
1417910_PM_at	Ccna2	3.71	5.09	3.39	4.88
1455768_PM_at	Npc2	7.00	7.77	6.60	8.09
1419985_PM_s_at	Ccdc69	8.37	8.51	3.48	4.95
1424278_PM_a_at	Birc5	3.34	4.98	3.14	4.61
1448182_PM_a_at	Cd24a	5.96	6.38	6.54	8.01
1450971_PM_at	Gadd45b	7.06	7.10	9.59	11.06
1438169_PM_a_at	Frmd4b	3.84	4.73	4.94	6.41
1418872_PM_at	Abcb1b	5.58	6.87	5.87	7.33
1416503_PM_at	Lxn	5.62	6.53	5.74	7.19
1454694_PM_a_at	Top2a	4.61	6.08	4.66	6.11
1429488_PM_at	Zdhhc21	6.98	7.14	5.68	7.13
1420394_PM_s_at	Gp49a /// Lilrb4	4.80	5.87	5.79	7.23
1424638_PM_at	Cdkn1a	3.83	3.91	4.98	6.42
1436574_PM_at	1700029I01Rik	5.47	7.03	4.16	5.60

1424067_PM_at	Icam1	6.65	7.43	8.67	10.11
1416802_PM_a_at	Cdca5	3.39	5.12	3.82	5.25
1419599_PM_s_at	Ms4a6d	4.99	7.24	3.88	5.31
1422527_PM_at	H2-DMa	7.85	8.96	6.90	8.33
1418930_PM_at	Cxcl10	3.43	3.15	8.17	9.59
1455104_PM_at	Mxd1	4.86	5.38	6.05	7.47
1436479_PM_a_at	Dpp7	4.93	6.14	3.35	4.76
1449874_PM_at	Ly96	2.95	4.57	2.92	4.31
1436576_PM_at	Fam26f	8.16	9.33	7.90	9.29
1421998_PM_at	Tor3a	4.72	5.13	4.59	5.97
1426179_PM_a_at	Twsg1	2.95	3.17	3.56	4.93
1418649_PM_at	Egln3	3.38	3.72	3.22	4.59
1421317_PM_x_at	Myb	7.87	9.54	9.20	10.57
1438009_PM_at	Hist1h2ad	9.58	10.33	8.33	9.69
1421679_PM_a_at	Cdkn1a	3.44	3.80	4.82	6.19
1460314_PM_s_at	<small>Hist1h3a // Hist1h3b // Hist1h3c // Hist1h3d // Hist1h3e // Hist1h3f // Hist1h3g // Hist1h3h // Hist1h3i // Hist1h3j // Hist1h3k // Hist1h3l</small>	6.59	6.54	5.13	6.49
1450194_PM_a_at	Myb	7.38	9.03	8.70	10.06
1422601_PM_at	Serpibn9	6.08	6.57	5.50	6.85
1453228_PM_at	Stx11	3.99	4.77	5.81	7.16
1419821_PM_s_at	Idh1	4.49	5.28	4.43	5.77
1454953_PM_at	Rnf157	3.73	4.45	5.24	6.58
1429400_PM_at	Clcn5	5.03	5.47	5.03	6.37
1455899_PM_x_at	Socs3	10.25	10.65	8.96	10.29
1448656_PM_at	Cacnb3	3.88	4.73	4.21	5.54
1423393_PM_at	Clic4	8.16	8.66	8.96	10.29
1454169_PM_a_at	Epsti1	8.74	9.97	6.42	7.75
1442183_PM_at	Gm13238	5.43	6.50	4.20	5.53
1449925_PM_at	Cxcr3	4.80	6.07	4.28	5.60
1440400_PM_at	Rnf157	3.27	4.28	5.93	7.23
1450291_PM_s_at	Ms4a4c	8.13	10.00	7.21	8.51
1448314_PM_at	Cdc2a	3.49	4.67	3.27	4.56
1417976_PM_at	Ada	5.91	7.04	6.03	7.32
1416910_PM_at	Dnajc15	9.17	10.50	8.26	9.54
1417909_PM_at	Serpinc1	3.09	3.24	4.45	5.74
1438606_PM_a_at	Clic4	7.87	8.42	8.68	9.97
1416309_PM_at	Nusap1	3.46	4.38	3.33	4.62
1417601_PM_at	Rgs1	7.61	8.22	8.61	9.89
1434372_PM_at	AW112010	10.74	11.81	9.99	11.27
1422948_PM_s_at	<small>Hist1h4a // Hist1h4b // Hist1h4c // Hist1h4d // Hist1h4e // Hist1h4f // Hist1h4g // Hist1h4h // Hist1h4i // Hist1h4j // Hist1h4k // Hist1h4l // Hist1h4m // LOC100041230</small>	8.11	8.26	7.40	8.68
1450886_PM_at	Gsg2	5.26	5.71	5.10	6.38
1416832_PM_at	Slc39a8	4.84	5.46	4.57	5.85
1422979_PM_at	Suv39h2	2.51	2.61	3.09	4.36
1429049_PM_at	Nuak2	5.51	6.16	5.57	6.83
1449852_PM_a_at	Ehd4	3.59	4.30	4.24	5.50
1456212_PM_x_at	Socs3	9.49	9.86	8.23	9.49
1455106_PM_a_at	Ckb	5.97	6.67	4.56	5.81
1417457_PM_at	Cks2	6.64	7.92	9.17	10.43

1427429_PM_at	Csf2	2.54	2.75	2.80	4.05
1443207_PM_at	---	3.14	3.35	4.12	5.37
1434428_PM_at	D330028D13Rik	4.86	8.47	4.89	6.14
1446670_PM_at	---	6.87	7.37	5.62	6.86
1424775_PM_at	Oas1a	7.62	8.31	6.64	7.89
1416702_PM_at	Serpini1	6.11	6.89	5.55	6.80
1416576_PM_at	Socs3	7.89	8.30	6.67	7.92
1431776_PM_at	Tob2	3.79	3.54	3.90	5.14
1419147_PM_at	Rec8	3.11	5.66	2.81	4.04
1439680_PM_at	Tnfsf10	6.13	5.99	3.59	4.82
1455790_PM_at	E2f2	5.93	7.55	3.46	4.69
1436847_PM_s_at	Cdca8	4.36	5.90	5.65	6.88
1428306_PM_at	Ddit4	9.15	9.56	5.91	7.14
1436201_PM_x_at	Mbp	2.52	3.12	4.34	5.57
1428468_PM_at	3110043O21Rik	5.42	5.00	6.07	7.27
1448148_PM_at	Grn	5.19	6.17	4.36	5.56
1441242_PM_at	---	6.10	6.38	4.87	6.07
1448143_PM_at	Aldh2	6.79	7.88	5.87	7.06
1426774_PM_at	Parp12	6.51	6.82	6.56	7.75
1426073_PM_at	Twsg1	2.78	2.70	3.31	4.50
1418764_PM_a_at	Bpnt1	5.36	5.49	5.70	6.88
1429863_PM_at	Lonrf3	2.60	2.83	3.20	4.38
1449168_PM_a_at	Akap2	4.51	4.54	4.45	5.62
1427202_PM_at	4833442J19Rik	7.10	8.09	7.17	8.34
1456956_PM_at	Ikzf2	4.85	4.82	4.75	5.92
1419297_PM_at	H2-Oa	5.65	6.53	3.55	4.72
1436026_PM_at	Zfp703	6.15	6.48	7.31	8.48
1421920_PM_a_at	Ccr9	5.19	9.25	3.34	4.50
1450476_PM_at	Cnr2	5.14	6.85	3.38	4.54
1418104_PM_at	Nrip3	2.38	2.47	3.16	4.32
1439790_PM_at	Serpib9	3.92	4.21	3.45	4.61
1436297_PM_a_at	Grina	7.91	8.60	6.63	7.78
1419598_PM_at	Ms4a6d	3.64	5.91	2.83	3.98
1416957_PM_at	Pou2af1	6.03	7.44	5.70	6.85
1424339_PM_at	Oasl1	6.15	6.31	5.24	6.39
1423392_PM_at	Clic4	4.63	5.45	5.54	6.69
1417698_PM_at	Gtf2f1	7.97	8.74	7.84	8.99
1419042_PM_at	ligp1	8.12	9.16	11.22	12.36
1415673_PM_at	Psph	7.60	8.27	6.14	7.28
1456006_PM_at	Bcl2l11	6.27	6.62	7.75	8.88
1452837_PM_at	Lpin2	5.56	6.56	5.47	6.61
1448449_PM_at	Ripk3	6.69	7.70	6.83	7.96
1424140_PM_at	Gale	4.56	5.53	4.88	6.01
1452123_PM_s_at	Frmd4b	4.14	5.01	5.45	6.58
1448303_PM_at	Gpnmb	3.72	5.11	3.73	4.85
1436419_PM_a_at	1700097N02Rik	7.18	8.31	4.23	5.36
1449363_PM_at	Atf3	3.58	3.50	4.90	6.02



1440866_PM_at	Eif2ak2	5.41	5.41	5.36	6.48
1424118_PM_a_at	Spc25	3.47	4.48	3.74	4.86
1448944_PM_at	Nrp1	4.62	6.47	3.84	4.95
1425685_PM_at	Akr1b10	4.84	5.21	4.02	5.13
1426550_PM_at	Sidt1	8.09	8.69	6.31	7.42
1453003_PM_at	Sorl1	7.03	7.51	4.75	5.86
1429816_PM_at	Armc3	5.34	6.29	3.14	4.24
1455908_PM_a_at	Scpep1	5.08	5.95	4.61	5.71
1452836_PM_at	Lpin2	6.26	7.27	5.98	7.08
1431734_PM_a_at	Dnajb4	5.52	5.87	5.78	6.87
1420197_PM_at	Gadd45b	2.81	2.87	3.80	4.89
1440047_PM_at	Socs1	6.37	6.15	7.79	8.88
1451310_PM_a_at	Ctsl	6.57	7.92	7.64	8.73
1436545_PM_at	Dtx4	5.18	6.57	4.97	6.06
1457673_PM_at	6820431F20Rik	4.13	4.07	6.56	7.65
1439040_PM_at	Cenpe	2.53	3.15	2.52	3.60
1456036_PM_x_at	Gsto1	6.37	6.66	5.76	6.84
1457445_PM_at	Trps1	6.00	5.81	6.02	7.10
1417654_PM_at	Sdc4	3.36	3.86	4.65	5.74
1437621_PM_x_at	Gm7669 /// Gm7901 /// Gm9347 /// Phgdh	8.70	9.14	7.08	8.16
1418932_PM_at	LOC100046232 /// Nfil3	3.62	3.62	4.81	5.89
1420671_PM_x_at	Ms4a4c	8.98	10.68	8.24	9.31
1452646_PM_at	Trp53inp2	3.34	4.08	5.45	6.52
1415822_PM_at	Scd2	8.19	9.17	5.60	6.66
1426258_PM_at	Sorl1	8.00	8.47	6.22	7.29
1426334_PM_a_at	Bcl2l11	2.62	3.05	4.36	5.42
1421688_PM_a_at	Ccl1	3.62	3.39	4.05	5.11
1423775_PM_s_at	Prc1	3.96	4.44	4.10	5.16
1418696_PM_at	Ttc36	2.98	3.05	3.81	4.87
1417911_PM_at	Ccna2	3.85	5.04	3.68	4.74
1448471_PM_a_at	Ctla2a	6.13	6.18	6.72	7.78
1452352_PM_at	Ctla2b	2.75	3.41	3.24	4.30
1429171_PM_a_at	Ncapg	2.39	3.23	2.39	3.45
1460042_PM_at	Slc23a3	2.45	2.30	3.60	4.66
1454714_PM_x_at	Gm7669 /// Gm7901 /// Gm9347 /// Phgdh	8.70	9.15	7.23	8.29
1449530_PM_at	Trps1	6.10	6.05	6.80	7.86
1456471_PM_x_at	<small>10527627 /// Gm1337 /// Gm1385 /// Gm7669 /// Gm7901 /// Gm8206 /// Gm8341 /// Gm8310 /// Gm8251 /// Gm8347 /// LOC320761 /// LOC320806 /// Phgdh</small>	7.39	7.83	5.95	7.00
1455990_PM_at	Kif23	6.69	8.75	5.12	6.18
1427364_PM_a_at	Odc1	4.80	5.00	7.41	8.46
1451828_PM_a_at	Acsl4	4.62	5.16	6.20	7.25
1436011_PM_at	Elmo2	4.60	4.66	5.59	6.64
1456228_PM_x_at	Mbp	2.79	3.13	4.65	5.69
1448943_PM_at	Nrp1	5.36	6.96	4.34	5.38
1422814_PM_at	Aspm	2.33	3.15	2.72	3.76
1418346_PM_at	Insl6	6.56	6.86	6.46	7.50
1451611_PM_at	Pla2g16	7.62	8.62	6.37	7.41
1435659_PM_a_at	Tpi1	6.79	7.16	5.47	6.51

1426665_PM_at	Katnb1	5.78	6.41	4.98	6.02
1450626_PM_at	Manba	6.93	7.66	5.51	6.55
1445597_PM_s_at	Pla2g16	8.84	9.67	7.58	8.62
1444152_PM_at	Cugbp2	7.06	7.40	5.67	6.71
1455065_PM_x_at	Gnpda1	7.73	8.77	6.62	7.66
1435517_PM_x_at	Ralb	4.87	5.54	4.05	5.08
1451776_PM_s_at	Hopx	6.40	6.79	5.03	6.06
1424724_PM_a_at	D16Ertd472e	7.41	7.47	7.11	8.14
1416343_PM_a_at	Lamp2	5.12	6.69	7.19	8.22
1441164_PM_at	---	2.77	3.55	2.80	3.83
1457944_PM_at	---	6.46	6.61	7.05	8.08
1424050_PM_s_at	Fgfr1	3.89	4.19	3.42	4.45
1449109_PM_at	Socs2	3.98	4.21	4.27	5.29
1435028_PM_at	Wdr7	6.69	7.26	5.08	6.11
1423717_PM_at	Ak3	6.87	7.13	5.43	6.46
1417458_PM_s_at	Cks2	6.97	8.30	9.82	10.85
1416344_PM_at	Lamp2	6.83	8.02	8.72	9.75
1452540_PM_a_at	<small>Hist1h2bc /// Hist1h2be /// Hist1h2b /// Hist1h2bm /// Hist1h2bp /// LOC100046213 /// LOC656522 /// RP23-38E20.1</small>	7.20	7.11	7.74	8.76
1420980_PM_at	Pak1	4.24	4.32	3.95	4.97
1450790_PM_at	Tg	2.59	2.32	2.98	4.00
1417491_PM_at	Ctsb	4.66	5.15	4.96	5.98
1420805_PM_at	Myl10	4.50	6.84	3.52	4.54
1434537_PM_at	Slco3a1	7.08	7.32	6.00	7.02
1425684_PM_at	Akr1b10	6.26	6.51	5.41	6.42
1428027_PM_at	---	2.71	2.75	5.99	7.00
1438910_PM_a_at	Stom	3.79	3.87	3.83	4.84
1428562_PM_at	2210403K04Rik	3.74	5.07	7.00	8.01
1450272_PM_at	Tnfsf8	7.53	7.58	6.78	7.78
1452927_PM_x_at	Tpi1	7.53	7.86	6.28	7.29
1430985_PM_at	1810027O10Rik	7.30	8.64	6.22	7.23
1418084_PM_at	Nrp1	4.66	6.15	3.81	4.81
1421529_PM_a_at	Txnrd1	7.74	7.95	7.30	8.30
1416619_PM_at	4632428N05Rik	9.77	9.25	8.97	7.97
1441415_PM_at	---	6.34	6.03	7.28	6.28
1436735_PM_at	Nsun3	5.67	5.60	5.52	4.51
1434353_PM_at	Sfmbt2	7.15	6.03	5.00	4.00
1445103_PM_at	---	2.89	2.91	3.77	2.75
1425532_PM_a_at	Bin1	7.74	7.38	6.76	5.74
1428288_PM_at	Klf9	4.92	4.64	6.13	5.11
1423389_PM_at	Smad7	9.29	8.75	9.08	8.05
1444299_PM_at	A430093F15Rik	7.92	8.05	7.15	6.11
1423161_PM_s_at	Spred1	3.04	2.72	4.29	3.26
1451899_PM_a_at	Gtf2ird1	6.43	5.23	6.34	5.30
1456851_PM_at	---	7.96	7.39	7.23	6.19
1448960_PM_at	Cxhc5	5.01	4.54	6.90	5.86
1417522_PM_at	Fbxo32	5.92	5.23	5.04	3.99
1455252_PM_at	Tsc1	7.43	6.62	7.55	6.49

1420754_PM_at	Ttf1	7.44	6.50	7.29	6.24
1455220_PM_at	Frat2	8.48	7.65	8.57	7.51
1449520_PM_at	Ttc28	7.66	6.81	7.48	6.41
1428122_PM_s_at	Fam125b	6.44	5.14	5.29	4.22
1439141_PM_at	Gpr18	9.36	8.39	9.40	8.33
1436892_PM_at	Spred2	5.76	5.24	7.12	6.04
1416107_PM_at	Nsg2	10.43	9.74	7.51	6.43
1451763_PM_at	Cnga1	5.98	5.46	4.56	3.47
1436329_PM_at	Egr3	7.90	7.66	9.68	8.59
1423059_PM_at	Ptk2	4.86	3.69	4.59	3.50
1429491_PM_s_at	Rif1	8.62	7.68	8.46	7.37
1440952_PM_at	---	8.49	7.80	7.59	6.48
1424838_PM_at	A330049M08Rik	5.68	6.14	6.28	5.17
1434027_PM_at	Rcan3	7.41	7.45	8.40	7.28
1460242_PM_at	Cd55	7.94	7.97	6.79	5.67
1423162_PM_s_at	Spred1	3.99	3.09	5.53	4.41
1434581_PM_at	2410066E13Rik	6.54	5.71	6.00	4.88
1438823_PM_at	---	3.96	3.61	6.28	5.15
1437584_PM_at	---	10.04	9.59	9.47	8.34
1427766_PM_at	<small>A130082M07Rik /// EG667677 /// ENMUSG00000076763 /// ENMUSG00000076765 /// Gm11046 /// LOC100444026 /// LOC677567 /// Tc4</small>	7.23	7.17	6.76	5.62
1435545_PM_at	BC032203	7.59	6.59	5.58	4.43
1426147_PM_s_at	Cldn10	5.23	4.93	4.85	3.70
1438712_PM_at	Dennd2d	7.61	7.17	6.35	5.20
1443553_PM_at	---	3.75	3.47	5.75	4.57
1448328_PM_at	Sh3bp2	5.43	5.71	7.24	6.06
1434891_PM_at	Ptgfrn	5.62	3.90	4.69	3.51
1460188_PM_at	Ptpn6	9.67	9.65	8.57	7.38
1450450_PM_at	Rcan3	5.87	6.00	6.75	5.56
1457549_PM_at	---	8.32	7.88	8.06	6.86
1460116_PM_s_at	Spred1	2.30	2.14	3.51	2.29
1436499_PM_at	Sgms1	8.87	8.76	8.13	6.90
1418762_PM_at	Cd55	8.21	7.96	7.75	6.52
1418836_PM_at	Qprt	6.49	6.10	6.63	5.40
1424032_PM_at	Hvcn1	9.01	9.12	7.02	5.78
1438805_PM_at	Ccnd3	8.72	8.37	8.67	7.42
1421594_PM_a_at	Sytl2	6.87	6.07	5.34	4.09
1454967_PM_at	---	7.14	6.67	8.15	6.89
1426576_PM_at	Sgms1	7.12	6.88	6.41	5.14
1437390_PM_x_at	Stx1a	7.77	6.86	8.18	6.91
1454745_PM_at	Arhgap29	7.10	6.59	6.93	5.66
1435265_PM_at	---	5.39	4.85	5.99	4.72
1446567_PM_at	---	7.91	7.36	7.48	6.20
1428682_PM_at	Zc3h6	6.77	6.31	5.55	4.25
1442437_PM_at	---	5.87	5.25	5.37	4.07
1423571_PM_at	S1pr1	10.22	9.83	10.09	8.79
1458802_PM_at	Hivep3	7.25	7.33	9.45	8.13
1418989_PM_at	Ctse	7.93	6.92	6.47	5.14

1423756_PM_s_at	Igfbp4	10.70	8.19	6.03	4.69
1448458_PM_at	Top2b	5.12	4.92	5.31	3.97
1454646_PM_at	Tcp11l2	8.79	8.40	8.53	7.15
1441286_PM_at	Gm13948	5.77	5.86	5.46	4.06
1419136_PM_at	Akr1c18	3.24	2.83	4.33	2.92
1434403_PM_at	Spred2	7.35	6.71	8.58	7.11
1423689_PM_a_at	Gpsm1	4.17	2.58	5.10	3.63
1448747_PM_at	Fbxo32	8.05	7.26	7.67	6.18
1448576_PM_at	Il7r	9.79	9.64	9.65	8.14
1451516_PM_at	Rhebl1	6.48	5.39	5.86	4.36
1436491_PM_at	Gm11346	9.57	9.11	8.17	6.67
1456509_PM_at	1110032F04Rik	4.81	3.79	4.65	3.11
1416630_PM_at	Id3	7.84	7.57	7.04	5.41
1421840_PM_at	Abca1	7.35	5.74	6.35	4.70
1448575_PM_at	Il7r	11.65	11.47	11.51	9.86
1417930_PM_at	Nab2	6.77	6.58	7.34	5.69
1433977_PM_at	Hs3st3b1	7.07	5.92	6.94	5.19
1436585_PM_at	BB182297	5.88	4.42	6.34	4.37