

Supplementary Information

The TSC-mTOR pathway regulates macrophage polarization

Vanessa Byles^{1,3}, Anthony J. Covarrubias^{1,3}, Issam Ben-Sahra¹, Dudley W. Lamming², David M. Sabatini², Brendan D. Manning¹, and Tiffany Horng¹

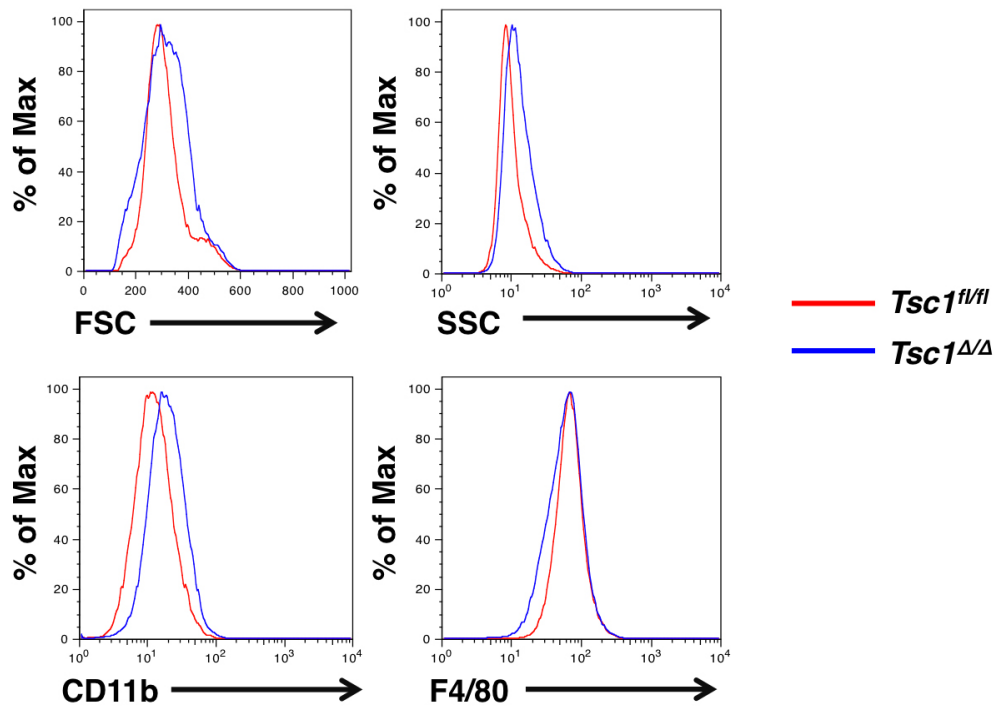
¹Department of Genetics & Complex Diseases, Harvard School of Public Health, Boston, Massachusetts

²Whitehead Institute for Biomedical Research, Cambridge, MA 02142; Department of Biology, MIT, Cambridge, MA 02139; Howard Hughes Medical Institute, MIT, Cambridge, MA 02139; Broad Institute of Harvard and MIT, Seven Cambridge Center, Cambridge, MA 02142; The David H. Koch Institute for Integrative Cancer Research at MIT, Cambridge, MA 02139

³These authors contributed equally to this work

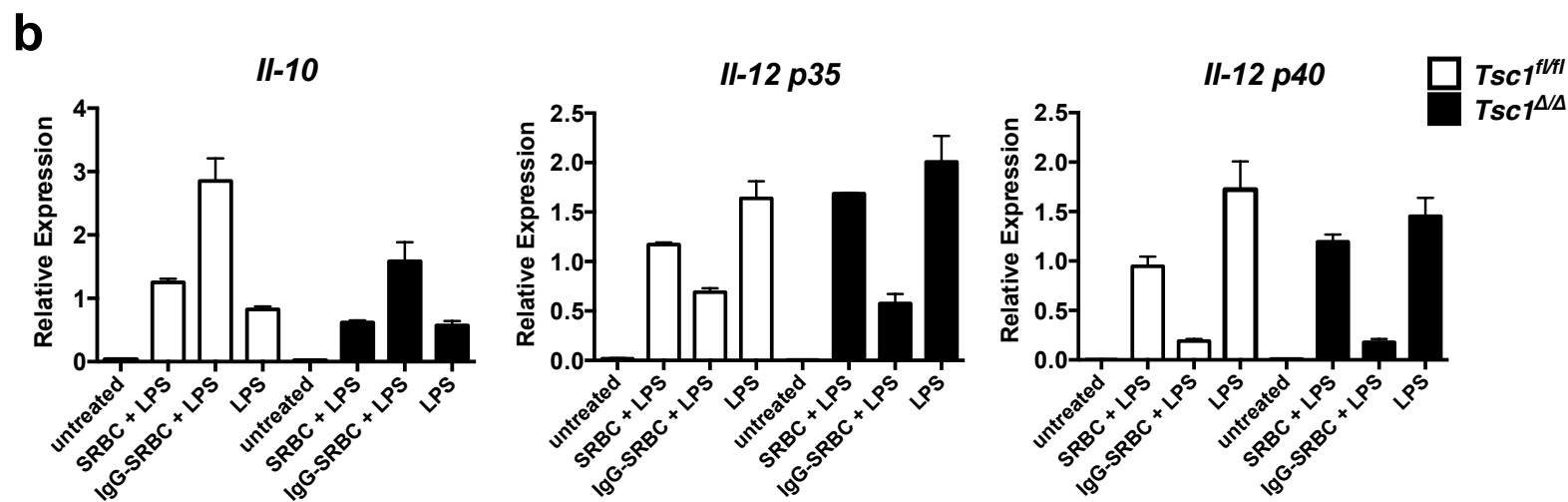
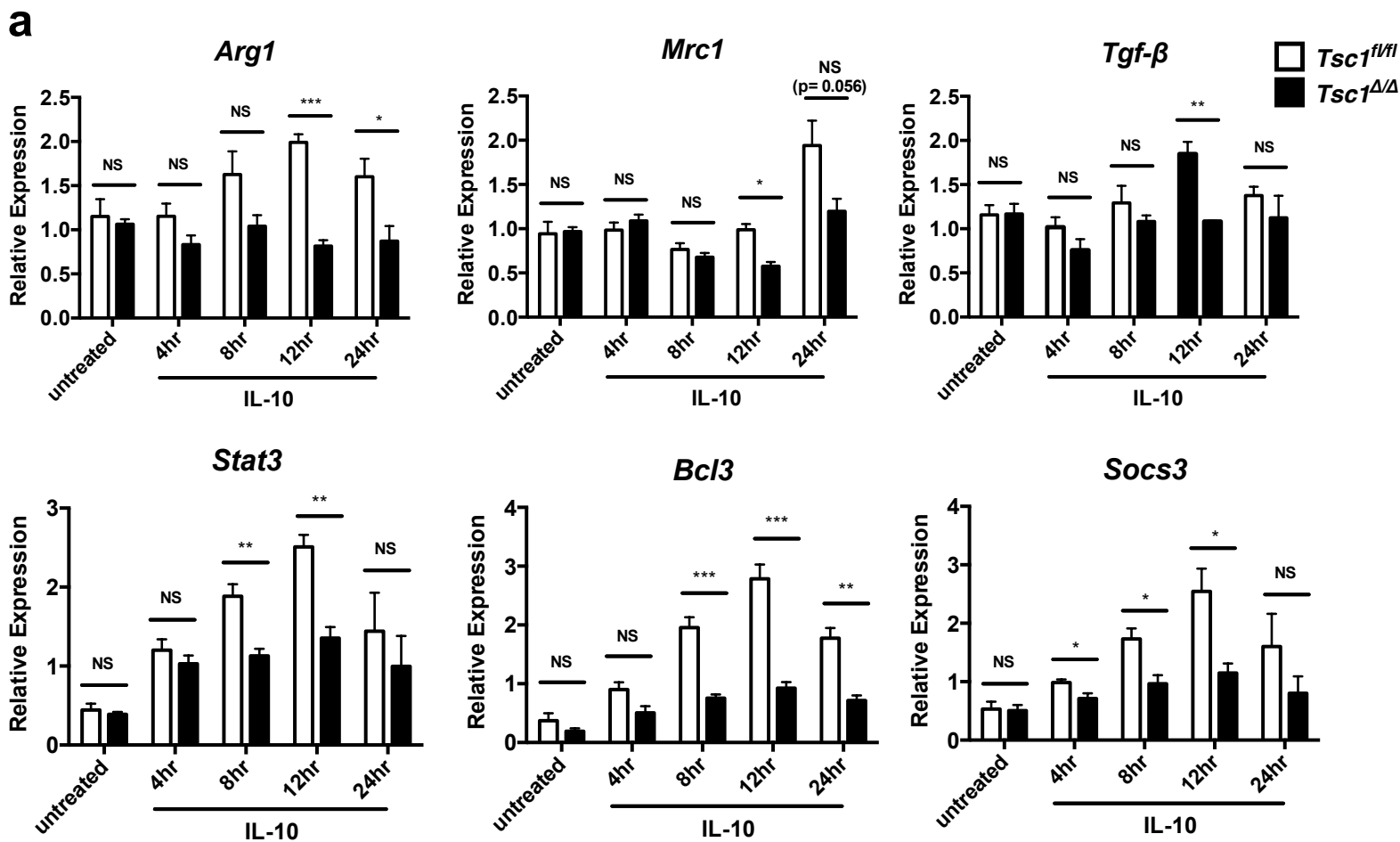
Corresponding author:
Tiffany Horng
Harvard School of Public Health
655 Huntington Ave, II-115
Boston, MA 02115
Ph: (617) 432-7526
F: (617) 432-5236
thorng@hsph.harvard.edu

Supplementary Figures S1-S9
Supplementary Table S1



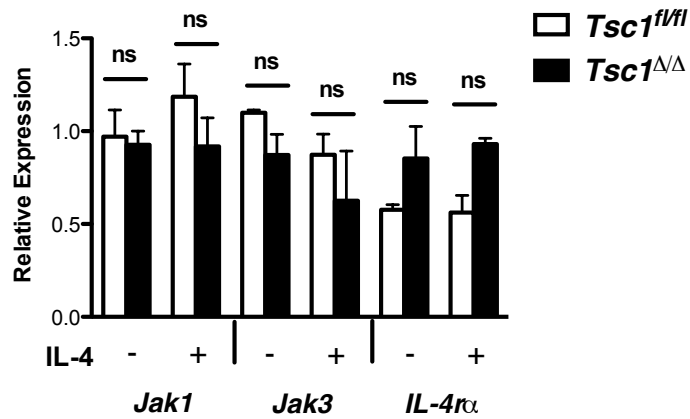
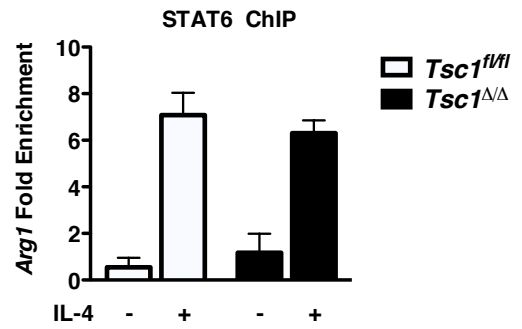
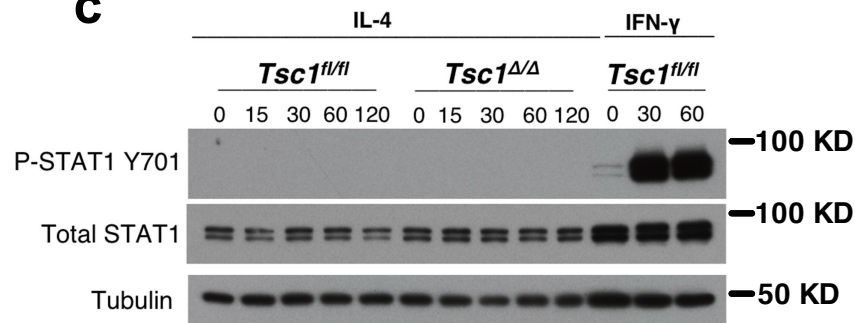
Supplementary Figure S1. Increased Cell Size and Granularity in *Tsc1^{Δ/Δ}* BMDMs

Flow cytometry analysis of *Tsc1^{fl/fl}* and *Tsc1^{Δ/Δ}* BMDMs stained with macrophage markers CD11b and F4/80. Histogram plots are shown as % of maximum. Red= *Tsc1^{fl/fl}* and Blue= *Tsc1^{Δ/Δ}*



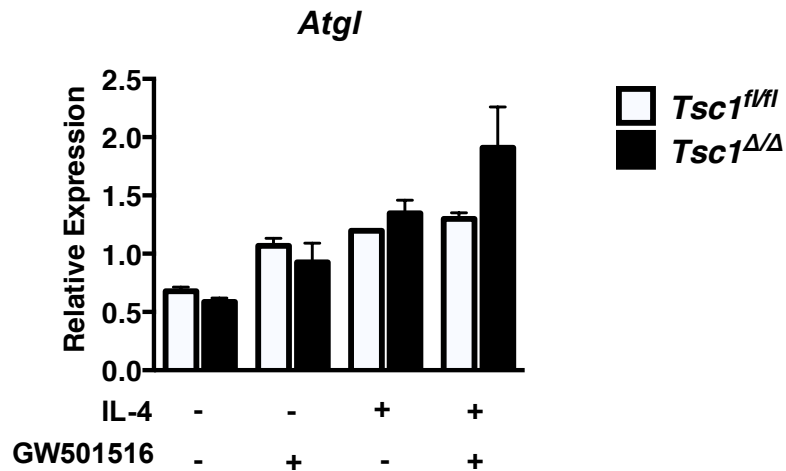
Supplementary Figure S2. Impaired M2c but not M2b Polarization in *Tsc1 ^{Δ/Δ}* BMDMs

a. M2c gene expression in *Tsc1^{fl/fl}* and *Tsc1 ^{Δ/Δ}* BMDMs stimulated with 10ng/ml IL-10 for the indicated times (n=2 representative experiments). **b.** M2b gene expression of *Tsc1^{fl/fl}* and *Tsc1 ^{Δ/Δ}* BMDMs stimulated simultaneously with LPS and non-opsonized (SRBC) or opsonized SRBC (IgG-SRBC) for 6h (Data representative of 3 independent experiments). Graphs are shown as mean \pm SEM, *p<0.05, **p<0.01, ***p<0.001 determined by Student's t-tests.

a**b****c**

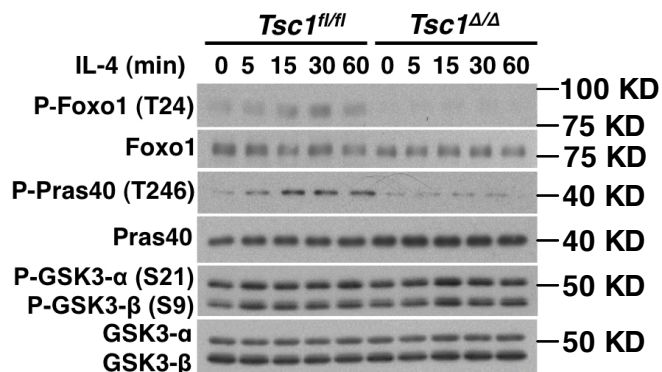
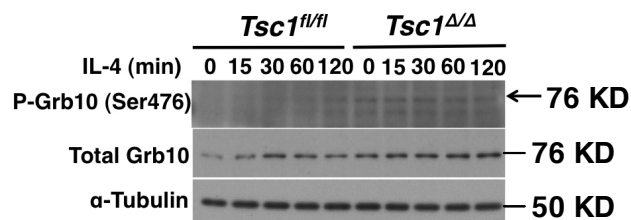
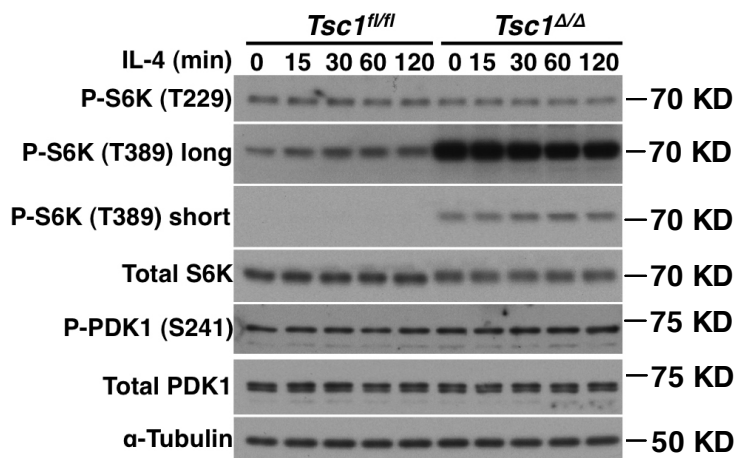
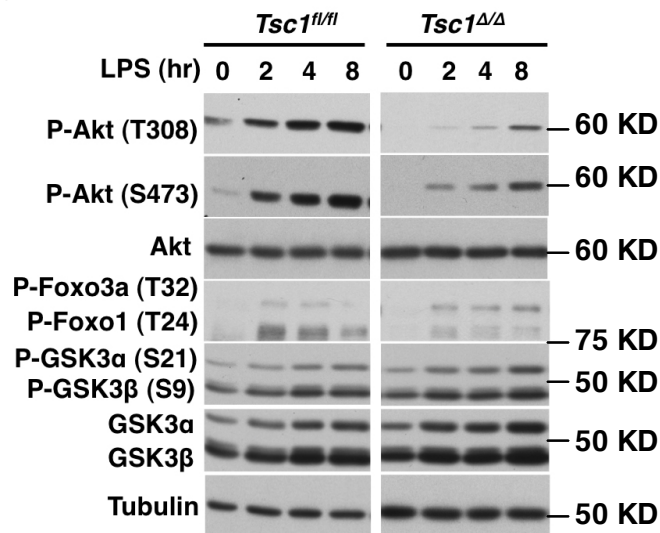
Supplementary Figure S3. The JAK-STAT6 Pathway is Intact in *Tsc1^{Δ/Δ}* BMDMs

a. Gene expression of components of the IL-4R-JAK-STAT6 pathway (n=2 representative experiments). **b.** STAT6 chromatin immunoprecipitation. Data is shown as fold enrichment for the *Arg1* promoter and is representative of 3 independent experiments. **c.** IL-4 fails to activate STAT1 in BMDMs. Immunoblot analysis of STAT1 activation in lysates from *Tsc1^{fl/fl}* and *Tsc1^{Δ/Δ}* BMDMs stimulated with IL-4 or IFN (control) for the indicated time points. Graphs are shown as mean \pm SEM.



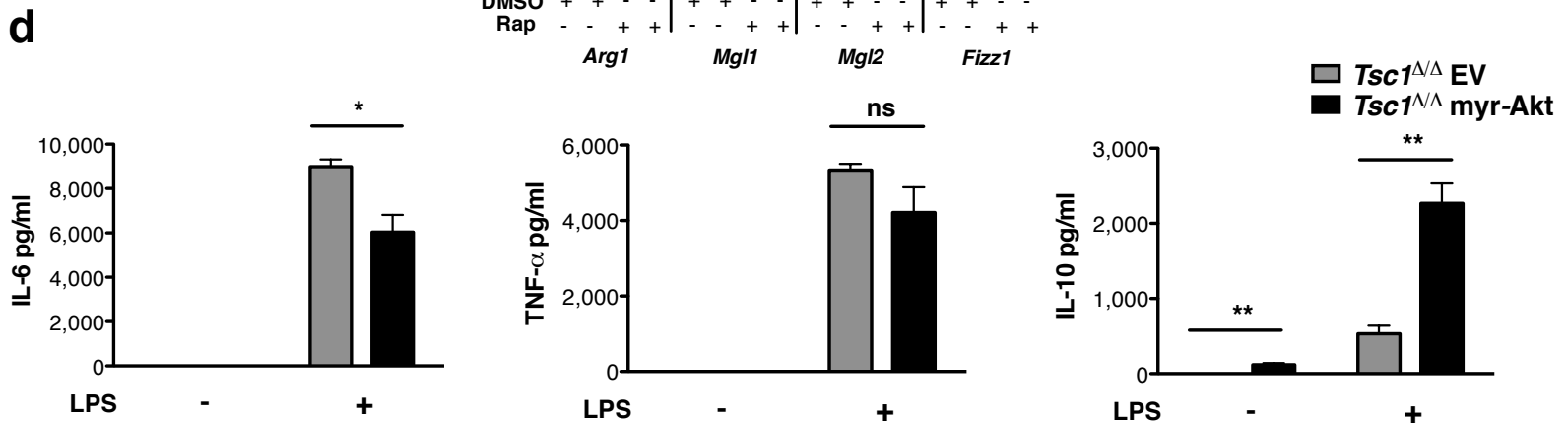
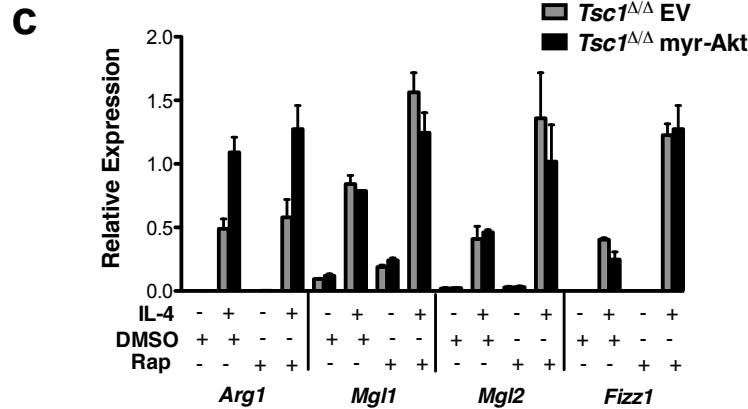
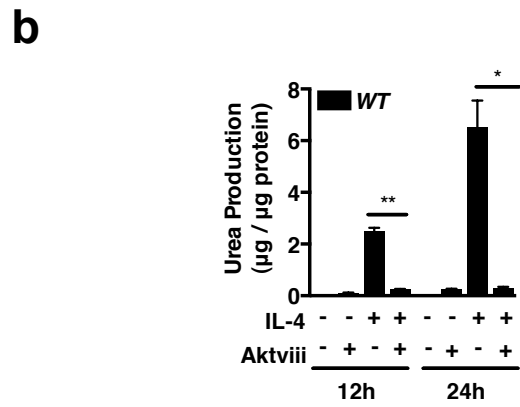
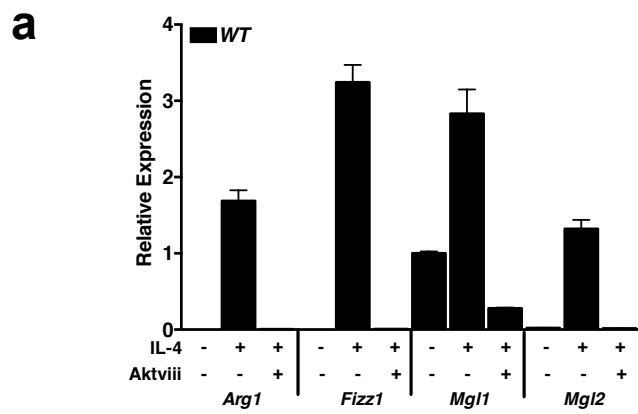
Supplementary Figure S4. PPAR δ Activity is Normal in *Tsc1^{Δ/Δ}* BMDMs

Gene expression of *Atgl* in *Tsc1^{fl/fl}* and *Tsc1^{Δ/Δ}* BMDMs stimulated with IL-4 in the presence or absence of GW501516. DMSO vehicle was used as control. Data is shown as mean \pm SEM (n=2 representative experiments).

a**b****c****d**

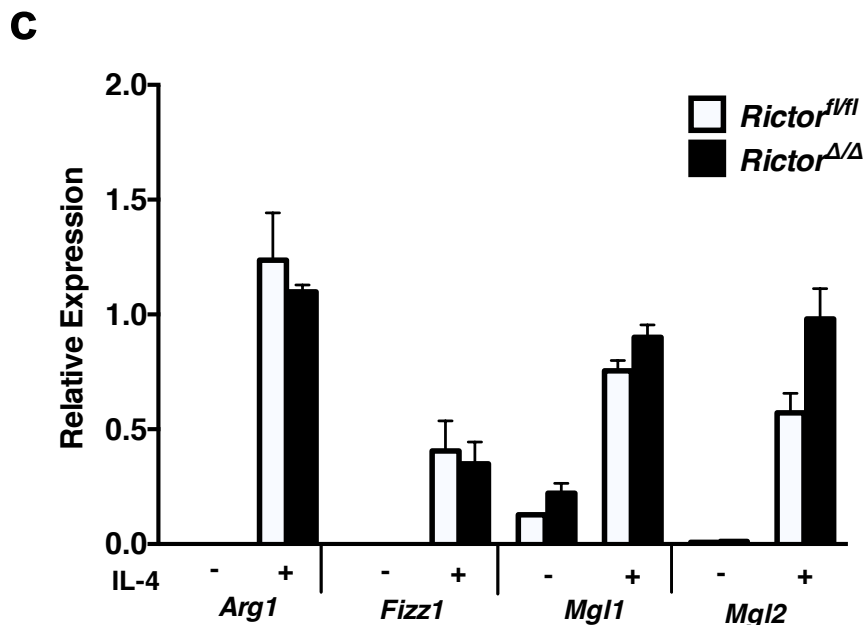
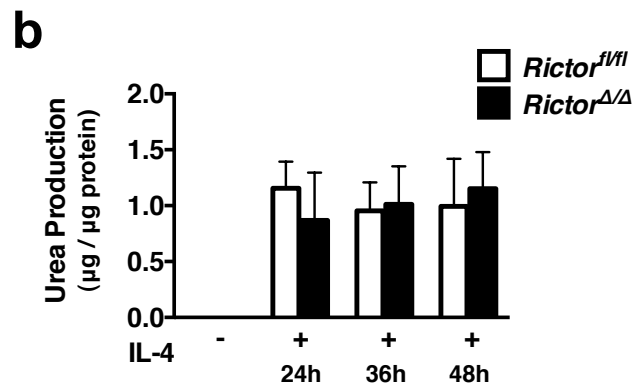
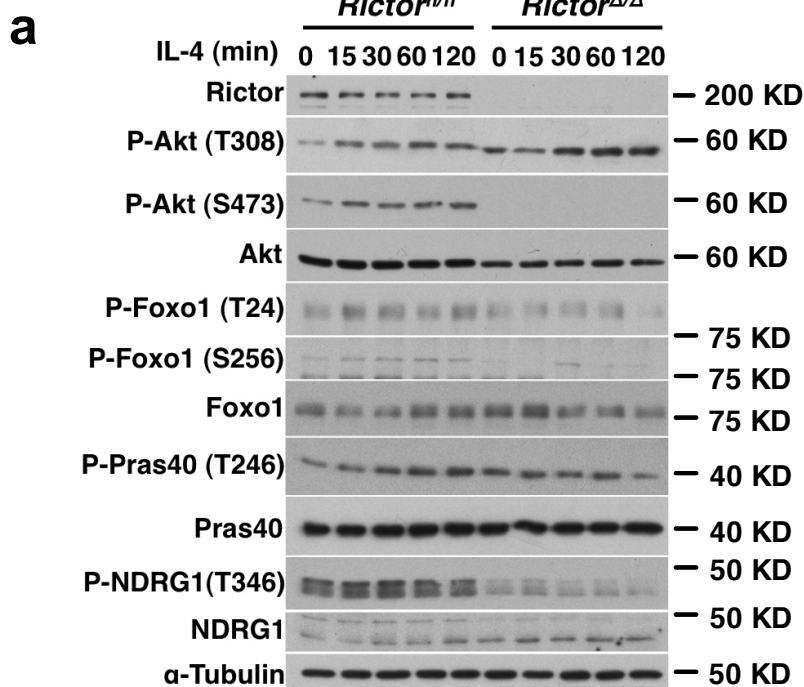
Supplementary Figure S5. IL-4 and LPS-Induced Akt Activation is Attenuated in *Tsc1^{Δ/Δ}* BMDMs

a-c. Immunoblots of *Tsc1^{fl/fl}* and *Tsc1^{Δ/Δ}* BMDMs stimulated with IL-4 as indicated. **a.** Analysis of Akt targets. **b.** Elevated levels of total and phosphorylated Grb10 in *Tsc1^{Δ/Δ}* BMDMs. **c.** Analysis of phosphorylated and total PDK1. Phosphorylation of PDK1 is not inducible, and levels of phosphorylated and total PDK1 are normal in *Tsc1^{Δ/Δ}* BMDMs. **d.** Immunoblot analysis of Akt signaling in lysates from *Tsc1^{fl/fl}* and *Tsc1^{Δ/Δ}* BMDMs stimulated with LPS as indicated.



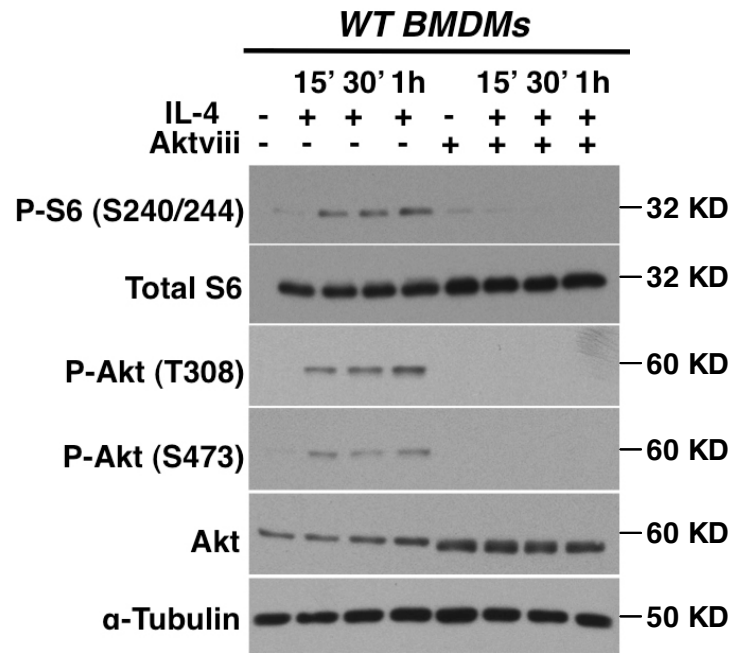
Supplementary Figure S6. Akt Signaling is Critical for Polarization in *Tsc1 Δ/Δ* BMDMs

a-b. WT BMDMs were pretreated with Aktviii for 1h before IL-4 stimulation. **a.** M2 gene expression was measured after 24h of IL-4 stimulation (representative of 3 independent experiments). **b.** Urea production normalized to total protein (n=3). **c.** M2 gene expression in myr-Akt *Tsc1 Δ/Δ* BMDMs and EV *Tsc1 Δ/Δ* BMDMs pretreated with rapamycin or DMSO for 30min and stimulated with IL-4 for 12h (n=2 representative experiments). **d.** Cytokine concentration in supernatants from myr-Akt *Tsc1 Δ/Δ* BMDMs and EV *Tsc1 Δ/Δ* BMDMs treated with LPS for 6h (n=3). Graphs are shown as mean \pm SEM, *p<0.05, **p<0.01 determined by Student's t-tests.



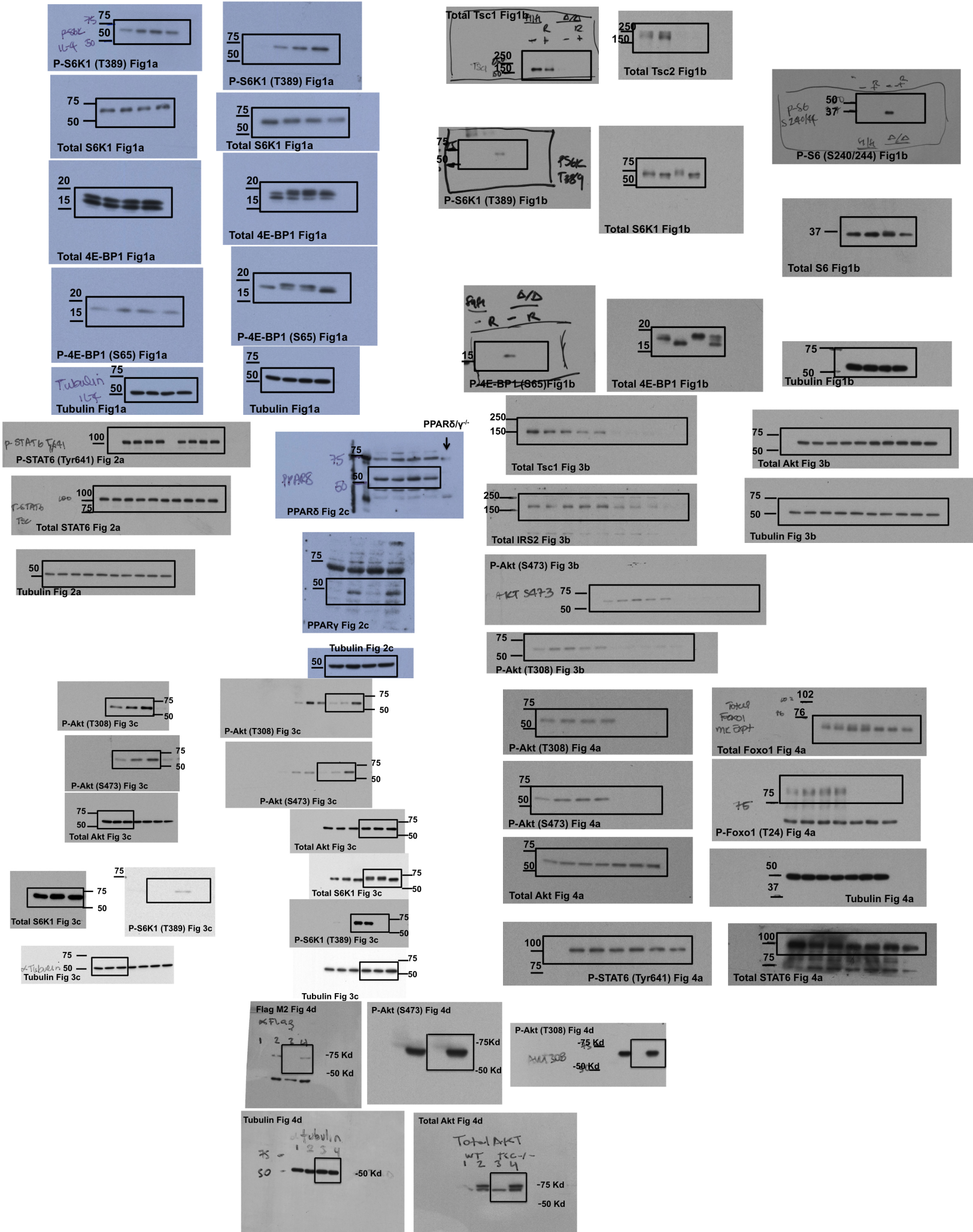
Supplementary Figure S7. Intact M2 Polarization in *Rictor^{Δ/Δ}* BMDMs

a. Reduced phosphorylation of the Akt target Foxo1, but not Pras40 in *Rictor^{Δ/Δ}* BMDMs. Immunoblot analysis of lysates from *Rictor^{fl/fl}* and *Rictor^{Δ/Δ}* BMDMs stimulated with IL-4 as indicated. **b.** Urea production normalized to total protein after stimulation with IL-4 as indicated (n=2 representative experiments). **c.** M2 gene expression analysis in *Rictor^{fl/fl}* and *Rictor^{Δ/Δ}* BMDMs stimulated with IL-4 for 24h (n=2 representative experiments). Graphs are shown as mean ± SEM.



Supplementary Figure S8. Akt Signaling is Critical for mTORC1 Activation in BMDMs

Immunoblot analysis of WT BMDMs pretreated with Aktviii for 1h before IL-4 stimulation for the indicated times. Reduced phosphorylation of S6 on S240/244 indicates attenuated mTORC1 activity in the presence of Aktviii.



Supplementary Figure S9. Full Blots for Main Text Figures

Supplementary Table S1. qPCR Primer Sequences.

| Gene | Primer Sequence (5'-3') |
|--------------------------------|--|
| <i>Arg-1</i> | F: ACC TGG CCT TTG TTG ATG TCC CTA R: AGA GAT GCT TCC AAC TGC CAG ACT |
| <i>Mgl-1</i> | F: TGC AAC AGC TGA GGA AGG ACT TGA R: AAC CAA TAG CAG CTG CCT TCA TGC |
| <i>Mgl-2</i> | F: GCA TGA AGG CAG CTG CTA TTG GTT R: TAG GCC CAT CCA GCT AAG CAC ATT |
| <i>Fizz-1</i> | F: TCC AGC TGA TGG TCC CAG TGA ATA R: ACA AGC ACA CCC AGT AGC AGT CAT |
| <i>Ym1</i> | F: AGA AGG GAG TTT CAA ACC T R: GTC TTG CTC ATG TGT GTA AGT GA |
| <i>Mrc1</i> | F: TGG GCT ACA GGA GAA CCC AAC TTT R: GCA GTG GCA TTG ATG CTG CTG TTA |
| <i>Pgc1β</i> | F: CTT CCG TTG GCC CAG ATA C R: CTG CTG GGC CTC TTT CAG TA |
| <i>Fabp4</i> | F: ATG AAA TCA CCG CAG ACG ACA GGA R: TGT GGT CGA CTT TCC ATC CCA CTT |
| <i>Cd36</i> | F: TCA TGC CAG TCG GAG ACA TGC TTA R: AAC TGT CTG TAC ACA GTG GTG CCT |
| <i>Il-6</i> | F: ACA AAG CCA GAG TCC TTC AGA GAG R: TTG GAT GGT CTT GGT CCT TAG CCA |
| <i>Tnf-α</i> | F: TCT CAG CCT CTT CTC ATT CCT GCT R: AGA ACT GAT GAG AGG GAG GCC ATT |
| <i>Il-10</i> | F: GCT CTT ACT GAC TGG CAT GAG R: CGC AGC TCT AGG AGC ATG TG |
| <i>Tgfβ</i> | F: GTG CGG CAG CTG TAC ATT GAC TTT R: TGT GTT GGT TGT AGA GGG CAA GGA |
| <i>Stat3</i> | F: TGT GAT GCC TCC TTG ATC GTG ACT R: CCA CAG GAT TGA TGC CCA AGC ATT |
| <i>Bcl3</i> | F: TCG CAA TTA TGA AGG GCT CAC TGC R: TGC ACC ATG TTC AGG CTG TTG TTC |

| | |
|--------------------------------|--|
| Socs3 | F: TAG ACT TCA CGG CTG CCA ACA TCT R: GGA GCT AGT CCC GAA GCG AAA TCT |
| Il-12 p35 | F: TGC CCT CCT AAA CCA CCT CAG TTT R: TTT CTC TGG CCG TCT TCA CCA TGT |
| Il-12 p40 | F: GGC TGG TGC AAA GAA ACA TGG ACT R: AGA GAC GCC ATT CCA CAT GTC ACT |
| Atg1 | F: TGG AAC CAA AGG ACC TGA TGA CCA R: AGA TGC TAC CCG TCT GCT CTT TCA |
| Cd68 | F: TTC TGC TGT GGA AAT GCA AG R: CAA TGA TGA GAG GCA GCA AG |
| Jak1 | F: GAT AAG GAG TAC TAC ACA GTC AAG R: CAG ACA TCA GAG GCG ATA TAA A |
| Jak3 | F: CTT GAT GAG CCA AGT ATC CTA C R: TGC TCC TAG ATA CAC AAA TTC C |
| Il4ra | F: AGA TGC AAG CCT TCA GAA TCT GGG R: CGT GGA TGC CCA AGA ACA CAC TAT |
| Pparγ | F: TCA TGA CCA GGG AGT TCC TC R: CAG GTT GTC TTG GAT GTC CTC |
| Pparδ | F: AGC ACA TCT ACA ACG CCT ACC TGA R: TCG ATG TCG TGG ATG ACA AAG GGT |
| Hprt | F: TTT CCC TGG TTA AGC AGT ACA GCC C R: TGG CCT GTA TCC AAC ACT TCG AGA |