Supplementary Figure Legend

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2 Supporting Figure 1. Gene expression by differentially polarized RAW and MSRC2 cells. 3 4 (A) qPCR analysis of polarization-related genes arg1 (arginase) and nos2 (iNOS) in RAW 264.7 5 cells cultured for 2h in either plain media (M0), LPS (M1), or IL-4 and IL-13 (M2). Expression 6 levels were calculated via the ddCT method and normalized to HPRT expression, then 7 normalized to expression of M0 stimulated RAW cells. Data are mean±SEM for n=3. (B) qPCR 8 analysis of msr1 (SR-AI) gene expression by RAW cells cultured for 2h in either plain media 9 (M0), LPS (M1), or IL-4 and IL-13 (M2). Expression levels were calculated as in A, data are 10 mean±SEM for n=3. (C) qPCR analysis of emr1 (F4/80) expression by RAW and MSRC2 cells 11 cultured for 2h in either plain media (M0), LPS (M1), or IL-4 and IL-13 (M2). Expression levels were calculated as in A, data are mean±SEM for n=3; *P<0.05, ***P<0.0005. 12 13 14 Supporting Figure 2. RAW cells, but not MSRC2 cells, exhibit a phagocytosis profile 15 consistent with M2 polarization. (A) Flow staining of MSRC2 and WT RAW cells following exposure to AdOVA and incubation with PE-stained carboxy-modified beads. Macrophages 16 17 were gated as live singlets with positive CFSE uptake, images are representative of 3

independent experiments. (B) Quantification of A, showing the percent of cells containing

fluorescence corresponding to the presence of 2+ beads as mean±SEM for n=3 replicates,

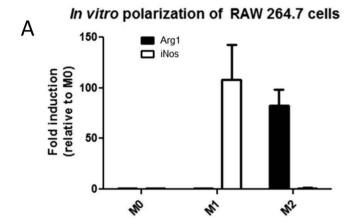
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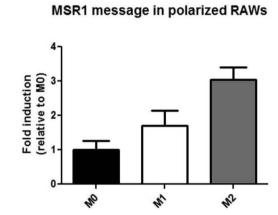
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**P<0.005.

- Supporting Figure 3. Quantification of Western blot results. Densitometry analysis of phosphorylated mTOR (**A**) and Mertk (**B**) protein in RAW and MSRC2 cells cultured for 2h in either plain media (M0), LPS (M1), or IL-4 and IL-13 (M2). Data are presented as the ratio of intensity of the phospho-mTOR or phospho-Mertk bands divided by the total mTOR or total
- 26 Mertk bands respectively, normalized to the expression of actin.





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