

1 **Supplementary Figure Legend**

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3 **Supporting Figure 1. Gene expression by differentially polarized RAW and MSRC2 cells.**

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
12 were calculated as in A, data are mean±SEM for n=3; *P<0.05, ***P<0.0005.

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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
16 exposure to AdOVA and incubation with PE-stained carboxy-modified beads. Macrophages
17 were gated as live singlets with positive CFSE uptake, images are representative of 3
18 independent experiments. (B) Quantification of A, showing the percent of cells containing
19 fluorescence corresponding to the presence of 2+ beads as mean±SEM for n=3 replicates,
20 **P<0.005.

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22 **Supporting Figure 3. Quantification of Western blot results.** Densitometry analysis of
23 phosphorylated mTOR (**A**) and Mertk (**B**) protein in RAW and MSRC2 cells cultured for 2h in
24 either plain media (M0), LPS (M1), or IL-4 and IL-13 (M2). Data are presented as the ratio of
25 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.





