

LAPTM5 is a Positive Regulator of Pro-inflammatory signaling pathways in Macrophages

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Supplementary Figure S1. Endogenous expression of LPTM5 in macrophages. (A) Confocal fluorescence analysis of RAW264.7 cells transfected with control or LPTM5 siRNA. At 72h post transfection, cells were fixed and immunostained with anti-LPTM5 antibody. Bar, 5 μ m. (B) Knockdown of LPTM5 using shRNA. Western blot analysis of RAW264.7 cell lines stably expressing control or LPTM5 shRNA. Cell lysates were immunoblotted with anti-LPTM5 or actin antibodies as a control for protein loading

Supplementary Figure S2. LPTM5 does not co-localize with transferrin. RAW264.7 cells were serum-starved for 2h in DMEM and subsequently incubated with 50 μ g/ml Alexa 488-labeled transferrin at 37°C. After 60min uptake, excess transferrin was washed away and cells were fixed and immunostained with antibodies towards LPTM5. Representative confocal images are shown. The merged panel shows an overlay of the two channels. Bar, 5 μ m. Quantification of co-localization between LPTM5 and transferrin was performed on randomly selected cells using Volocity 5.4.1 software and expressed as Colocalization Coefficient M2 = 0.221, r(20), p>0.10 (corresponding to the degrees of freedom (r) and level of significance (p)).

Supplementary Figure S3. A20 is upregulated in LPTM5-deficient RAW264.7 cells. Stable RAW264.7 cells expressing control or LPTM5 shRNA were stimulated with (A) 1 μ g/ml LPS or (B) 50 ng/ml TNF α for the indicated time points. Cells were lysed and analyzed by immunoblotting with antibodies towards A20 and actin as a control for protein loading. Note increased basal A20 expression in RAW264.7 cells expressing shRNA against LPTM5 compared to control cells, indicated by the asterisks. Panels on the right show quantification of basal A20 protein levels relative to actin from control (white column) and LPTM5 knockdown (black column) RAW264.7 cells. The graph indicates the increase of A20 protein in LPTM5 knockdown relative to control cells and is presented as mean \pm SD of three independent experiments.

Figure S1

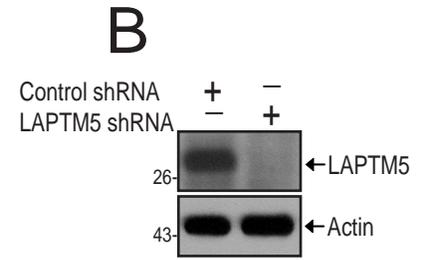
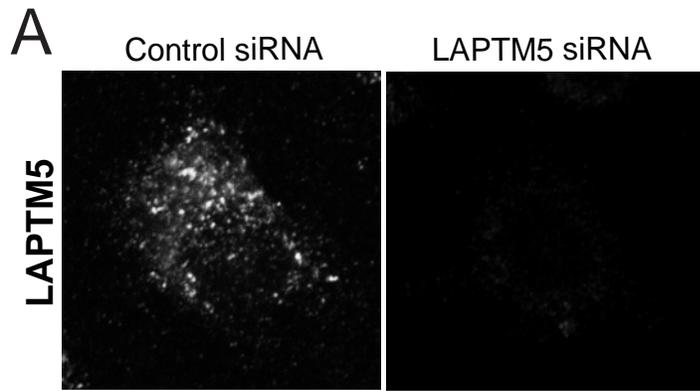


Figure S2

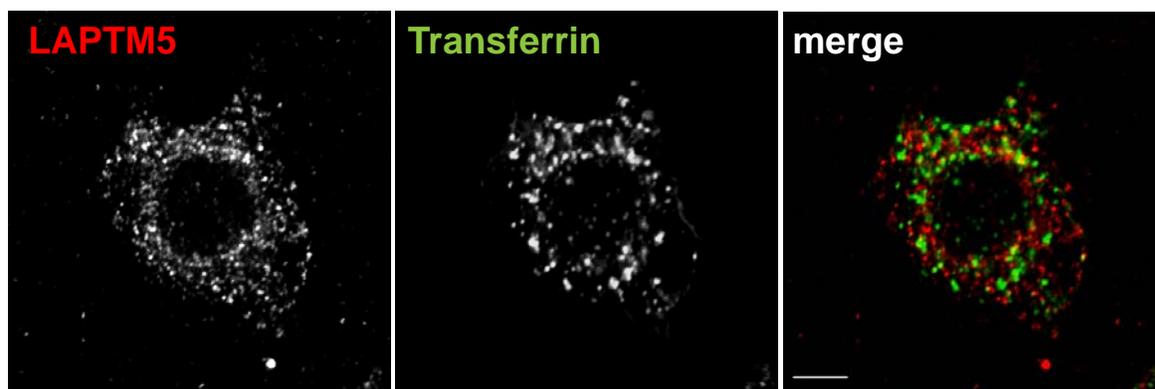
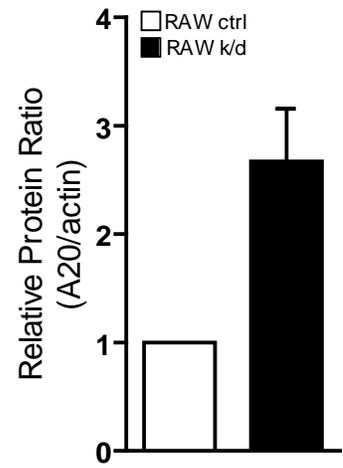
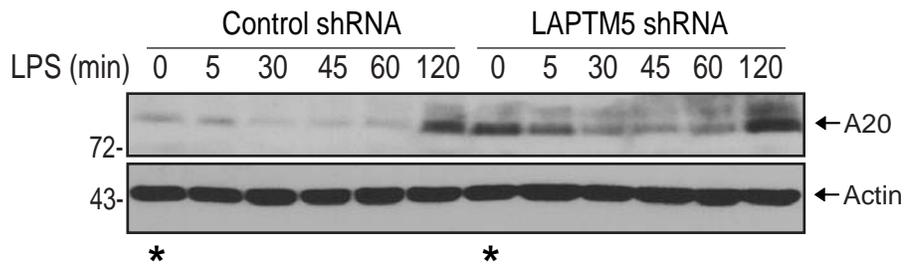


Figure S3

A



B

