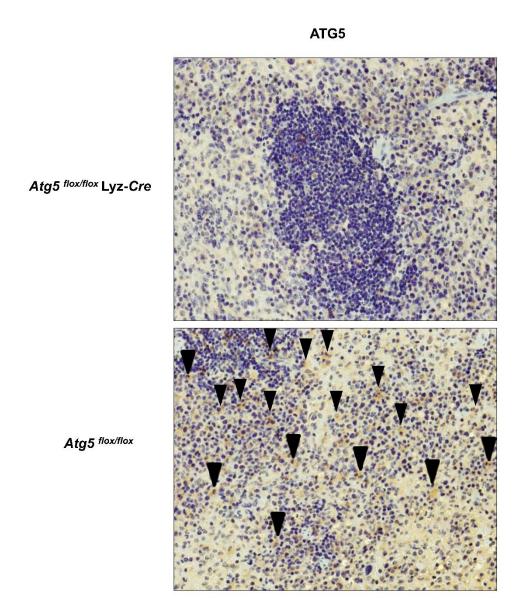
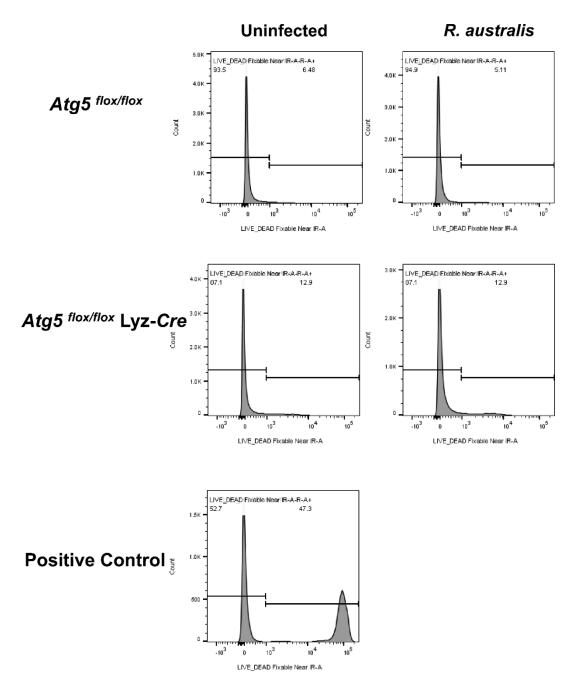
Supplementary FIG 1



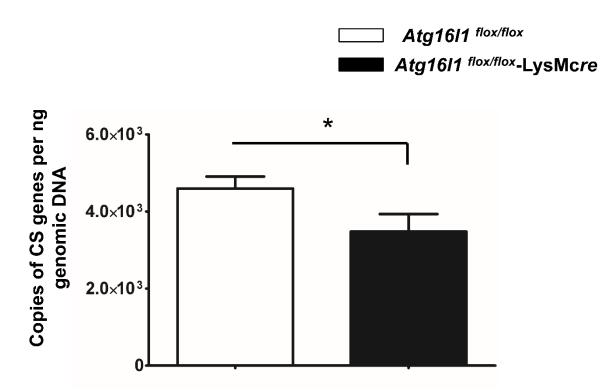
Supplementary FIG 1.Immunohistochemical staining of ATG5 in *Atg5* flox/flox and *Atg5* flox/flox Lyz-*Cre* mice. Spleen of infected *Atg5* flox/flox Lyz-*Cre* and *Atg5* flox/flox mice was processed for immunohistochemical analysis of ATG5 using specific antibodies directed against ATG5. ATG5 was stained as brown (shown as black arrow heads).

Supplementary FIG 2



Supplementary FIG 2. Viability of *R. australis*-infected BMMs of *Atg5* flox/flox and *Atg5* flox/flox Lyz-*Cre* mice. BMMs of *Atg5* flox/flox and *Atg5* flox/flox Lyz-*Cre* mice were infected with *R. australis* at an MOI of 2. At 48 h p.i., Cells were collected and stained for viability using LIVE/DEAD® Fixable Near-IR Dead Cell Stain Kit (Life Technologies, Grand Island, NY) in accordance with the manufacturer's protocol. Positive control was composed of half alive and half dead BMMs of *Atg5* flox/flox Lyz-*Cre* mice. Flow cytometry was performed on 30,000 cells with a FACSCalibur laser cytometer (Becton-Dickinson, BD Biosciences, San Jose, CA). Data analysis on the entire ungated cell population was performed using FlowJo software. Data represented two independent experiments.

Supplementary FIG 3



Supplementary FIG 3. Significantly increased concentrations of *R. australis* in Atg16l1 flox/flox BMMs compared to Atg16l1 flox/flox -LysMCre BMMs in vitro. BMMs of Atg16l1 flox/flox -LysMCre and Atg16l1 flox/flox mice were infected with R. australis at an MOI of 2. Rickettsial concentration in these primary mouse macrophages was evaluated by quantitative real-time PCR at 48 h p.i.. Number of citrate synthase (CS) gene copies per ng of genomic DNA represents the quantity of rickettsiae. *, p<0.05.