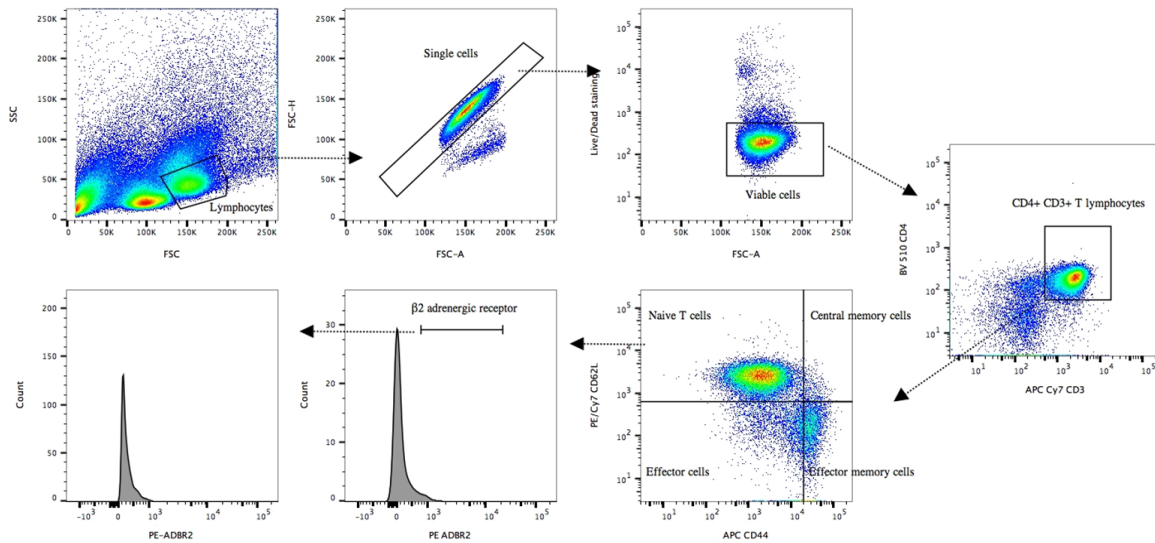


# Supplementary Figures

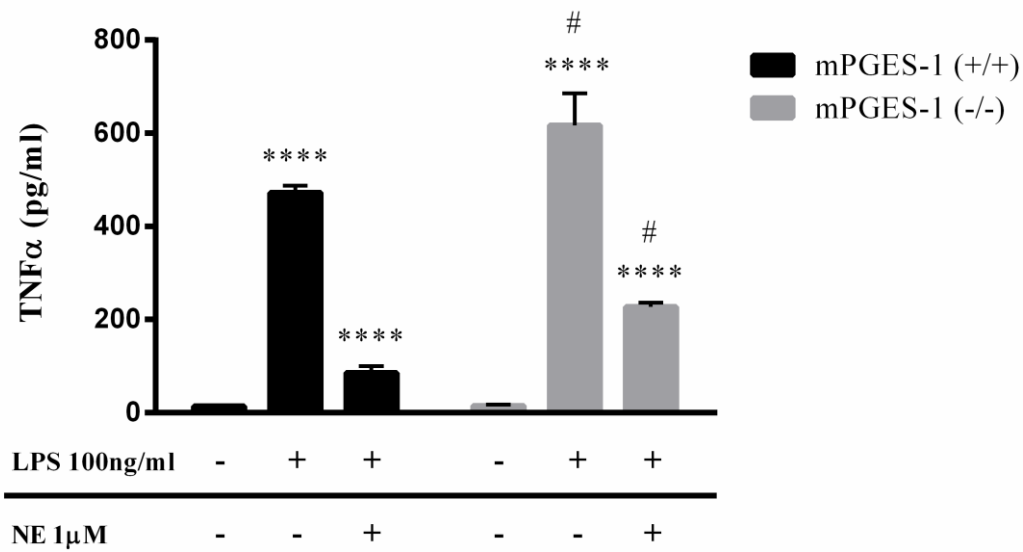
Figure A.



**Fig A. Gating strategy for CD4<sup>+</sup> CD3<sup>+</sup> T lymphocytes and their different subsets.**

Lymphocytes were gated using SSC-A and FSC characteristics followed by stepwise exclusion of non-single cells and non-viable cells. In viable, single CD4 T lymphocytes (CD4<sup>+</sup> CD3<sup>+</sup>) CD44 and CD62L were used to distinguish between different CD4+ T cell subsets. Corresponding  $\beta_2$  expression on these subsets were measured as %  $\beta_2^+$  cells of parent population or mean fluorescence intensity using histograms.

**Figure B**



**Fig B. Cytokine profiling of activated mPGES-1 (-/-) splenocytes in response to NE stimulation.** Primary splenocyte cultures established from mPGES-1 (+/+) and (-/-) mice were pre-treated with norepinephrine (NE) at 1μM concentration for 30 mins followed by LPS (100ng/ml) activation. Cell supernatants were analyzed for cytokine production following 3 hours of treatment. (\*\*\*\*p<0.0001; LPS versus LPS+NE). (# p<0.05; mPGES-1(+/+) versus (-/-) within LPS+NE treatment; student's T-test). Each sample was run as duplicates during the assay and values are represented as mean ±SEM from 3 independent experiments. Statistical analysis was done using One-way ANOVA unless otherwise indicated.

Figure C.

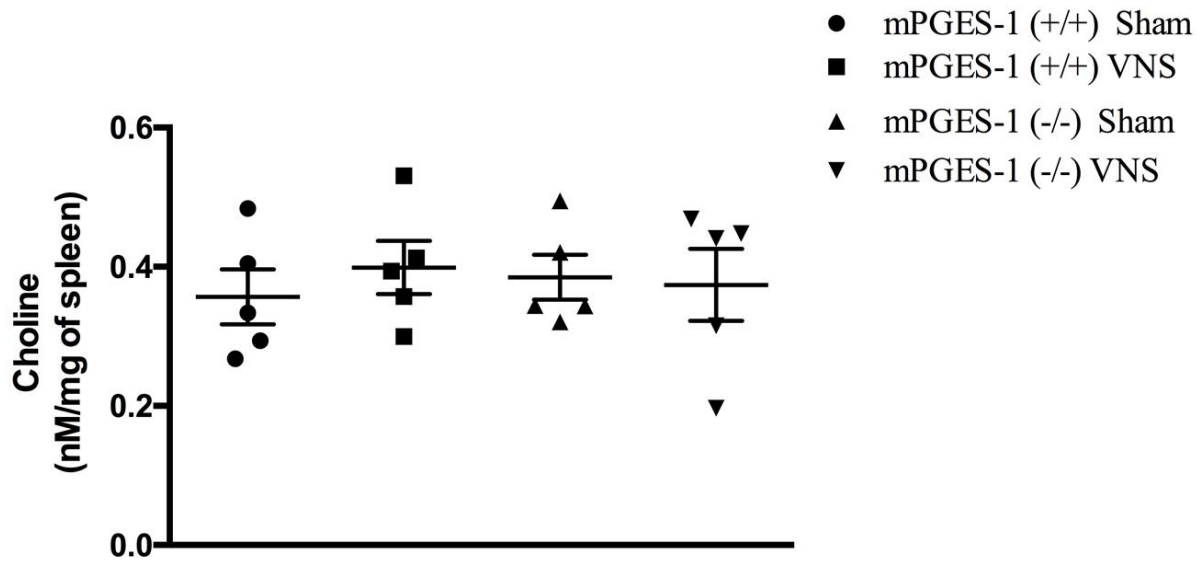


Fig C. Choline measurement in mouse spleen following SHAM and VNS treatment.

mPGES-1 (+/+) and (-/-) mice underwent SHAM and VNS for 30 minutes and spleens were homogenized for choline measurement. No differences were observed between both treatment groups and mPGES-1 depletion does not affect the choline levels. Samples were run as duplicates during the choline assay and values are represented as mean  $\pm$ SEM from 5 individual animals per treatment group ( $p > 0.05$ ; One-way ANOVA)