

Supplemental Material to:

**Linda M. Stephenson, Brian C. Miller, Aylwin Ng,
Jason Eisenberg, Zijiang Zhao, Ken Cadwell,
Daniel B. Graham, Noboru N. Mizushima, Ramnik Xavier,
Herbert W. Virgin, Wojciech Swat**

**Identification of *Atg5*-dependent transcriptional changes
and increases in mitochondrial mass in *Atg5*-deficient
T lymphocytes**

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www.landesbioscience.com/journals/autophagy/article/8133

Figure S1

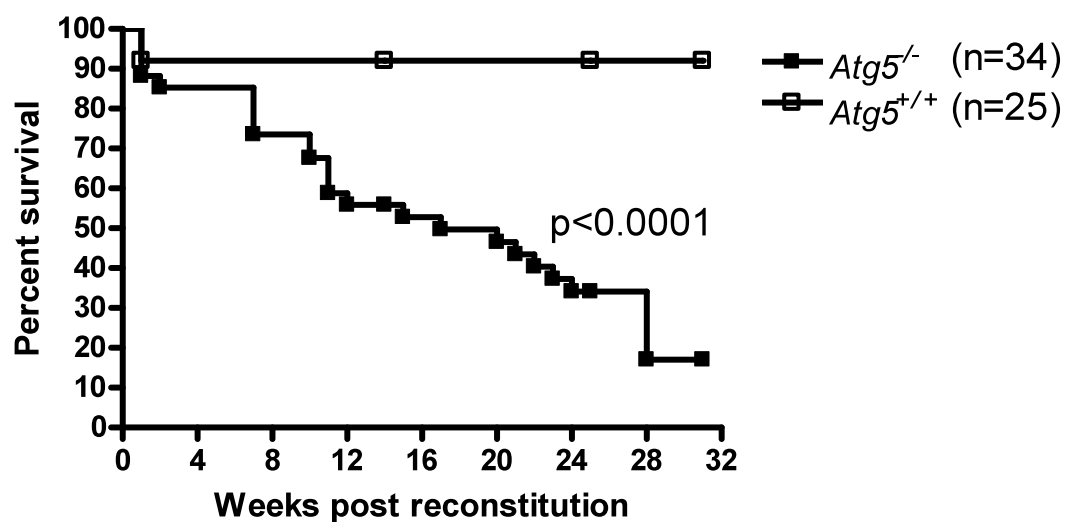
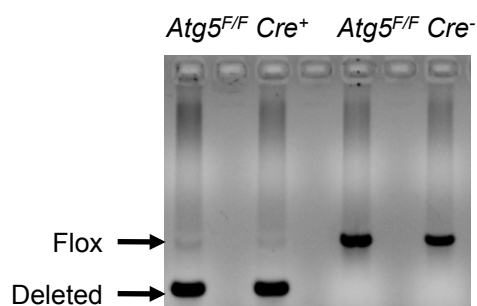
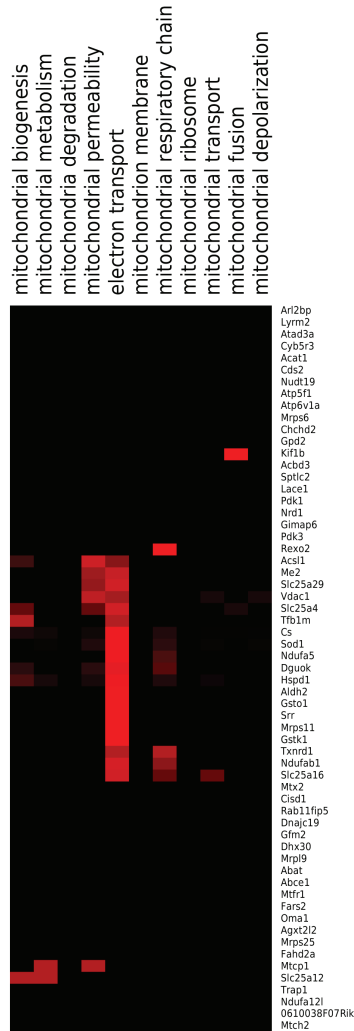


Figure S2



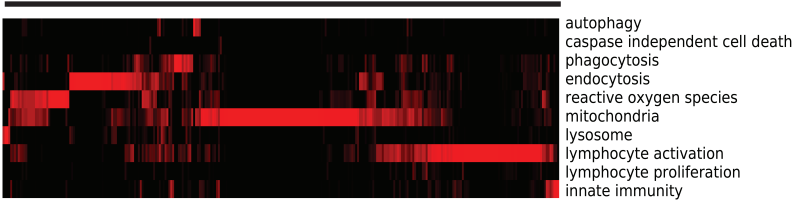
A

Literature co-citation analysis
using terms associated with
mitochondrial functions



B

292 of 699 differentially expressed genes have
PubMed citations for the terms searched



Supplemental Figure 1. *Atg5*^{-/-} chimeras have decreased survival compared with *Atg5*^{+/+} chimeras. The survival of Atg5 chimeric mice was recorded up to 31 weeks post-irradiation and stem cell reconstitution. The p value between the *Atg5*^{-/-} and *Atg5*^{+/+} survival curves is shown. Data pooled from 4 independently generated batches of reconstituted chimeras.

Supplemental Figure 2. Recombination of the *Atg5*^{lox} allele in *Atg5*^{F/F} *Cre*⁺ thymocytes. PCR analysis performed on DNA isolated from thymocyte of *Atg5*^{F/F} *Cre*⁺ and *Atg5*^{F/F} *Cre*⁻ mice to determine recombination of the *Atg5*^{lox} locus. Two mice are shown of each genotype. Representative gel from 4 independent experiments.

Supplemental Figure 3. Matrices generated from literature co-citation analysis of differentially expressed genes and citation terms associated with mitochondrial and cellular functions. (A) Mitochondrial functions and (B) general cellular processes or specific T-lineage immune functions, displayed as a heatmap. The intensity of red on the heatmap denotes the extent to which each gene is co-cited with each specific term in the PubMed database.