# Supplementary information for Mitochondria are Secreted in Extracellular Vesicles When Lysosomal Function is Impaired

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FCCP

DMSO

**Supplementary Figure 1. EV release is increased in MEFs when lysosomal function is compromised. a.** Scheme of EV isolation from conditioned media using differential centrifugation or immunoaffinity purification. The immunoaffinity purification method uses an antibody cocktail against tetraspanin surface markers CD9, CD63 and CD81. **b.** Quantification of proteins in small EV fraction from Fig. 1a (Alix n=7; Tsg101 n=8, CD81 n=4 biologically independent experiments). **c.** Representative Western blot and quantification of large EVs isolated using affinity capture (n=4 biologically independent experiments). **d.** Evaluation of cell death in MEFs after 48 h of Baf A1 treatment (5 nM) (n=5 biologically independent experiments). **e.** Representative Western blot of proteins in WCL and EVs secreted by WT MEFs after treatment with FCCP (10  $\mu$ M) for 24 h. Quantification of large and small EVs (Tsg101 n=4; CD81 n=5, MTCO1 n=5 biologically independent experiments)). **f.** Venn diagrams showing the overlap of proteins identified in large EVs with those reported in Vesiclepedia and ExoCarta databases. Data are mean±SEM. P values shown are by two-sided Student's t-test. Source data are provided as a Source Data file.



**Supplementary Figure 2. EV release is increased in hearts when lysosomal function is compromised. a.** Scheme of EV isolation from heart tissue (Created using BioRender.com). **b.** Western blot analysis and quantification of total EVs isolated from mouse plasma (n=4 biologically independent experiments). **c.** Quantification of proteins in small EV fraction from Fig. 1e (n=4 biologically independent experiments). **d.** Western blot analysis and quantification of CD63, CD81, Tim23, and Mito-Dendra2 protein levels in EV fractions isolated from heart tissue using immunoaffinity capture (n=4 biologically independent experiments). Data are mean±SEM. ns = not significant. P values shown are by two-sided Student's t-test. Source data are provided as a Source Data file.



**Supplementary Figure 3. EV release is increased in Rab7-deficient cells. a.** Quantification of proteins in small EV fraction from Fig. 2a (Alix n=7; Tsg101 n=6, CD81 n=5 biologically independent experiments). **b**. Representative Western blot and quantification of CD81 (EV marker, n=4 biologically independent experiments), Calreticulin (ER, n=3 biologically independent experiments) and MnSOD (mitochondria, n=4 biologically independent experiments) in whole cell lysates (WCL) and EV fractions from WT and *Rab7<sup>-/-</sup>* MEFs. **c.** Representative Western blot and quantification of proteins in WCL and large EVs (n=4 biologically independent experiments). Cells were infected with Ad-β-gal or Ad-Rab7 for 24h. **d.** Representative Western blot and quantification of proteins in WCL and large EVs (n=4 biologically independent experiments). Cells were infected with Ad-β-gal or Ad-Rab7 for 24h. **d.** Representative Western blot and quantification of proteins in WCL and large EVs (n=4 biologically independent experiments). Cells were infected with Ad-β-gal or Ad-Rab7 for 24h. **d.** Representative Western blot of proteins in whole cell lysates (WCL) and large EVs (n=4 biologically independent experiments). Cells were infected with Ad-β-gal or Ad-Rab7T22N for 36h. **e**. Representative Western blot of proteins in whole cell lysates (WCL) and large EV fraction from *Rab7<sup>-/-</sup>* MEFs after treatment with GW4869 (2.5μM, 48hr). Quantification of proteins in the large EV fraction (n=4 biologically independent experiments). **f**. Quantification of CD81 and Tom20 levels in large EVs after proteinase K digestion (n=4 biologically independent experiments). Data are mean±SEM. P values shown are by two-sided Student's t-test. Source data are provided as a Source Data file.











**Supplementary Figure 4. Endosomal proteins are increased in** *Rab7<sup>-/-</sup>* **MEFs. a.** Representative western blot and quantification of various EV proteins in WT and *Rab7<sup>-/-</sup>* MEFs (CD81/Gapdh and Tsg101/Gapdh: WT and *Rab7<sup>-/-</sup>* (n=8 biologically independent experiments); Alix/Gapdh: WT (n=8 biologically independent experiments) and *Rab7<sup>-/-</sup>* (n=6 biologically independent experiments). **b.** Representative Western blot and quantification of endosomal proteins in WT and *Rab7<sup>-/-</sup>* MEFs (n=8 biologically independent experiments). **b.** Representative Western blot and quantification of endosomal proteins in WT and *Rab7<sup>-/-</sup>* MEFs (n=8 biologically independent experiments). **c.** Co-localization between CD81 (red) and Cytochrome *c* (green) in WT and *Rab7<sup>-/-</sup>* infected with the indicated adenovirus for 24 h. Scale bar = 20µm. Mander's overlap coefficient (n=3 independent experiments with a total of 90 cells). **d.** Co-localization analysis between CD63 (green) and Hsp60 (red, mitochondria) in WT +/- Baf A1 (50nM for 24hr) and *Rab7<sup>-/-</sup>* MEFs. Scale bar = 20µm. Mander's overlap coefficient (n=3 independent experiments with a total of 90 cells). Data are mean±SEM. ns = not significant. P values shown are by two-sided Student's t-test (a, b) or ANOVA with Tukey's post-hoc testing (**c, d**). Source data are provided as a Source Data file.



Supplementary Figure 5. EV secretion can occur independently of autophagosome formation and mitophagy. a. Western blotting for Atg5, LC3, and p62 protein levels in WT and  $Atg5^{-/-}$  MEFs. Atg5 exists in a complex with Atg12 in WT MEFs (representative from n=3 independent experiments). b. Quantification of proteins after Atg7 knockdown (n=3 biologically independent experiments). c. qPCR analysis for Pink1 transcript levels in MEFs infected with *control* or *Pink1* shRNA lentiviral particles (2 MOI, n=3). d. Representative Western blots of whole cell lysate (WCL) and extracellular vesicle fractions for CD81 and MTCO1. The EVs were obtained by differential centrifugation of proteins in large EVs (n=3 biologically independent experiments). f. LysoTracker Red staining of cells and quantification of fluorescence intensity (n=3 independent experiments with a total of 90 cells scored). g. Representative images of co-localization between GFP-LC3 (green) and Cytochrome c (red) in WT and  $Rab7^{-/-}$  MEFs +/- Baf A1 (50nM for 24hr). Mander's correlation coefficient (n=3 independent experiments with a total of 90 cells shown are by two-sided Student's t-test (c, e) or ANOVA with Tukey's post-hoc testing (g). Source data are provided as a Source Data file.



**Supplementary Figure 6. Effect of Rab27a knockdown in** *Rab7*-deficient cells. **a.** Western blot for Rab27 levels in WT MEFs after treatment with BafA1 (5nM, 24h, n=4 biologically independent experiments). **b.** Western blot for Rab27 in WT mouse hearts 24 h after administration of chloroquine (CQ) (n=4 biologically independent animals). **c.** Confirmation of Rab27 knockdown in *Rab7*<sup>-/-</sup> MEFs. Representative western blot and quantification of Rab27 protein levels in WT and *Rab7*<sup>-/-</sup> MEFs after infection with control or Rab27a shRNA lentivirus (n=6 biologically independent experiments). **d.** Quantification of proteins in small EV fractions from Fig. 5e (Alix n=6; Tsg101, CD81 n=5 biologically independent experiments). **e.** Representative Western blots and quantification of mitochondrial proteins in MEFs after Rab27 knockdown (n=7 biologically independent experiments). **f.** Analysis of mRNA levels by qPCR in WT MEFs (n=4 biologically independent experiments). Data are mean±SEM. All sample numbers represents biologically independent experiments. ns = not significant. P values shown are by two-sided Student's t-test. Source data are provided as a Source Data file.



**Supplementary Figure 7. Characterization of** *Rab7<sup>tiff</sup>MCM* mice. a. Mice were injected with tamoxifen (40 mg/kg) for 5 days and all phenotyping conducted at 28 days post-tamoxifen treatment. b. Echocardiographic analysis of ventricular function and structure 28 days post-tamoxifen treatment. Percent (%) ejection fraction (EF) and left ventricular internal dimension in systole (LVED; s) . *MCM* (n=8 biologically independent animals), *Rab7<sup>tif</sup> MCM* (n=11 biologically independent animals), *Rab7<sup>tif</sup> MCM* (n=11 biologically independent animals). c. Assessment of mitochondrial respiration using isolated mitochondria from *Rab7<sup>tif</sup> and Rab7<sup>tif</sup> MCM* hearts show no differences in maximal respiration rates (FCCP uncoupled) with substrates for complex I (pyruvate/malate and palmitoyl carnitine/malate) or II (succinate/rotenone) (*n* = 4 biologically independent samples). Data are mean±SEM. ns = not significant. Statistical test performed by ANOVA with Tukey's post-hoc testing (b) or two-sided Student's t-test (c). Source data are provided as a Source Data file.



**Supplementary Figure 8. Characterization of EV release in hearts**. **a.** Representative Western blots and quantification of proteins in large EV fractions isolated from hearts using immunoaffinity purification (CD63, Tim23: *Rab7<sup>i/f</sup>* and *Rab7<sup>i/f</sup> MCM* (n=4 biologically independent animals); CD81: *Rab7<sup>i/f</sup>* and *Rab7<sup>i/f</sup> MCM* (n=5 biologically independent animals)). **b.** Representative Western blots of protein levels in large EV fractions from WT and *MCM* heart tissue at D28 post-tamoxifen injection. **c.** Quantification of proteins in large EV fractions (WT and *MCM*, n=5 biologically independent animals). **d.** Nano particle tracking analysis (NTA) to assess size distribution of large EVs isolated from *Rab7<sup>i/f</sup>* and *Rab7<sup>i/f</sup> MCM* heart tissue. **e.** Mitochondrial DNA (mtDNA) content in large EVs preparations (D-loop n=5, Cyt B n=4 biologically independent animals). **f.** Representative Western blots and quantification of proteins in total (large and small) EV fractions isolated from plasma using immunoaffinity purification (CD63 n=8, CD81 n=9 biologically independent animals). **g.** Gating strategy for the macrophage content analysis and quantification of cardiac macrophage content in *Rab7<sup>i/f</sup> MCM* hearts analyzed by flow cytometry (n=6 biologically independent animals). Data are mean±SEM. ns = not significant. P values shown are by two-sided Student's t-test. Source data are provided as a Source Data file.



**Supplementary Figure 9. Release of small EVs is unchanged in aged (24 months) or Lamp2-deficient hearts. a.** Quantification of proteins in small EV fractions from Fig. 8a (n=4 biologically independent animals). **b.** Quantification of CD63 (n=4 biologically independent animals) and CD81 (n=6 biologically independent animals) proteins in small EV fractions from Fig. 8d. Data are mean±SEM. ns=not significant. Statistical analysis performed by two-sided Student's t-test. Source data are provided as a Source Data file.