

Expanded View Figures

Figure EV1. MAPL SUMOylates ABCD3 and its absence alters ABCD3 complex assembly.

- A Representative immunoblot of total lysate and streptavidin-beads pull-down fractions of HEK-293T-REX cells expressing Flag-MAPL-BirA or Flag-BirA (as control) incubated with biotin and probed for ABCD3, DRP1, MAPL and beta-actin.
- B Quantification from three independent experiments of ABCD3 signals of the heavy membrane SIM-beads elution fraction (Fig 1C). * $P < 0.05$ in an unpaired two-tailed t-test.
- C qRT-PCR investigating ABCD3 mRNA levels in livers of four different mice (2-month-old males).
- D Representative immunoblots from whole-cell liver extracts probed for ABCD3, DRP1 and Mfn2 (Left panel). ABCD3, Drp1 and Mfn2 signals were quantified ($n = 3$ for each genotype, right panel).
- E Representative immunoblots from whole-cell liver extracts probed for PEX14, ACOX1 and SCP2 (Left panel). PEX14, ACOX1 and SCP2 signals were quantified ($n = 3$ for each genotype, right panel).

Data are represented as mean $\pm$ SEM.

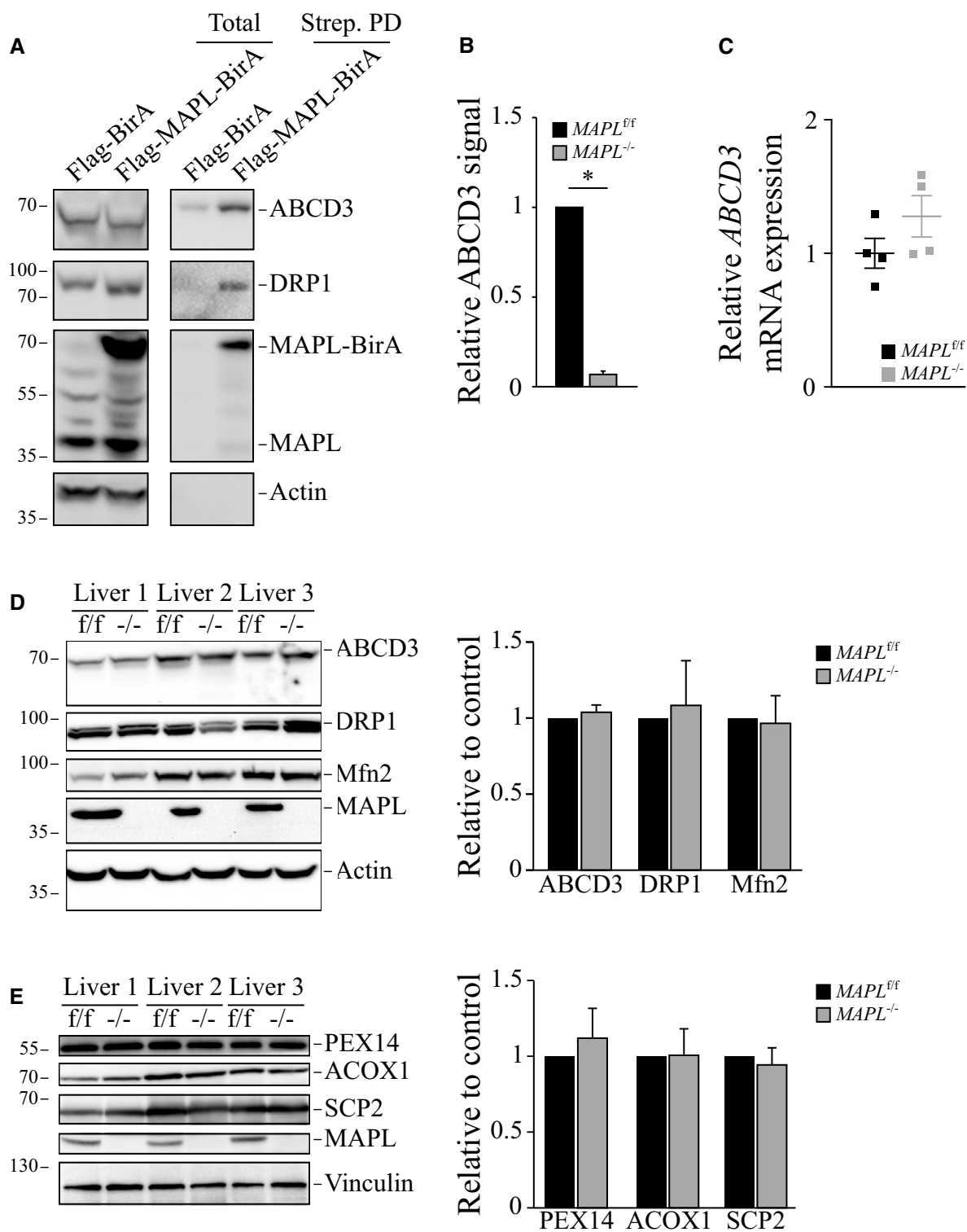


Figure EV1.

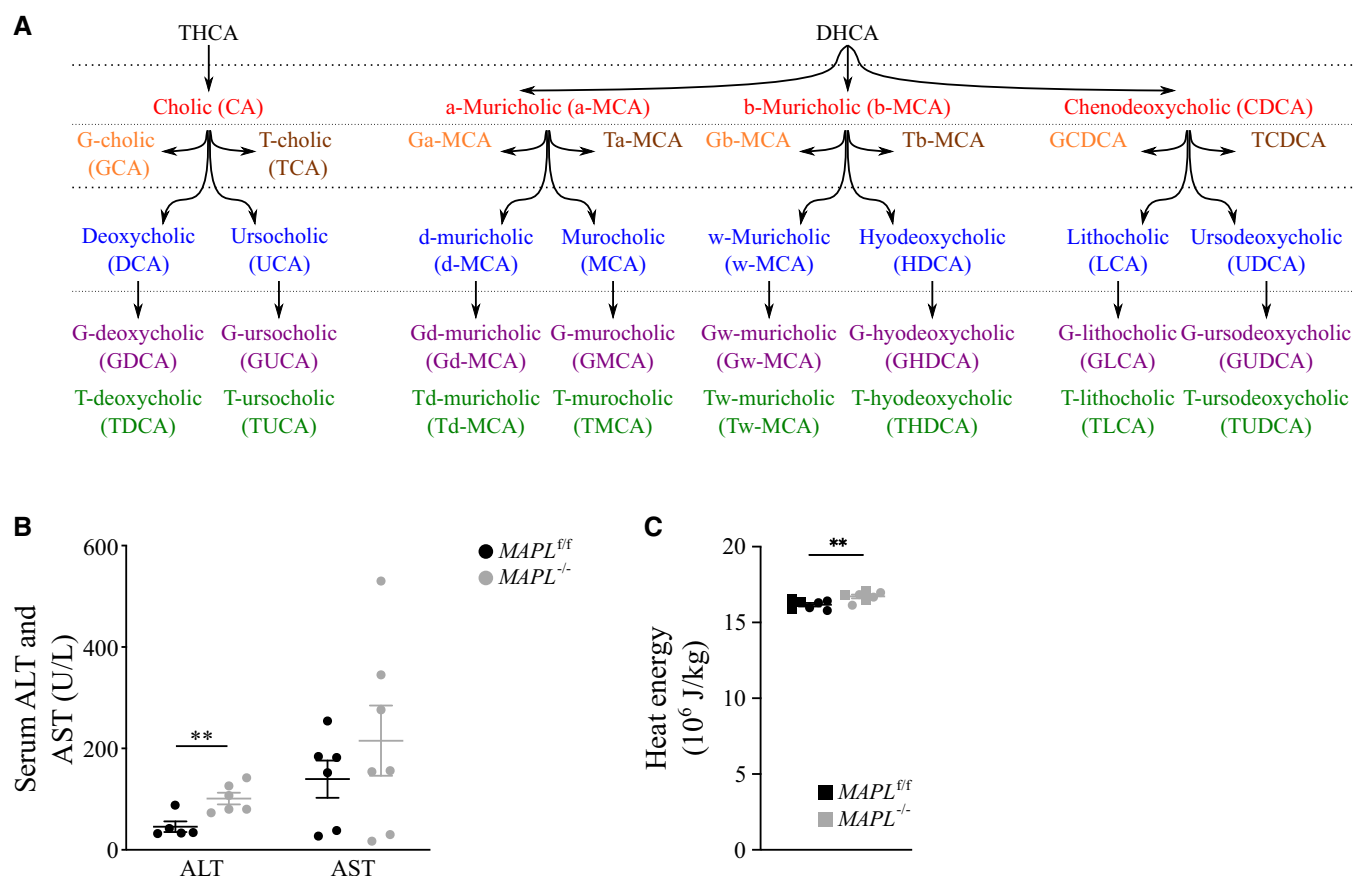


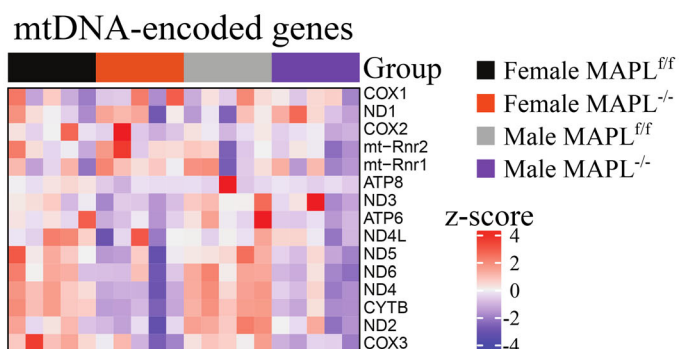
Figure EV2. Loss of MAPL alters bile acid synthesis.

- A A model depicting the generation of the main murine bile acids from THCA and DHCA (tri- and dihydroxycholestanoic acids) (G: glyco-; T: Tauro-).
- B Levels of alanine transaminase (ALT) and aspartate transaminase (AST), markers of liver damage, were quantified and plotted from serum collected from 6-month-old females ($n = 5$ and $n = 6$ for *MAPL^{fl/fl}* and *MAPL^{-/-}*, respectively for ALT quantification and $n = 6$ and $n = 7$ for *MAPL^{fl/fl}* and *MAPL^{-/-}*, respectively for AST quantification). ** $P < 0.01$ in an unpaired two-tailed t -test.
- C The amount of energy present in 3-month animal feces has been measured using an oxygen bomb calorimeter, as described in material and methods ($n = 7$ with four females and three males for each genotype). ** $P < 0.01$ in an unpaired two-tailed t -test.

Data are represented as mean $\pm$ SEM. Squares represent males whereas circles represent females.

Figure EV3. Organellar gene expression in MAPL-deficient mice.

From liver RNAseq ($N = 5$ per sex and genotype). Volcano plots (left panels) and heat maps (right panels) of peroxisomal (top panels) and mitochondrial (bottom panels) gene expression in the liver of *MAPL^{-/-}* mice.



EV4

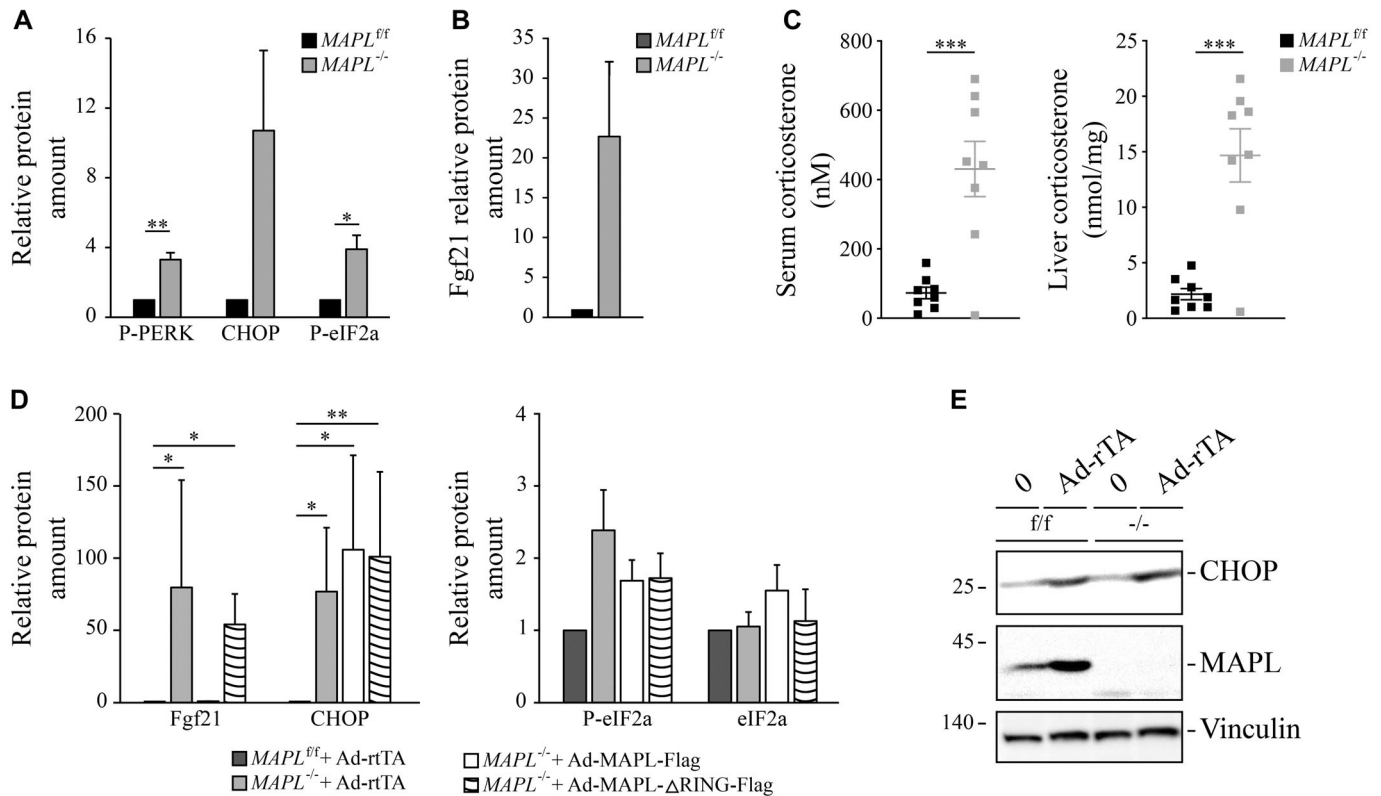


Figure EV4. Loss of MAPL leads to hepatic ER stress, eIF2 α activation and Fgf21 expression.

- A Quantification of phospho-PERK, CHOP and phospho-eIF2 α signals presented in Fig 6A and B immunoblots ($n = 3$ different mouse for each genotype). * $P < 0.05$, ** $P < 0.01$ in an unpaired two-tailed t-test.
- B Quantification of Fgf21 signals presented in Fig 6E ($n = 3$ for each genotype).
- C Circulating and liver corticosterone levels were quantified as described in materials and methods ($n = 8$ for each genotype, 2-month-old males). *** $P < 0.001$ in an unpaired two-tailed t-test.
- D Quantification of FGF21, CHOP, phospho-eIF2 α and eIF2 α signals presented in Fig 6J ($n = 5, 5, 3$ and 4 for MAPL^{f/f} + rtTA, MAPL^{-/-} + rtTA, MAPL^{-/-} + MAPL-Flag and MAPL^{-/-} + MAPL- Δ RING-Flag, respectively, 2–3 months old). * $P < 0.05$, ** $P < 0.01$ in a Kruskal-Wallis test.
- E Representative immunoblot of CHOP protein expression of livers isolated from MAPL^{f/f} and MAPL^{-/-} animals injected or not with the empty virus rtTA.

Data are represented as mean $\pm$ SEM. Squares represent males whereas circles represent females.

Figure EV5. MAPL^{-/-} female mice are lean, resistant to weight gain and develop spontaneous hepatocellular carcinoma.

- A Female mice fed with normal chow were weighed weekly; MAPL^{f/f} ($n = 7$) and MAPL^{-/-} ($n = 10$).
- B MAPL^{f/f} and MAPL^{-/-} ($n = 8$ for each genotype) females were fed with normal chow for 5 months (dotted lines), or for 3 months followed by 2 months of 60% fat chow (solid lines, diet change indicated by HFD arrow).
- C Body weight (g, left panel) and length (cm, right panel) of 7-month-old MAPL^{f/f} ($n = 4$ for body weight and $n = 3$ for length) and MAPL^{-/-} ($n = 2$) female mice. *** $P < 0.01$ in an unpaired two-tailed t-test.
- D Wet weight of organs including liver, kidney, gastrocnemius muscle, epididymal white fat (WAT) and interscapular brown fat (BAT) isolated from 7-month-old MAPL^{f/f} ($n = 4$) or MAPL^{-/-} ($n = 2$) female mice.
- E White adipose tissue gene expression performed by qRT-PCR. $N = 3$, in triplicate, 2-month-old MAPL^{f/f} and MAPL^{-/-} male mice.
- F Insulin tolerance test in MAPL^{f/f} ($n = 8$), MAPL^{-/-} ($n = 8$) female mice. Insulin (0.5 U/kg) was injected intraperitoneally after a 2 h fast and blood glucose was measured at indicated times. Repeated Measures 2way ANOVA analysis for a difference between genotypes $P = 0.0135$.
- G Glucose tolerance test in MAPL^{f/f} ($n = 8$), MAPL^{-/-} ($n = 8$) female mice. Glucose (2 g/kg) was injected intraperitoneally after an overnight fast and blood glucose was measured at indicated times. Repeated Measures 2way ANOVA for a difference between genotypes $P = 0.0903$.
- H Liver glycogen measured enzymatically in male MAPL^{f/f} ($n = 3$), MAPL^{-/-} ($n = 3$) mice. ** $P < 0.01$ in an unpaired two-tailed t-test.
- I Survival curve of MAPL^{f/f} ($n = 11$) and MAPL^{-/-} ($n = 10$) female mice.
- J Cancer-free survival curve of MAPL^{f/f} ($n = 11$) and MAPL^{-/-} ($n = 10$) female mice.

Data are represented as mean $\pm$ SEM. Squares represent males whereas circles represent females.

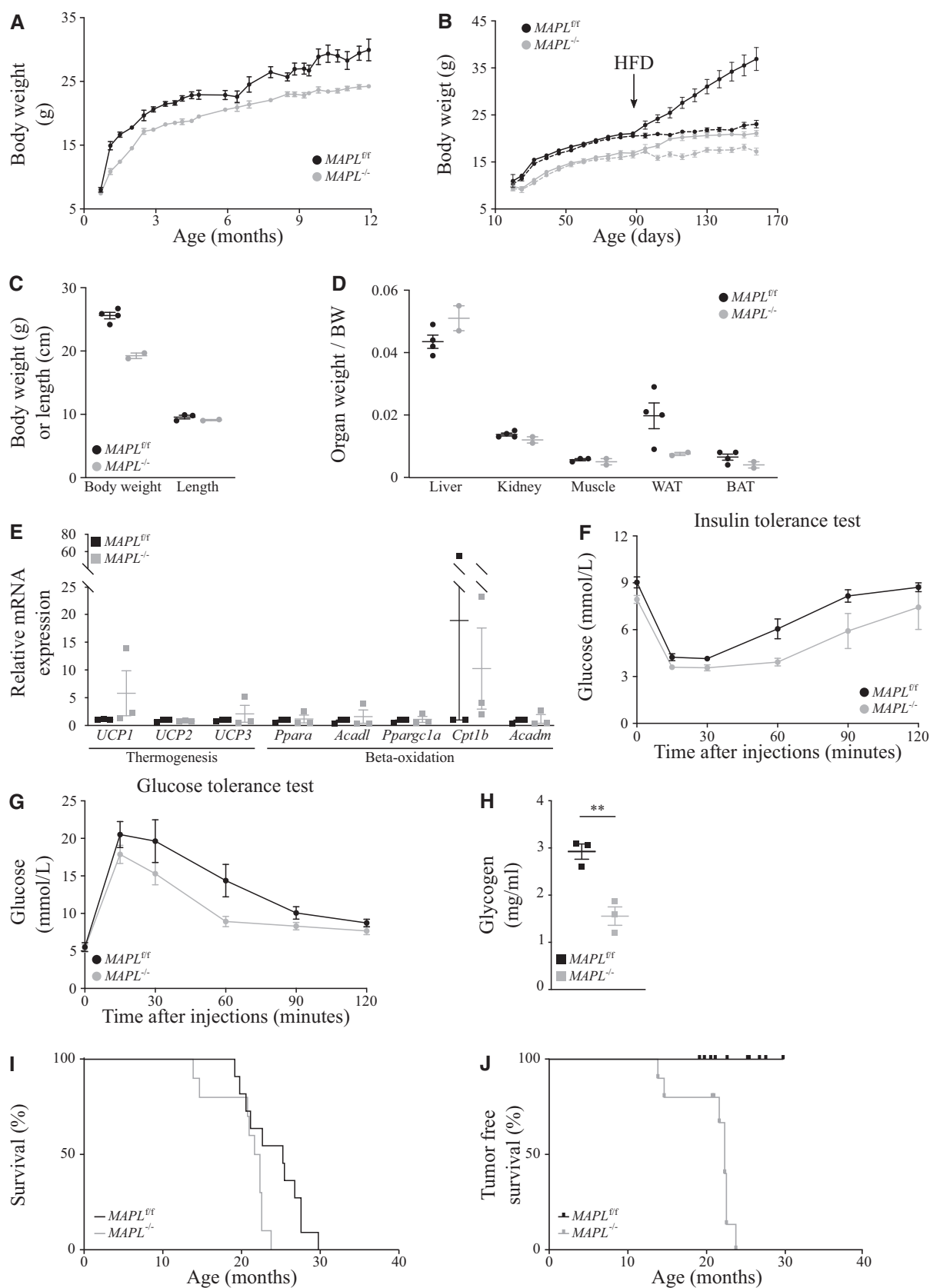


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