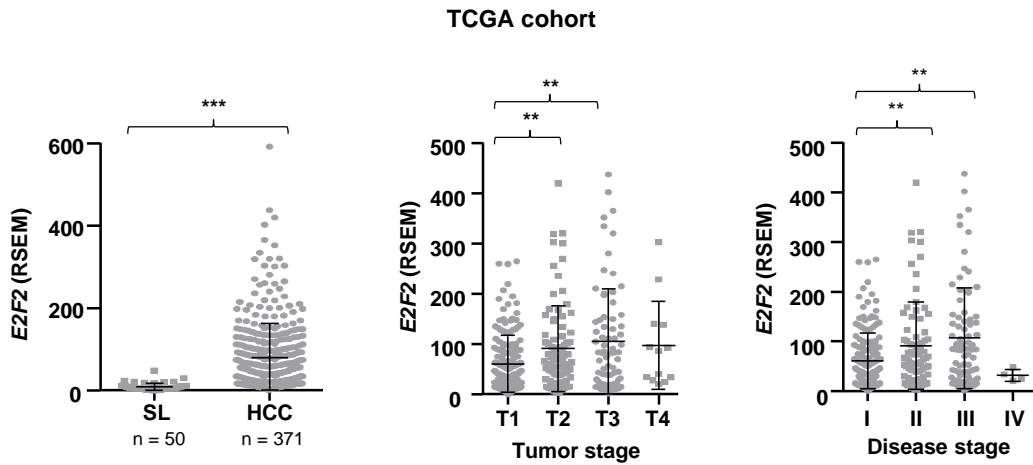
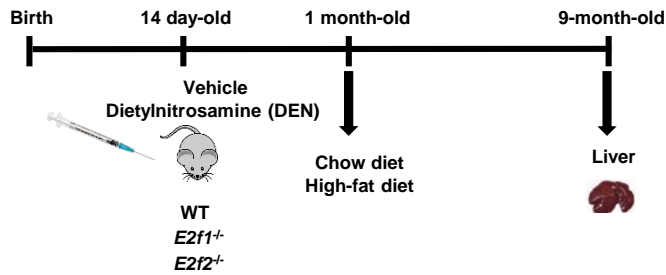


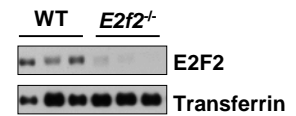
A



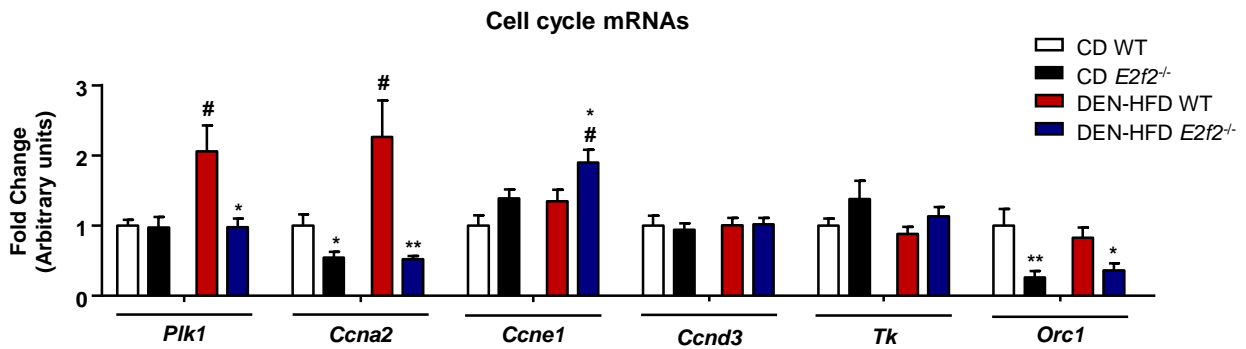
B



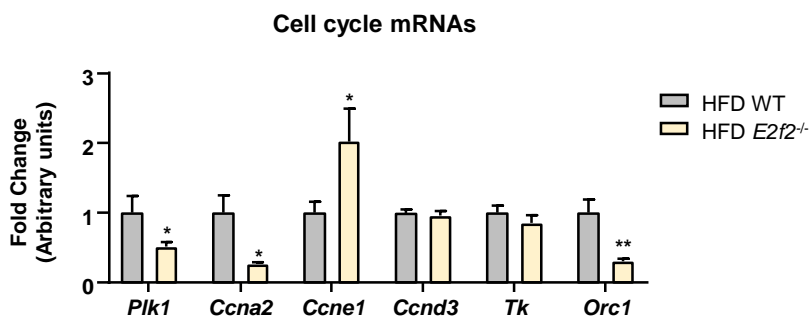
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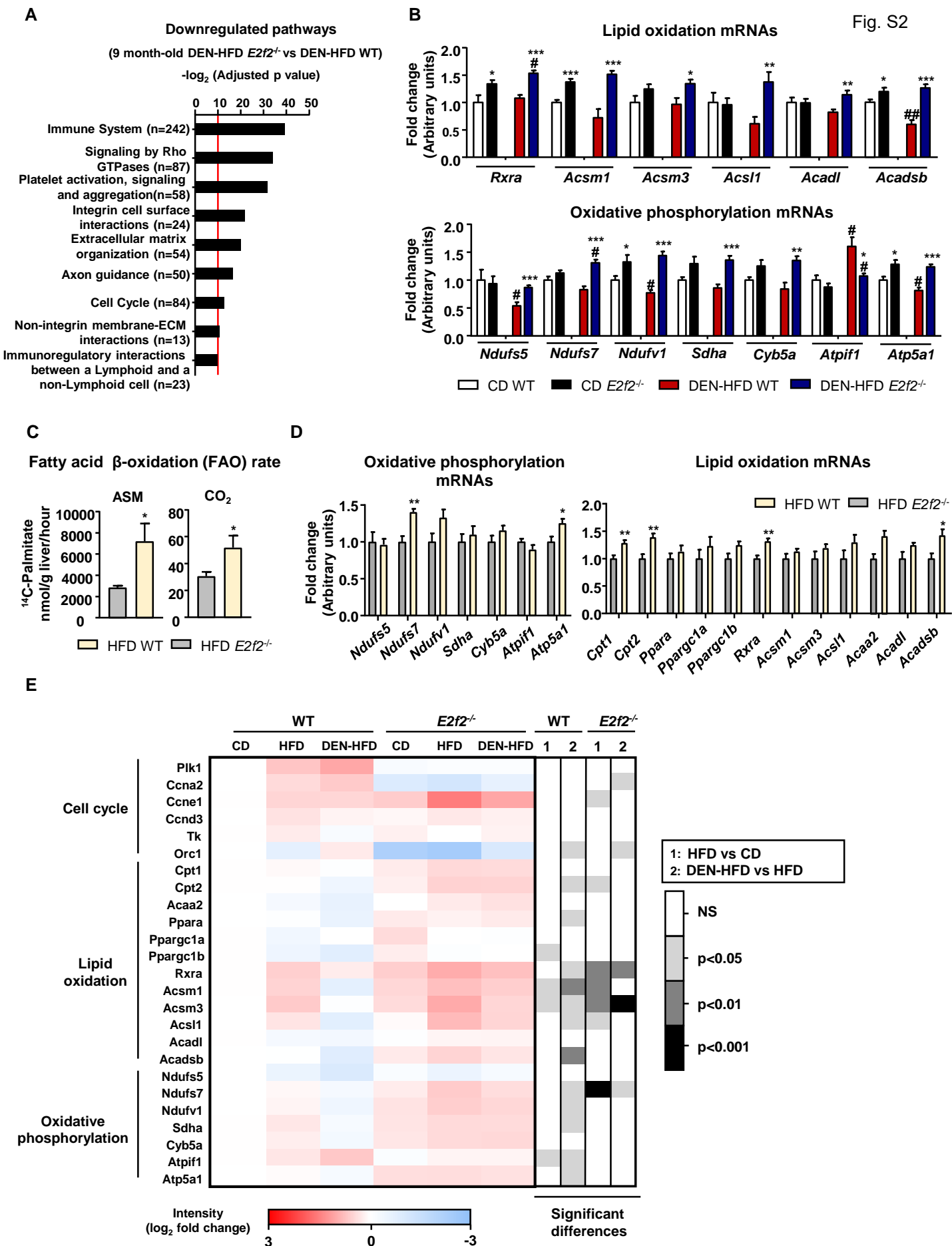


D

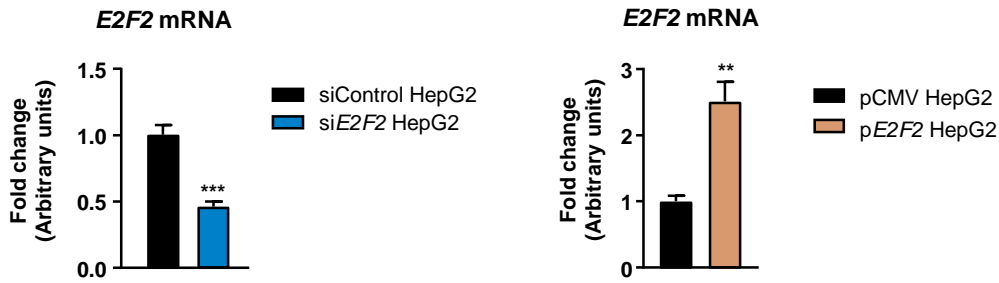


E

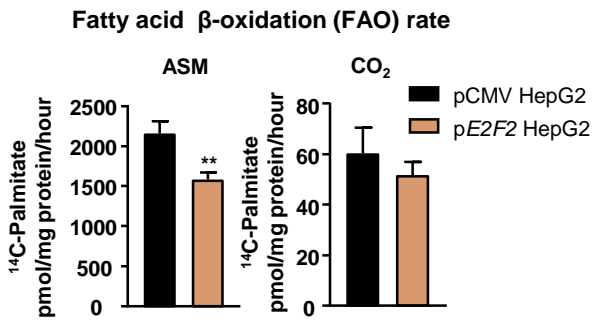




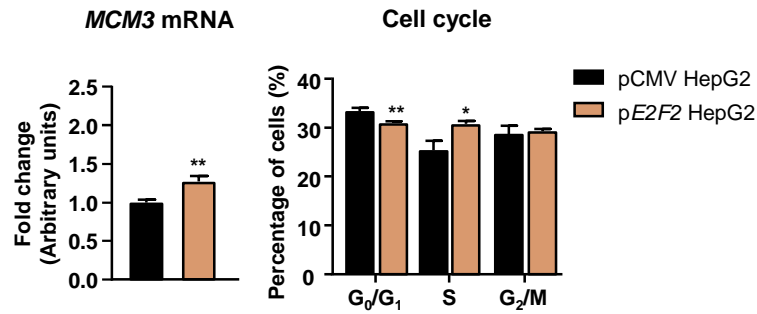
A



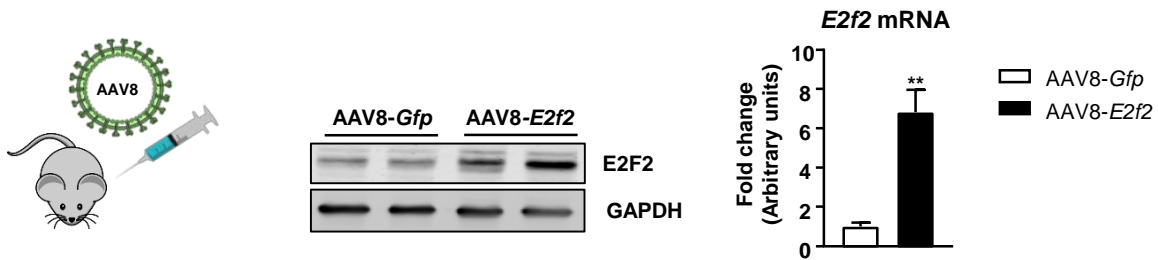
B



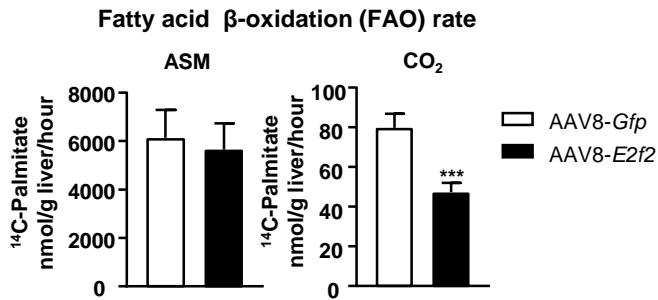
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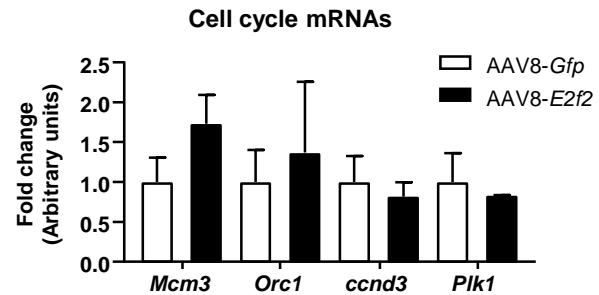
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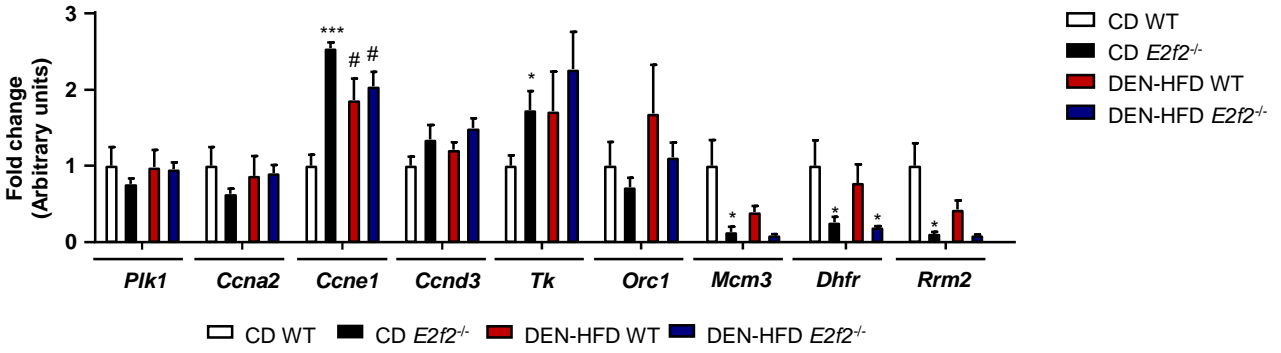
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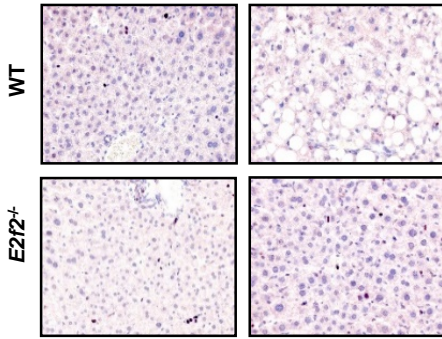
F



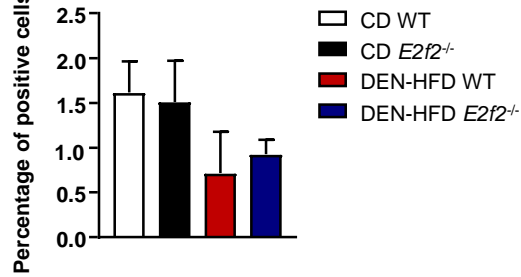
A Cell cycle mRNAs



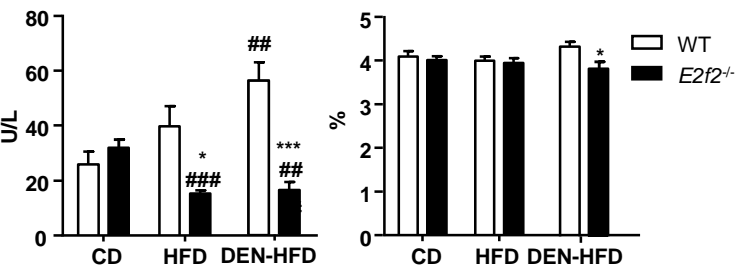
B CD DEN-HFD



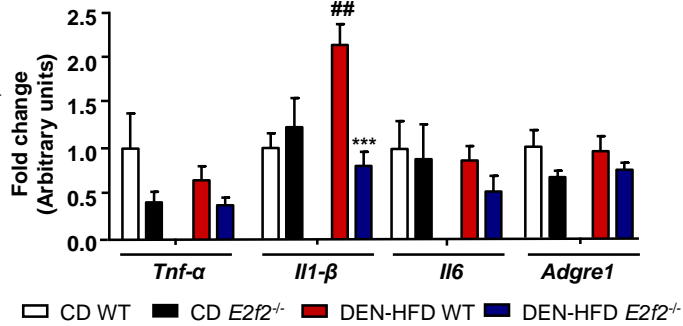
Ki67 levels



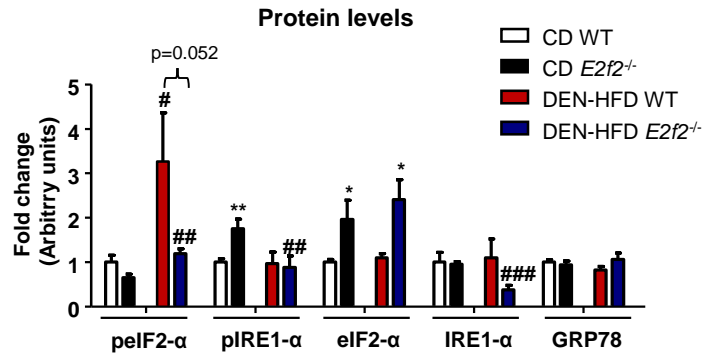
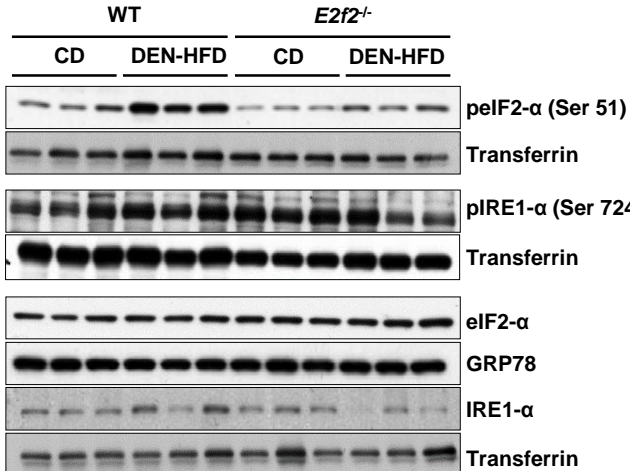
C ALT Liver weight to body weight

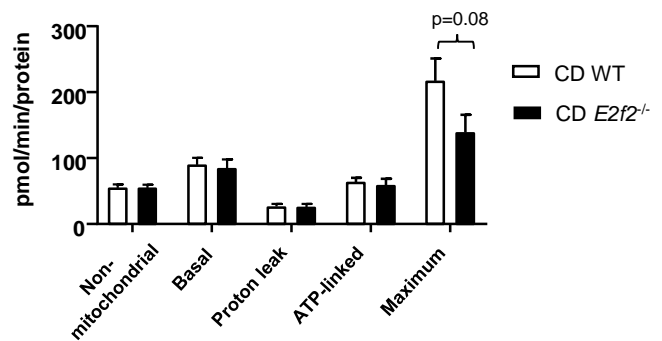
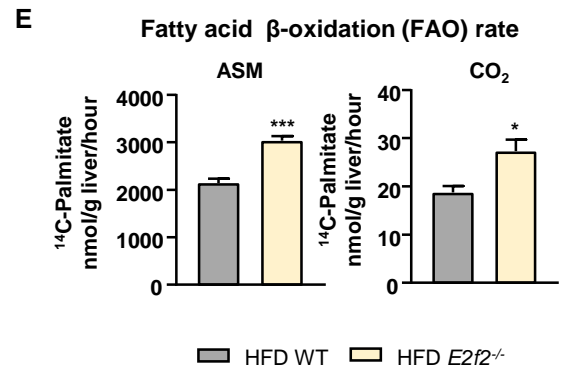
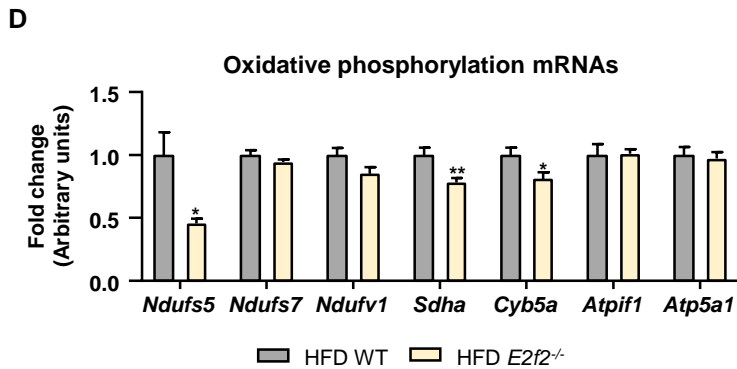
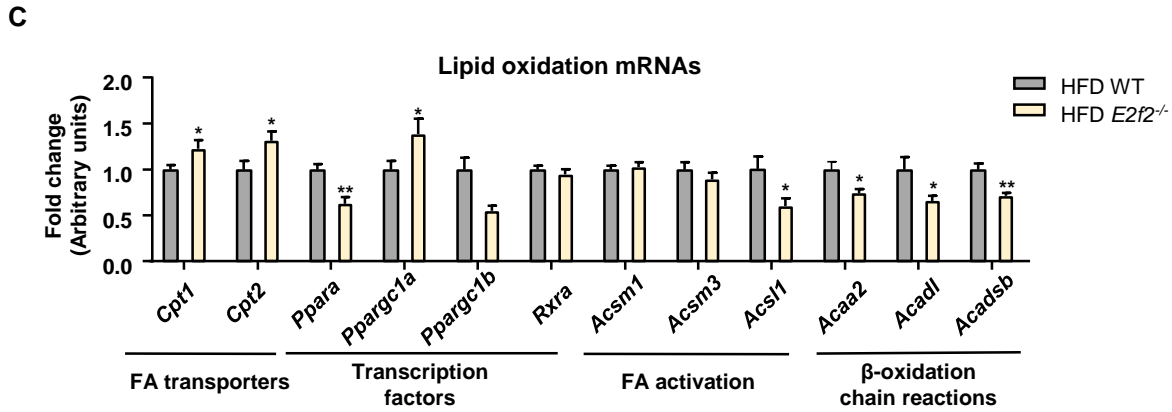
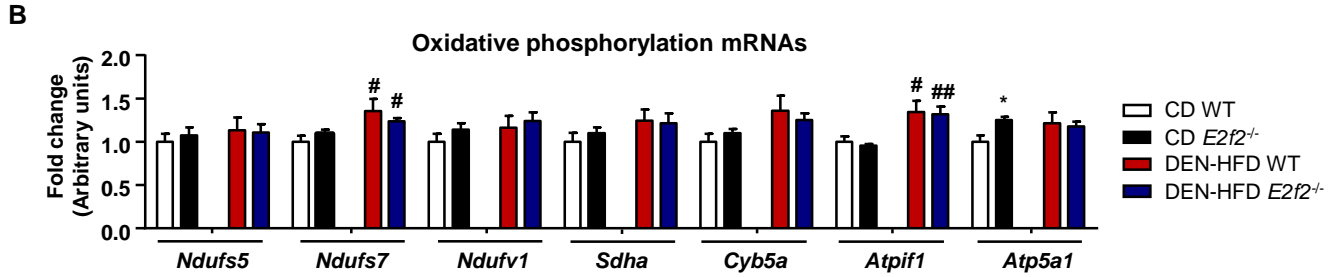
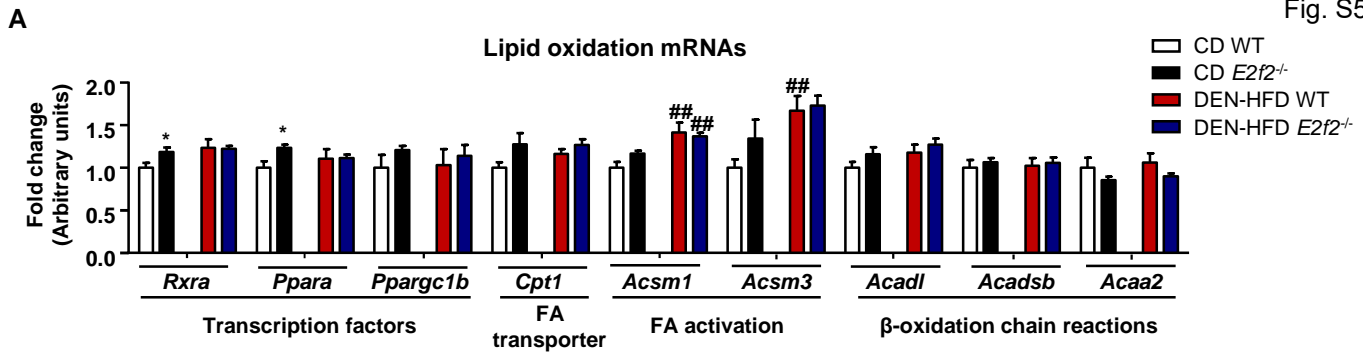


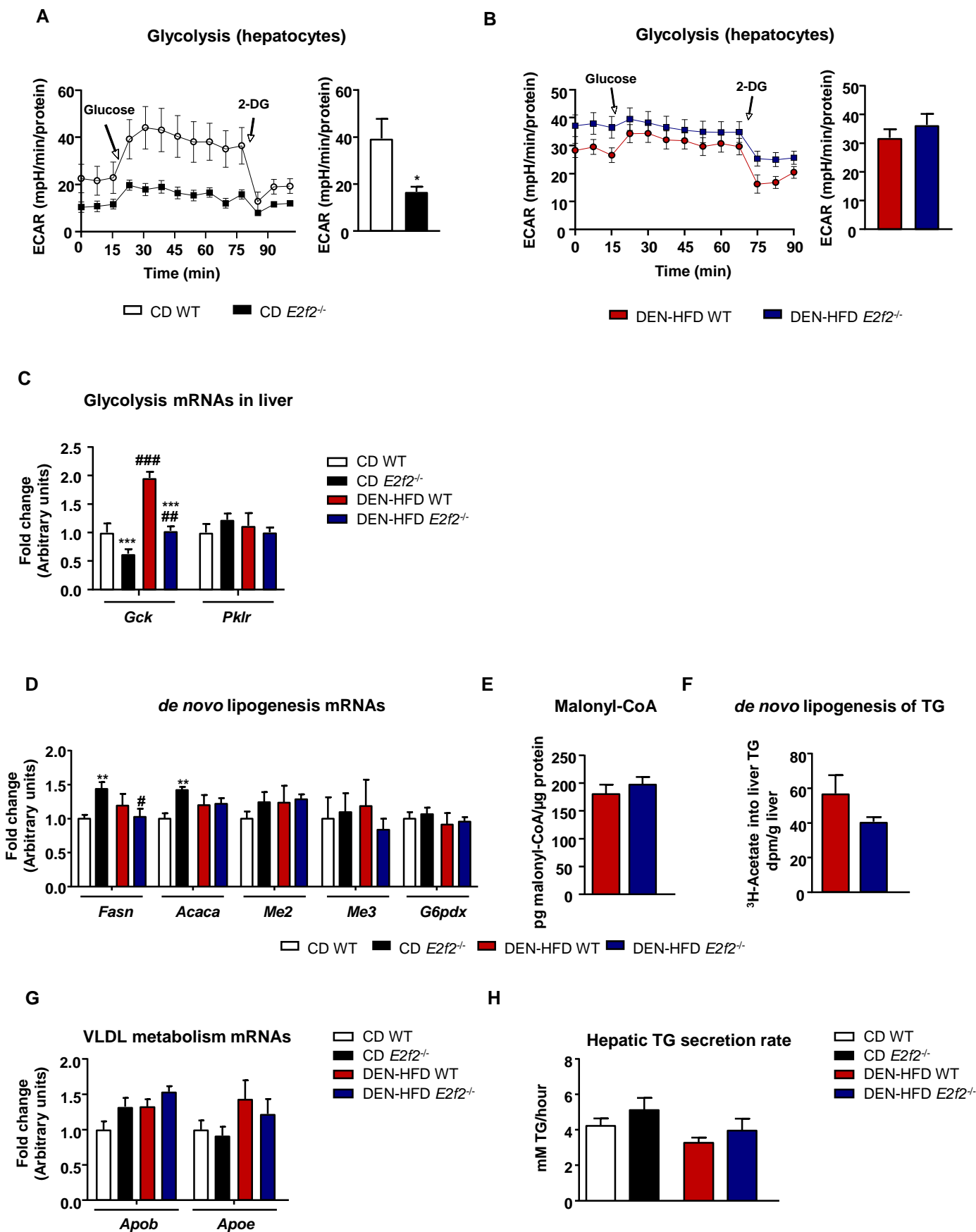
D Inflammation mRNAs



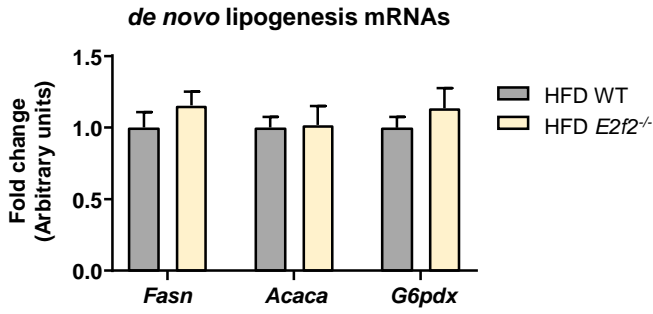
E WT *E2f2*^{-/-}



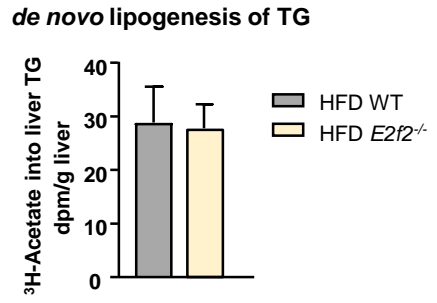




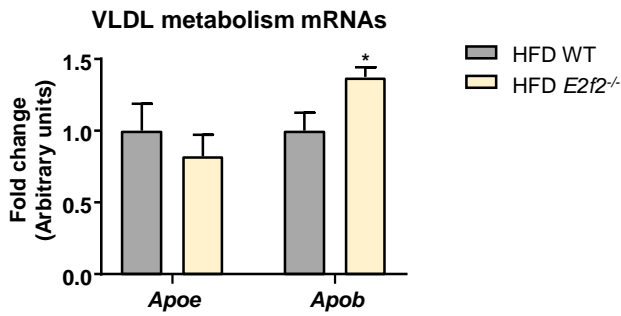
A



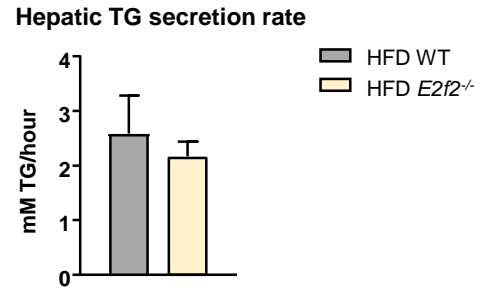
B



C

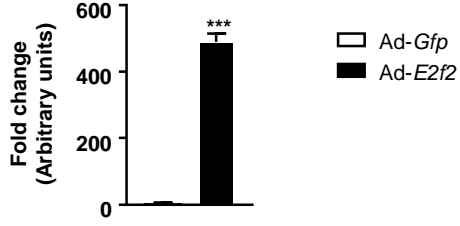
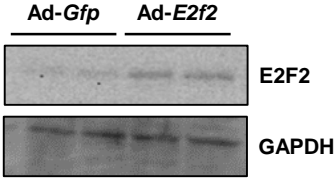


D



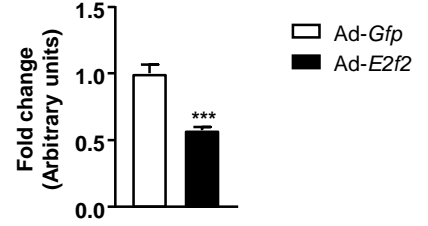
A

E2f2 mRNA (hepatocytes)



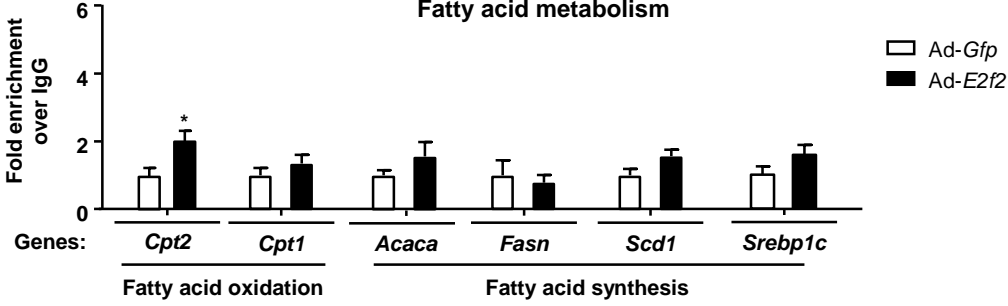
B

Cpt2 mRNA (hepatocytes)



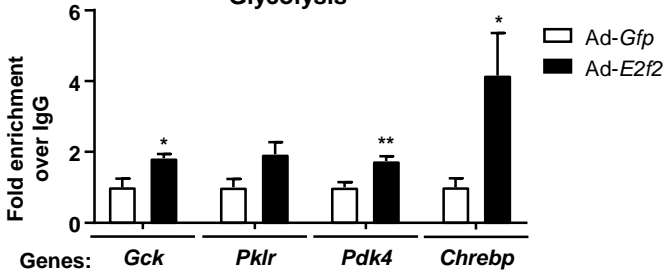
C

Ad-*E2f2* ChIP (hepatocytes)
Fatty acid metabolism



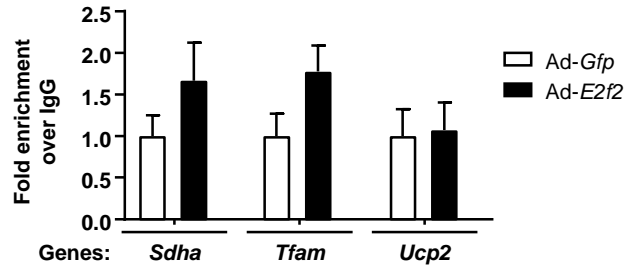
D

Ad-*E2f2* ChIP (hepatocytes)
Glycolysis

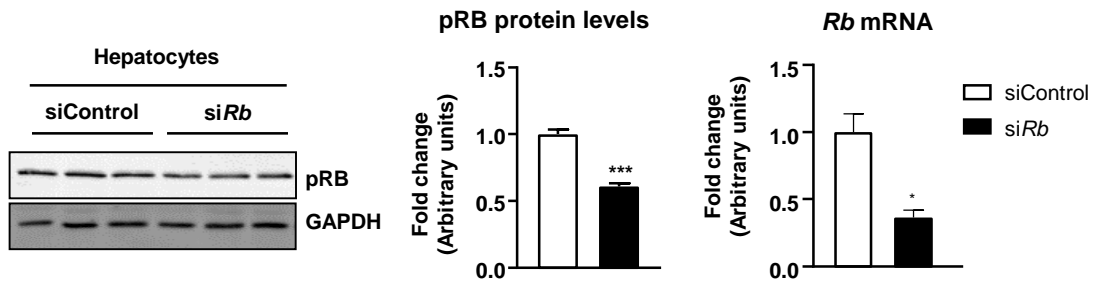


E

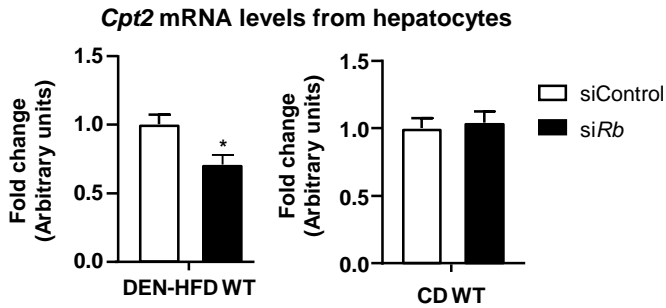
Ad-*E2f2* ChIP (hepatocytes)
Mitochondria



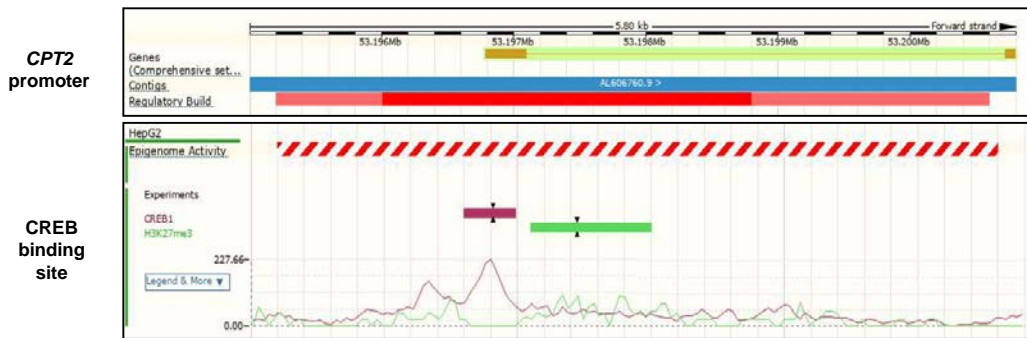
A



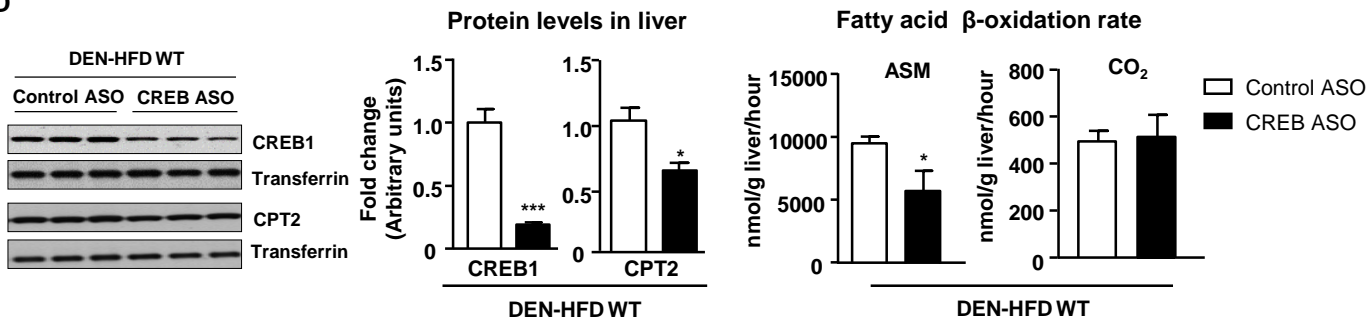
B



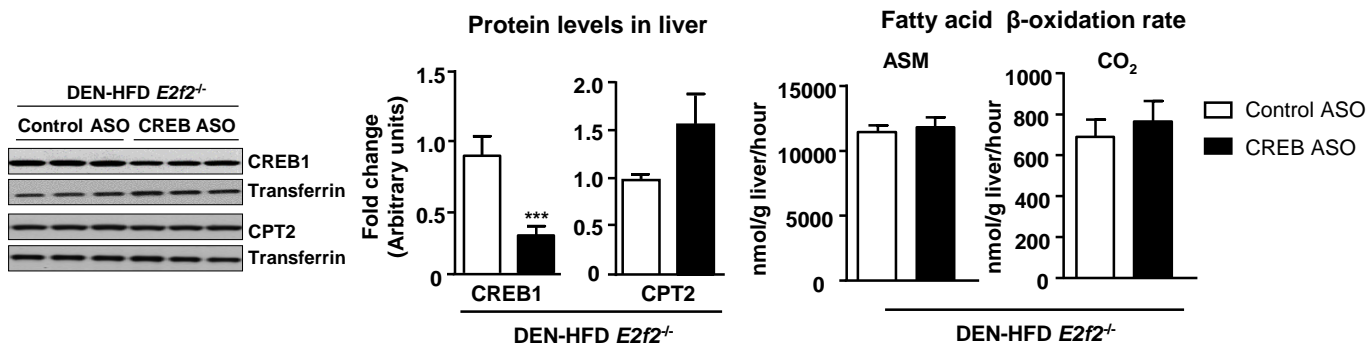
C



D



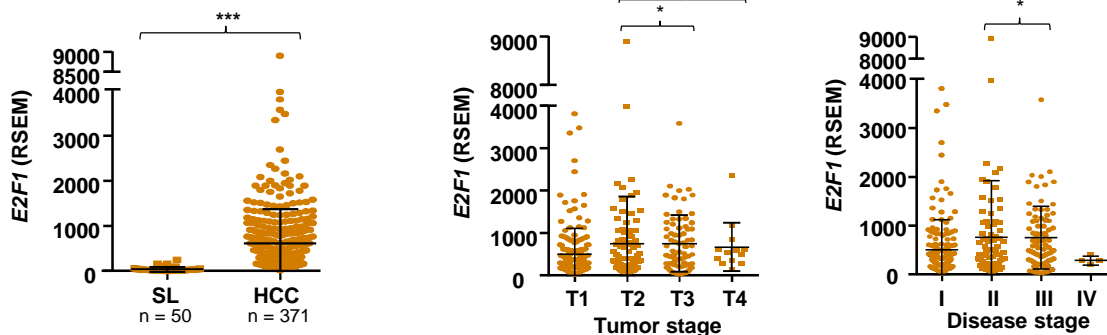
E



A

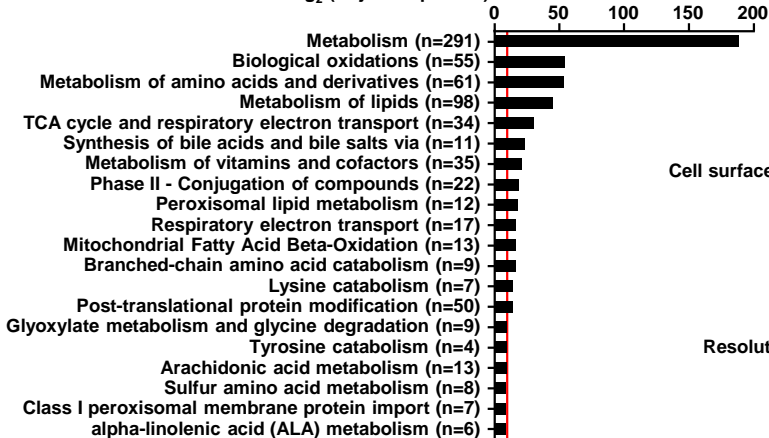
TCGA cohort

Fig. S10



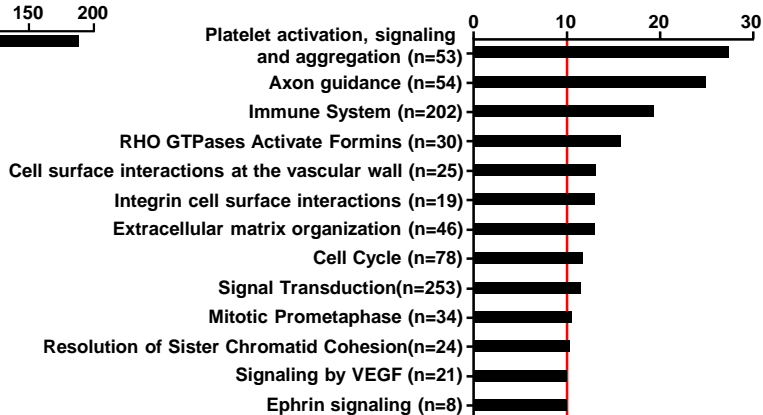
B

Upregulated pathways

(9 month-old DEN-HFD *E2f1*^{+/+} vs DEN-HFD WT)-log₂ (Adjusted p value)

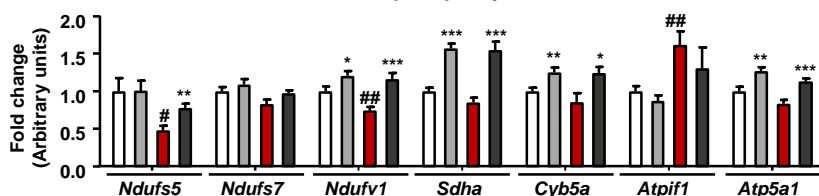
C

Upregulated pathways

(9 month-old DEN-HFD *E2f1*^{-/-} vs DEN-HFD WT)-log₂ (Adjusted p value)

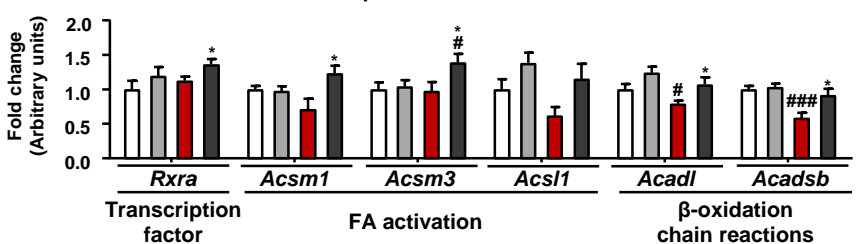
D

Oxidative phosphorylation mRNAs

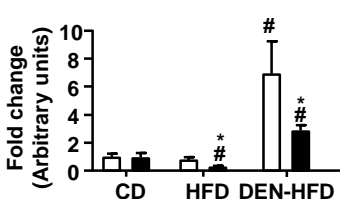


E

Lipid oxidation mRNAs



F

3 month-old *E2f1* mRNA

G

9 month-old *E2f1* mRNA