



Supplementary information, Fig. S14. RNA-seq analysis of cardiomyocytes from adult *Lrp6*-CKO and *Cre*^{+/-} hearts. a Gene ontology (GO) classification of differentially expressed genes in the *Cre*^{+/-} and *Lrp6*-CKO cardiomyocytes. Significant GO terms are shown. **b** Scatter plot for KEGG enrichment results: *Cre*^{+/-} cardiomyocytes versus *Lrp6*-CKO cardiomyocytes. *n* = 2-3 mice per group.