

Supplementary Materials

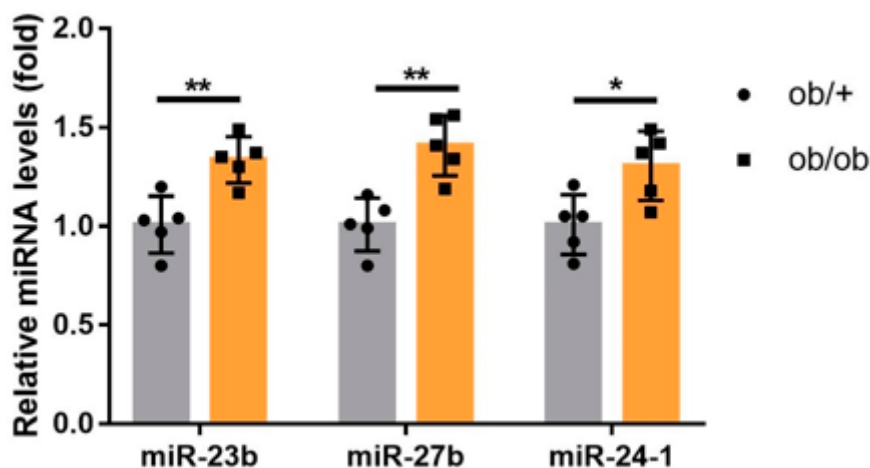


Figure S1. Expression pattern of the miR-23b cluster. miR-23b, miR-27b, and miR-24-1 were increased in the liver of ob/ob mice ($n = 4-5$). Data are represented as mean \pm SEM. Significance was determined by Student's t -test analysis, * $p < 0.05$, ** $p < 0.01$.

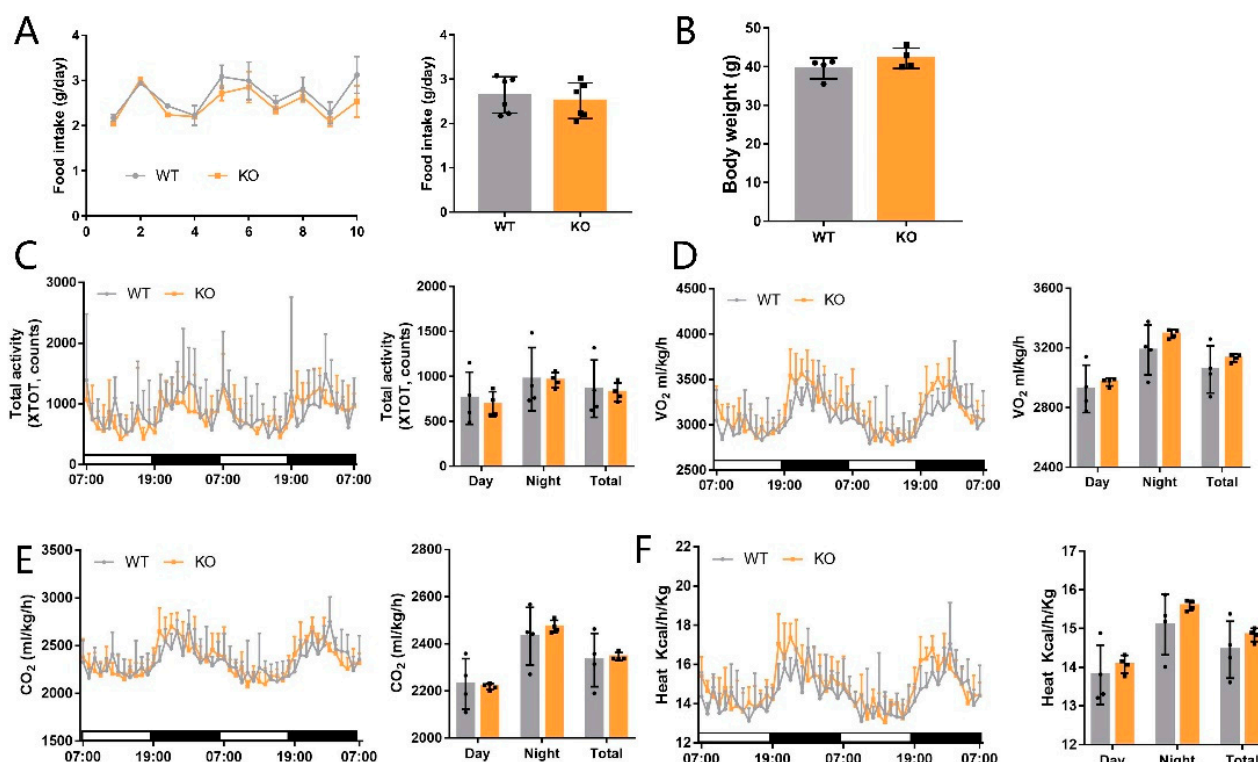


Figure 2. Overall metabolic alterations in knockout mice. (A) Continuous monitoring of food intake for 10 days in WT and KO mice fed an HFD ($n = 6$). Right panel, average food intake per mouse. (B) Body weights of KO and WT mice before metabolic cage experiments. (C–F) Total activity (XTOT), Oxygen consumption (VO₂), CO₂ production (VCO₂), and heat production were measured during 2 successive days using metabolic cages. The daytime and night-time averages are on the right panel. Mice were feed under HFD ($n = 4v4$). Data are represented as mean \pm SD. Significance was determined by Student's t test analysis.

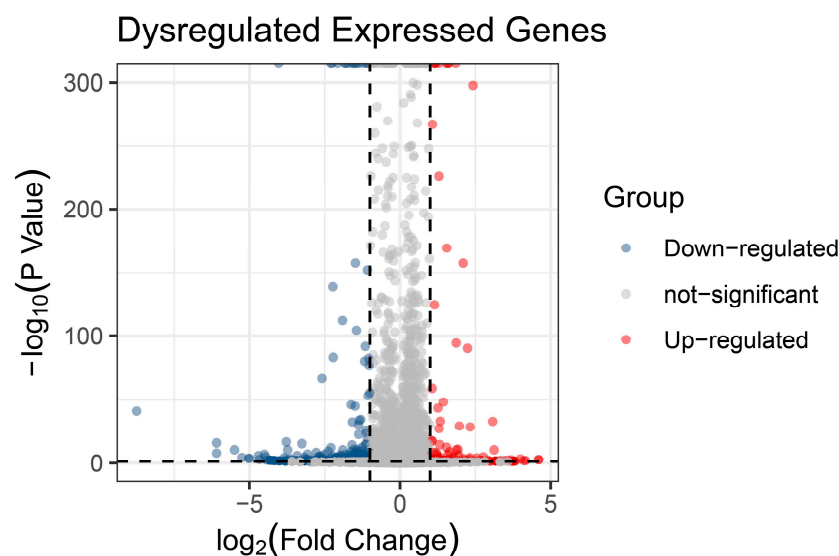


Figure S3. Liver transcriptome profiles after ablation of miR-23b cluster. Dysregulated expressed genes in miR23b cluster-deficient liver tissue compared with WT mice under HFD. The threshold is set to $|\text{Log}_2 \text{FC}| > 1$ and $Q\text{-value} < 0.05$. Red dots represent upregulated and blue dots represent downregulated.

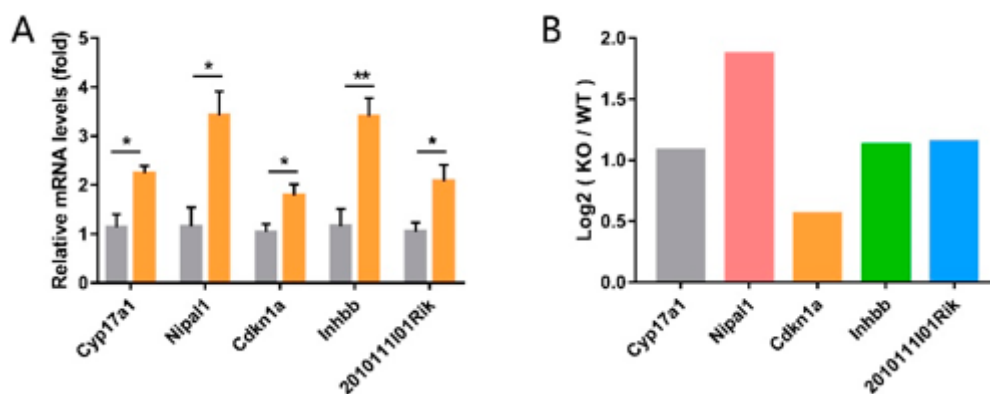


Figure S4. Validation of RNA-seq results. (A) qPCR analysis of mRNA levels from RNA-seq data with relatively high fold changes, high expression, and low p -values ($n = 4$). (B) $\text{Log}_2 \text{FC}$ of validated genes from transcriptomics. Data are represented as mean \pm SEM. Significance was determined by Student's t test analysis, * $p < 0.05$, ** $p < 0.01$.

Table S1. Primer sequences for quantitative real-time PCR.

Target	Forward Primer (5'-3')	Reverse Primer (5'-3')
N209	GCTGTGTTGTCAAGAGTGCA	CAGCAGGGACAAAGAACCAC
T1348	CCAGAGACATCCCAACCCAT	AGACAGGCATTCTCACTGCT
<i>Pklr</i>	GATCCGAAGTTCCGGACAAGG	ATGAGCCCGTCGTCAATGTAG
<i>Cyp17a1</i>	GGCACTGCATCACGATAA	TCCGAAGGGCAAATAACT
<i>Nipal1</i>	GTCAGAGTCGCTGCCTTATCC	TGCAAGAGACCCTTCTTTTGAG
<i>Cdkn1a</i>	CCTGGTGATGTCCGACCTG	CCATGAGCGCATCGCAATC
<i>Inhbb</i>	CTTCGTCTCTAATGAAGGCAACC	CTCCACCACATTCCACCTGTC
2010111101Rik	GTCAACATGGCAGGCTACAG	GGGCTTCATTTCTGTCCAGC
β -actin	CATCCGTAAAGACCTCTATGCCAAC	ATGGAGCCACCGATCCACA
<i>Gapdh</i>	AGGTCGGTGTGAACGGATTTG	TGTAGACCATGTAGTTGAGGTCA
miR-23b-3p	GGCAGACAATCACATTGCCA	TATGGTTGTTACGACTCCTTCAC
miR-27b-3p	CTCGTCAATTCACAGTGGCTAAG	TATGGTTTTGACGACTGTGTGAT
miR-24-3p	CCGGTGCCTACTGAGCTGATAT	TATCCTTCTTCACGACTCCTTCAC
U6	CAGCACATATACTAAAATTGGAACG	ACGAATTTGCGTGTCATCC