

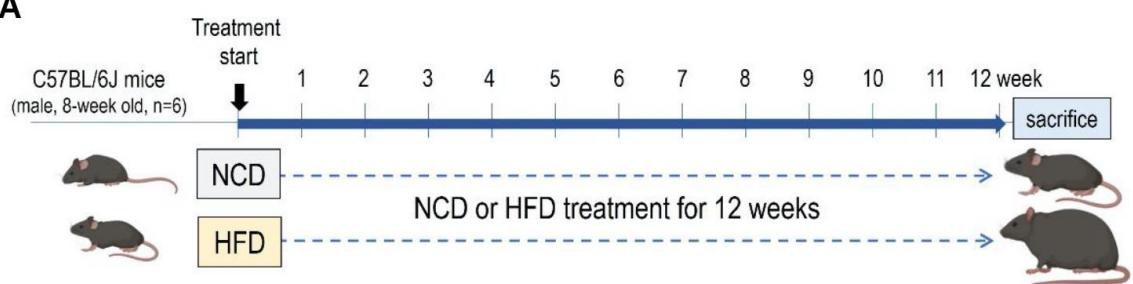
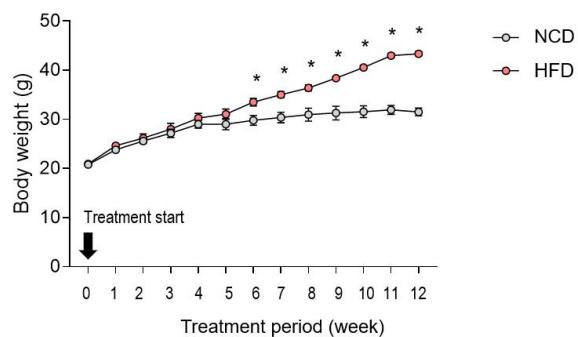
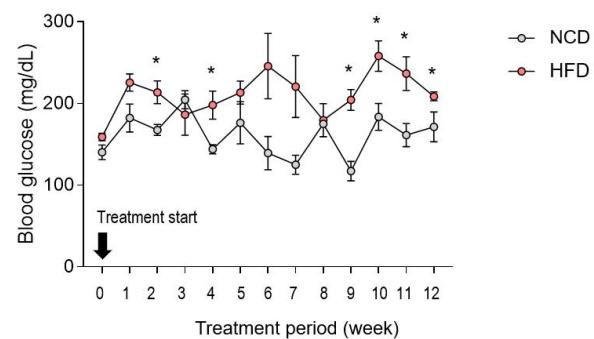
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Figure S1 Metabolic changes of NCD and HFD model mice. (A) Schematic of the experimental plan. (B) Body weight measurement of NCD and HFD fed mice. (C) Blood glucose measurements of NCD and HFD fed mice. Statistical analysis was conducted to determine the difference between NCD and HFD model mice using a Mann-Whitney test (non-parametric *t*-test). *, P<0.05. NCD, normal chow diet mice group; HFD, high-fat diet mice group.

Table S1 Primer list

| Gene name | Direction | Seq. information |
|----------------|-----------|-------------------------|
| mTGF beta | Forward | CTCCCGTGGCTCTAGTGC |
| | Reverse | GCCTTAGTTGGACAGGATCTG |
| m α SMA | Forward | AACGCCTCCGCTGCC |
| | Reverse | CGATGCCGCTGACTCC |
| mCOL1A1 | Forward | GCTCCTCTTAGGGGCCACT |
| | Reverse | CCACGTCTCACCATGGGG |
| mPDGFR | Forward | TTCCAGGAGTGATACCAGCTT |
| | Reverse | AGGGGGCGTGATGACTAGG |
| mTIMP1 | Forward | GCAACTCGGACCTGGTCATAA |
| | Reverse | CGGCCCCGTGATGAGAAACT |
| mGDF15 | Forward | AGCCGAGAGGACTCGAACTCAG |
| | Reverse | GGTGACGCGGAGTAGCAGCT |
| mGFRAL | Forward | TTCCTGGCTGTTACGTTAAC |
| | Reverse | GCCATTGACATCAATCAAGCA |
| mIL-1 β | Forward | AAGATCGACACATTGATTCCAGC |
| | Reverse | CATGGTTGAAGTGAATTCCCAC |
| mIL-6 | Forward | CTTCGACGTGACAGACGCT |
| | Reverse | GCAGGGGCAGTGAAACTTATT |
| mTNF α | Forward | CAGCACACTCGATATGGACCA |
| | Reverse | CCTCGGGCTCAGGATAGTCT |
| mIL-10 | Forward | AGACCAAGGTGTCACAAGGC |
| | Reverse | CCAAGGAGTTTTCCGTTAGC |
| mCox-2 | Forward | AAGCCTCTCCAACCTCTCC |
| | Reverse | GCTGGCAAAGAATGCAAAC |
| mCCI2 | Forward | CATCTGCCCTAACGGTCTTCAG |
| | Reverse | GGTTGTGAAAAGGTAGTGG |
| mL32 | Forward | TCTGGTGAAGCCCCAAGATGG |
| | Reverse | CTCTGGGTTCCGCCAGT |
| mBax | Forward | TGAAGACAGGGGCCCTTTG |
| | Reverse | AATTGCCGGAGACACTCG |
| mBCL-2 | Forward | GTCGCTACCGTCGTGACTTC |
| | Reverse | CAGACATGCACCTACCCAGC |
| mAco | Forward | ATGCACTACAGCGTCCATGA |
| | Reverse | AATTCTACCAATCTGGCTGCAC |
| mCyp2e1 | Forward | CTTGCAGAACAGAGACCA |
| | Reverse | ATGCACTACAGCGTCCATGA |

Table S1 (continued)

| Gene name | Direction | Seq. information |
|-------------|-----------|------------------------------|
| mCyp4a10 | Forward | CAACTTGCCCCATGATCACACA |
| | Reverse | CATCCTGCAGCTGATCCTTC |
| mNox2 | Forward | GAAAACCTCTGGGTCAAGCACT |
| | Reverse | ATTCGACACACTGGCAGCA |
| mGCL-c | Forward | GTTATGGCTTGAGTGCTGCAT |
| | Reverse | ATCACTCCCCAGCGACAATC |
| mGpx-1 | Forward | CCAGGAGAATGGCAAGAATGA |
| | Reverse | TCTCACCAATTCACTTCGCACTT |
| mSod2 | Forward | TCTCACCAATTCACTTCGCACTT-3 |
| | Reverse | GGTGGCGTTGAGATTGTTCA-3 |
| mG6pdh | Forward | CTGGAACCGCATCATCGTGGAG-3 |
| | Reverse | CCTGATGATCCAAATTCACTAAATAG-3 |
| mG6pase | Forward | CGACTCGCTATCTCCAAGTGA |
| | Reverse | GTTGAACCAGTCTCCGACCA |
| mPEPCK | Forward | AACTGTTGGCTGGCTCTC |
| | Reverse | GAACCTGGCGTTGAATGC |
| mNRF2 | Forward | TCTTGAGTAAGTCGAGAAGTGT |
| | Reverse | GTTGAAACTGAGCGAAAAAGGC |
| mPGC-1alpha | Forward | TATGGAGTACATAGAGTGTGCT |
| | Reverse | CCACTTCAATCCACCCAGAAAG |
| m c-Fos | Forward | CTGTCGCTCTCTAGTGCCAAC |
| | Reverse | CCTCCTGACACGGTCTTCAC |
| mPSD95 | Forward | GGTAACTCAGGTCTGGCTTC |
| | Reverse | CACTGCAGCTGAATGGGTCA |
| mSyp | Forward | ACATGCAAGGAACTGAGGGA |
| | Reverse | CCAGGTTCAGGAAGCCAAAC |

Table S1 (continued)