

Supplemental Data

Figure S1

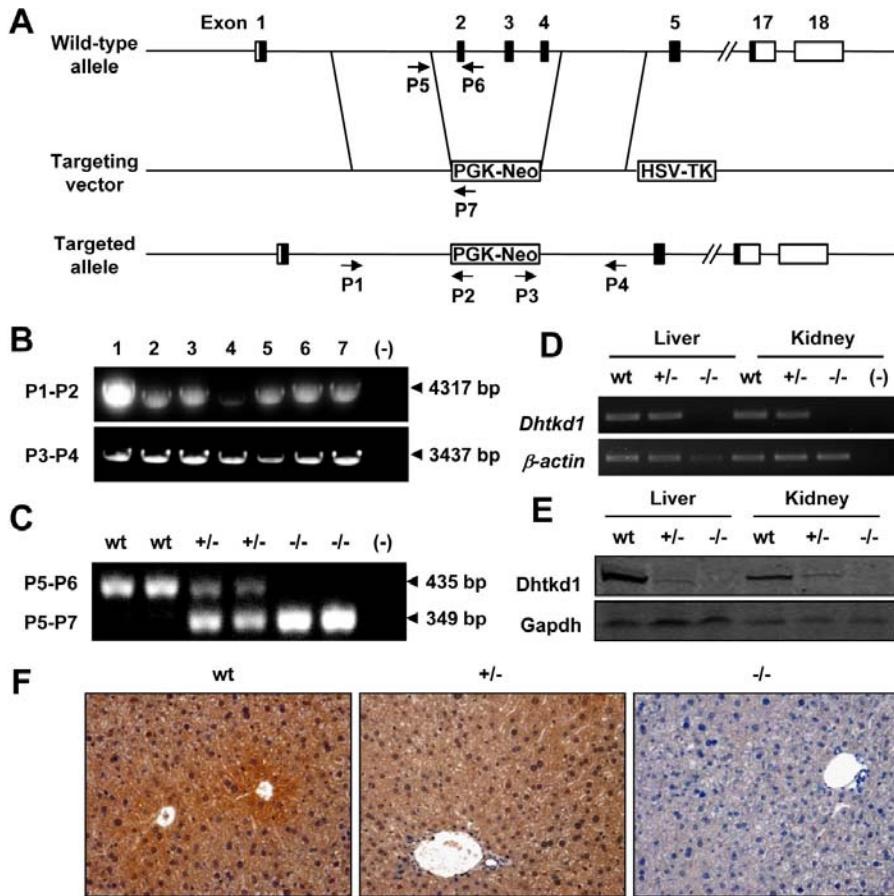


Figure S1. Targeted deletion of *Dhtkd1* in mice. (A) Schematic representation of *Dhtkd1* gene knock-out strategy. The targeting vector was designed to delete the genomic region from exon 2 to exon 4. Exons and introns are represented by boxes and horizontal lines, respectively. The coding region is shown in black. Fragments amplified by PCR for genotyping are indicated by arrows. (B) PCR analysis of recombinant ES cell clones. Genomic DNA extracted from ES cell clones was amplified using 5'-external and 3'-external primers as shown in panel. Homologous recombination events yielded a 4.3-kb or 3.4-kb fragments, respectively. (C) PCR genotyping of progenies from heterozygous matings. Wild-type (wt), heterozygous (*Dhtkd1*^{+/−}) and homozygous (*Dhtkd1*^{−/−}) mice were identified by PCR amplification of the fragments specific for either *Dhtkd1* wild-type allele (435 bp) or the targeted allele (349 bp). (D) RT-PCR analysis of *Dhtkd1* mRNA expression in liver and kidney of wt, heterozygous, and homozygous mice. *β-actin* was used as loading control. (E) RT-PCR analysis of *Dhtkd1* mRNA expression in liver and kidney of wt, heterozygous, and homozygous mice. *Gapdh* was used as loading control. (F) Immunohistochemistry (IHC) analysis of *Dhtkd1* protein expression in liver tissue of wt, heterozygous, and homozygous mice. Brown staining indicates positive expression; blue staining indicates hematoxylin counterstain.

(349 bp). (D) RT-PCR analysis for *Dhtkd1* mRNA in liver and kidney with primer e1/e3. The 418-bp *Dhtkd1*-specific fragment is absent only in *Dhtkd1*^{-/-} mice. *Actb* is used as a loading control. (E) Western blot analysis shows that Dhtkd1 protein is disappeared in liver and kidney of *Dhtkd1*^{-/-} mice. (F) Immunohistochemical staining for Dhtkd1 in liver. Magnification, 200×.

Figure S2

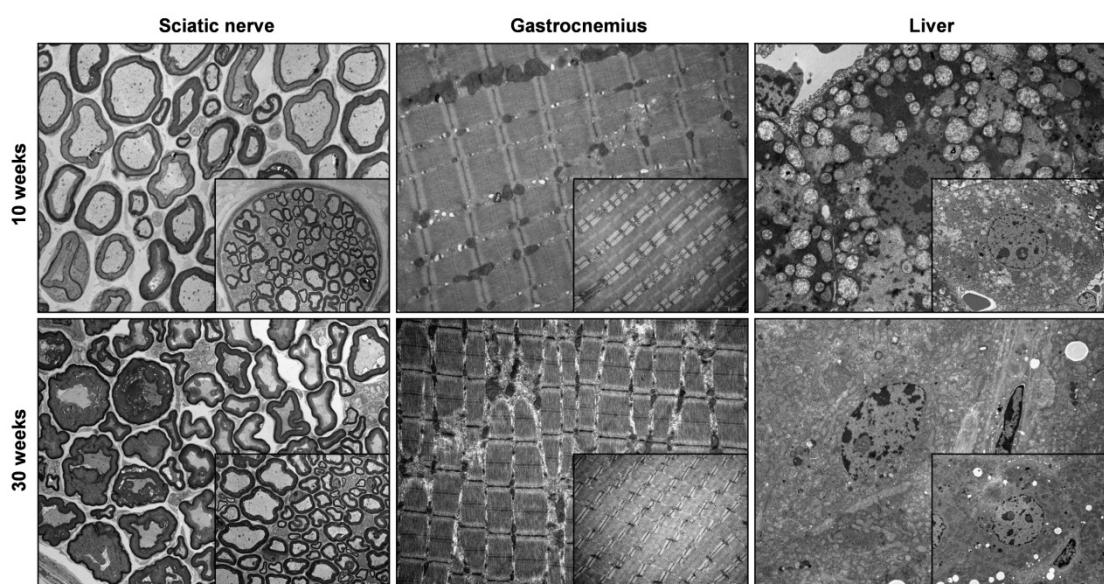


Figure S2. Aggravation of peripheral nerve, muscle and liver changes with the age of mice lacking *Dhtkd1*. Electron micrographs of cross-sections of sciatic nerves, gastrocnemius muscle and liver in the young adulthood (10 weeks old) and middle age (30 weeks old) of *Dhtkd1*^{-/-} mice are shown. The inserts indicate morphology of wt tissues.

Figure S3

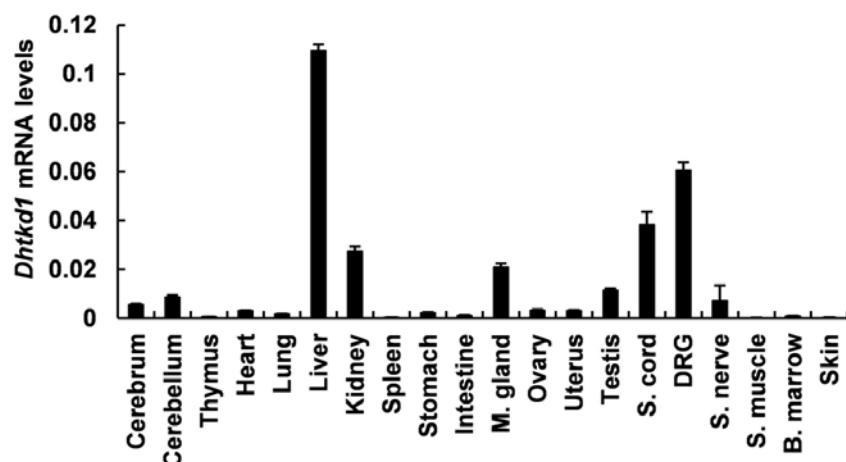


Figure S3. *Dhtkd1* expression profile in mice. Real-time PCR analysis for *Dhtkd1* mRNA expression.

Values are shown as means \pm S.D. M. gland, mammary gland; S. cord, spinal cord; DRG, dorsal root ganglion; S. nerve, sciatic nerve; S. muscle, skeletal muscle; B. marrow, bone marrow.

Figure S4

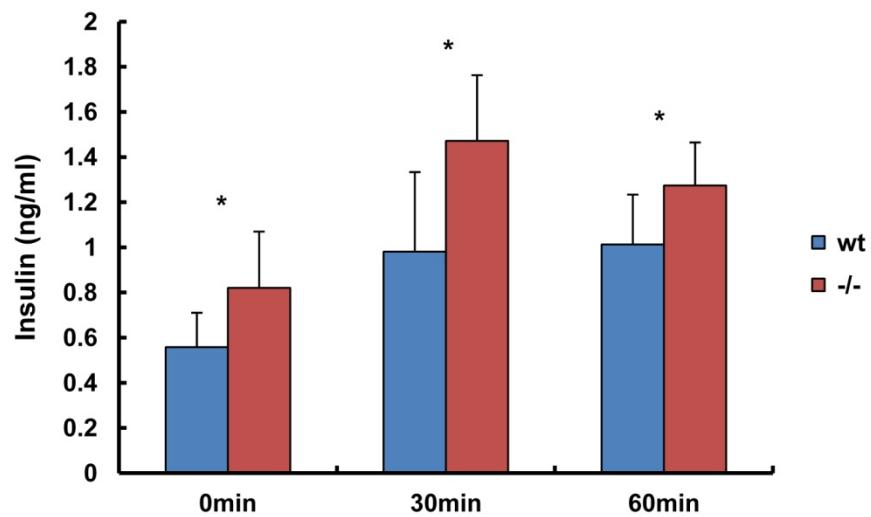


Figure S4. Insulin levels during GTT, related to Fig3G. During the GTT, insulin levels of different time

points were determined by ELISA. Values are shown as means \pm S.D.

Figure S5

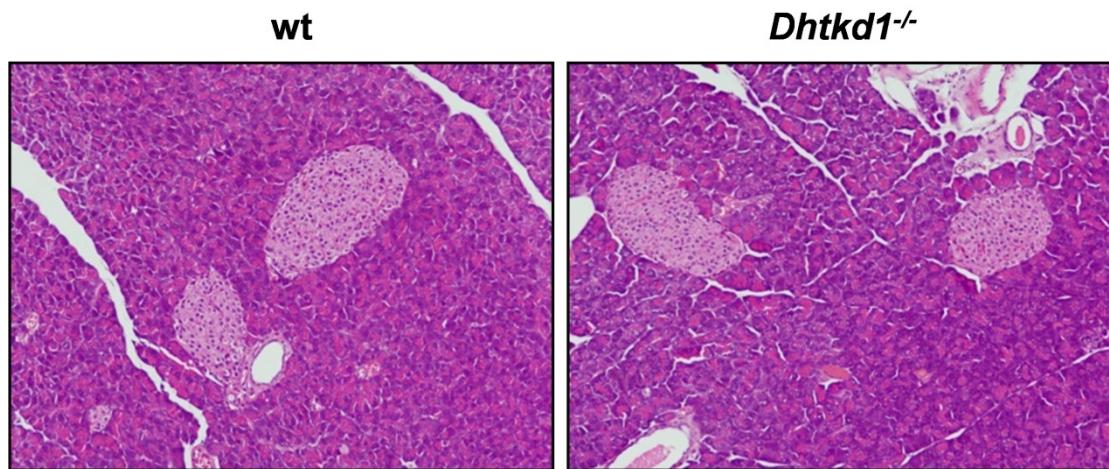


Figure S5. Comparison of islet structure between wt and *Dhtkd1*^{-/-} mice. H&E staining of tissue and cellular morphology in wt and *Dhtkd1*^{-/-} pancreas and islets, showing no significant structural difference between two genotypes.

Figure S6

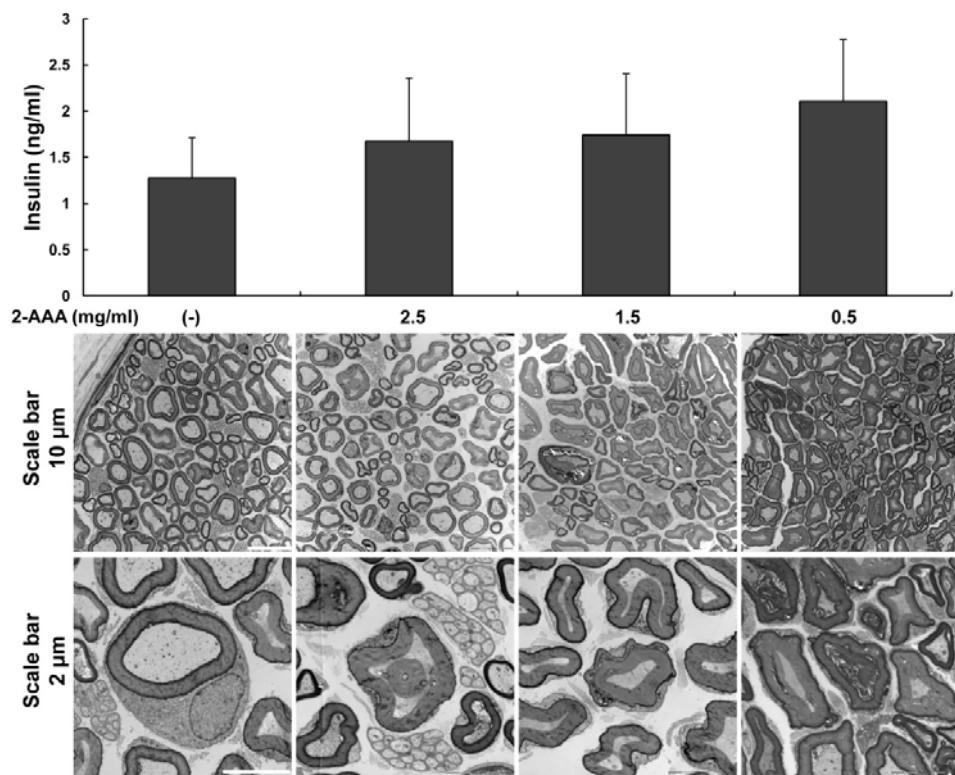


Figure S6. Insulin levels and pathological changes (TEM) of sciatic nerve.

Table S1. The primers used in this study

P1	GAAAAATGGCGATCCTCAAG
P2	GGGGAACCTCCTGACTAGGG
P3	TCGCCTTCTTGACGAGTTCT
P4	CTCTGAGCAGACCTGGGAAC
P5	GTTGGCTTGGTGTGCGTC
P6	AAGGCTCTCCTGCCAGTTC
P7	GGTAGAATTGACGACCTGC
Dhtkd1-e1	ATTCTCCACGGCGTCTT
Dhtkd1-e3	CTCTCTCTCCTGGCTCTGAA
Dhtkd1	GCTGCTGCGGTTATTGGTG
	CAAGGCAAGTGTCTCTGCG
Gjb1	GCACGTAGCTCACCAACAG
	TGATGACATAGGTCCACCACA
Mpz	TACAGTGACAACGGCACTTC
	GCAGTACCGAATCAGGTAGAAGA
Mbp	GGCCAGTAAGGATGGAGAGAT
	CCTCTGAGGCCGTCTGAGA
Pmp22	ATGGACACACGACTGATCTCT
	CAGCCATTGCTCACTGATGA
Prx	GGCGGCAAAGAAGGAATC
	CAAAGAACACACGGGCACT

Srebp2	CAGGTGCAGACGGTACAGG
	CGTGGTCAACACAAGGGAATC
Fbxo32	CAGCTTCGTGAGCGACCTC
	GGCAGTCGAGAAGTCCAGTC
MuRF1	GTGTGAGGTGCCTACTTGCTC
	GCTCAGTCTTCTGTCCCTGGAA
Glut2	TCAGAAGACAAGATCACCGGA
	GCTGGTGTGACTGTAAGTGGG
Gck	AGACGAAACACCAGATGTATTCC
	GAAGCCCTTGGTCCAGTTGAG
Pcx	AATGTCCGGCGTCTGGAGTA
	ACGCACGAAACACTCGGAT
Ins1	CACTTCCTACCCCTGCTGG
	ACCACAAAGATGCTGTTGACA
Ins2	GCTTCTTCTACACACCCATGTC
	AGCACTGATCTACAATGCCAC
Syt14	TGAGAAAAGGATTCGGCGACT
	CTGGCACAGGTTCGATCACT
Ucn3	AAGCCTCTCCCACAAGTTCTA
	GAGGTGCGTTGGTTGTCATC
Glp1r	ACGGTGTCCCTCTCAGAGAC
	ATCAAAGGTCCGGTTGCAGAA

Abcc8	TGAGCATTGGAAGACCCTCAT
	CAGCACCGAAGATAAGTTGTCA
Kcnj11	AAGGGCATTATCCCTGAGGAA
	TTGCCTTCTTGGACACGAAG
Pclo	AAGAGTTGGATAGTAGTCAGGCT
	ACCTAACGTGTCCGTAGTTCT
Noc2	AGCAGAGAGGCTAGACATCCT
	AGACACTGGGAGAGACCGTT
Pcsk1	CTTCGCCTCTTTGCGTTT
	TCCGCCGCCATTCAATTAAAC
Erol1b	ACCCTGAGCTTCCTCTCAAGT
	AAAGGACATGGTCGTTTCAGATT
Slc30a8	AGCCACCAAGATGTACGCC
	CTTGCTTGCTCGACCTGTT
Egr2	GCCAAGGCCGTAGACAAAATC
	CCACTCCGTTCATCTGGTCA
S100b	TGGTTGCCCTCATTGATGTCT
	CCCATCCCCATCTCGTCC
Bdnf	TTACCTGGATGCCGAAACAT
	TGACCCACTCGCTAATACTGTC
Ngf	TGATCGGCGTACAGGCAGA
	GCTGAAGTTAGTCCAGTGGG

β -actin	GGCTGTATTCCCCCTCCATCG
	CCAGTTGGTAACAATGCCATGT