

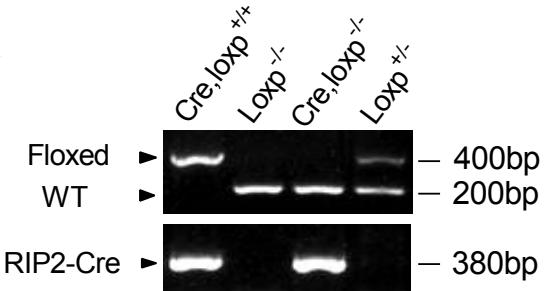
Supplemental Materials

Molecular Biology of the Cell

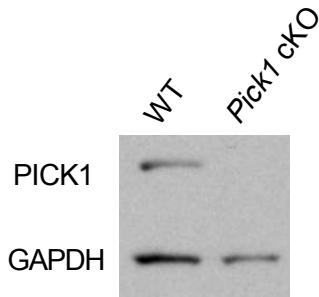
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Figure S1

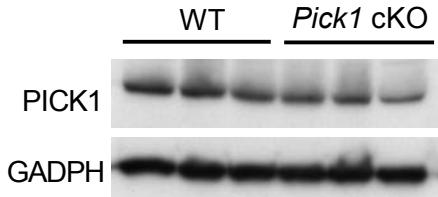
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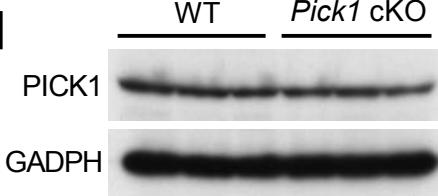
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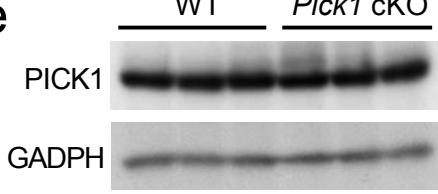
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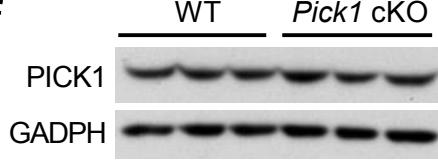
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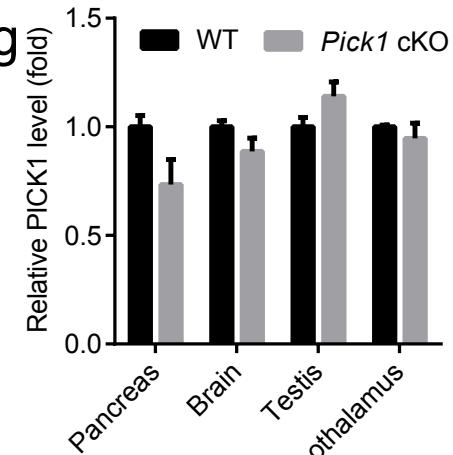
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f



g



h

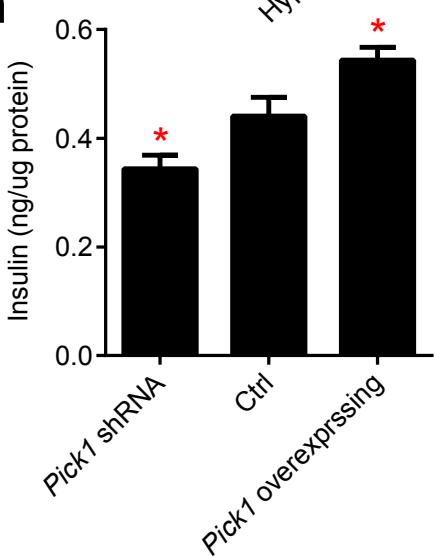


Fig. S1.(a) Representative PCR genotyping results of the indicated genotypes. (b) Western blot analysis of isolated islets from WT and cKO mice aged 4 months old. (c-f) Western blot analysis of PICK1 in the pancreas, brain, testis and hypothalamus from 5-month-old control and PICK1 cKO animals. (g) Relative PICK1 protein level was measured and normalized to that of GAPDH, n=3 per group. (h) Evaluation of insulin in INS1E cells with reduced or overexpressed PICK1. Data are represented as the mean ± SEM. *P<0.05; two sided student's t-test.

Figure S2

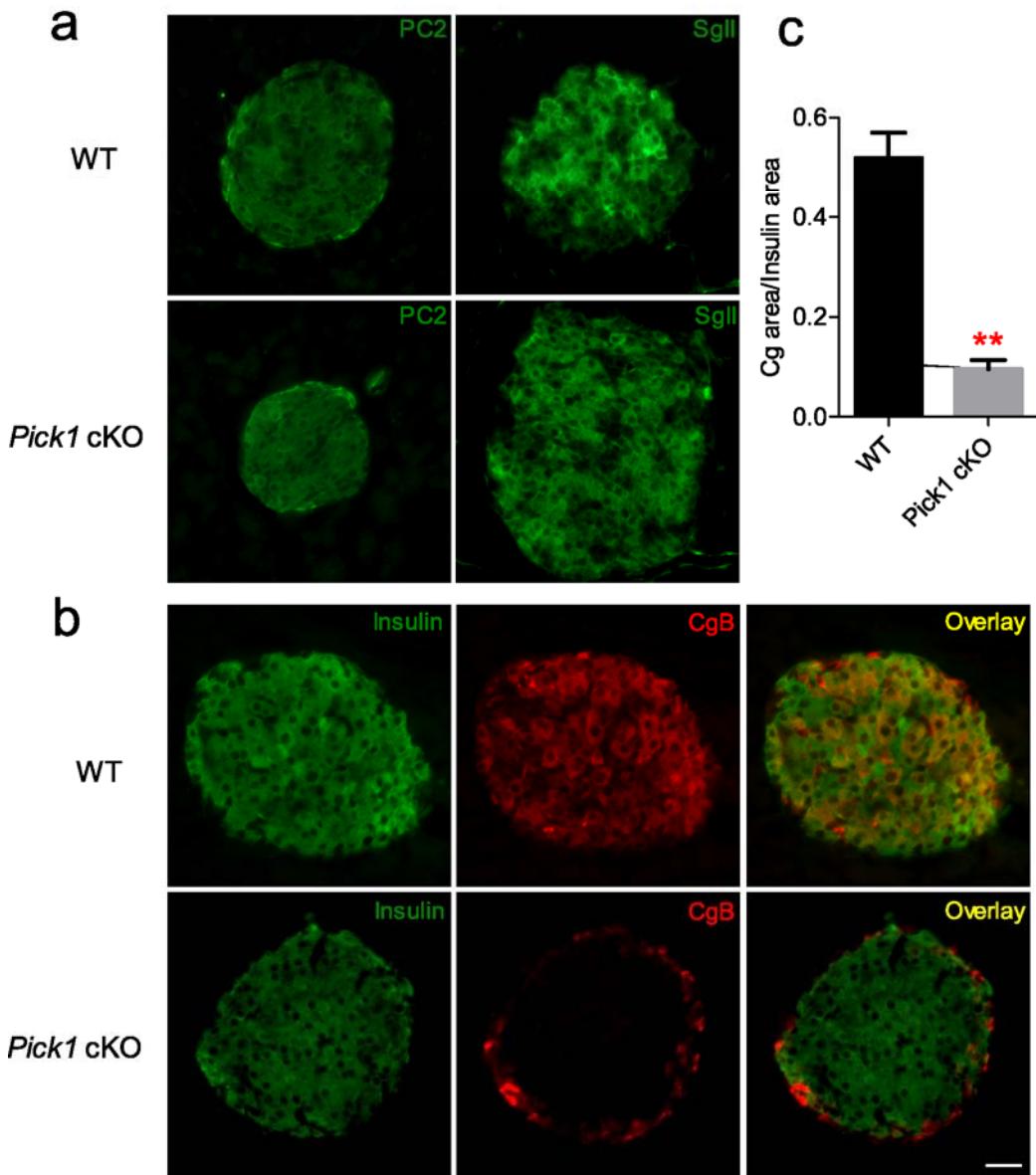


Fig. S2.(a-b) Pancreatic sections from 5-month-old mice of the indicated genotypes were stained with antibodies against PC2 (a, left) or SgII (a, right), insulin (b, green) and CgB (b, red). Scale bar represents 50 μ m. SgII, secretogranin II. (c) Quantification analysis of the CgB/insulin area. 20 islets were calculated per group. Data are represented as the mean \pm SEM. **P<0.01; two sided student's t-test.

Table S1. Q-PCR Primer Sequence

Gene	Forward	Reverse
<i>Pdx1</i>	TTTGAAGTCAGTCAGTTGCTCCTT	CCTTCAACCCCTCTTGTATT
<i>Mafa</i>	AGGAGGTCATCCGACTGAAACAGA	ATTTCCTTGTACAGGTCCGCT
<i>Ins1</i>	TGGCTTCTTCTACACACCCAAGTC	ACTGATCCACAATGCCACGCTTCT
<i>Ins2</i>	GTGGCTTCTTCTACACACCCATGT	GCACTGATCTACAATGCCACGCTT
<i>Pax6</i>	ACAACGACGAAAGAGAGG	ACAAAGACACCACCAAGC
<i>NeuroD1</i>	TCAGCATCAATGGCAACTTC	AAGATTGATCCGTGGCTTG
<i>CHOP</i>	CCACCCACCTGAAAGCAGAA	AGGTGAAAGGCAGGGACTCA
<i>sXbp1</i>	CTGAGTCCGAATCAGGTGCAG	GTCCATGGGAAGATGTTCTGG
<i>Xbp1</i>	TGGCCGGGTCTGCTGAGTCCG	GTCCATGGGAAGATGTTCTGG
<i>ATF4</i>	GGGTTCTGTCTCCACTCCA	AAGCAGCAGAGTCAGGCTTC
<i>Pc1/3</i>	AATGGGCGGCGGAGAT	CCAAAAGGTCATACCCAGTTC
<i>Pc2</i>	GCTGGACCAGCCTTATGA	ATGTCGATCAGTTGAGGCATGT
<i>Cpe</i>	CTGATCCACAGCACCCGAAT	GCGATGCGGCTTCTCA
<i>Actin</i>	CTGAATGGCCCAGGTCTGA	CCCTGGCTGCCTAACAC
<i>Gapdh</i>	GGTGAAGGTGGTGTGAACG	CTCGCTCCTGGAAGATGGTG