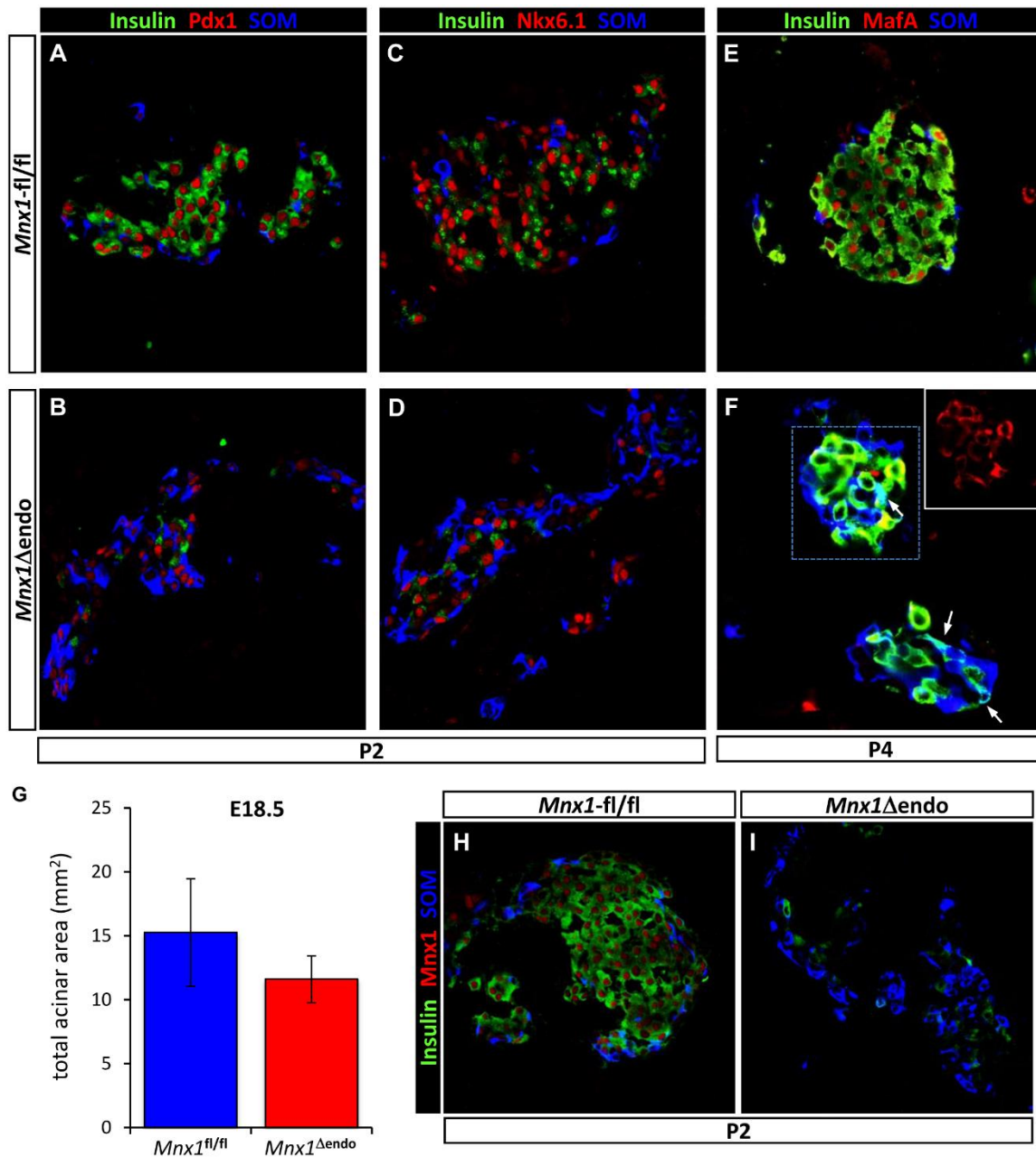
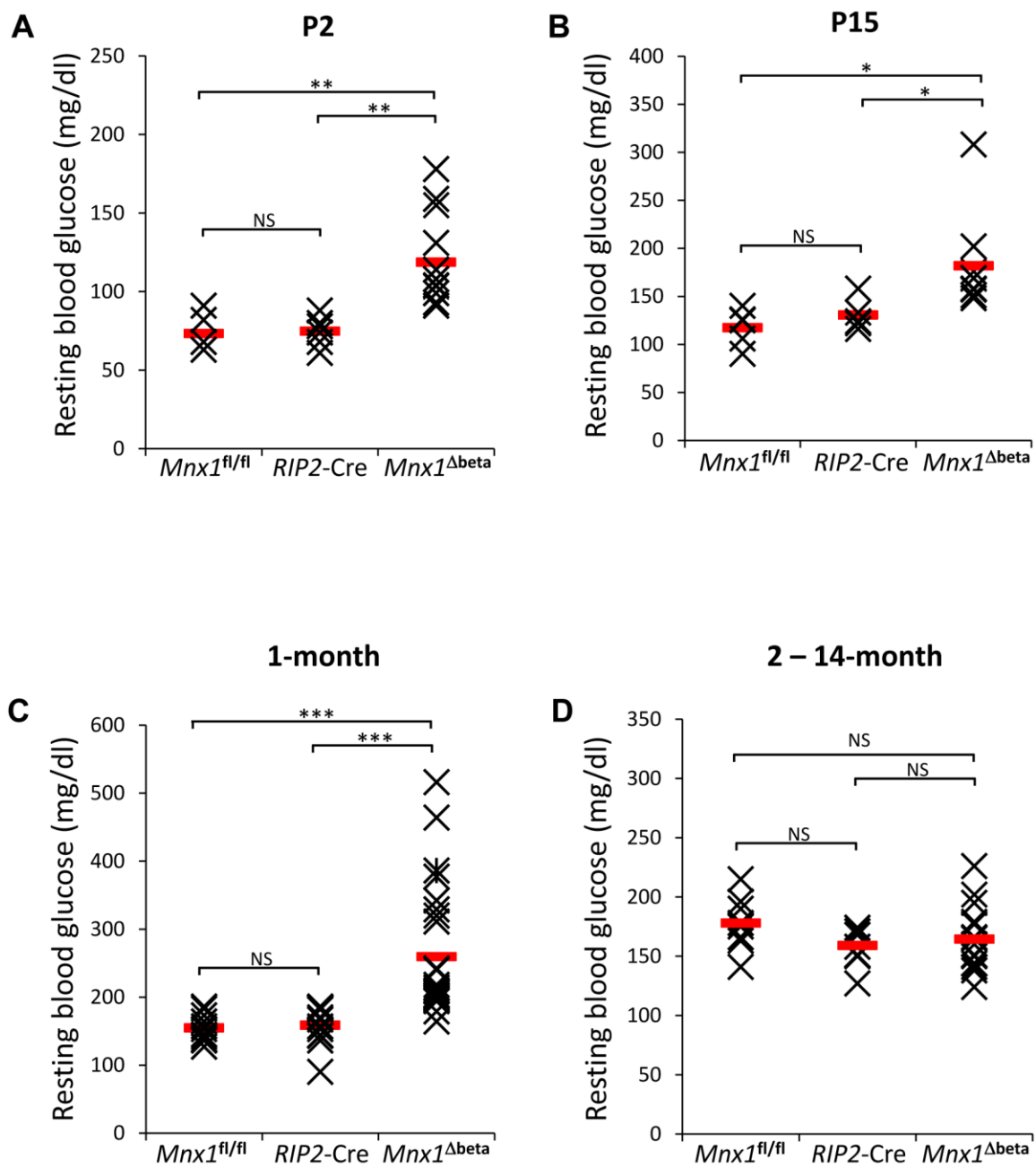


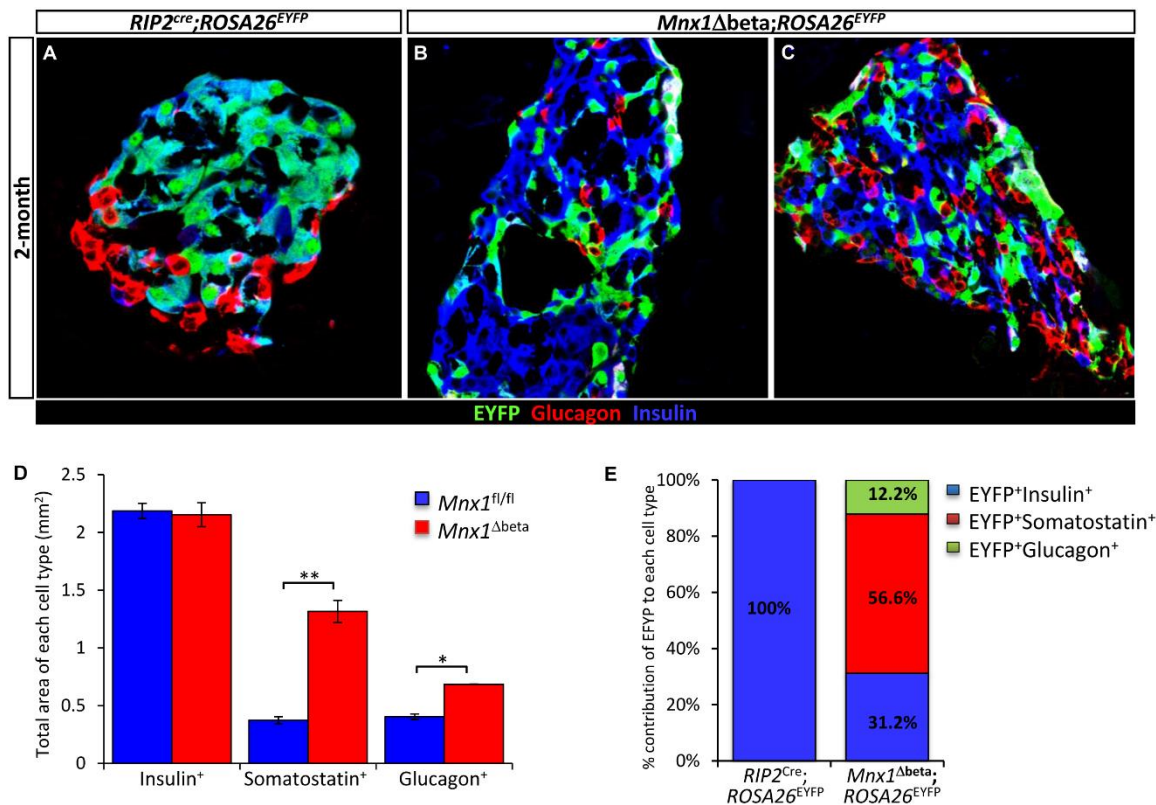
Supplementary Fig. 1 Verification of *Mnx1* floxed allele using germ-line activated Cre line (*EIIA^{Cre}*). (A) Schematic showing generation of *Mnx1* global null mice by crossing *Mnx1^{fl/fl}* mice to the germ line deleter *EIIA^{Cre}*. (B, C) Gross morphology of E18.5 gut show the absence of dorsal pancreas (demarcated in red dotted line), whereas ventral pancreas (delineated by blue dotted line) develop normally in the *Mnx1^{null}* mice. We are not able to test for the presence of the truncated protein with the currently available specific Mnx1 antibody using immunofluorescence, because this antibody was raised against the C-terminal region of Mnx1, which is absent when exon 3 is deleted. Nevertheless, the *EIIA^{Cre};Mnx1^{FL/+}*, *Ngn3-Cre;Mnx1^{FL/+}* and *RIP2-Cre;Mnx1^{FL/+}* mice are normal and do not exhibit any defects in motor neuron function, pancreatic endocrine differentiation and β -cell function, suggesting that the truncated protein is either not made or is degraded, not exhibiting any dominant negative effects.



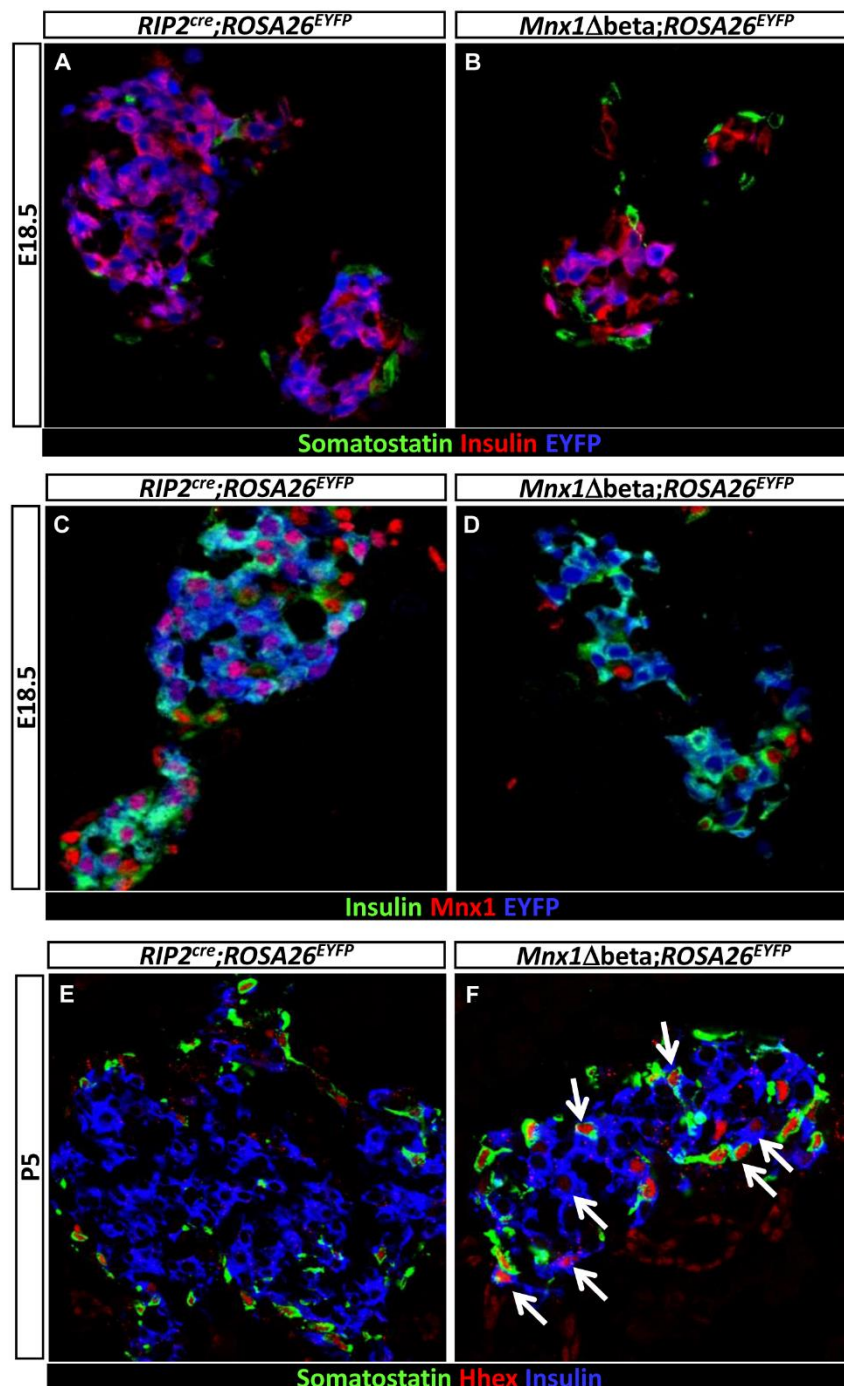
Supplementary Fig. 2 The remaining β cells in *Mnx1*^{Δendo} are not mature. Immunofluorescence analysis show that the remaining β cells in the *Mnx1*^{Δendo} expressed (A, B) Pdx1, and (C, D) Nkx6.1 as in control. (E, F) But MafA protein become localized to the cytoplasm compared to the nucleus localization in control β cell, indicating that these mutant β cells are immature. Insulin⁺somatostatin⁺ cells (arrow) were MafA⁻ indicating the departure of this cell type from β cells. (G) Total acinar area were not changed significantly in *Mnx1*^{Δendo}. (H, I) The remaining β cells in the *Mnx1*^{Δendo} mutants do not express Mnx1.



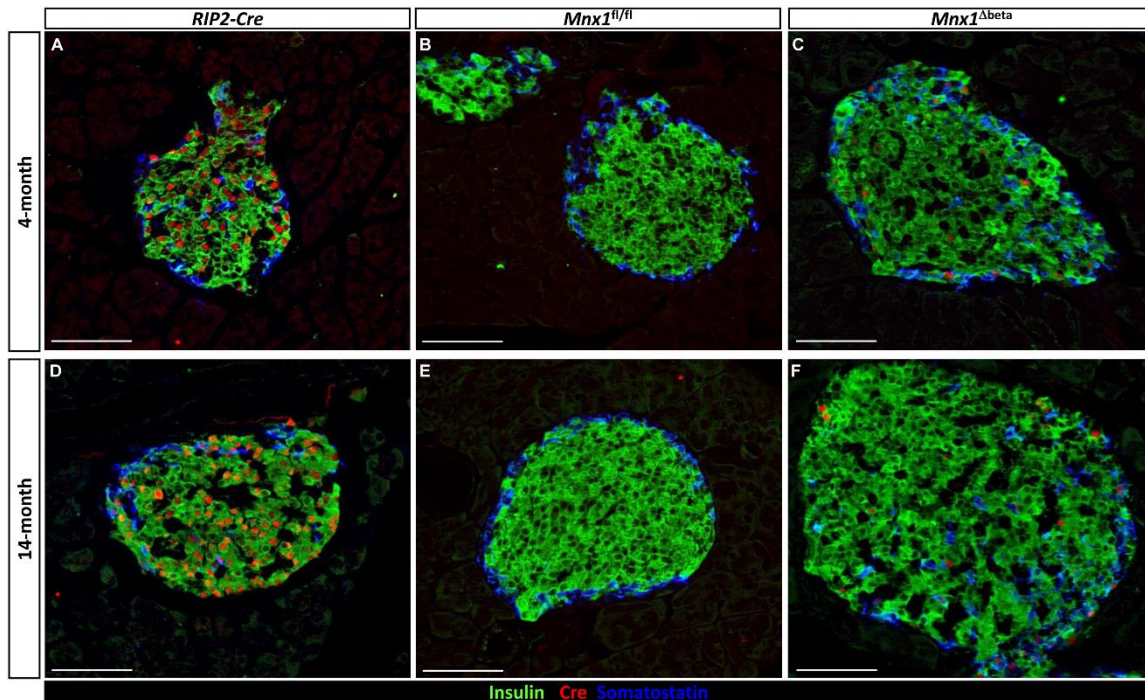
Supplementary Fig. 3 Resting blood glucose of *Mnx1^{Δbeta}* mutants improved with age. Measurement of resting blood glucose of *Mnx1^{fl/fl}*, *RIP2-Cre* and *Mnx1^{Δbeta}* mice at (A) P2, (B) P15, (C) 1-month old, and (D) 2 – 14-month old show a slightly elevated blood glucose at early age but blood glucose level improved with age. * $p < 0.05$, ** $p < 0.005$, *** $p < 0.001$



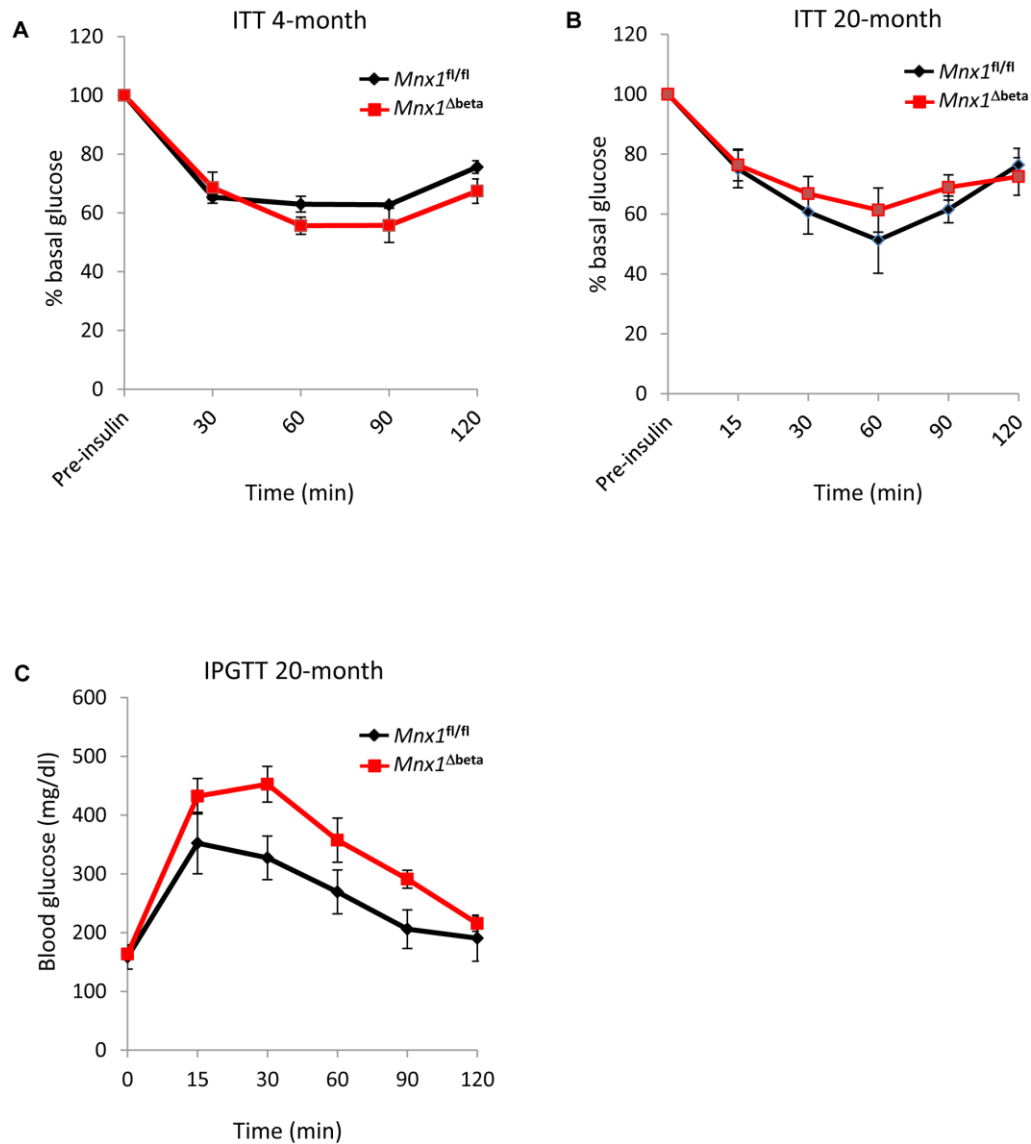
Supplementary Fig. 4 β-to-α transdifferentiation was also observed in *Mnx1^{Δbeta}* mutants, albeit at lower frequency. (A, B, C) The presence of EYFP⁺Glucagon⁺ cells in *Mnx1^{Δbeta}* islets indicate β-to-α transdifferentiation also occurred when *Mnx1* function is deleted in β cells. (D) Quantitative analysis of total area of each hormone⁺ cell types show that β-cell numbers were restored at 4-month old *Mnx1^{Δbeta}*, concomitant with increased of δ and α-cell numbers; (E) Quantitative analysis show that the percentage contribution of EYFP⁺ cell in each of the β, δ, and α cell compartments in *RIP2^{Cre}* and *Mnx1^{Δbeta}* mutants.



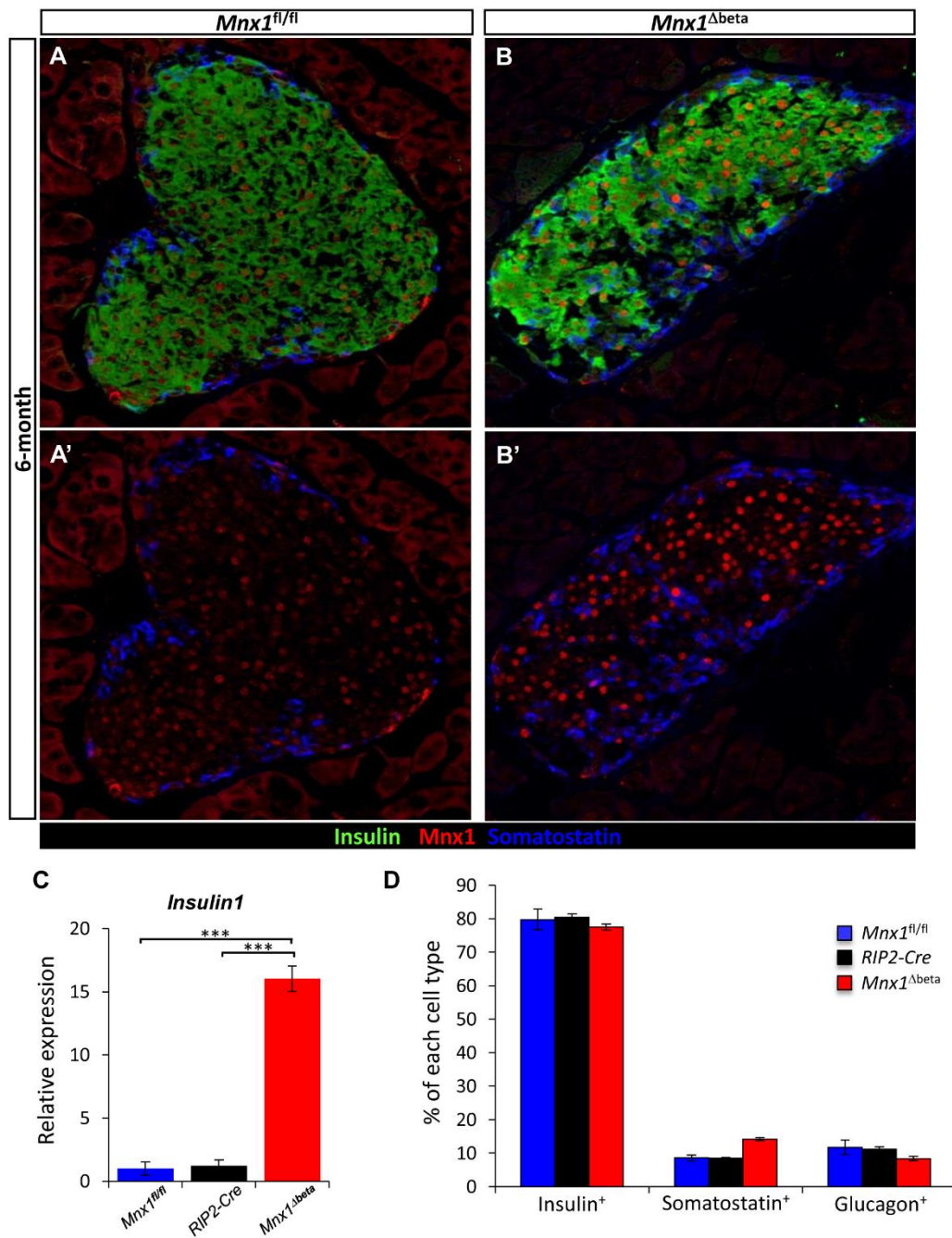
Supplementary Fig. 5 Initiation of β -to- δ cell transdifferentiation in $Mnx1^{\Delta\beta}$ mutants was observed at P5 but not at E18.5. (A, B) Immunofluorescence analysis with EYFP, insulin and somatostatin show the absence of EYFP⁺somatostatin⁺ cells at E18.5 in $Mnx1^{\Delta\beta}$ mice, indicating no β -to- δ transdifferentiation at this stage. (C, D) All EYFP⁺ cells are $Mnx1^{-}$, indicating $Mnx1$ is efficiently deleted in the $Mnx1^{\Delta\beta}$ β cells at E18.5. (E, F) The presence of Hhex⁺insulin⁺ and Hhex⁺insulin⁺somatostatin⁺ (white arrows) cells in the $Mnx1^{\Delta\beta}$ mutants at P5 indicate the initiation of β -to- δ transdifferentiation.



Supplementary Fig. 6 Escaper β cells in *Mnx1^{Δbeta}* mutants were devoided of Cre. Immunolabeling of Cre showed that majority of the escaper β cells that repopulated the islet in *Mnx1^{Δbeta}* do not produce Cre recombinase at (A, B, C) 4 months and, (D, E, F) 14 months, compared to the RIP2-Cre control islets. It is noteworthy that even in RIP2-Cre islets, 15-20% of insulin⁺ β cells do not produce Cre recombinase, consistent with the 85% recombination efficiency in the islet in this allele. Scale bar, 25 μ m.



Supplementary Fig. 7 Glucose clearance defects in *Mnx1^{Δbeta}* mutants were not caused by peripheral insulin resistance. (A) Insulin tolerance test on *Mnx1^{fl/fl}* and *Mnx1^{Δbeta}* indicate that *Mnx1^{Δbeta}* mice do not have peripheral insulin resistance at 6-month old, but (B) developed mild insulin resistance and (C) glucose intolerance at 20-month old. ITT, Insulin tolerance test; IPGTT, intraperitoneal glucose tolerance test.



Supplementary Fig. 8 *Mnx1* protein level and insulin mRNA expression were upregulated in the β cells of *Mnx1*^{Δbeta} mutants. (A, A', B, B') Immunolabeling of *Mnx1* show that *Mnx1* protein was significantly induced in *Mnx1*^{Δbeta} β cells at 6-month old. (C) qRT-PCR data showed that *Insulin1* mRNA expression is highly upregulated in the remaining Cre⁻ β cells at 6-month old. (D) Quantitative analysis of β , δ , and α cell fraction showing the percentage of β cells within islets of *Mnx1*^{Δbeta} is comparable to control *Mnx1*^{fl/fl} and *RIP2-Cre* at 14-month old.

Supplementary Table S1: List of antibodies used

Primary antibodies				
Antigen	Species	Dilution	Staining type	Source
Mnx1	Rabbit	1:5000	IF	Samuel Pfaff (Salk Institute)
Mnx1	Mouse	1:500	TSA	DSHB
Ptf1a	Goat	1:1000	TSA	Chris Wright
Sox9	Rabbit	1:1000	IF	Chemicon
Pax6	Rabbit	1:800	TSA	Covance
GFP	Rabbit	1:500 1:1000	IF TSA	Clontech
Menin	Goat	1:500	IF	Bethyl
Insulin	Guinea Pig	1:1000	IF	Linco
Insulin-A	Goat	1:250	IF	Santa Cruz
Glucagon	Guinea Pig	1:1000	IF	Linco
Glucagon	Rabbit	1:1000	IF	Linco
Somatostatin	Goat	1:1000	IF	Santa Cruz
Pancreatic Polypeptide	Guinea Pig	1:1000	IF	Linco
Cpal	Goat	1:250	IF	BD Bioscience
E-cadherin	Mouse	1:500	IF	BD Bioscience
Nkx6.1	Rabbit	1:1000	IF	BCBC
MafA	Rabbit	1:1000	TSA	Bethyl
GLUT2	Rat	1:200	IF	Alpha Diagnostic
Pdx1	Rabbit	1:1000	IF	Chris Wright (Vanderbilt)
Ki67	Rabbit	1:500	IF	Sigma
Ngn3	Guinea Pig	1:2000	TSA	Maike Sander (UCSD)
Hhex	Rabbit	1:300	IF	Clifford Bogue (Yale University)
Cre	Rabbit	1:500	IF	Novagen

Secondary antibodies			
Antigen	Conjugation	Dilution	Source
Rabbit/Guinea pig/ Goat/Mouse/Chicken	Cy3	1:300	Jackson ImmunoResearch
Rabbit/Guinea pig/ Goat/Mouse/Chicken	Cy2	1:300	Jackson ImmunoResearch
Rabbit/Guinea pig/Goat/Mouse	Cy5	1:300	Jackson ImmunoResearch
Rabbit/Guinea pig/ Goat/Mouse/Chicken	Biotinylated	1:1000	Vector Laboratories

Supplementary Table S2: Primers used in qRT-PCRs

Primer name	Sequence
<i>GAPDH</i>	Forward: AACTTTGGCATTGTGGAAGG
	Reverse: GGATGCAGGGATGATGTTCT
<i>Insulin1</i>	Forward: CAGCAAGCAGGTCATTGTTT
	Reverse: GGGACCACAAAGATGCTGTT
<i>Mnx1</i>	Forward: AAGCGTTTTGAGGTGGCTAC
	Reverse: CCATTTCAATCGGCGGTTCT
<i>Cdkn2a</i>	Forward: GGGATGATGGACTTTTGAGG
	Reverse: TCTGGCTTCTAAGAGAAGATCTAA
<i>Bmi1</i>	Forward: AAACCAGACCACTCCTGAACA
	Reverse: TCTTCTTCTTCTCATCTCATTTTGA
<i>Cdkn1a</i>	Forward: GCTTGGATGTCAGCGGGA
	Reverse: CAGAGTTTGCCTGAGACCCA
<i>CDK4</i>	Forward: CGAGCGTAAGATCCCCTGCT
	Reverse: CGAGCGTAAGATCCCCTGCT
<i>CDK6</i>	Forward: TGCGAGTGCAGACCAAGTGG
	Reverse: AGGTCTCCAGGTGCCTCAGC
<i>CyclinD1</i>	Forward: CCCTCGGTGTCCTACTTCAA
	Reverse: GGGGATGGTCTCCTTCATCT
<i>Menin</i>	Forward: ACCCACTCACCTTTATCACA
	Reverse: ACATTTTCGGTTGCGACAGT
<i>Insulin2</i>	Forward: GGCTTCTTCTACACACCCAT
	Reverse: CCAAGGTCTGAAGGTCACCT