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## **Reporting Summary**

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Statistics	
	es, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a Confirmed	
The exact sam	uple size (n) for each experimental group/condition, given as a discrete number and unit of measurement
A statement of	on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	test(s) used AND whether they are one- or two-sided ests should be described solely by name; describe more complex techniques in the Methods section.
X A description	of all covariates tested
X A description	of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	ion of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	thesis testing, the test statistic (e.g. $F$ , $t$ , $r$ ) with confidence intervals, effect sizes, degrees of freedom and $P$ value noted sexact values whenever suitable.
For Bayesian	analysis, information on the choice of priors and Markov chain Monte Carlo settings
For hierarchic	al and complex designs, identification of the appropriate level for tests and full reporting of outcomes
Estimates of e	effect sizes (e.g. Cohen's $d$ , Pearson's $r$ ), indicating how they were calculated
	Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.
Software and o	code
Policy information abo	ut <u>availability of computer code</u>
Data collection	Imaging: MetaMorph v7, Electrophysiology: HEKA Patchmaster v2
Data analysis	MetaMorph v7, ImageJ v1.52 with custom analysis routines (Gandasi Nat Comm 2014), Origin Pro v2018
	om algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers. deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.
Data	
<ul><li>Accession codes, un</li><li>A list of figures that</li></ul>	ut <u>availability of data</u> include a <u>data availability statement</u> . This statement should provide the following information, where applicable: ique identifiers, or web links for publicly available datasets have associated raw data restrictions on data availability
We have no restrictions of	on sharing the raw data. However, we are not aware of a practical way to make the raw data (600GB) available through public repositories.
Field-speci	fic reporting
Please select the one b	elow that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.
<b>X</b> Life sciences	Behavioural & social sciences Ecological, evolutionary & environmental sciences
For a reference copy of the d	ocument with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf

	nces study design sclose on these points even when the disclosure is negative.
Sample size	No sample size calculation was performed, and sample size was instead based on previous experience with single cells and human islets (Gandasi et al Cell Metab 2018 DOI: 10.1016/j.cmet.2017.12.017).
Data exclusions	No data were excluded.
Replication	At least 2 biological repetitions for each experiment, meaning islets from different donors on different days. All repetitions at repetition were successful and included in the statistical analysis.
Randomization	Not relevant. When comparing eg drug effects, islet samples were divided onto several coverslips that were treated differently only during data acquisition.
Blinding	Blinding was not formally used, but data files from individual donors were processed together and without knowledge of the treatment group (except for diabetic status, which was known on arrival of islet samples).
Reportin	g for specific materials, systems and methods
	ion from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, sted is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.
Materials & ex	perimental systems Methods

## **Antibodies**

x

Antibodies used

Involved in the study

**x** Eukaryotic cell lines

Clinical data

Palaeontology

Animals and other organisms

Human research participants

**x** Antibodies

Mouse anti-Glucagon (K79bB10) Sigma G2654, dilution 1/1500 Rabbit anti-Glugacon DAKO (Uppsala) A0565, dilution 1/200 Guinea Pig anti-Insulin DAKO (Uppsala) A0564, dilution 1/200 Rabbit anti-Somatostatin DAKO (Uppsala) A0566, dilution 1/200 Rabbit anti-SSTR2 (UMB1) abcam ab134152, dilution 1/100

n/a | Involved in the study

Flow cytometry

MRI-based neuroimaging

ChIP-seq

X

X

Goat anti-Rabbit Alexa-546 Invitrogen A11035, dilution 1/500 Goat anti-Mouse Alexa-555 Invitrogen A21424, dilution 1/500 Goat anti-Guinea Pig Alexa-555 Invitrogen A21435, dilution 1/500 Goat anti-Rabbit Alexa-488 Invitrogen A11034, dilution 1/500

Validation

All primary antibodies are tested and characterized as specific in human tissues, by the manufacturers, and are widely cited.

 $Mouse\ anti-Glucagon\ (Sigma\ G2654):\ IHC\ in\ fixed\ sections\ of\ pancreas\ from\ human,\ www.sigmaaldrich.com/catalog/product/$ sigma/g2654, antibodyregistry.org/AB\_259852

anti-glucagon (DAKO A0565): antibodyregistry.org/AB\_10013726, IHC: www.proteinatlas.org/ENSG00000115263-GCG/antibody

anti-insulin (DAKO A0564): antibodyregistry.org/AB\_10013624, www.citeab.com/antibodies/3382917-a0564-insulin

anti-somatostatin (DAKO A0566): antibodyregistry.org/AB\_2688022, IHC: www.proteinatlas.org/ENSG00000157005-SST/

anti SSTR2 (abcam ab134152): manufacturer information: reacts with mouse, rat, human, tested in human pancreas and SSTR2 ko mice. Fischer et al Clin Endocrinol Metab 93:4519-24 (2008), antibodyregistry.org/ab134152

### Human research participants

Policy information about studies involving human research participants

Population characteristics

Is lets were from 68 non-diabetic donors (42 male, 26 female, age 59.0+/-12.9, BMI 26.5+/-4.3, HbA1c 5.5+/-0.3; mean+/-SD) and 21 type-2 diabetic donors (13 male, 8 female, age 60.4+/-8.7, BMI 29.7+/-6.3, HbA1c 6.6+/-0.7; mean+/-SD).

Sections were from 5 ND (2 male, 3 female, age 63.8+/-6.2, BMI 25.4+/-4.9, HbA1c 6.3+/-2.1) and 5 type-2 diabetic donors (1 male, 4 female, age 62.4+/-5.9, BMI 33.9+/-2.2, HbA1c 7.6+/-0.4; mean+/-SD).

Recruitment

Cadaveric donor pancreata deemed unsuitable for transplantation were obtained and processed by the islet isolation cores. No selection criteria were applied.

Ethics oversight

Human pancreatic islets and pancreas sections were obtained from the Nordic Network for Clinical Islet Transplantation Uppsala (ethical approved by Uppsala Regional Ethics Board) or the ADI Isletcore at the University of Alberta (ethical approval by Alberta Human Research Ethics Board, Pro00001754) and with the donor families' written informed consent. The study was approved by the Uppsala Regional Ethics Board (2006/348).

Note that full information on the approval of the study protocol must also be provided in the manuscript.