

Supplementary Figure 11. The characterization data for reserpine .

Reserpine had high purity (>90%).

Reserpine was purchased as a dry powder from Calbiochem Novabiochem Novagen.

HPLC, LC-MS (ESI) and ¹H-NMR profiles are shown.

Supplementary Figure 12. The characterization data for Tetrabenazine.

Tetrabenazine had high purity (>90%).

Tetrabenazine was purchased as a dry powder from Tocris Bioscience.

HPLC, LC-MS (ESI) and ¹H-NMR profiles are shown.

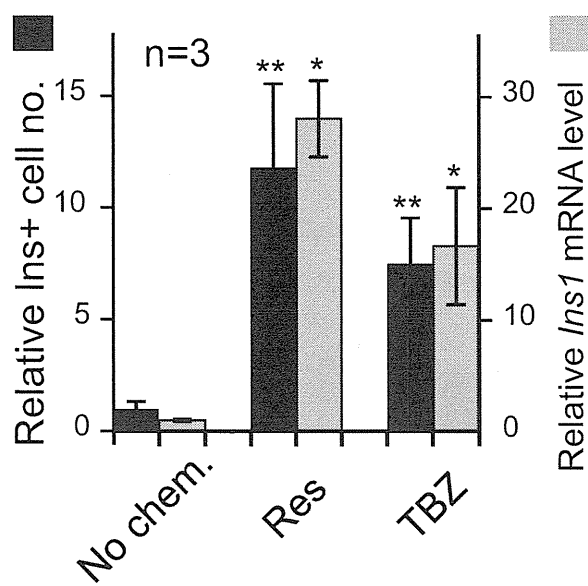
Supplementary Figure 13. The characterization data for dBu-cAMP.

dBu-cAMP had high purity (>90%).

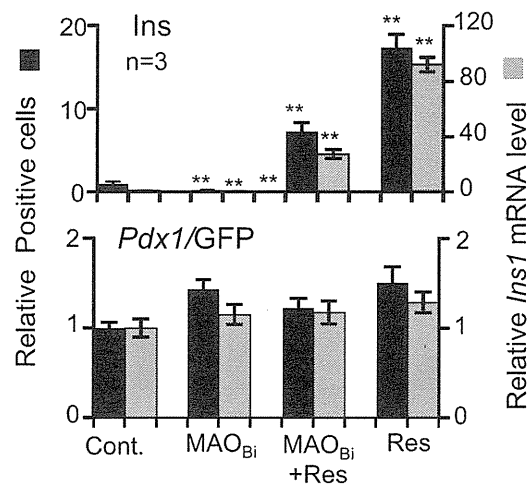
dBu-cAMP was purchased as a dry powder from BIOMOL International. HPLC,

LC-MS (ESI) and ¹H-NMR profiles are shown.

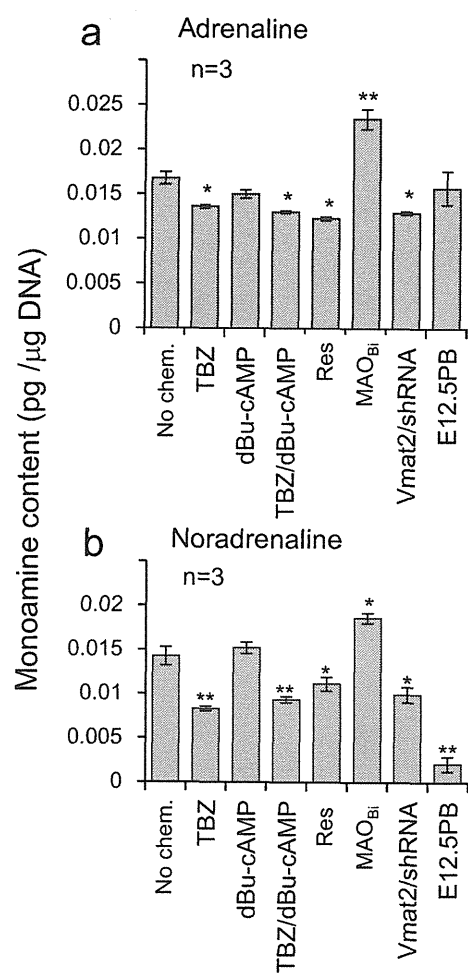
Supplementary Figure 1



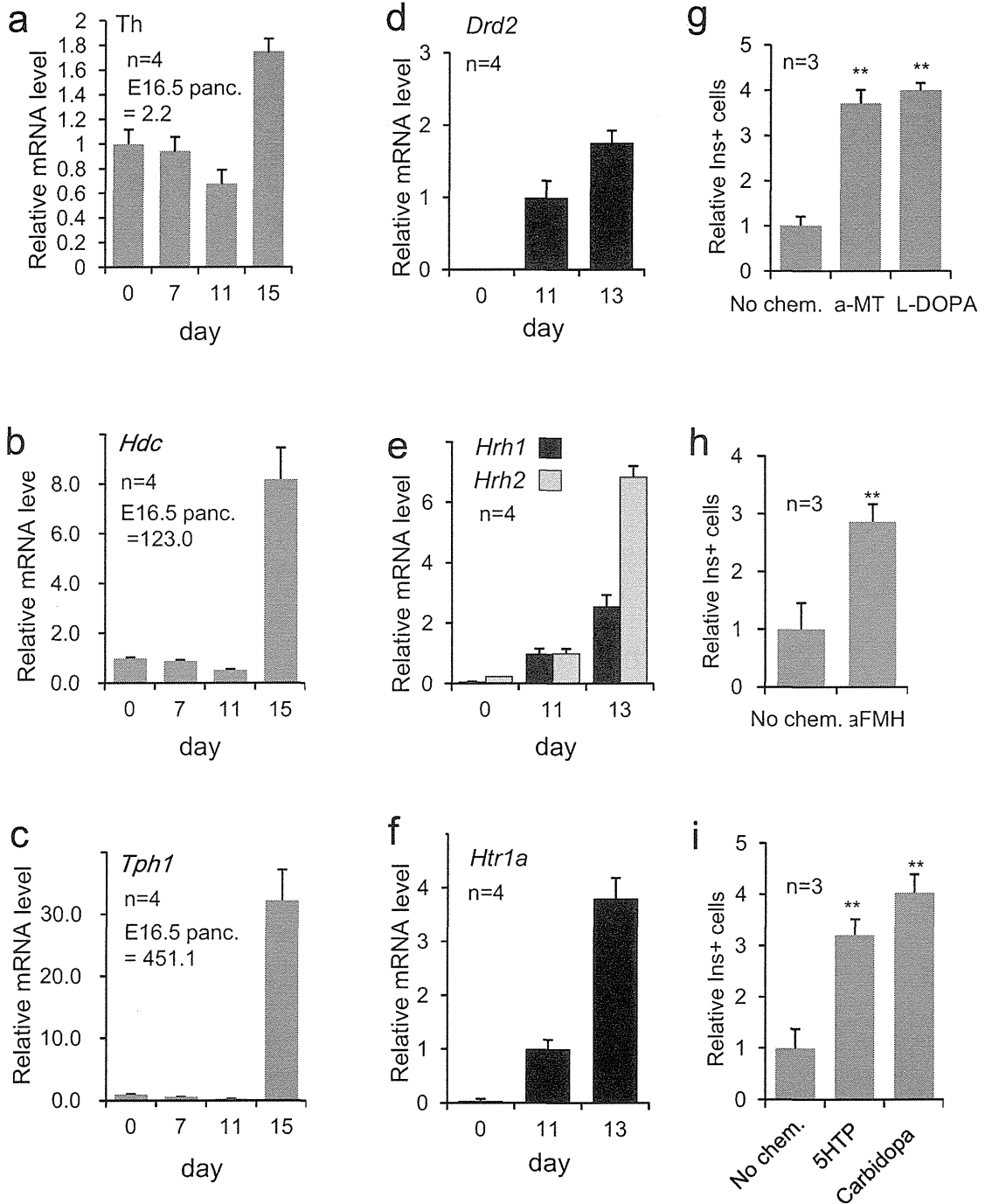
Supplementary Figure 2



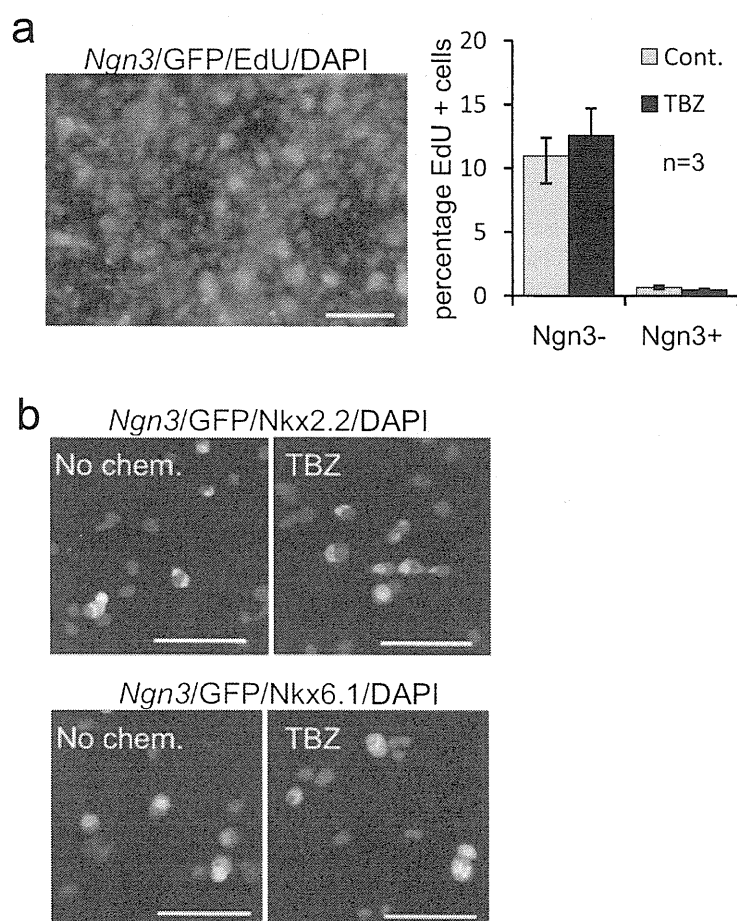
Supplementary Figure 3



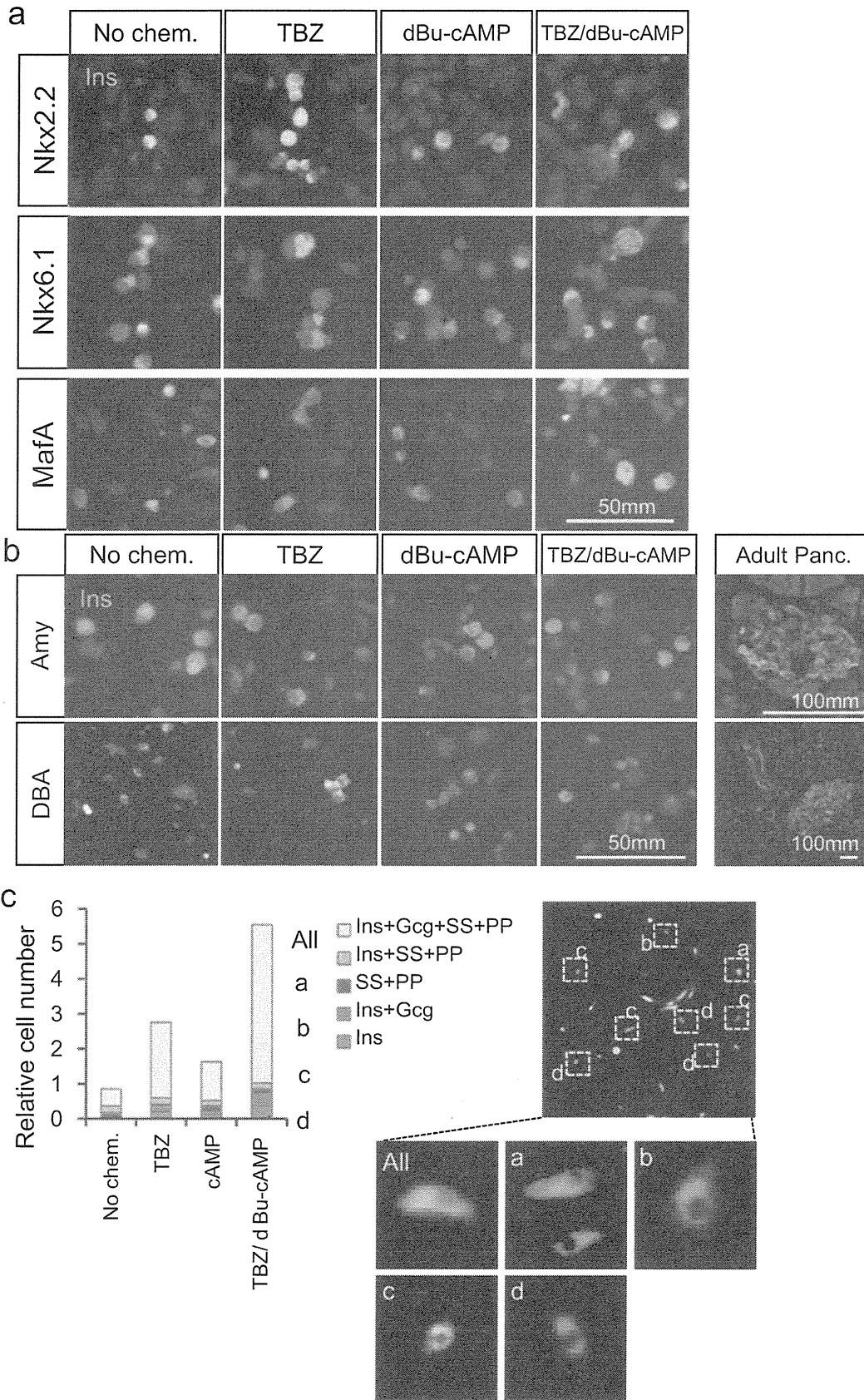
Supplementary Figure 4



Supplementary Figure 5



Supplementary Figure 6

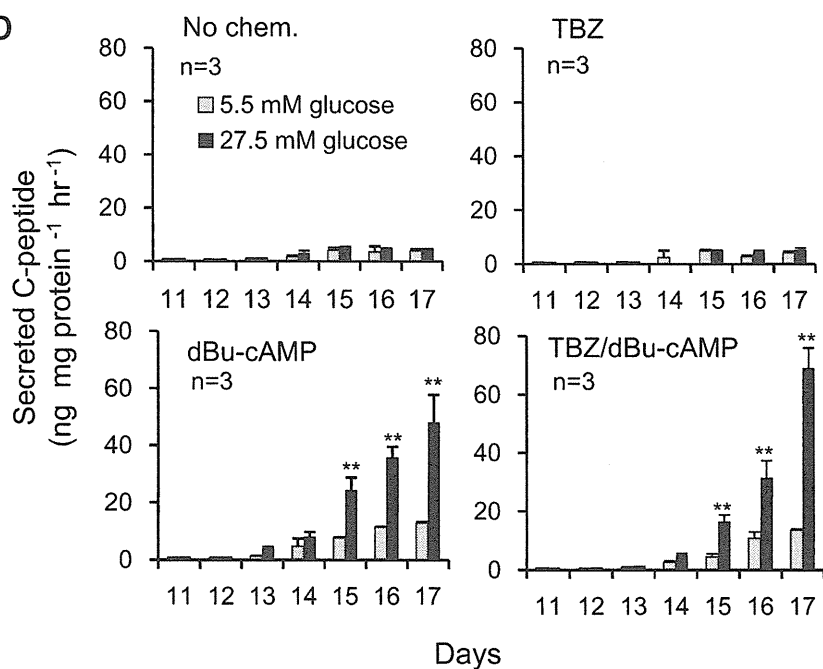


Supplementary Figure 7

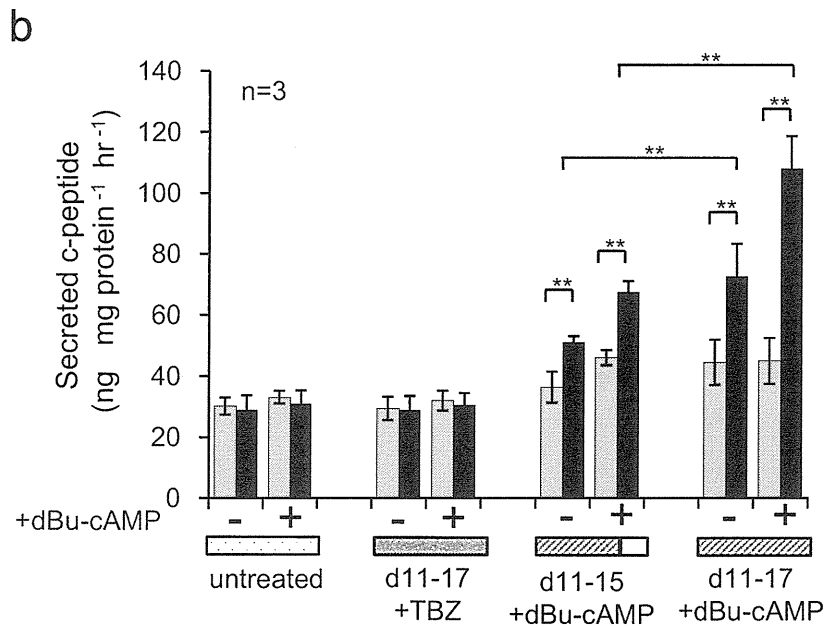
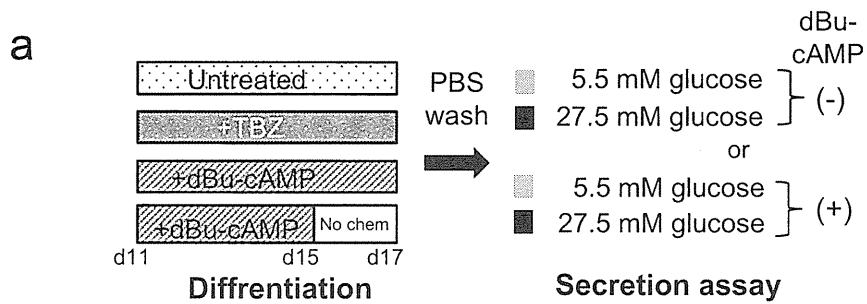
a

	C-peptide (ng/ well, $\sim 1 \times 10^7$ ES cells)	C-peptide (ng/ μ g DNA)	Folds
No chem.	326 \pm 76	8.5 \pm 2.0	1
TBZ	1432 \pm 152	48.8 \pm 5.2	5.7
dBu-cAMP	604 \pm 11	22.9 \pm 0.4	2.7
TBZ/dBu-cAMP	1670 \pm 119	68.9 \pm 4.9	8.1
10 islets	16707 \pm 829 ng	118.7 \pm 5.9 ng	

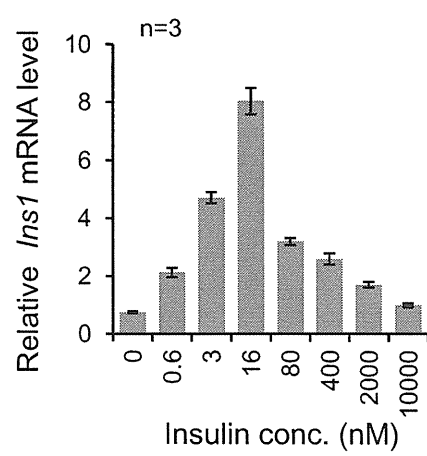
b



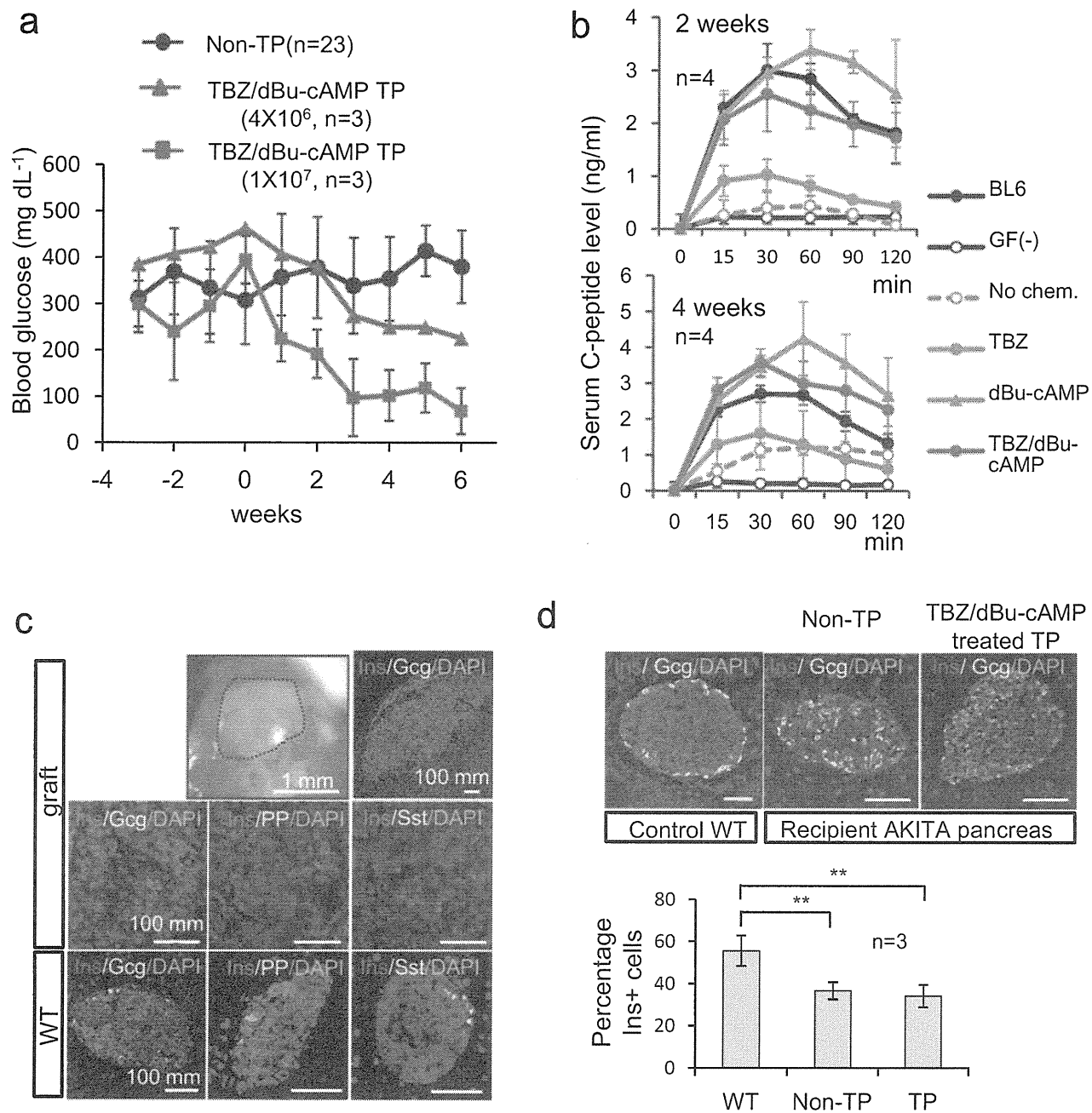
Supplementary Figure 8



Supplementary Figure 9

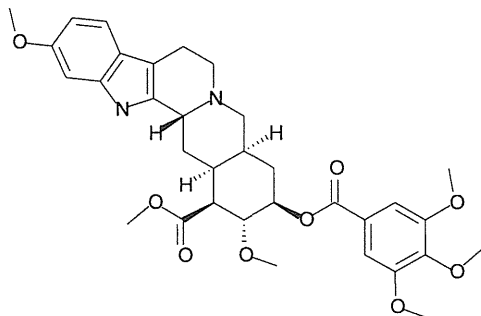


Supplementary Figure 10

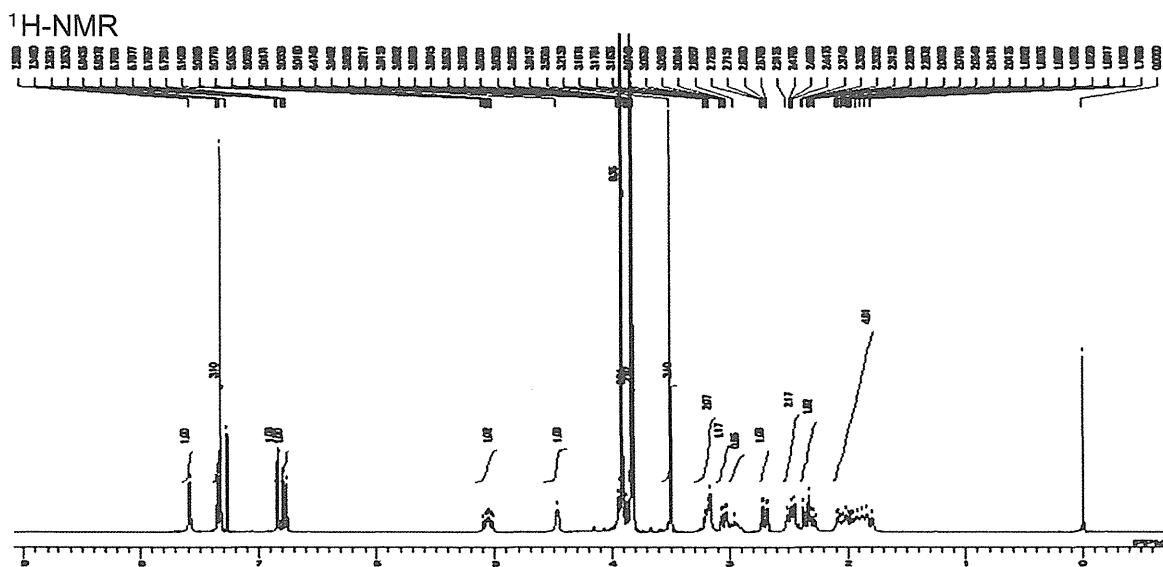
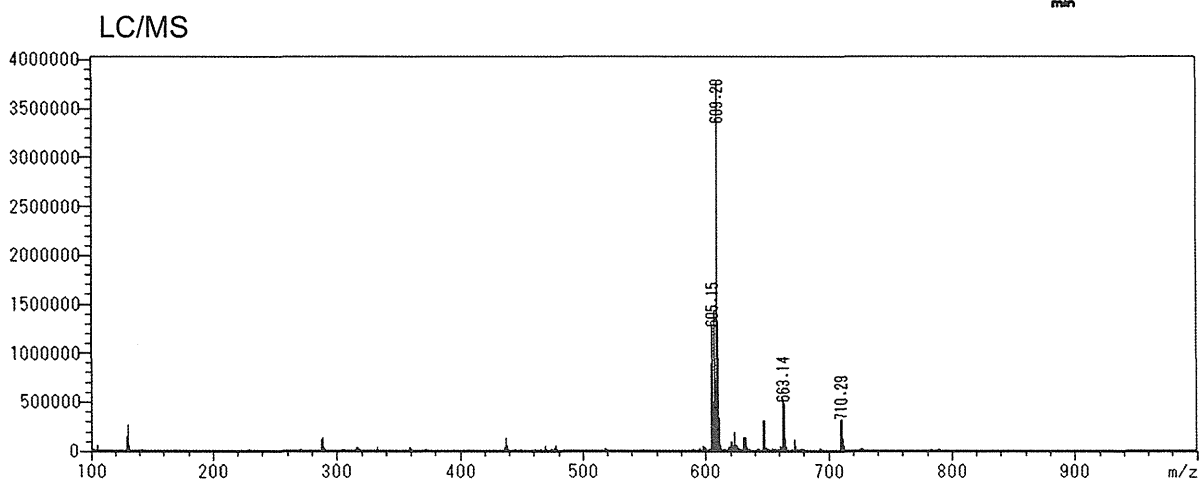
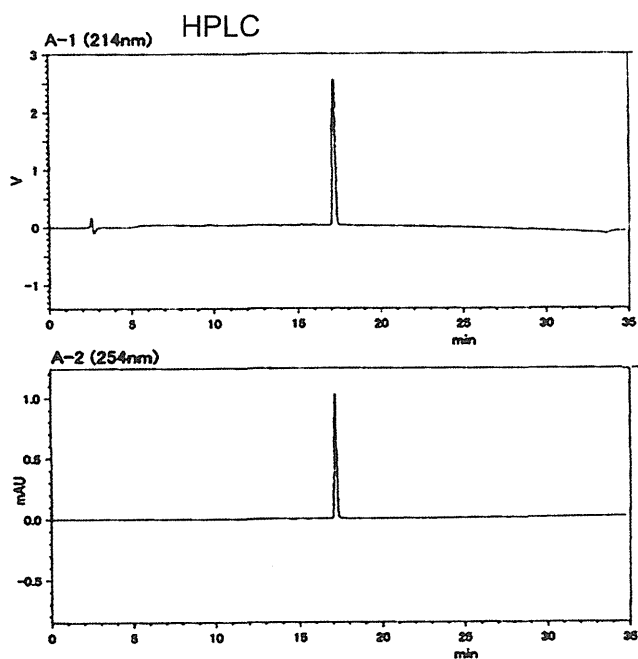


Supplementary Figure 11

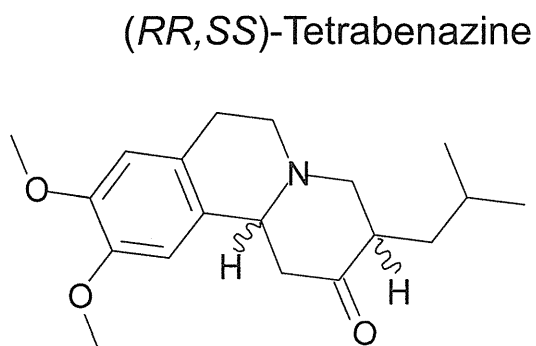
Reserpine



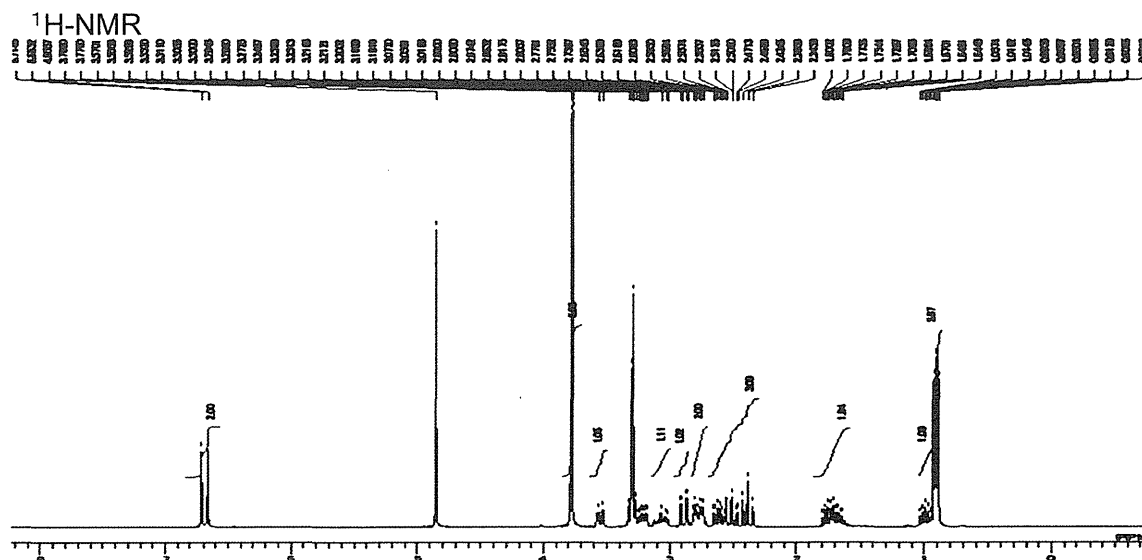
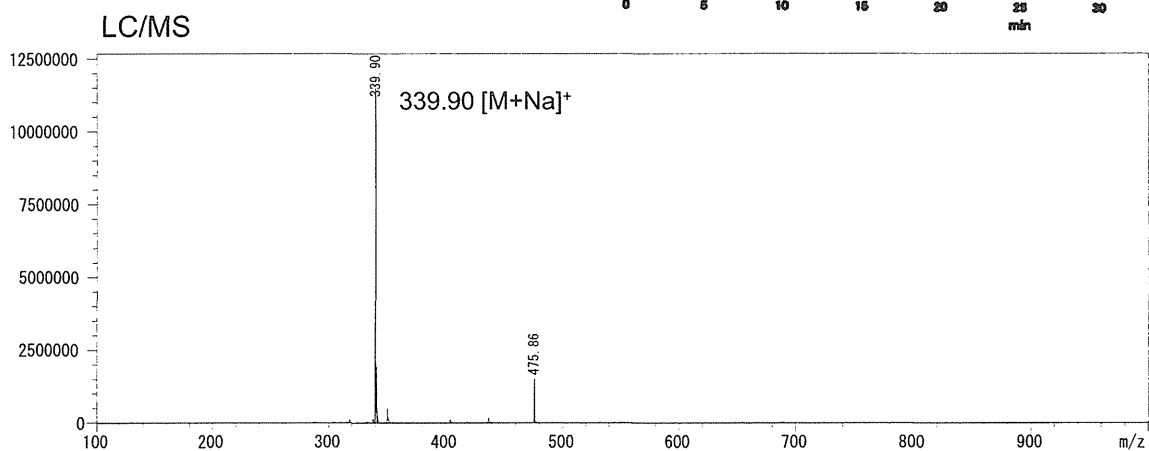
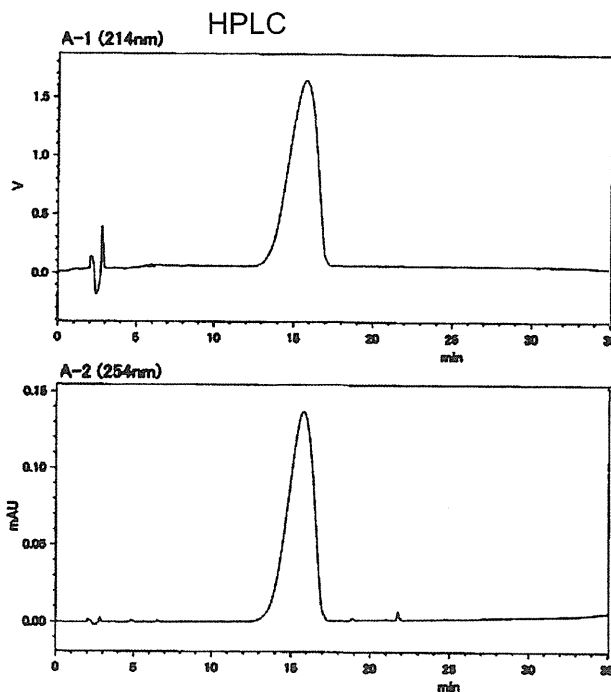
Chemical Formula: $C_{33}H_{40}N_2O_9$
Molecular Weight; 608.68



Supplementary Figure 12

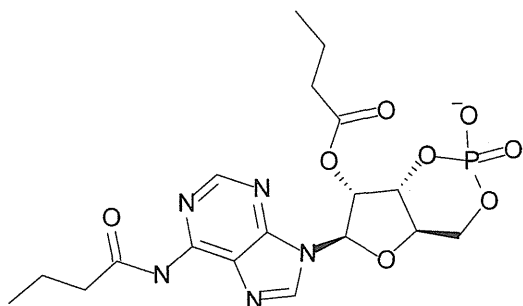


Chemical Formula: $C_{19}H_{27}NO_3$
 Molecular Weight; 317.42

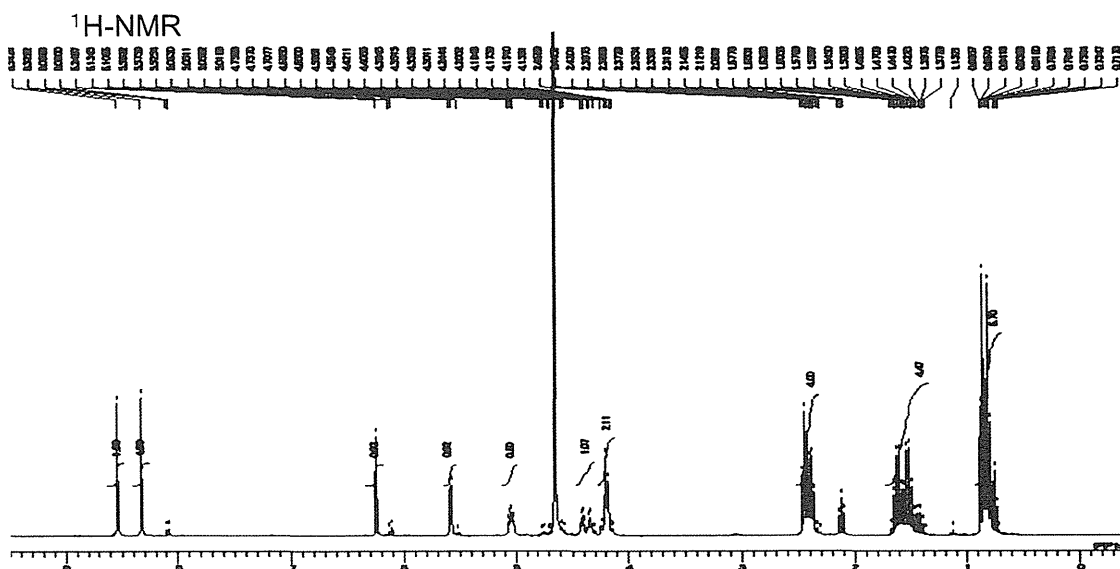
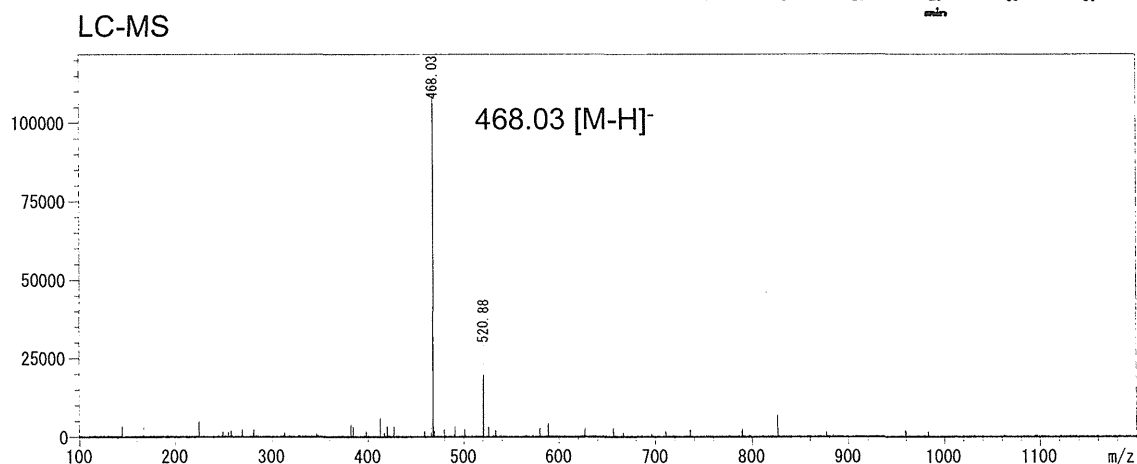
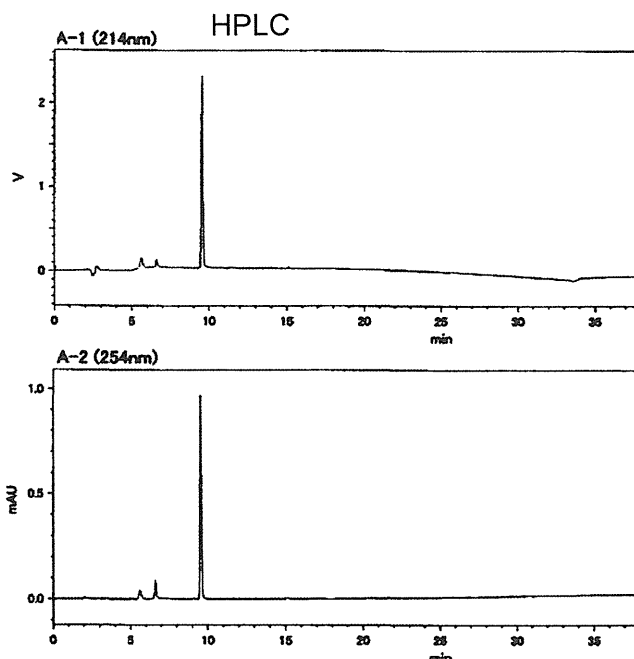


Supplementary Figure 13

N⁶,2'-O-Dibutyryladenine
3',5'-cyclic monophosphate



Chemical Formula: C₁₆H₂₄N₅O₈P
Molecular Weight; 469.39



Supplementary Table 1. Small molecule screening data

Category	Parameter	Description
Assay	Type of assay	cell-based (differentiating mouse ES cells)
	Target	Mouse Insulin 1 protein expression level
	Primary measurement	Counting the number of Insulin+ cells detected by immunocytochemistry analysis.
	Key reagents	Monoclonal anti-insulin Antibody K36AC10 (SIGMA)-
	Assay protocol	Described in this manuscript, partially referring to previous report report (doi: 10.1242/jcs.066886)
	Additional comments	
Library	Library size	1120
	Library composition	biologically active known compounds arrayed as single compounds in dimethyl sulfoxide (DMSO)
	Source	Prestwick Chemical Library
	Additional comments	
Screen	Format	96 well plates
	Concentration(s) tested	2 μ M compounds and 1% DMSO
	Plate controls	10% pdx1/GFP positive cells should be appeared on d11 before chemical screening.
	Reagent/ compound dispensing system	manual
	Detection instrument and software	ImageXpress Micro scanning system and MetaXpress cellular image analysis software (Molecular Devices, Japan)
	Assay validation/QC	CV value = 0.36 \pm 0.0447
	Additional comments	Pdx1/GFP fluorescence detection as a criteria for continuing the assay
Post-HTS analysis	Hit criteria	>2-fold increase in the proportion of Insulin+ cells
	Hit rate	0.1% (10/1120)
	Confirmation of hit purity and structure	HPLC, LC-MS (ESI), and 1H-NMR profiles
	Additional comments	In 2 nd screen, dose-dependencies were examined

Supplementary Table 2 PCR primers used to detect mouse gene expressions

Gene	Forward primer	Reverse primer
<i>Abcc8</i>	CCTGCAGCCAGACATAGACA	GCTCTGGTCTTGCCTGTTC
<i>β-actin</i>	GTGATGGTGGGAATGGGTCA	TTTGATGTCACGCACGATTTC
<i>Drd2</i>	GAGAAGGCTTTGCAGACCAC	AGGACAGGACCCAGACAATG
<i>Gk</i>	TAT GAA GAC CGC CAA TGT GA	TTT CCG CCA ATG ATC TTT TC
<i>Glut2</i>	ACAGAGCTACAATGCAACGTGG	CAACCAGAATGCCAATGACGAT
<i>Hdc</i>	GATCAGATTCTACCTGTGG	GTGTACCATCATCCACTTGG
<i>Hrh1</i>	AAGTGGTTTCTCCACGGTTG	CTTTTTGGAGAAGGCGAGTG
<i>Hrh2</i>	AAGCTTCAGGTGTGGAGTGG	ATTGCCACTGCCAGAGATTC
<i>Htr1a</i>	TTGGAACTACTTTGGGTTATGG	ATTGTCAATTTCTTTGGTGAGTG
<i>Ins1</i>	CAGCCCTTAGTGACCAGCTA	ATGCTGGTGCAGCACTGATC
<i>MafA</i>	TTTCCTCGGCAGCGTCCACTTGTA	GGGGGTTCCTCCGGGTTTTCTAAT
<i>Ngn3</i>	CTGCGCATAGCGGACCACAGCTTC	CTTCACAAGAAGTCTGAGAACACCAG
<i>Nkx6.1</i>	TACTTGGCAGGACCAGAGAG	CGCTGGATTTTGTGCTTTTTTC
<i>Pdx1</i>	CCAAAACCGTCGCATGAAGTG	CTCTCGTGCCCTCAAGAATTTTC
<i>Snap25</i>	GAGAACCTGGAGCAGGTGAG	GGGGGTGACTGACTCTGTGT
<i>Sox17</i>	GAACAGTTGAGGGGCTACAC	GTTTAGGGTTTCTTAGATGC
<i>Th</i>	GGTATACGCCACGCTGAAGG	TAGCCACAGTACCGTTCCAGA
<i>Tph1</i>	CACAGTTCAGATCCCCTCTACA	GAACGTGGCCTAGGAGTTCA
<i>Vmat2</i>	TGTGAAGTCTGGTGTGTTAAGG	CATCACATCACAAAGGCATCC

