

1 **Paired box 6 programs essential exocytotic genes in the regulation of glucose-**  
2 **stimulated insulin secretion and glucose homeostasis**

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16 **Running title:** PAX6 replenishment improves glucose-stimulated insulin secretion and glucose  
17 homeostasis

18

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20 box 6, type 2 diabetes.

## 1 **Abstract**

2 The Paired box 6 (PAX6) transcription factor is crucial for normal pancreatic islet development  
3 and function. Heterozygous mutations of *PAX6* are associated with impaired insulin secretion  
4 and early-onset diabetes mellitus in human. Yet the molecular mechanism of PAX6 in  
5 controlling insulin secretion in human beta-cells and its pathophysiological role in Type 2  
6 diabetes (T2D) remain ambiguous. We investigated the molecular pathway of PAX6 in the  
7 regulation of insulin secretion and the potential therapeutic value of PAX6 in T2D by using  
8 human pancreatic beta-cell line EndoC- $\beta$ H1, the *db/db* mouse model and primary human islets.  
9 Through loss- and gain-of-function approaches, we uncovered a novel mechanism by which  
10 PAX6 modulates glucose-stimulated insulin secretion (GSIS) through a cAMP response element-  
11 binding protein (CREB)/Munc18-1/2 pathway. Moreover, under diabetic conditions, beta-cells  
12 and pancreatic islets displayed dampened PAX6/CREB/Munc18-1/2 pathway activity and  
13 impaired GSIS, which were reversed by PAX6 replenishment. Notably, adeno-associated virus-  
14 mediated PAX6 overexpression in *db/db* mouse beta-cells led to a sustained alleviation of  
15 glycemic perturbation *in vivo*. Our study highlights the pathophysiological role of PAX6 in T2D-  
16 associated beta-cell dysfunction in human, and suggests clinical translation of PAX6 gene  
17 transfer in preserving and restoring beta-cell function and T2D management.

## 1 **Abbreviations**

2	AAV	Adeno-associated virus
3	ATF1	Activating transcription factor 1
4	AUC	Area under curve
5	CaMK	Ca <sup>2+</sup> /calmodulin-dependent protein kinase
6	CRE	cAMP response element
7	CREB	cAMP response element-binding protein
8	CREM	cAMP response element modulator
9	CRTC2	CREB-regulated transcription coactivator 2
10	FSK	Forskolin
11	GIP	Glucose-dependent insulinotropic polypeptide
12	GLP-1	Glucagon-like peptide-1
13	GSIS	Glucose-stimulated insulin secretion
14	HGPA	High glucose and palmitic acid
15	IBMX	Isobutylmethylxanthine
16	IPGTT	Intraperitoneal glucose tolerance test
17	IRS-2	Insulin receptor substrate-2
18	ITT	Insulin tolerance test
19	KSIS	Potassium-stimulated insulin secretion
20	PAX6	Paired box 6
21	PKA	Protein kinase A
22	shRNA	Short hairpin RNA
23	SM proteins	Sec1/Munc18 (SM) proteins
24	SNAP25	Synaptosomal-associated membrane protein 25 kDa
25	SNARE	Soluble <i>N</i> -ethylmaleimide-sensitive factor attachment protein receptor
26	STX1	Syntaxin-1
27	SYB2	Synaptobrevin 2
28	T2D	Type 2 diabetes

## 1 **Introduction**

2 Type 2 diabetes (T2D), the predominant form of diabetes, remains one of the leading causes of  
3 mortality and morbidity worldwide. T2D develops when insulin secretion is defective to  
4 compensate for the increased insulin demand elicited by insulin resistance. Physiologic and  
5 genetic data suggest that pancreatic beta-cell failure is the central event leading to the  
6 progression of insulin resistance to hyperglycemia and overt diabetes (1, 2). Thus, great efforts  
7 have been dedicated to the protection of beta-cells from dysfunction and death in T2D research  
8 as the pathological pathways driving beta-cell failure remain poorly characterized.

9 Preservation of beta-cell function and mass can be achieved by maintaining optimal expression  
10 and activity of key transcription factors that control the expression of genes essential for normal  
11 beta-cell differentiation, function and survival. Defining the roles and signaling network of these  
12 transcription factors in beta-cells provides significant insights to decipher T2D pathogenesis and  
13 more importantly to develop effective approaches for T2D management. In this regard, paired  
14 box 6 (PAX6) is a transcription factor pivotal for the normal development of pancreas. PAX6  
15 deletion greatly reduces alpha- and beta-cell mass with dramatic increase in ghrelin expression, a  
16 gut hormone transiently expressed in fetal pancreas (3-5). Previous studies revealed that PAX6 is  
17 an activator of several beta-cell genes involved in insulin synthesis, glucose sensing, insulin  
18 secretion and thereby maintaining mature beta-cell function in rodents (6, 7). Correspondingly,  
19 PAX6 deficiency in adult murine islet cells causes progressive hyperglycemia and glucose  
20 intolerance with reduced insulin production and secretion (5, 8, 9). In human, clinical studies  
21 revealed that humans heterozygous for *PAX6* show glucose intolerance and early-onset diabetes  
22 associated with impaired insulin secretion (10-12). These observations strongly suggest the  
23 essential regulatory role of PAX6 in insulin secretion not only in rodents but also in human.

1 However, evidence regarding the role of PAX6 in islet function and the pathophysiology of  
2 human T2D remains fragmentary.

3 cAMP response element (CRE)-binding protein (CREB) has emerged as another key  
4 transcription factor modulating beta-cell proliferation and survival. Glucose is a natural activator  
5 of CREB. Upon glucose stimulation, glycolysis and mitochondrial oxidative metabolism lead to  
6 increased ATP/ADP ratio and closure of ATP-sensitive K<sup>+</sup> channels, followed by membrane  
7 depolarization and Ca<sup>2+</sup> influx. Ca<sup>2+</sup> sensors then trigger exocytosis of insulin granules (13). The  
8 ATP generated would be converted into cAMP by adenylyl cyclase; elevated cAMP levels  
9 enhance glucose-stimulated insulin secretion (GSIS) through protein kinase A (PKA)-dependent  
10 and -independent mechanisms (14). On the other hand, catalytic subunits of activated PKA  
11 would translocate to the nucleus and phosphorylate CREB at serine 133 (Ser133) (15). Ca<sup>2+</sup>  
12 influx also promotes CREB phosphorylation by activating Ca<sup>2+</sup>/calmodulin-dependent protein  
13 kinase (CaMK) IV. Phosphorylation of Ser133 is a prerequisite for CREB-mediated transcription  
14 (16). Activated CREB binds to a conserved CRE that was first documented as an eight-base-pair  
15 palindrome, TGACGTCA with a higher conservation at the 5' end (TGACG). The CRE also  
16 occurs as a half-site motif (TGACG/CGTCA) (17, 18). CREB-dependent gene regulation is  
17 essential to govern beta-cell proliferation and survival (19, 20). Disruption of CREB activity in  
18 beta-cells induces apoptosis, glucose intolerance and ultimately overt diabetes (21, 22). Whilst  
19 most of the previous studies illustrated the role of CREB signaling in beta-cell survival, however,  
20 being a transcription factor activated by a variety of insulin-secreting stimuli including glucose,  
21 incretin hormones such as glucagon-like peptide-1 (GLP-1) and glucose-dependent  
22 insulinotropic polypeptide (GIP) (23), the role of CREB in insulin secretion remains obscure.

1 Exocytosis plays a critical role in cellular processes as it is the final but common step of  
2 neurotransmission and hormone secretion. Sec1/Munc18 (SM) proteins and soluble *N*-  
3 ethylmaleimide-sensitive factor attachment protein receptor (SNARE) proteins are the two key  
4 protein families of exocytosis. Three conserved SNARE proteins including syntaxin-1 (STX1),  
5 synaptosomal-associated protein 25 (SNAP25) and synaptobrevin 2 (SYB2) are involved to  
6 promote granule exocytosis by forming a heterotrimeric complex (24, 25). SM proteins are  
7 essential regulators of SNARE protein-mediated vesicle docking or fusion events, acting as high  
8 affinity binding partners for syntaxin proteins (26). There are three plasma membrane-localized  
9 SM proteins (Munc18-1/2/3) expressed in mammalian cells which were found to play specific  
10 roles in beta-cell insulin secretion and to modulate glucose homeostasis. Depletion of Munc18-1,  
11 Munc18-2 or Munc18-3 causes defective GSIS in mouse and human islets whereas  
12 overexpression of either Munc18-1 or Munc18-2 potentiates GSIS in human islets (27-29). On  
13 the other hand, down-regulation of SM proteins has been observed in pancreatic islets of insulin-  
14 resistant rats and T2D patients (28, 30, 31), suggesting a positive association between islet  
15 Munc18 expression level and islet function in both rodents and human. While the pivotal  
16 function of these proteins in insulin secretion is irrefutable, the regulatory mechanism of SM  
17 proteins, especially their diminution during T2D remains ambiguous. Therefore, deciphering the  
18 mechanism that regulates SM protein expression or activity is of paramount importance to  
19 understand the pathophysiology of T2D, paving the way for generation of innovative anti-  
20 diabetic therapy.

21 In this study, we uncovered a novel mechanism by which PAX6 regulates human beta-cell  
22 insulin secretion through modulation of CREB signaling pathway, a direct transcriptional  
23 regulator of exocytotic proteins Munc18-1 and Munc18-2. Suppression of this pathway under

1 diabetic condition results in defective insulin secretion; while restoration of PAX6 expression  
2 rescues CREB activation and Munc18-1/2 down-regulation, enhances insulin-secreting capacity  
3 and ultimately ameliorates glucose homeostasis.

#### 4 **Results**

5 **PAX6 regulates Munc18-1/2 expression in human pancreatic beta-cells.** Exocytosis is the  
6 final and critical step of insulin secretion. To elucidate the role of PAX6 in insulin exocytosis,  
7 the expression levels of several core exocytotic proteins were first measured under conditions  
8 with PAX6 manipulated. Lentivirus-mediated PAX6 overexpression in EndoC- $\beta$ H1 cells  
9 increased mRNA levels of *MUNC18-1* and *MUNC18-2* by ~3-fold and 2-fold over the respective  
10 control levels (Fig. 1A) without altering the mRNA levels of other selected exocytotic proteins  
11 including *MUNC18-3*, *MUNC13-1*, *SNAP25*, *SYB2* and *STX1A*. Consistently, short hairpin RNA  
12 (shRNA)-mediated *PAX6* knockdown significantly decreased mRNA levels of *MUNC18-1* and  
13 *MUNC18-2* (Fig. 1B; ~50% and 70% of control group respectively) without affecting the mRNA  
14 levels of some other exocytotic proteins. In addition, we examined Munc18-1 and Munc18-2  
15 protein expression under PAX6 overexpression and knockdown conditions which confirmed the  
16 mRNA expression findings (Fig. 1C and 1D). These results showed that PAX6 is a positive  
17 regulator of Munc18-1/2 expression and loss of PAX6 would lead to reduced expression of  
18 Munc18-1/2.

19 **PAX6 regulates GSIS through modulation of Munc18-1/2 expression.** We next investigated  
20 the role of PAX6/Munc18-1/2 axis in insulin secretion. PAX6 knockdown in EndoC- $\beta$ H1 cells  
21 led to a decrement in protein levels of both Munc18-1 and Munc18-2, which were reversed by  
22 individual overexpression of either Munc18-1 or Munc18-2 without cross-affecting the other one

1 (Fig. S2A and S2B). Although insulin content was lower in cells with PAX6 knockdown, there  
2 was no significant difference in basal insulin secretion (Fig. 2A and S3A). However, GSIS was  
3 greatly suppressed in the PAX6 knockdown group when compared with the control group.  
4 Notably, individual overexpression of Munc18-1 or Munc18-2 did not exert significant effect on  
5 GSIS, but mitigated the defective GSIS caused by PAX6 knockdown (Fig. 2A). Perfusion of  
6 EndoC- $\beta$ H1 cells with high glucose resulted in a burst of insulin secretion (first-phase) which  
7 declined and was followed by a slow increase (second-phase) in insulin secretion. PAX6  
8 knockdown markedly reduced GSIS in both first- and second-phases and overexpression of  
9 Munc18-1 or Munc18-2 tended to improve them (Fig. 2B). To examine whether reduced insulin  
10 secretion could be the result of a distal signaling defect, we measured potassium-stimulated  
11 insulin secretion (KSIS) in PAX6 knockdown cells. Similar to GSIS, KSIS was significantly  
12 impaired by PAX6 knockdown. Importantly, overexpression of Munc18-1 or Munc18-2 largely  
13 restored the KSIS defect (Fig. 2B). To further elucidate the interaction between PAX6 and  
14 Munc18-1/2 in regard to insulin secretion, we first generated stable Munc18-1 or Munc18-2  
15 knockdown in EndoC- $\beta$ H1 cells, then overexpressed PAX6 in these cells. As shown in Fig. S2C  
16 and S2D, reduced Munc18-1 and Munc18-2 protein levels by shRNA knockdown were enhanced  
17 by PAX6 overexpression. For insulin secretion, Munc18-1 or Munc18-2 knockdown resulted in  
18 decreased GSIS compared to the control group. PAX6 overexpression alone had no effect on  
19 GSIS, but could fully rescue the reduced insulin secretion associated with Munc18-1 or Munc18-  
20 2 knockdown (Fig. 2C). Dynamic measurement of insulin secretion further demonstrated that  
21 Munc18-1 and Munc18-2 knockdown reduced GSIS in both first- and second-phases while  
22 PAX6 overexpression could relieve the defects. Knockdown of Munc18-1 or Munc18-2 also  
23 suppressed KSIS where PAX6 overexpression tended to restore it (Fig. 2D). In addition, PAX6



1 knockdown also diminished insulin secretion induced by Exendin-4, a GLP-1 analog (Fig. S3B).  
2 This result further validated the essential role of PAX6 in insulin exocytosis, the final and  
3 common step of stimulated insulin secretion.

4 **PAX6 regulates Munc18-1/2 expression via modulation of CREB signaling.** The above  
5 results demonstrated that PAX6 modulated the expression of exocytotic proteins Munc18-1 and  
6 Munc18-2, which in turn affected insulin secretion. But how PAX6 regulates Munc18-1/2  
7 expression is unclear. To examine whether both Munc18-1 and Munc18-2 are the direct targets  
8 of PAX6, we checked if the promoter regions of Munc18-1/2 contain PAX6 consensus binding  
9 site (TTCACGCATGAGTG) but did not identify any putative binding sites. Luciferase reporter  
10 assay using luciferase vectors carrying human *MUNC18-1* and *MUNC18-2* promoters with GFP  
11 or PAX6-overexpressing vector co-transfection also showed that PAX6 itself was unable to  
12 activate *MUNC18-1* and *MUNC18-2* promoters (Fig. S4A), suggesting that the regulation might  
13 depend on other factors. Several putative CREs had been identified in both mouse and human  
14 *MUNC18-1* and *MUNC18-2* promoters (Table S2). Luciferase reporter assay showed that CREB  
15 overexpression, raising phosphorylated and total CREB levels (Fig. S1D), drastically enhanced  
16 luciferase activity (Fig. 3A), indicating the transactivation of both *MUNC18-1* and *MUNC18-2*  
17 promoters by CREB. Of note, deletion of the putative CREs in both human *MUNC18-1* and  
18 *MUNC18-2* promoters reduced the transactivation by CREB overexpression, confirming the  
19 functionality of those CREs. Moreover, glucose and forskolin (FSK), the CREB activators,  
20 increased the promoter activity with effects weakened by ablation of CREs (Fig. S4C).  
21 Accordingly, chromatin immunoprecipitation assay showed that there was a significant  
22 enrichment of phospho-CREB to the CREs in both human *MUNC18-1* and *MUNC18-2*  
23 promoters (Fig. 3B), which revealed the physical bindings of phospho-CREB to both promoters.

1 CREB activation mediated by FSK or isobutylmethylxanthine (IBMX) greatly enhanced the  
2 binding of phospho-CREB to *MUNC18-1* and *MUNC18-2* promoters (Fig. S4B). Functionally,  
3 FSK and IBMX induced Munc18-1 and Munc18-2 expression at both mRNA and protein levels  
4 whereas CREB knockdown down-regulated their expression levels (Fig. S5). These data  
5 unravelled CREB as a direct transcriptional regulator of Munc18-1/2. To determine whether  
6 PAX6 modulates Munc18-1/2 expression via CREB signaling, regulation of CREB signaling by  
7 PAX6 was examined. As shown in Fig. 3C and S6A, PAX6 knockdown reduced total CREB  
8 protein level and the mRNA expression of some of the key CREB family members including  
9 *CREB* and *CREM* while PAX6 overexpression enhanced their mRNA levels. Blunted glucose-  
10 induced CREB phosphorylation (Ser133) and transcription of CREB target genes, *cFOS* and  
11 *NR4A1* (32) were observed in cells with PAX6 knockdown (Fig. 3C and S6B). Concomitantly,  
12 glucose-stimulated ATP and cAMP production as well as  $Ca^{2+}$  influx, which are the substrates  
13 for CREB activation, were suppressed by PAX6 knockdown (Fig. 3D, 3E and S6C), likely  
14 resulting from reduced expression of Glut1 (Fig. S8A), the predominant glucose transporter in  
15 human beta-cells (33). In particular, overexpression of CREB or CREB-regulated transcription  
16 coactivator 2 (CRTC2) prevented the down-regulation of both Munc18-1 and Munc18-2 caused  
17 by PAX6 knockdown (Fig. 3F, S7A and S7C) whereas CREB inhibition (666-15) prevented their  
18 up-regulation induced by PAX6 overexpression (Fig. 3G and S7B). These results indicated that  
19 the regulation of Munc18-1/2 expression by PAX6 is mostly CREB-dependent. For insulin  
20 secretion, CREB or CRTC2 overexpression preserved GSIS impaired by PAX6 knockdown (Fig.  
21 3H). Collectively, these observations provided evidence that PAX6 regulated CREB signaling  
22 which in turn modulated Munc18-1/2 expression and insulin secretion at the functional level.

1 **Repressed PAX6/CREB/Munc18 axis in pancreatic beta-cells under diabetic condition**  
2 **leads to blunted insulin secretion.** As reduced GSIS is a hallmark of beta-cell dysfunction in  
3 T2D, we examined the potential participation of PAX6/CREB/Munc18 axis in T2D-associated  
4 defective GSIS by treating EndoC- $\beta$ H1 cells or primary mouse islets with high glucose and  
5 palmitic acid (HGPA) which mimicked diabetic condition. Long-term (>48 h) HGPA treatment  
6 evidently attenuated protein levels of PAX6, Munc18-1 and Munc18-2 in both EndoC- $\beta$ H1 cells  
7 and mouse islets (Fig. 4A); while these were alleviated by PAX6 restoration in cells (Fig. 4B and  
8 S7D). The involvement of CREB signaling was also investigated. Results showed that HGPA  
9 treatment prominently decreased Glut1 expression, inhibited glucose-induced ATP and cAMP  
10 production, Ca<sup>2+</sup> influx as well as CREB Ser133 phosphorylation in EndoC- $\beta$ H1 cells, all of  
11 which could be restored by PAX6 overexpression (Fig. S8B, 4C, 4D, S6D and 4E). As a  
12 downstream effector of PAX6, direct CREB or CRT2 overexpression was able to counteract  
13 Munc18-1 and Munc18-2 down-regulation led by HGPA treatment (Fig. 4F and S7D),  
14 confirming the regulatory role of CREB in Munc18-1 and Munc18-2 expression under diabetic  
15 condition. For insulin secretion, 72 h-incubation of EndoC- $\beta$ H1 cells in HGPA considerably  
16 suppressed GSIS while individual overexpression of PAX6, Munc18-1, Munc18-2, CREB or  
17 CRT2 ameliorated the defect to varying extents (Fig. 4G). These data indicated that repressed  
18 PAX6/CREB/Munc18 axis in beta-cells contributed to compromised beta-cell function in  
19 diabetic condition.

20 **Beta-cell specific PAX6 overexpression preserves CREB/Munc18-1/2 signaling pathway**  
21 **and GSIS in *db/db* mouse islets.** The above results evidently illustrated the modulatory role of  
22 PAX6/CREB/Munc18 axis in beta-cell insulin secretion. We then substantiated their role in a  
23 real physiological situation using *db/db* mice, a rodent model of T2D. In agreement with the

1 observation in EndoC-βH1 cells under HGPA condition, isolated *db/db* mouse islets exhibited  
2 marked diminution in PAX6, Munc18-1 and Munc18-2 protein expression compared to *db/+*  
3 control mice (Fig. 5A). To elucidate the *in vivo* effects of PAX6 restoration in beta-cell function,  
4 8-week-old *db/db* mice were injected with adeno-associated virus (AAV) overexpressing GFP  
5 control or PAX6 driven by the rat insulin promoter. Immunofluorescence staining showed that  
6 most of the exogenous PAX6 was expressed in beta-cells but not in alpha-cells with minimal  
7 effect on other internal organs (Fig. 5B, S9A and S9B). Western blotting further showed the  
8 presence of exogenous PAX6 in pancreatic islets 6 weeks after gene delivery (Fig. S9C and S9D).  
9 Of note, AAV-PAX6 counteracted the down-regulation of both Munc18-1 and Munc18-2 as well  
10 as Glut2 in *db/db* mouse islets (Fig. 5C and S8C). Glucose-induced ATP and cAMP production  
11 was abolished in *db/db* mouse islets compared to *db/+* control which was elevated by AAV-  
12 PAX6 (Fig. 5D and 5E). Concurrently, *db/db* mouse islets displayed attenuated CREB Ser133  
13 phosphorylation upon high glucose stimulation which was again augmented by PAX6  
14 overexpression (Fig. 5F). For insulin-secreting capacity, isolated *db/db* mouse islets had  
15 diminished GSIS which was substantially potentiated by AAV-PAX6 (Fig. 5G). Dynamic  
16 measurement of insulin release showed a significant decline in both first- and second-phases of  
17 GSIS in *db/db* mouse islets compared to *db/+* mouse islets while AAV-PAX6 prominently  
18 enhanced the first-phase of insulin secretion. KSIS was also weakened in *db/db* mouse islets, and  
19 this was largely recovered by AAV-PAX6 (Fig. 5H).

20 **Beta-cell specific PAX6 overexpression improves glucose homeostasis in *db/db* mice.** *Db/db*  
21 mouse islets displayed repressed PAX6/CREB/Munc18 axis correlated with impaired GSIS,  
22 which is fundamental to deteriorated glucose homeostasis in T2D, we hence examined the  
23 potential anti-diabetic effect of AAV-PAX6 in *db/db* mice *in vivo*. Strikingly, intraperitoneal

1 glucose tolerance test (IPGTT) showed that AAV-PAX6 improved glucose tolerance in *db/db*  
2 mice 2 weeks after injection and the effect was sustained for at least 6 weeks (Fig. 6A).  
3 Measurement of insulin secretion during IPGTT revealed that the early phase of insulin secretion  
4 was lost in *db/db* mice but was regained by AAV-PAX6 treatment (Fig. 6B). Hyperglycemia was  
5 also progressively alleviated without affecting weight gain (Fig. 6C and 6D). Given that  
6 peripheral insulin resistance is another factor contributing to glucose intolerance in T2D, insulin  
7 tolerance test (ITT) was performed to evaluate the effect of AAV-PAX6 on insulin sensitivity.  
8 As shown in Fig. 6E, insulin sensitivity in *db/db* mice was not influenced by beta-cell specific  
9 PAX6 overexpression, indicating that the moderation of glucose intolerance was mostly due to  
10 the potentiation of insulin secretion at those time points. Collectively, these results provided  
11 compelling evidence that restoration of the PAX6/CREB/Munc18 axis in islets promoted GSIS  
12 which in turn ameliorated glucose metabolism in diabetic mice.

13 **Beta-cell specific PAX6 overexpression restores CREB/Munc18-1/2 signaling pathway and**  
14 **GSIS in T2D human islets.** To determine whether reduction of islet PAX6 and Munc18-1/2  
15 expression might occur in human T2D, we first determined the protein expression levels of  
16 PAX6 and Munc18-1/2 in islets from normal and T2D donors. In accordance with the results  
17 observed in *db/db* mice, the expression levels of PAX6, Munc18-1 and Munc18-2 were  
18 markedly decreased in T2D islets (Fig. 7A). AAV-PAX6 led to the expression of exogenous  
19 PAX6 mostly in beta-cells (Fig. S10A and S10B), noticeably restored the protein levels of  
20 Munc18-1 and Munc18-2 as well as Glut1 in T2D islets (Fig. 7B and S8D). Glucose-induced  
21 ATP and cAMP generation was dampened in T2D islets which was elevated by AAV-PAX6 (Fig.  
22 7C and 7D). For CREB signaling, by contrast with the robust induction of Ser133  
23 phosphorylation in normal islets, glucose had only modest effect on T2D islets which was

1 augmented by AAV-PAX6 (Fig. 7E). Functionally, GSIS was attenuated in T2D islets but could  
2 be mitigated by AAV-PAX6 (Fig. 7F). Dynamic measurement of insulin release showed an  
3 absence of the first-phase and a weakened second-phase of insulin release upon glucose  
4 challenge in T2D islets compared to normal islets; whereas a clear biphasic pattern of insulin  
5 release was regained by AAV-PAX6. The reduced K<sub>SI</sub>S in T2D islets was also enhanced by  
6 AAV-PAX6 (Fig. 7G). Overall, these results corroborated the existence of dampened  
7 PAX6/CREB/Munc18 axis in human T2D islets; while replenishment of PAX6 reversed those  
8 defects and potentiated GSIS.

## 9 **Discussion**

10 Impaired insulin secretion is a major feature of beta-cell failure determining the progression of  
11 insulin resistance to hyperglycemia and overt T2D, which has become a global public health  
12 threat. Therefore, gaining a better understanding of the molecular mechanism underlying beta-  
13 cell dysfunction is of paramount importance because it provides new insights into potential  
14 effective intervention strategies for T2D. Insulin secretion is a dynamic process and its regulation  
15 is highly complex, resulting from a combinatorial network of transcription factors. PAX6 is one  
16 of the major factors determining adult beta-cell function. PAX6 ablation in adult murine islet  
17 cells attenuates insulin production and secretion resulting in hyperglycemia and glucose  
18 intolerance (5, 7, 8). Mechanistic studies performed in rodent beta-cells demonstrated that PAX6  
19 modulates the proximal signaling of GSIS including glucose sensing, ATP production and Ca<sup>2+</sup>  
20 dynamics (6, 8). Most of the prior studies looked into the physiological roles of PAX6 in rodent  
21 beta-cells, however, whether PAX6 exerts the same functions in human beta-cells has not been  
22 demonstrated, there is also no direct evidence showing the involvement of PAX6 in human T2D.  
23 In this study, we made several novel observations concerning the role of PAX6 in beta-cell

1 function and T2D progression. Firstly, we provided *in vitro* and *ex vivo* evidence showing for the  
2 first time that PAX6 regulates both proximal and distal signaling of GSIS in human beta-cells by  
3 controlling glucose transport, ATP and cAMP production, Ca<sup>2+</sup> influx, CREB activation and  
4 expression of exocytotic proteins Munc18-1 and Munc18-2. Secondly, we found that the  
5 PAX6/CREB/Munc18-1/2 axis is dampened in human pancreatic islets under T2D condition,  
6 unraveling the pathophysiological role of this axis in human diabetes. Thirdly, we showed that  
7 AAV-mediated replenishment of PAX6 prominently improved beta-cell function and alleviated  
8 glycemic disturbance in *db/db* mice *in vivo*.

9 Exocytosis is the final but the most critical step of insulin secretion. Various nutrients or stimuli  
10 trigger insulin secretion through different pathways but all converge on exocytosis to accomplish  
11 secretion. Cognate SM proteins are indispensable to execute insulin secretion in mouse and  
12 human islets (27-29). We have provided new evidence illustrating the regulatory role of PAX6 in  
13 the distal signaling of GSIS by modulating the expression of exocytotic proteins Munc18-1/2 in  
14 human beta-cells. Importantly, the present findings are in line with data from earlier studies (8).  
15 Thus, interrogation of the RNA-Seq data from the study reveals that *Stxbp1* (Munc18-1) mRNA  
16 was reduced highly significantly (~1.6 fold) in adult beta-cell-selective *Pax6* KO mice (8),  
17 though *Stxbp2* (Munc18-2) expression was not changed at the mRNA level. Our *in vitro* study  
18 using human beta-cell line demonstrated that Munc18-1/2 knockdown suppressed both GSIS and  
19 KSIS while PAX6 overexpression largely restored the defects. Conversely, PAX6 knockdown  
20 diminished Munc18-1/2 expression and suppressed both GSIS and KSIS. Overexpression of  
21 Munc18-1/2 could only partially rescue the reduced GSIS. This partial rescue is likely because  
22 PAX6 modulates both proximal and distal mechanisms of GSIS and overexpression of Munc18-

1 1/2 could only restore the distal mechanism, but not the proximal signaling caused by PAX6  
2 knockdown.

3 We further demonstrated that CREB is the downstream mediator of PAX6 in controlling  
4 Munc18-1/2 expression. CREB is a transcription factor critical to maintain functional beta-cell  
5 mass by enhancing cell survival and preventing apoptosis. CREB specifically regulates the  
6 expression of crucial genes including Bcl-2, cyclins and insulin receptor substrate-2 (IRS-2) in  
7 beta-cells (19, 20, 22, 34). Although it is activated by a range of insulin-secreting stimuli such as  
8 glucose, GLP-1 and GIP (23), the role of CREB in insulin secretion remains elusive. In the  
9 present study, we uncovered Munc18-1 and Munc18-2 as the novel transcriptional targets of  
10 CREB. Putative CREs were identified in the promoter regions of both Munc18-1 and Munc18-2  
11 genes in mouse and human. Most putative CREs identified locate within 200 nucleotides  
12 upstream of the transcription start sites which are defined as the proximal promoter regions,  
13 where the CREs are most likely to be functional. Frequent occurrence of CREs also indicates that  
14 they are highly conserved across species (35). The full CREs existing in human (TGACGCGC)  
15 and mouse (TGACGCGT) Munc18-2 promoters had been recognized in previous studies (36,  
16 37); whilst the full CRE in human *MUNC18-1* promoter (TGACGCCA) has been shown to be a  
17 functional CRE (38). Chromatin immunoprecipitation study and luciferase reporter assay  
18 substantiated the physical binding and transactivation of human *MUNC18-1* and *MUNC18-2*  
19 promoters by CREB. Of note, our results showed that PAX6 knockdown in EndoC-βH1 cells  
20 reduced Glut1 expression, glucose-induced ATP/ cAMP generation and Ca<sup>2+</sup> influx, which  
21 would in turn lead to repressed CREB activation. Indeed, PAX6 knockdown *per se* reduced the  
22 expression levels of CREB and other CREB family members which would further attenuate  
23 CREB activity. However, whether PAX6 regulates CREB expression by a direct transcriptional



1 modulation or through other signaling pathways requires further investigation. Functionally,  
2 PAX6 knockdown and overexpression resulting in Munc18-1/2 down- and up-regulation were  
3 counteracted by CREB overexpression/ activation and inhibition respectively. Impaired GSIS led  
4 by PAX6 knockdown and HGPA treatment was also relieved by CREB/ CRT2 overexpression.  
5 Overall, apart from the reported regulatory role of PAX6 in the proximal signaling of GSIS, our  
6 results provide compelling evidence elucidating a previously unknown signaling network  
7 between PAX6 and CREB in the modulation of Munc18-1/2 expression, which underlies the role  
8 of PAX6 in the coordination of the proximal and distal signaling of GSIS.

9 Progressive deterioration of beta-cell function, particularly impairment of GSIS, is a central  
10 event driving T2D progression. To gain insights into the pathophysiological role of the  
11 PAX6/CREB/Munc18-1/2 axis in T2D-associated beta-cell dysfunction, EndoC- $\beta$ H1 cells were  
12 exposed to HGPA to mimic T2D condition. HGPA down-regulated the entire pathway and  
13 suppressed GSIS which were rescued by restoration of PAX6 expression, indicating that  
14 dysregulation of the PAX6/CREB/Munc18-1/2 would be one of the pathways driving impaired  
15 GSIS under diabetic condition. The therapeutic potential of PAX6 in T2D was further delineated  
16 using *db/db* mice which exhibited prominently defective insulin secretion and frank diabetes. In  
17 line with the aforementioned *in vitro* results, the PAX6/CREB/Munc18-1/2 axis was repressed in  
18 *db/db* mouse islets. As there is no known specific agonist for PAX6, we adopted the AAV  
19 approach to restore PAX6 expression. Non-integrative AAV vectors, derived from non-  
20 pathogenic viruses, persist for years as episomes in nuclei of non-dividing cells (39). Multiple  
21 clinical studies have provided promising evidence of AAV-mediated long-term expression of  
22 therapeutic proteins without causing significant adverse effects (40-42). Here, we took advantage  
23 of AAV vectors and their ability to mediate long-term production of target proteins to develop a

1 gene therapy strategy for T2D based on PAX6 transfer to beta-cells. AAV-PAX6 potentiated the  
2 CREB/Munc18-1/2 axis, markedly enhanced the first-phase of GSIS *in vivo* and *ex vivo* while  
3 ameliorating glucose homeostasis in *db/db* mice. Improvement in glucose homeostasis could be  
4 observed 2 weeks after AAV injection and was sustained for least 6 weeks. To further explore  
5 the relevance of these findings to humans, we compared the axis in human islets between normal  
6 and T2D donors. Strikingly, a similar phenomenon had been observed in human islets. The  
7 PAX6/CREB/Munc18-1/2 axis was significantly dampened in T2D islets with impaired insulin  
8 secretion and PAX6 replenishment in beta-cells rescued the pathway and amplified both GSIS  
9 and KSIS. Our results unveiled the molecular link between PAX6 and CREB in the regulation of  
10 Munc18-1/2 expression and insulin secretion in human islets, in which attenuated PAX6 and  
11 CREB activity may explain the decrease in Munc18-1/2 expression under diabetic conditions (28,  
12 30, 31). These observations further support the notion that PAX6 crucially determines beta-cell  
13 function in both mouse and human, while disruption of PAX6 expression or activity drives islet  
14 dysfunction and inflicts high risk to overt T2D progression.

15 Early and late phases of insulin secretion probably serve distinct physiological roles. Early phase  
16 response appears to efficiently suppress endogenous glucose production after meal while the late  
17 phase response predominately serves to enhance insulin-mediated glucose disposal in skeletal  
18 muscle and adipose tissue (43, 44). The progression from impaired glucose tolerance to frank  
19 diabetes is characterized by a dramatic decline in early insulin secretion as loss of early insulin  
20 secretion initially leads to postprandial hyperglycemia (45, 46). In this regard, by replenishing  
21 the upstream diminished modulator, AAV-PAX6 remarkably potentiated the first-phase of GSIS  
22 in diabetic mouse and human islets by targeting glucose transport, ATP/ cAMP generation and  
23 insulin exocytosis, thus restraining the deterioration of hyperglycemia in *db/db* mice. Although

1 AAV-PAX6 might not fully reverse T2D in mice, this strategy mitigated beta-cell dysfunction,  
2 one of the major contributors to T2D and demonstrated its potential in sustained alleviation of  
3 glycemic disturbance in T2D. Additionally, loss of beta-cell identity (dedifferentiation) is an  
4 emerging mechanism contributing to beta-cell dysfunction in T2D (47), while PAX6 has been  
5 demonstrated to maintain beta-cell identity and to repress alternative islet cell genes (7). Apart  
6 from the regulation of insulin secretion, further investigation on human beta-cell  
7 dedifferentiation is warranted to gain better understanding on the effects of PAX6 replenishment.  
8 It would also be important to investigate how PAX6 expression or activity is regulated.  
9 Identification of candidates that critically confer blunted PAX6 action in beta-cells under  
10 diabetic condition would aid the development of novel, effective therapy against T2D.

11 In summary, the present study advances the understanding of PAX6 as an indispensable  
12 transcription factor controlling GSIS in human pancreatic beta-cells through coordination of the  
13 proximal and distal signaling. Our results also shed light onto the pathophysiological role of  
14 PAX6 in T2D and provide new insights and scientific basis to support the future clinical  
15 translation of PAX6 gene transfer to overcome beta-cell failure in T2D.

## 16 **Materials and Methods**

17 **Study Design.** This study aimed to examine the molecular pathway of PAX6 in the regulation of  
18 insulin secretion, the pathophysiological role of PAX6 in beta-cell dysfunction and the potential  
19 therapeutic value of PAX6 in T2D. We performed loss- and gain-of-function studies using  
20 human EndoC- $\beta$ H1 cell line to investigate the regulatory role of PAX6 in GSIS and the  
21 molecular mechanism. Type 2 diabetic *db/db* mice and primary human islets from normal and  
22 T2D donors were used to study the link between PAX6 and T2D-associated beta-cell failure.

1 Beta-cell specific overexpression of PAX6 was achieved by AAV administration. Glucose  
2 homeostasis of *db/db* mice and their lean control was monitored by IPGTT and ITT with insulin-  
3 secreting capacity evaluated. Islets were harvested at the endpoint for mechanistic studies. Mice  
4 were randomly divided into different groups and no animals were excluded because of  
5 illness. All experimental procedures involving animals were approved by the Institutional  
6 Animal Care and Use Committee in at A\*STAR (Ref. No. 191479). All procedures involving  
7 human tissues were approved by A\*STAR Institutional Review Board (Ref. No. 2019-021). All  
8 experiments were performed in at least triplicate biological repeats.

9 **Cell culture.** The human EndoC- $\beta$ H1 cell line (Univercell-Biosolutions, Toulouse, France) was  
10 cultured as previously described (48). Cells were maintained under normal condition (5.6 mmol/l  
11 glucose) or treated with HGPA (25 mmol/l D-glucose and 0.3 mmol/l palmitic acid).

12 **Lentivirus-mediated gene overexpression and knockdown.** For gene overexpression, coding  
13 sequences expressing human *PAX6*, *MUNC18-1*, *MUNC18-2*, *CREB1* (OriGene, Rockville, MD,  
14 USA) and *CRTC2* (VectorBuilder, Santa Clara, CA, USA) were subcloned into the lentiviral  
15 pLV vector (Addgene, Cambridge, MA, USA) to generate overexpression constructs. For gene  
16 knockdown, shRNA sequences targeting human *PAX6*, *MUNC18-1*, *MUNC18-2* and *CREB1*  
17 (Table S1) were subcloned into the lentiviral pLKO.1 vector (Addgene) to generate knockdown  
18 constructs. Three shRNA sequences were used to knock down each of the gene with the  
19 sequence showing the highest knockdown efficiency chosen for downstream experiments (Fig.  
20 S1A-S1C). A non-targeting shRNA sequence was used as control. All DNA constructs were  
21 validated by sequencing. Viral particles were generated by transient transfection of HEK-293FT  
22 cells (American Type Culture Collection, Manassas, VA, USA) with the lentiviral plasmids and  
23 helper plasmids (pRSV-Rev, pCMV-VSVG and pMDLg/pRRE; Addgene) using lipofectamine

1 2000 reagent (Thermo Fisher Scientific, Rockford, IL, USA). The cell supernatant was collected,  
2 concentrated and titrated at 48 h post-transfection. EndoC-βH1 cells were infected with  
3 overexpressing or shRNA viral particles and were subjected to antibiotic selection to generate  
4 stable cell lines. For double infections, stable knockdown cell lines were first generated followed  
5 by overexpression. Viral particles carrying different selection markers (puromycin for  
6 knockdown and blasticidin for overexpression) were used.

7 **Animal models.** Male genetically diabetic C57BL/KsJ-db (*db/db*) and their age-matched, lean  
8 heterozygote littermates (*db/+*) were obtained from the Jackson Laboratory (Bar Harbor, ME,  
9 USA). Animals were fed *ad libitum* and maintained in a specific pathogen free facility with  
10 constant ambient temperature and a 12-h light/dark cycle.

11 **Administration of AAV.** The AAV8 carrying PAX6-flag-GFP (AAV-PAX6) or GFP only  
12 (AAV-Ctrl) driven by the rat insulin promoter was purchased from VectorBuilder. AAV-PAX6  
13 or AAV-Ctrl was injected intraperitoneally into *db/+* and *db/db* mice (8-week-old) at the dose of  
14  $1 \times 10^{11}$  viral genomes (vg) per mouse as previously described (49, 50). Mice were observed for  
15 6 weeks after AAV injection with pancreatic islets and organs isolated for analyses at the end  
16 point.

17 **In vivo glucose homeostasis.** Glucose tolerance was assessed by IPGTT. After 16 h fasting,  
18 mice were given 1 g/kg body weight of glucose (Sigma-Aldrich, St. Louis, MO, USA) by  
19 intraperitoneal injection. Blood glucose and serum insulin levels were measured at the indicated  
20 time points. The first- and second-phases of insulin secretion were defined as 0-15 and 15-60  
21 min respectively. For ITT, mice were injected with insulin (1 U/kg body weight; Sigma-Aldrich)  
22 after 4 h fasting and blood glucose levels were subsequently measured.

1 **Pancreatic islet isolation, primary culture and treatments.** Intact pancreatic islets were  
2 isolated from mice as previously described (51). Briefly, mouse pancreas was given an intra-  
3 ductal injection of collagenase P (Roche, Mannheim, Germany) in Hanks' balanced salt solution  
4 (Gibco, Grand Island, NY, USA). The pancreas was removed and digested in 37°C. After  
5 washing and gradient centrifugation, the islets were handpicked under a stereomicroscope.  
6 Isolated islets were further dispersed into single cells by trypsin digestion for ATP/ADP and  
7 cAMP measurement. Intact islets or dispersed cells were then cultured overnight for recovery in  
8 RPMI-1640 medium (Thermo Fisher Scientific) supplemented with 10% (vol/vol) fetal bovine  
9 serum (FBS; Gibco) and 1% (vol/vol) penicillin and streptomycin (Thermo Fisher Scientific).

10 **Human islets, culture and transduction.** Primary human islets from 4 normal and 4 T2D  
11 donors were obtained from Prodo Laboratories, Inc. (Aliso Viejo, CA, USA). Human islets were  
12 cultured in PIM(S) complete media (Prodo Labs) and transduced with AAV-Ctrl or AAV-PAX6  
13 at a dose of 20 000 vg per islet cell.

14 **Quantitative real-time PCR.** Total RNA from cell samples was extracted using TRIzol reagent  
15 (Thermo Fisher Scientific) and subjected to reverse transcription using High-Capacity cDNA  
16 Reverse Transcription Kits (Applied Biosystems, Foster City, CA, USA). Gene expression was  
17 quantified by real-time PCR using SYBR™ Green PCR Master Mix (Applied Biosystems). The  
18 reactions were performed using an i-Cycler Thermal Cycler (Applied Biosystems). Relative gene  
19 expression was analyzed using the  $2^{-\Delta\Delta C_t}$  method and normalized relative to beta-actin. The  
20 sequences of primers used are listed in Table S1.

21 **Western blotting.** Proteins from isolated islets or organs were extracted using the RIPA Lysis  
22 and Extraction Buffer (Thermo Fisher Scientific). Proteins were separated by SDS-PAGE,

1 transferred to nitrocellulose membrane by iBlot™ 2 Gel Transfer Device (Thermo Fisher  
2 Scientific). After blocking, the membrane was incubated with antibodies against the following  
3 proteins: GAPDH (sc-32233; Santa Cruz Biotechnology Inc., Santa Cruz, CA, USA), phospho-  
4 CREB (9198; Cell Signaling, Danvers, MA, USA), CREB (9197; Cell Signaling), CRT2  
5 (ab109081; Abcam, Cambridge, MA, USA), PAX6 (ab197768; Abcam), Munc18-1 (ab109023;  
6 Abcam), Munc18-2 (ab97598; Abcam and GTX33529; Genetex Inc., Irvine, CA, USA), Glut1  
7 (ab40084; Abcam), Glut2 (ab54460; Abcam) or Flag (F1804; Sigma-Aldrich). After washing,  
8 the membrane was incubated with appropriate IR-Dye labeled secondary antibodies (LI-COR  
9 Biosciences, Lincoln, NE, USA). Labeled protein bands were visualized and quantitated by  
10 Odyssey® CLx Imager (LI-COR Biosciences).

11 **Insulin secretion assay.** For static insulin release, cells or size-matched isolated islets were pre-  
12 incubated in Krebs-Ringer bicarbonate buffer (KRBB; supplemented with 10 mmol/l HEPES  
13 and 2 mg/ml BSA) with 2.5 mmol/l glucose for 1 h followed by incubation in KRBB containing  
14 2.5 mmol/l or 16.7 mmol/l glucose for an additional 1 h, as described previously (51). Buffer was  
15 collected to measure insulin release by ELISA kits (Antibody and Immunoassay Services,  
16 University of Hong Kong, Hong Kong, China). For dynamic insulin release, cells or isolated  
17 islets were perfused with KRBB containing 2.5 mmol/l or 16.7 mmol/l glucose or 20 mmol/l  
18 KCl at a constant flow rate for the indicated periods of time. The first- and second-phases of  
19 insulin secretion were defined as 0-12 and 12-30 min respectively. Eluted fractions were  
20 collected to measure insulin release.

21 **Luciferase reporter assay.** DNA fragments corresponding to the promoter regions of human  
22 *MUNC18-1* and *MUNC18-2* genes were amplified by PCR from EndoC-βH1 cell genomic DNA  
23 using primers listed in Table S1. The PCR products were subcloned into the pGL3-basic

1 luciferase reporter vector (Promega, Madison, WI, USA). All constructs were validated by  
2 sequencing. EndoC-βH1 cells were co-transfected with the reporter/ control vector, a pRL-TK  
3 renilla luciferase vector (Promega) and a CREB-overexpressing/ control vector using  
4 Lipofectamine 2000 for 48 h. Cells were subjected to luciferase activity measurement by a dual-  
5 luciferase reporter assay kit (Promega) on a luminometer (Promega). The firefly luciferase  
6 activity was normalized against renilla luciferase activity.

7 **Chromatin immunoprecipitation.** EndoC-βH1 cells were fixed in 1% formaldehyde.  
8 Crosslinking reaction was stopped by glycine at 125 mmol/l final concentration. Cells were lysed  
9 and nuclear fraction was extracted for sonication to shear DNA fragments in the range of 200-  
10 600bp. Cell lysate was then subjected to immunoprecipitation with magnetic beads (Thermo  
11 Fisher Scientific) coated with anti-phospho-CREB antibody or normal rabbit IgG (Merck  
12 Millipore, Darmstadt, Germany) overnight at 4°C. The complex was washed and eluted with  
13 elution buffer (1% SDS and 0.1 M NaHCO<sub>3</sub>). The crosslinks were reversed by incubating the  
14 complex at 65°C overnight followed by DNA purification using QIAquick PCR Purification Kit  
15 (Qiagen, Hilden, Germany). The DNA fragments were analyzed by quantitative real-time PCR  
16 using primers targeting human *MUNC18-1* and *MUNC18-2* gene promoters. Primers targeting  
17 *GAPDH* region containing no CRE were used as negative control. All primers used are listed in  
18 Table S1.

19 **Measurement of intracellular ATP/ADP ratio and cAMP level.** EndoC-βH1 cells or  
20 dispersed islet cells were pre-incubated in KRBB with 2.5 mmol/l glucose for 1 h followed by  
21 stimulation with 2.5 mmol/l or 20 mmol/l glucose for 15 min. The relative ATP/ADP content  
22 and cAMP level were measured using the ADP/ATP ratio assay kit (Abcam) and cAMP  
23 immunoassay kit (Sigma-Aldrich) according to manufacturers' instructions.



1 **Immunofluorescence staining.** Isolated islets and organs were embedded and frozen. Cryostat  
2 sections were collected and fixed. Slides were incubated with mouse anti-flag (F1804; Sigma-  
3 Aldrich), guinea pig anti-insulin (PA1-26938; Thermo Fisher Scientific), rabbit anti-glucagon  
4 (ab92517; Abcam) and chicken anti-GFP (GFP-1020; Aves Labs, Tigard, OR, USA) antibodies.  
5 After washing, slides were probed with Alexa Fluor dye-conjugated secondary antibodies  
6 (Thermo Fisher Scientific). Slides were mounted with antifade mountant with DAPI (Thermo  
7 Fisher Scientific) before image acquisition. Digital images were acquired by a Ti-E  
8 fluorescence microscope (Nikon, Tokyo, Japan).

9 **Ca<sup>2+</sup> imaging.** EndoC-βH1 cells were transfected with the GCaMP7s vector (addgene) using  
10 Lipofectamine 2000 for 48 h. EndoC-βH1 cells were pre-incubated in KRBB with 2.5 mmol/l  
11 glucose for 1 h followed by stimulation with 20 mmol/l glucose. A Ti-E fluorescence microscope  
12 (Nikon) was used for live cell imaging. Image analysis was performed using the NIS-Elements  
13 (Nikon) software.

14 **Statistical analysis.** Data were expressed as means ± standard errors (SEMs). Comparisons  
15 between groups were analyzed by two-tailed Student's *t* test, one-way or two-way analysis of  
16 variance (ANOVA) followed by Tukey's *post hoc* test, where  $p < 0.05$  was considered statistically  
17 significant.

## 18 **Supplementary Materials**

19 **Fig. S1.** Western blotting showing shRNA knockdown efficiency and CREB overexpression in  
20 EndoC-βH1 cells.

21 **Fig. S2.** Western blotting showing protein expression of Munc18-1/2 in EndoC-βH1 cells with  
22 manipulation of PAX6 and Munc18-1/2.

- 1 **Fig. S3.** Effects of PAX6 knockdown on insulin content and Exendin-4-induced insulin secretion  
2 in EndoC-βH1 cells.
- 3 **Fig. S4.** Luciferase activity of human *MUNC18-1* and *MUNC18-2* promoters and chromatin  
4 immunoprecipitation assay showing binding of pCREB onto human *MUNC18-1* and *MUNC18-2*  
5 promoters.
- 6 **Fig. S5.** CREB phosphorylation, protein and mRNA levels of Munc18-1/2 in EndoC-βH1 cells  
7 after CREB activation or knockdown.
- 8 **Fig. S6.** PAX6's regulation of CREB family member gene expression, CREB target gene  
9 induction and Ca<sup>2+</sup> influx.
- 10 **Fig. S7.** Effects of CREB signaling regulation on Munc18-1/2 expression in EndoC-βH1 cells.
- 11 **Fig. S8.** Western blotting showing Glut1 and Glut2 protein expression.
- 12 **Fig. S9.** The expression of exogenous PAX6 in mice receiving AAV-PAX6.
- 13 **Fig. S10.** The expression of exogenous PAX6 in human islets with AAV-PAX6 treatment.
- 14 **Table S1.** Oligonucleotide sequences used in shRNA vectors construction, quantitative real-time  
15 PCR, luciferase vectors construction and ChIP assay.
- 16 **Table S2.** Sequences and positions of putative CREs in promoters of Munc18-1 and Munc18-2  
17 in both human and mouse.
- 18 **Table S3.** Human islet donor information.

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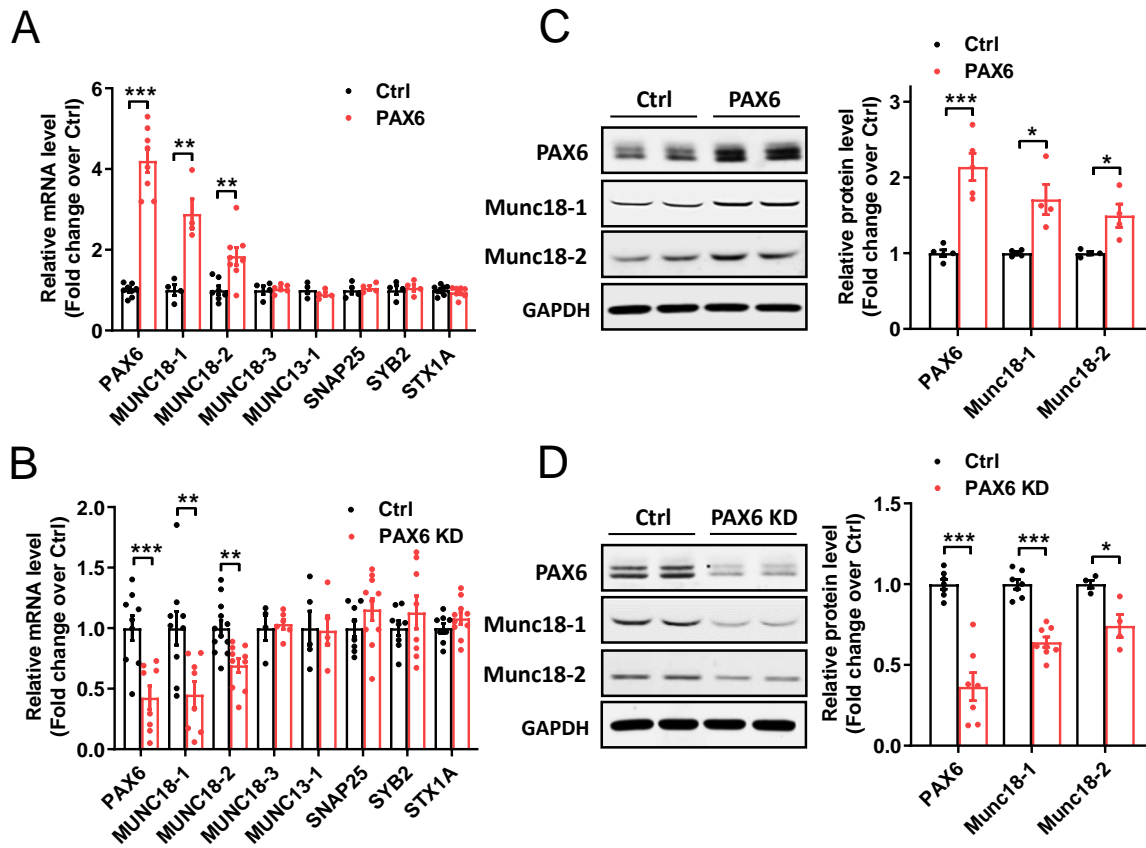
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8  
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20 W.N.L. performed experiments. A.K.K.T. edited and revised the manuscript. G.A.R. designed  
21 experiments, edited and revised the manuscript. W.H. designed the experiments, analyzed and  
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23 conflict of interest. **Data and materials availability:** All data associated with this study are  
24 present in the paper or the Supplementary Materials.

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1 **Figures**  
2

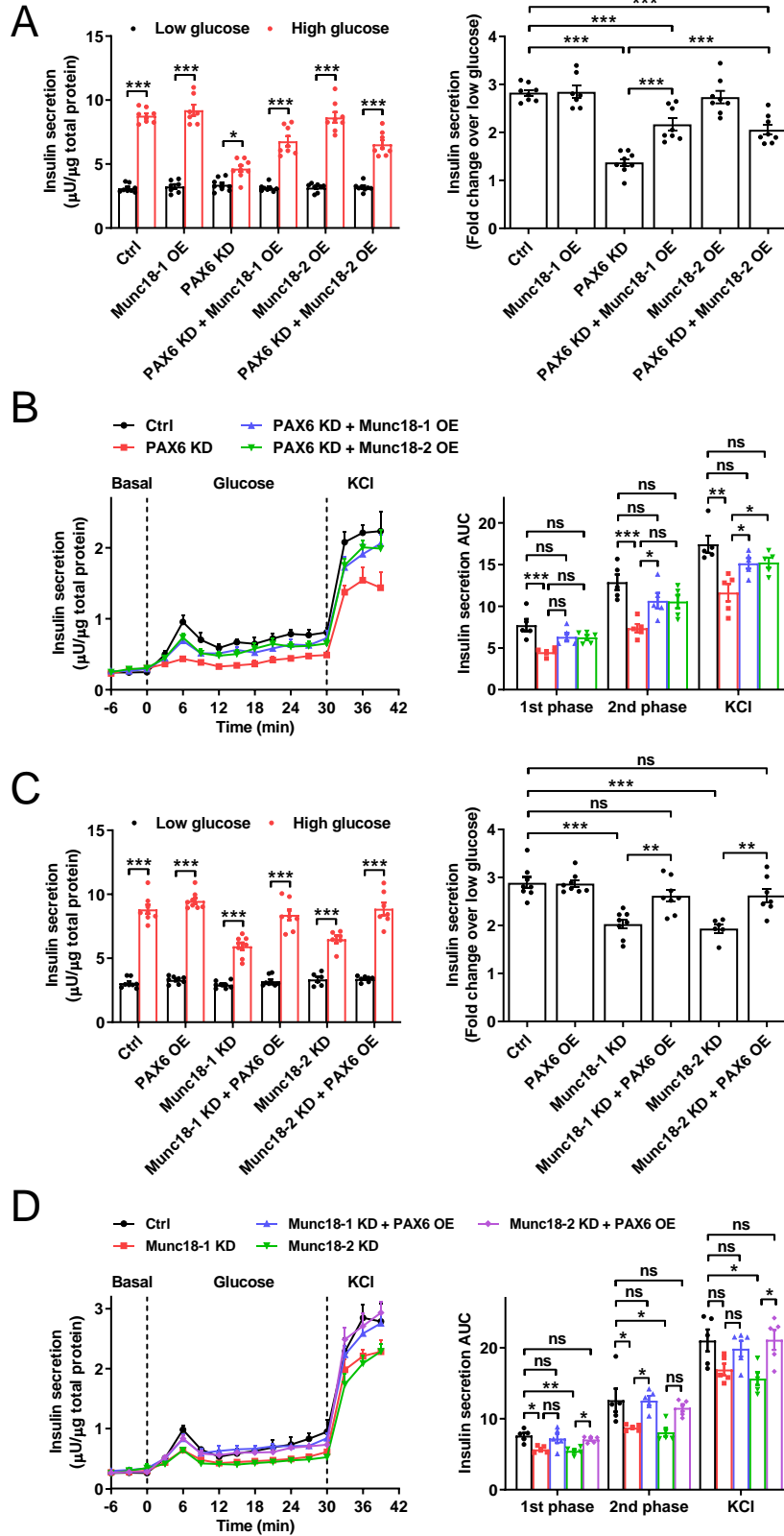
**Figure 1**



3

4 **Fig. 1.** PAX6 regulates Munc18-1/2 expression in human pancreatic beta-cells. mRNA  
5 expression levels of *PAX6* and exocytotic proteins in EndoC-βH1 cells with lentivirus-mediated  
6 PAX6 (A) overexpression and (B) knockdown (KD) ( $n=4-12$ ). Protein expression levels of PAX6  
7 and Munc18-1/2 in EndoC-βH1 cells with lentivirus-mediated PAX6 (C) overexpression and (D)  
8 knockdown. \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$  ( $n=4-8$ ). Data are means  $\pm$  SEMs.

Figure 2

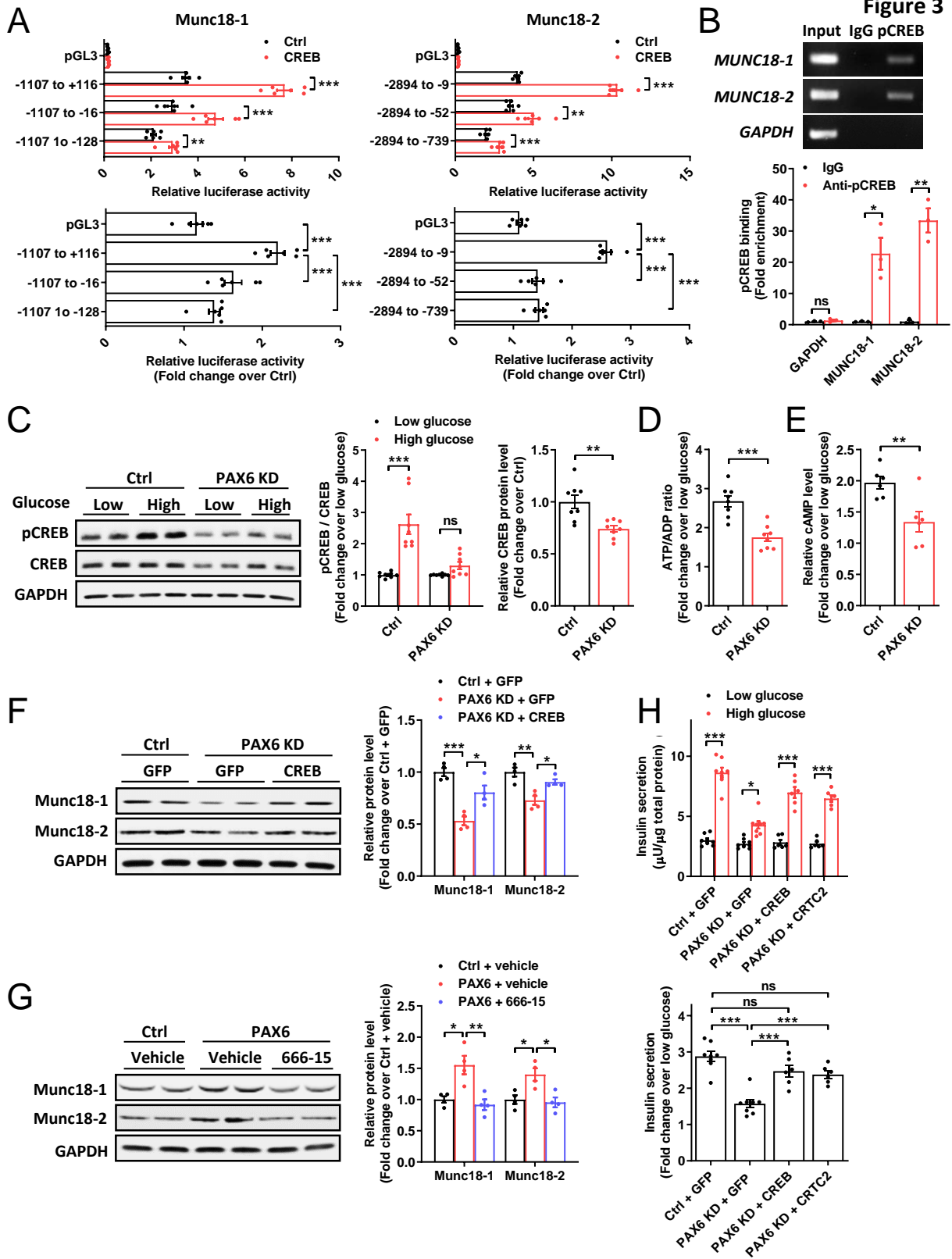


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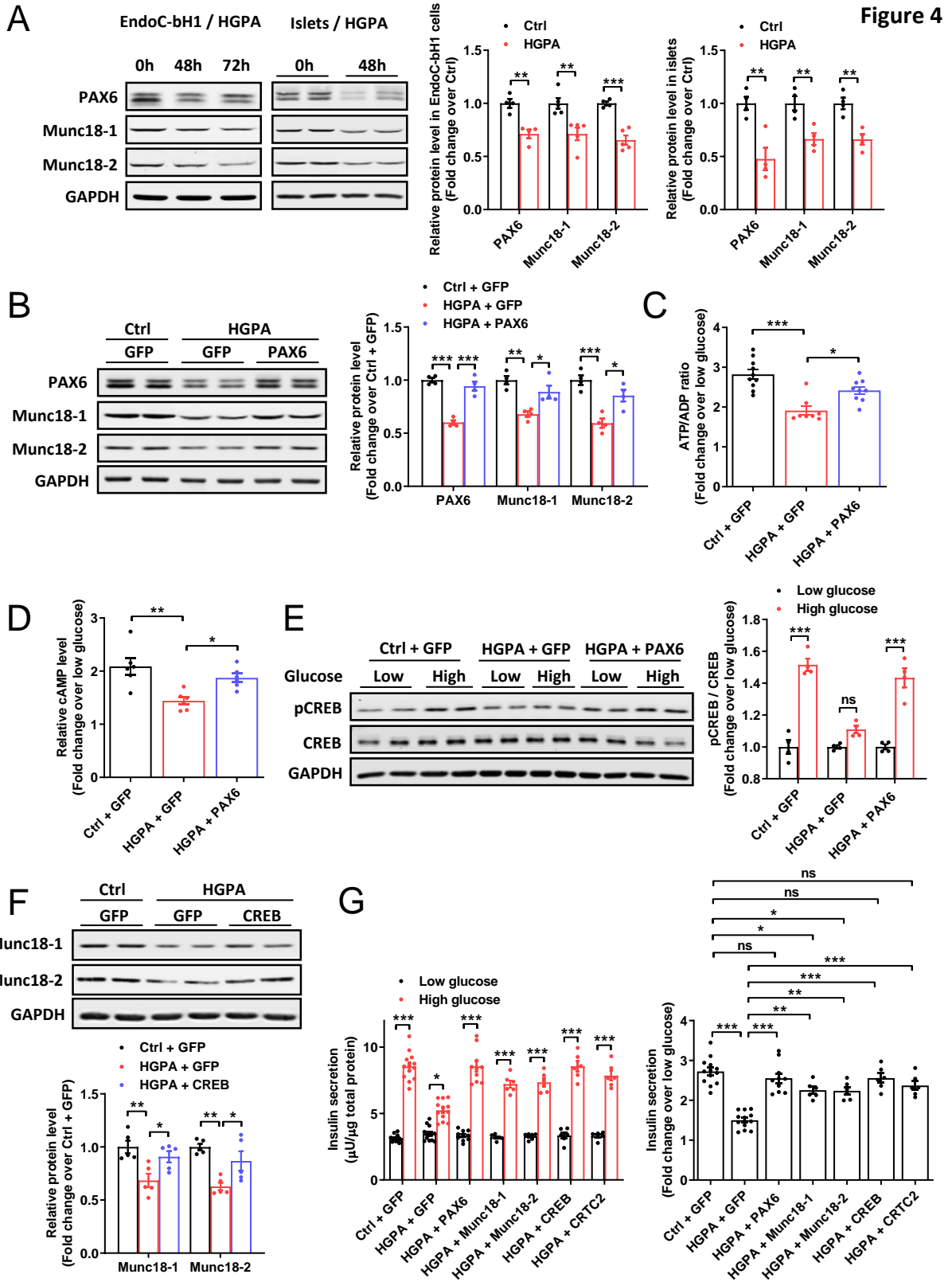
1 **Fig. 2.** PAX6 regulates GSIS through modulation of Munc18-1/2 expression. (A) Static insulin  
2 secretion of EndoC-βH1 cells was expressed as absolute amount (μU/μg total protein) and fold  
3 change ( $n=7-9$ ). (B) Dynamic insulin secretion of EndoC-βH1 cells in response to glucose (16.7  
4 mmol/l) and KCl (20 mmol/l) stimulation ( $n=5$ ). (C) Static insulin secretion of EndoC-βH1 cells  
5 was expressed as absolute amount (μU/μg total protein) and fold change ( $n=6-8$ ). (D) Dynamic  
6 insulin secretion of EndoC-βH1 cells in response to glucose (16.7 mmol/l) and KCl (20 mmol/l)  
7 stimulation ( $n=5$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ ; ns: non-significant. Data are means ±  
8 SEMs.



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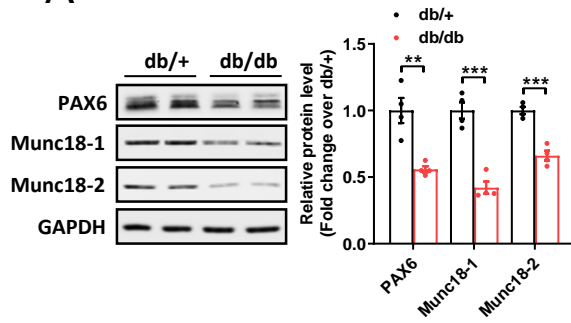
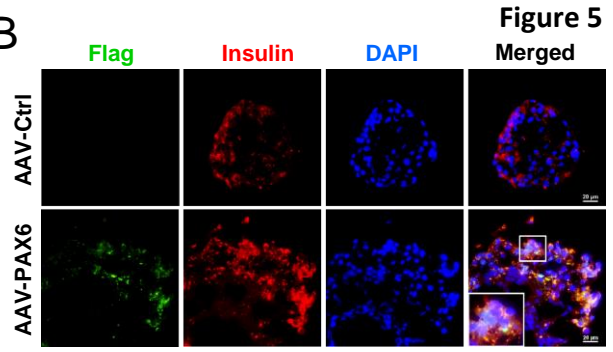
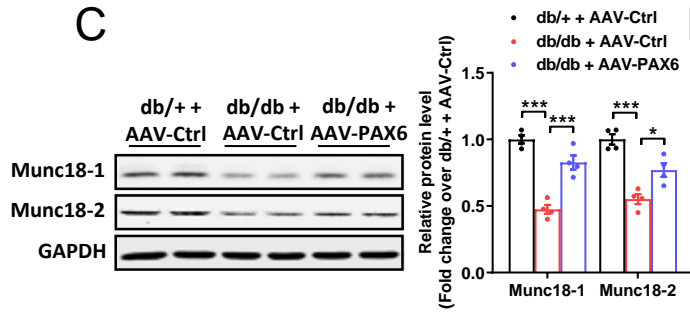
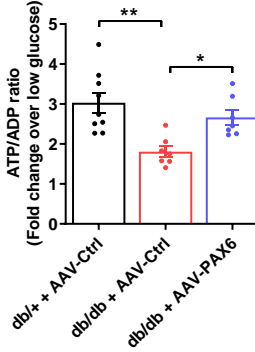
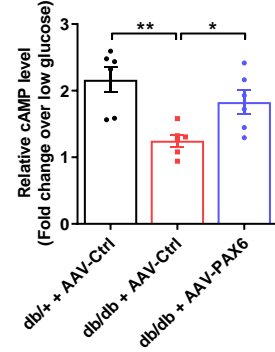
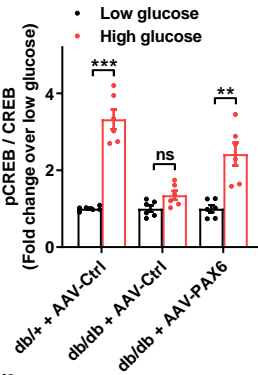
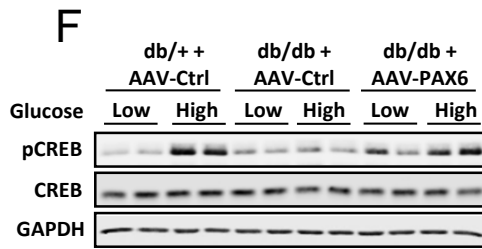
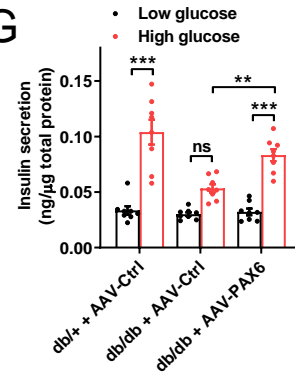
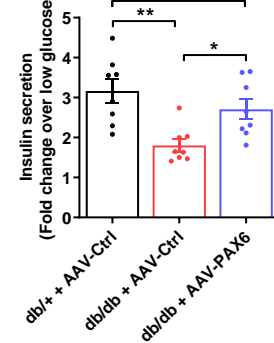
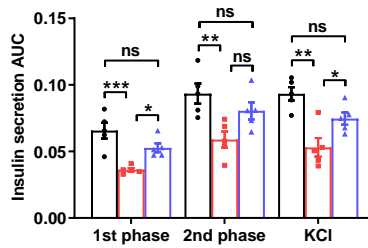
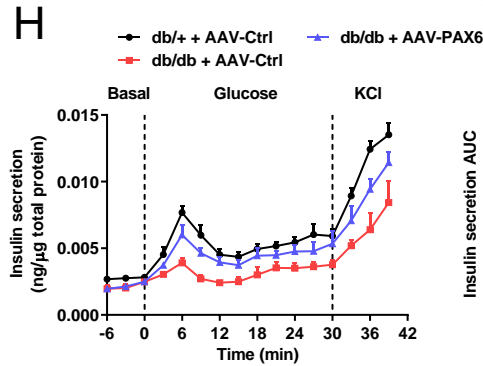
1 **Fig. 3.** PAX6 regulates Munc18-1/2 expression via modulation of CREB signaling. (A)  
2 Luciferase activity of human *MUNC18-1* and *MUNC18-2* promoters was measured in EndoC-  
3  $\beta$ H1 cells ( $n=6$ ). (B) Chromatin immunoprecipitation assay showing pull-down of *MUNC18-1*  
4 and *MUNC18-2* promoters by phospho-CREB antibody in EndoC- $\beta$ H1 cells ( $n=3$ ). (C)  
5 Phosphorylated and total CREB levels in EndoC- $\beta$ H1 cells with PAX6 knockdown after 15-min  
6 glucose (2.5 or 20 mmol/l) stimulation ( $n=8$ ). (D) ATP/ADP ratio ( $n=7-8$ ) and (E) cAMP level  
7 ( $n=6$ ) were measured in EndoC- $\beta$ H1 cells with PAX6 knockdown after 15-min glucose (2.5 or  
8 20 mmol/l) stimulation. (F) Protein expression of Munc18-1/2 in EndoC- $\beta$ H1 cells with PAX6  
9 knockdown and CREB overexpression ( $n=4$ ). (G) Protein expression of Munc18-1/2 in EndoC-  
10  $\beta$ H1 cells with PAX6 overexpression and treatment of CREB inhibitor 666-15 (40 nM) for 72 h  
11 ( $n=4$ ). (H) Static insulin secretion of EndoC- $\beta$ H1 cells was expressed as absolute amount ( $\mu$ U/ $\mu$ g  
12 total protein) and fold change ( $n=6$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ ; ns: non-significant. Data  
13 are means  $\pm$  SEMs.



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1 **Fig. 4.** Repressed PAX6/CREB/Munc18 axis in pancreatic beta-cells under diabetic condition  
2 leads to blunted insulin secretion. (A) Protein expression of PAX6 and Munc18-1/2 in EndoC-  
3  $\beta$ H1 cells and isolated mouse islets exposed to HGPA for the indicated periods of time ( $n=4-5$ ).  
4 (B) Protein expression of PAX6 and Munc18-1/2 in EndoC- $\beta$ H1 cells with PAX6 overexpression  
5 after 72-h exposure to normal or HGPA condition ( $n=4$ ). After 72-h exposure to normal or HGPA  
6 condition, (C) ATP/ADP ratio ( $n=8-10$ ), (D) cAMP level ( $n=6$ ), (E) phosphorylated and total  
7 CREB levels ( $n=4$ ) in EndoC- $\beta$ H1 cells with PAX6 overexpression were measured after 15-min  
8 glucose (2.5 or 20 mmol/l) stimulation. (F) Protein expression of Munc18-1/2 in EndoC- $\beta$ H1  
9 cells with CREB overexpression after 72-h exposure to normal or HGPA condition ( $n=5$ ). (G)  
10 Static insulin secretion of EndoC- $\beta$ H1 cells was expressed as absolute amount ( $\mu$ U/ $\mu$ g total  
11 protein) and fold change ( $n=6-11$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ ; ns: non-significant. Data  
12 are means  $\pm$  SEMs.

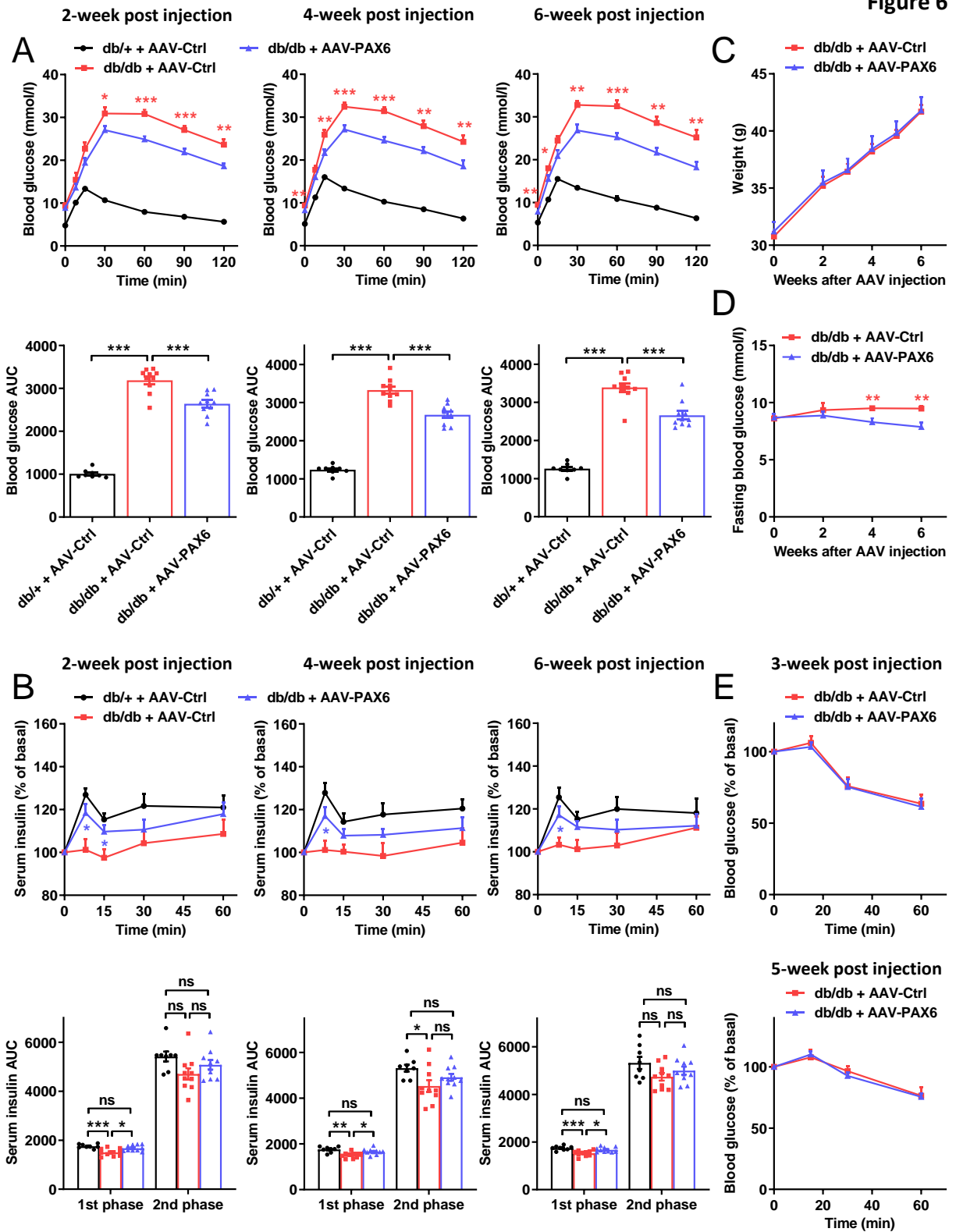
**A****B****Figure 5****C****D****E****F****G****H**

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1 **Fig. 5.** Beta-cell specific PAX6 overexpression preserves CREB/Munc18-1/2 signaling pathway  
2 and GSIS in *db/db* mouse islets. (A) Protein expression of PAX6 and Munc18-1/2 in isolated  
3 islets of *db/+* and *db/db* mice ( $n=4$ ). (B) Representative immunostaining of *db/db* mouse islets  
4 labeled for flag (green), insulin (red) and DAPI (blue) 6-week post AAV injection. Scale bar = 20  
5  $\mu\text{m}$ . (C) Protein expression of Munc18-1/2 in isolated islets of *db/+* and *db/db* mice with AAV-  
6 Ctrl or AAV-PAX6 injection ( $n=4$ ). (D) ATP/ADP ratio ( $n=7-9$ ) and (E) cAMP level ( $n=6$ ) in  
7 dispersed mouse islet cells after 15-min glucose (2.5 or 20 mmol/l) stimulation. (F)  
8 Phosphorylated and total CREB levels in isolated islets of *db/+* and *db/db* mice after 15-min  
9 glucose (2.5 or 20 mmol/l) stimulation ( $n=4$ ). (G) Static insulin secretion of isolated islets of  
10 *db/+* and *db/db* mice was expressed as absolute amount (ng/ $\mu\text{g}$  total protein) and fold change  
11 ( $n=8$ ). (H) Dynamic insulin secretion of isolated islets in response to glucose (16.7 mmol/l) and  
12 KCl (20 mmol/l) stimulation ( $n=5$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ ; ns: non-significant. Data  
13 are means  $\pm$  SEMs.

Figure 6



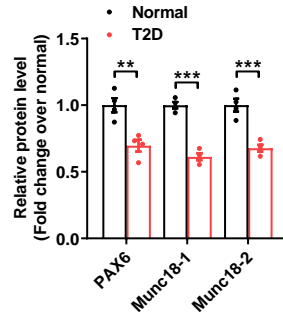
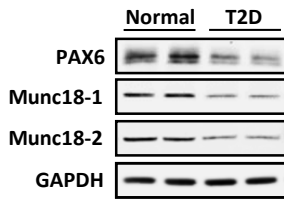
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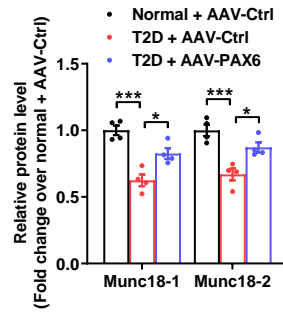
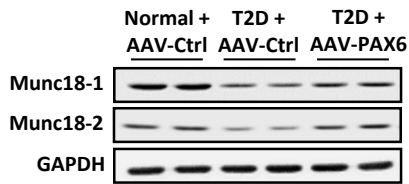


1 **Fig. 6.** Beta-cell specific PAX6 overexpression improves glucose homeostasis in *db/db* mice. (A)  
2 Glucose tolerance by IPGTT was performed 2-, 4- and 6-week post AAV injection with glucose  
3 profiles calculated as AUC. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$  versus *db/db* + AAV-PAX6. (B)  
4 Serum insulin levels during IPGTT were expressed as % of basal. \* $P < 0.05$  versus *db/db* + AAV-  
5 Ctrl. (C) Weight and (D) fasting blood glucose at the indicated time points after AAV injection.  
6 (E) ITT was performed 3- and 5-week post AAV injection and blood glucose levels were  
7 expressed as % of basal level. \* $P < 0.05$ ; \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ ; ns: non-significant ( $n=8-10$ ).  
8 Data are means  $\pm$  SEMs.

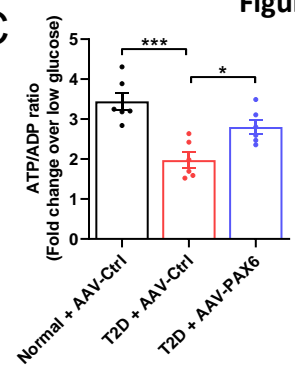
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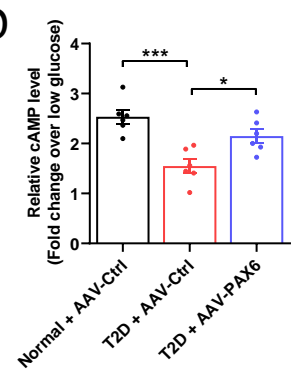
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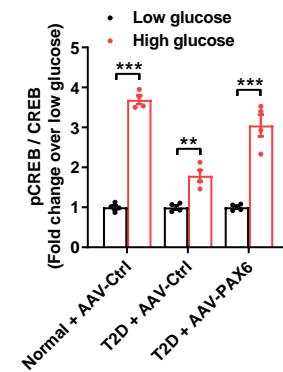
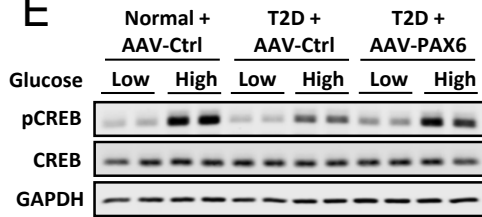
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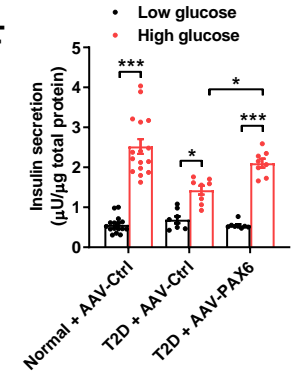
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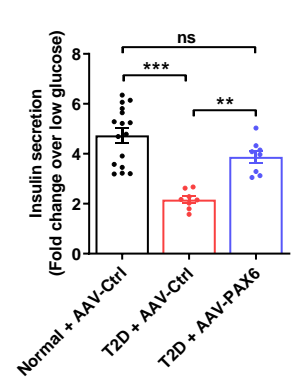
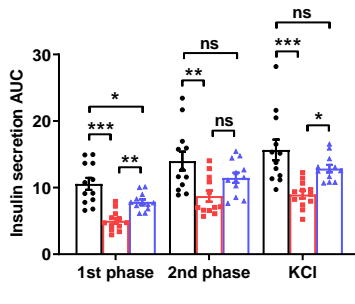
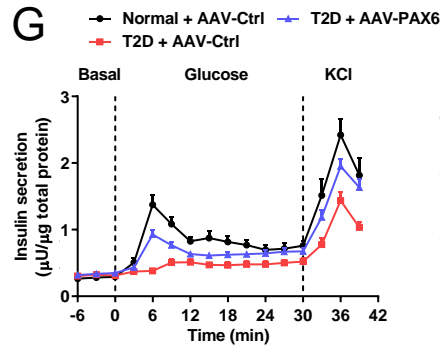
**E**



**F**



**G**



**Figure 7**

1 **Fig. 7.** Beta-cell specific PAX6 overexpression restores CREB/Munc18-1/2 signaling pathway  
2 and GSIS in T2D human islets. (A) Protein expression of PAX6 and Munc18-1/2 in normal and  
3 T2D human islets ( $n=4$ ). (B) Protein expression of Munc18-1/2 in human islets with AAV-Ctrl or  
4 AAV-PAX6 transduction ( $n=4$ ). (C) ATP/ADP ratio ( $n=6$ ) and (D) cAMP level ( $n=6$ ) in dispersed  
5 human islet cells after 15-min glucose (2.5 or 20 mmol/l) stimulation. (E) Phosphorylated and  
6 total CREB levels in human islets after 15-min glucose (2.5 or 20 mmol/l) stimulation ( $n=4$ ). (F)  
7 Static insulin secretion of human islets was expressed as absolute amount ( $\mu\text{U}/\mu\text{g}$  total protein)  
8 and fold change ( $n=8-16$ ). (G) Dynamic insulin secretion of human islets in response to glucose  
9 (16.7 mmol/l) and KCl (20 mmol/l) stimulation ( $n=12$ ).  $*P<0.05$ ;  $**P<0.01$ ;  $***P<0.001$ ; ns:  
10 non-significant. Data are means  $\pm$  SEMs.

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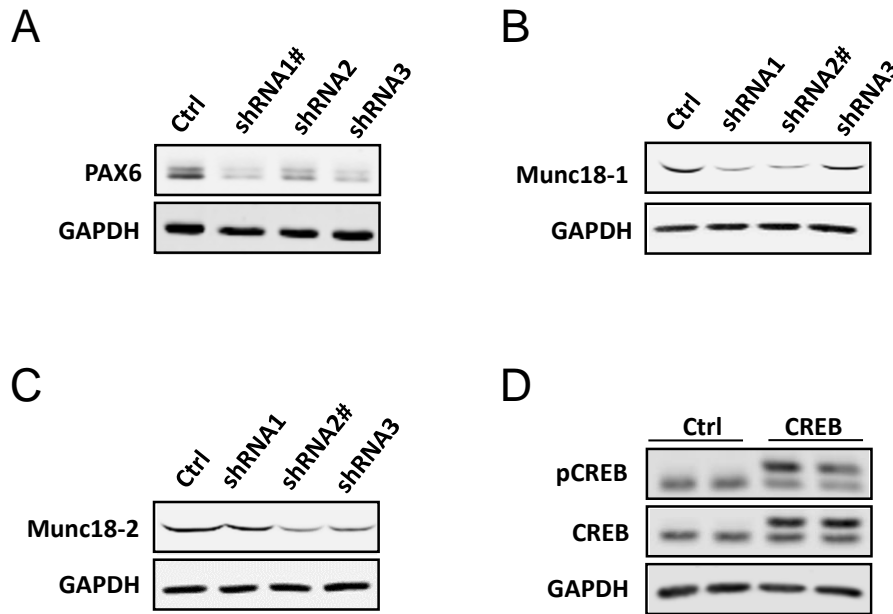
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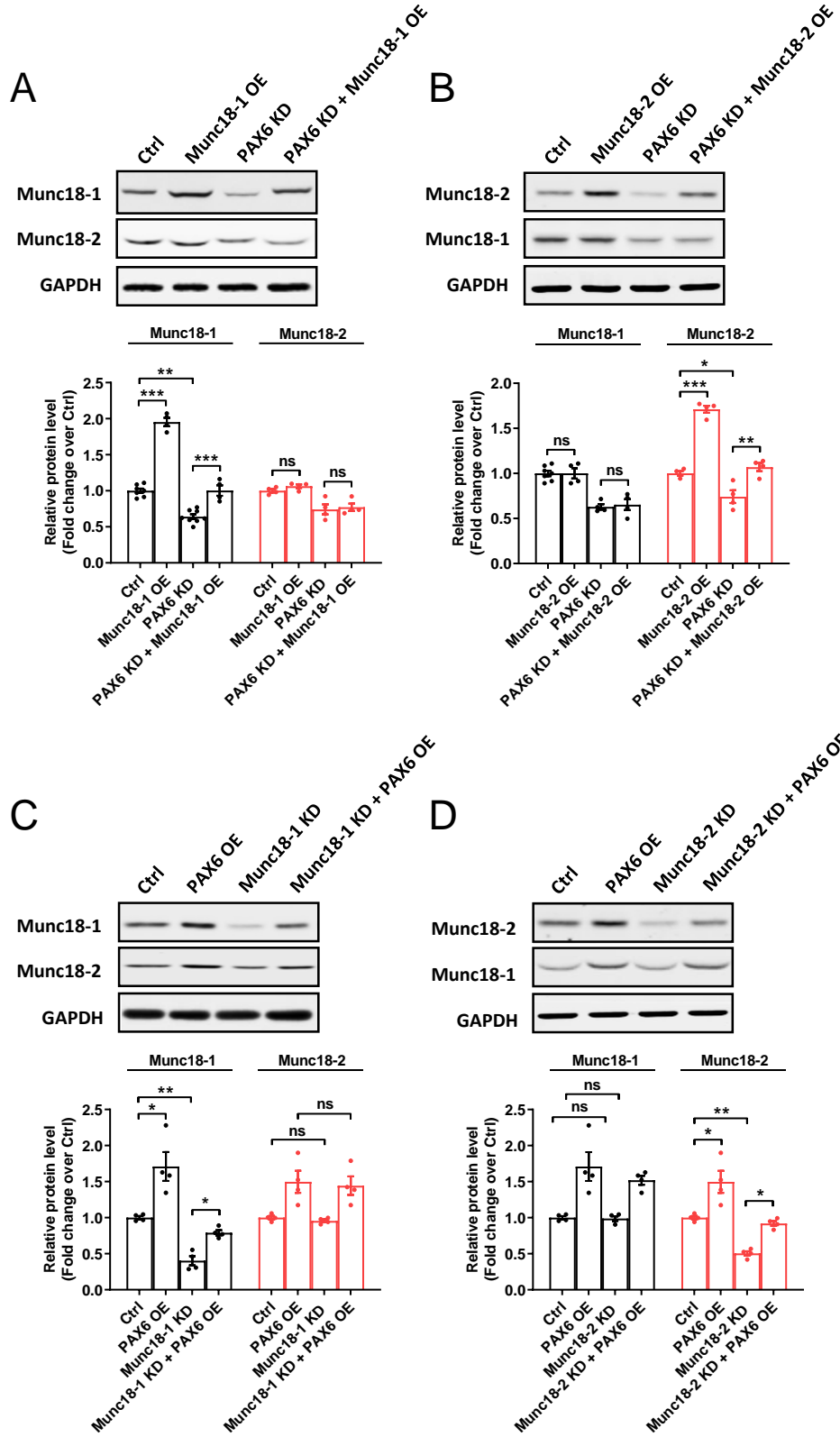
1 **Supplementary Materials**

**Figure S1**



2

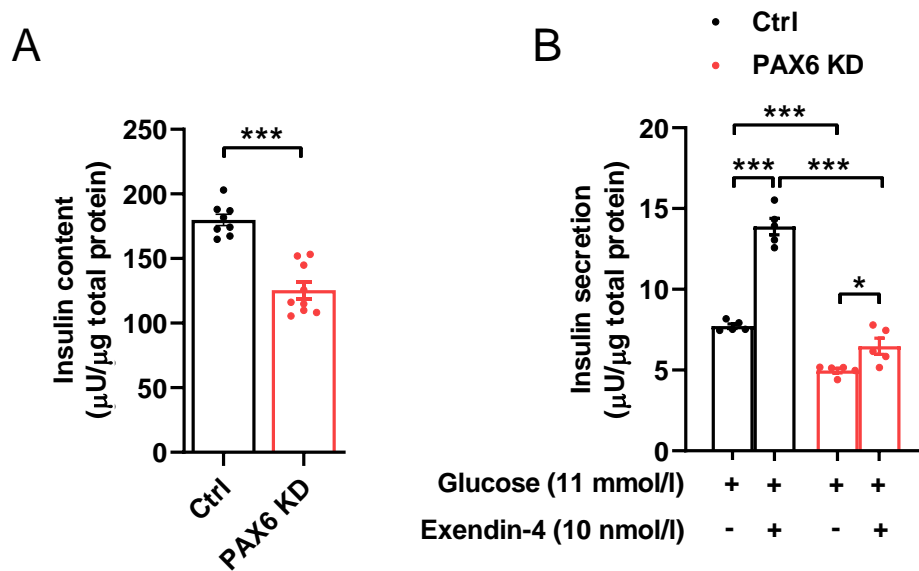
3 **Fig. S1.** Western blotting showing knockdown efficiency of three shRNA constructs targeting  
4 human (A) PAX6, (B) Munc18-1 and (C) Munc18-2 in EndoC-βH1 cells. Hashtags indicate the  
5 shRNA constructs chosen for downstream experiments. (D) Western blotting showing the  
6 increase in phosphorylated and total CREB in EndoC-βH1 cells with CREB overexpression.



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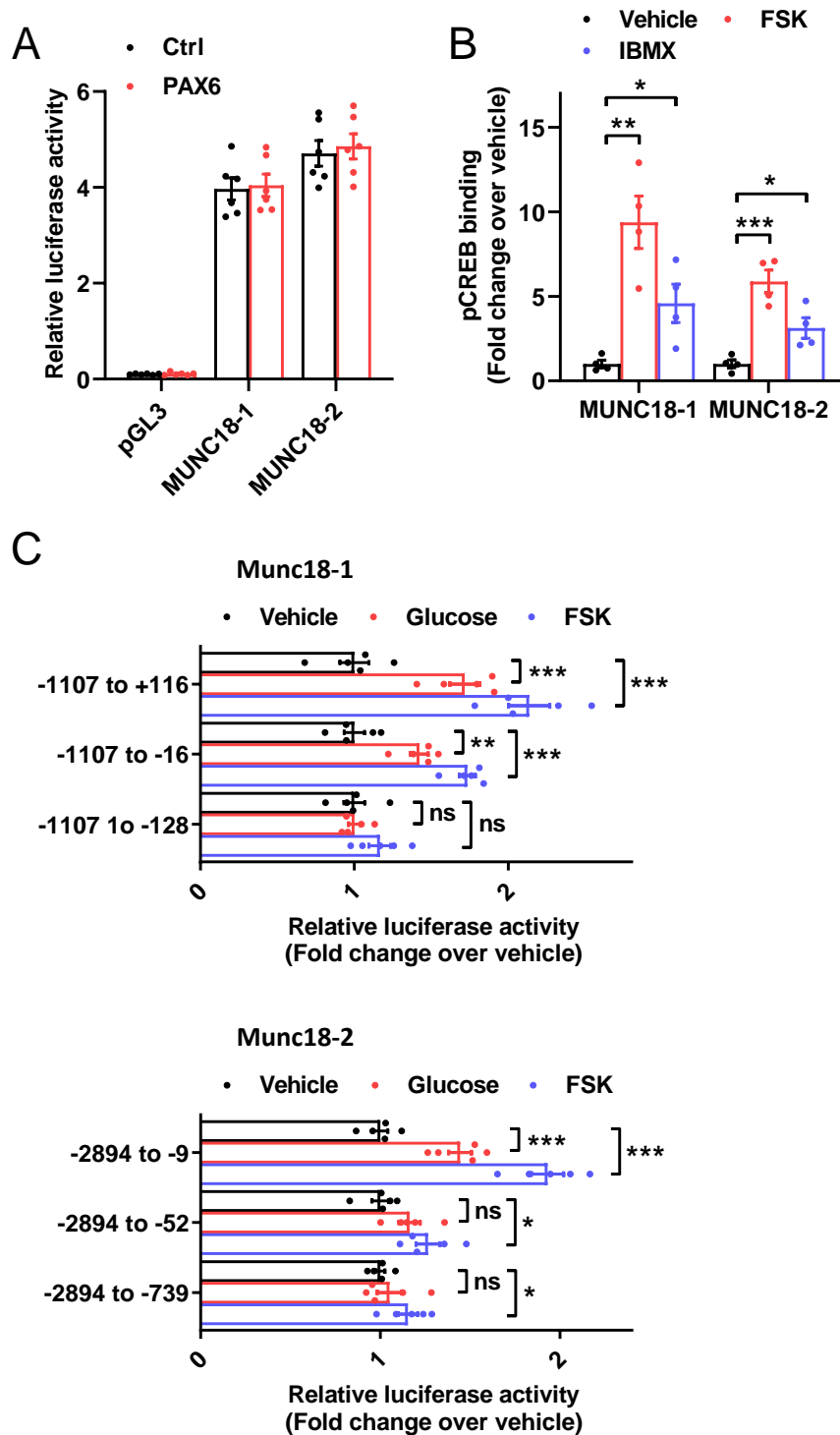
1 **Fig. S2.** Western blotting showing Munc18-1/2 expression in EndoC-βH1 cells under different  
2 conditions: (A) cells with Munc18-1 overexpression, PAX6 knockdown and PAX6 knockdown  
3 with Munc18-1 overexpression; (B) cells with Munc18-2 overexpression, PAX6 knockdown and  
4 PAX6 knockdown with Munc18-2 overexpression; (C) cells with PAX6 overexpression,  
5 Munc18-1 knockdown and Munc18-1 knockdown with PAX6 overexpression; (D) cells with  
6 PAX6 overexpression, Munc18-2 knockdown and Munc18-2 knockdown with PAX6  
7 overexpression ( $n=4-8$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ ; ns: non-significant. Data are means  $\pm$   
8 SEMs.



1

2 **Fig. S3.** Effects of PAX6 knockdown on insulin content and insulin secretion in EndoC-βH1  
 3 cells. (A) Insulin content was expressed as absolute amount (μU/μg total protein) ( $n=8-9$ ). (B)  
 4 Control or PAX6 knockdown cells were stimulated with exendin-4 (10 nmol/l) in the presence of  
 5 11 mmol/l glucose for 1 h. Insulin secretion was expressed as absolute amount (μU/μg total  
 6 protein) ( $n=5$ ). \* $P<0.05$ ; \*\*\* $P<0.001$ . Data are means  $\pm$  SEMs.

Figure S4



1

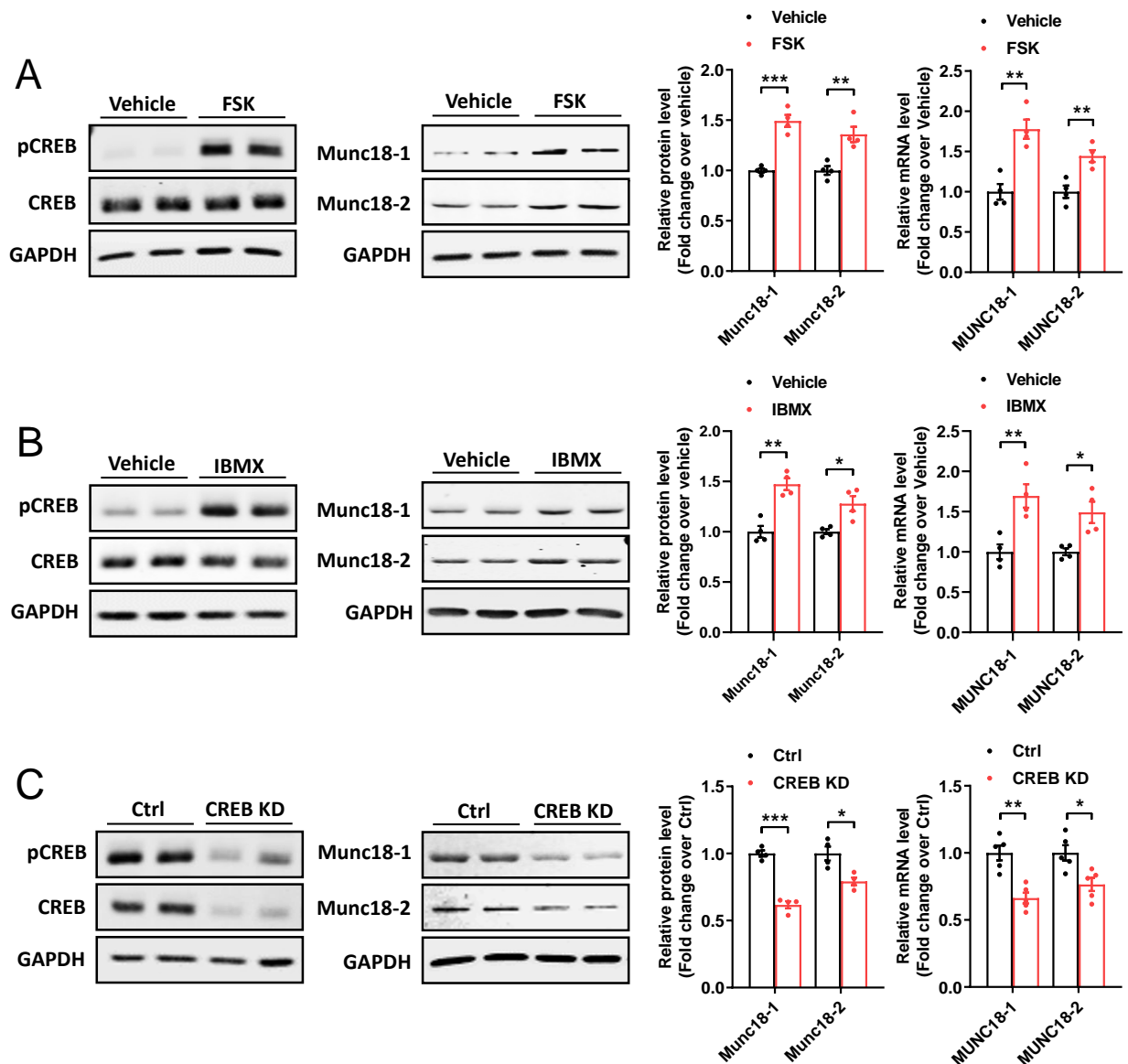
2 **Fig. S4.** (A) Luciferase activity of human *MUNC18-1* and *MUNC18-2* promoters was measured  
 3 in EndoC- $\beta$ H1 cells with or without PAX6 overexpression ( $n=6$ ). (B) Chromatin



1 immunoprecipitation assay showing increased pCREB binding onto *MUNC18-1* and *MUNC18-2*  
2 promoters after 2 h treatment of FSK (10  $\mu\text{mol/l}$ ) or IBMX (300  $\mu\text{mol/l}$ ) in EndoC- $\beta$ H1 cells  
3 ( $n=4$ ). (C) Luciferase reporter assay showing enhanced activity of human *MUNC18-1* and  
4 *MUNC18-2* promoters after 16 h treatment of glucose (20  $\text{mmol/l}$ ) or FSK (10  $\mu\text{mol/l}$ ) in EndoC-  
5  $\beta$ H1 cells ( $n=5$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ ; ns: non-significant. Data are means  $\pm$  SEMs.

6

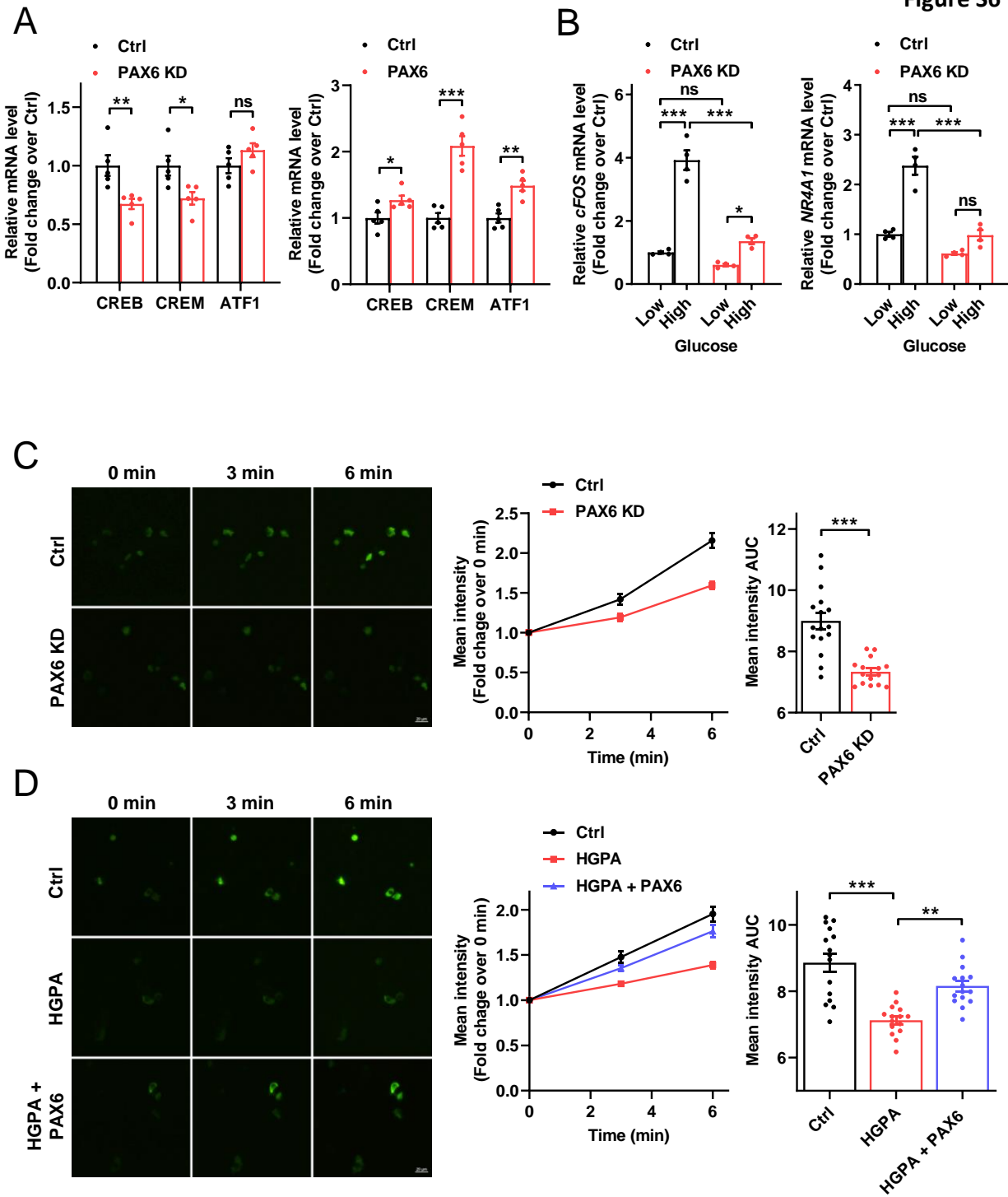
Figure S5



1

2 **Fig. S5.** Phosphorylation of CREB, Munc18-1 and Munc18-2 protein and mRNA levels in  
 3 EndoC-βH1 cells after 24-h treatment of (A) FSK (10 μmol/l) or (B) IBMX (300 μmol/l) (n=4).  
 4 (C) Phosphorylation of CREB, Munc18-1 and Munc18-2 protein and mRNA levels in control  
 5 and CREB knockdown cells (n=4-5). \*P<0.05; \*\*P<0.01; \*\*\*P<0.001. Data are means ± SEMs.

Figure S6

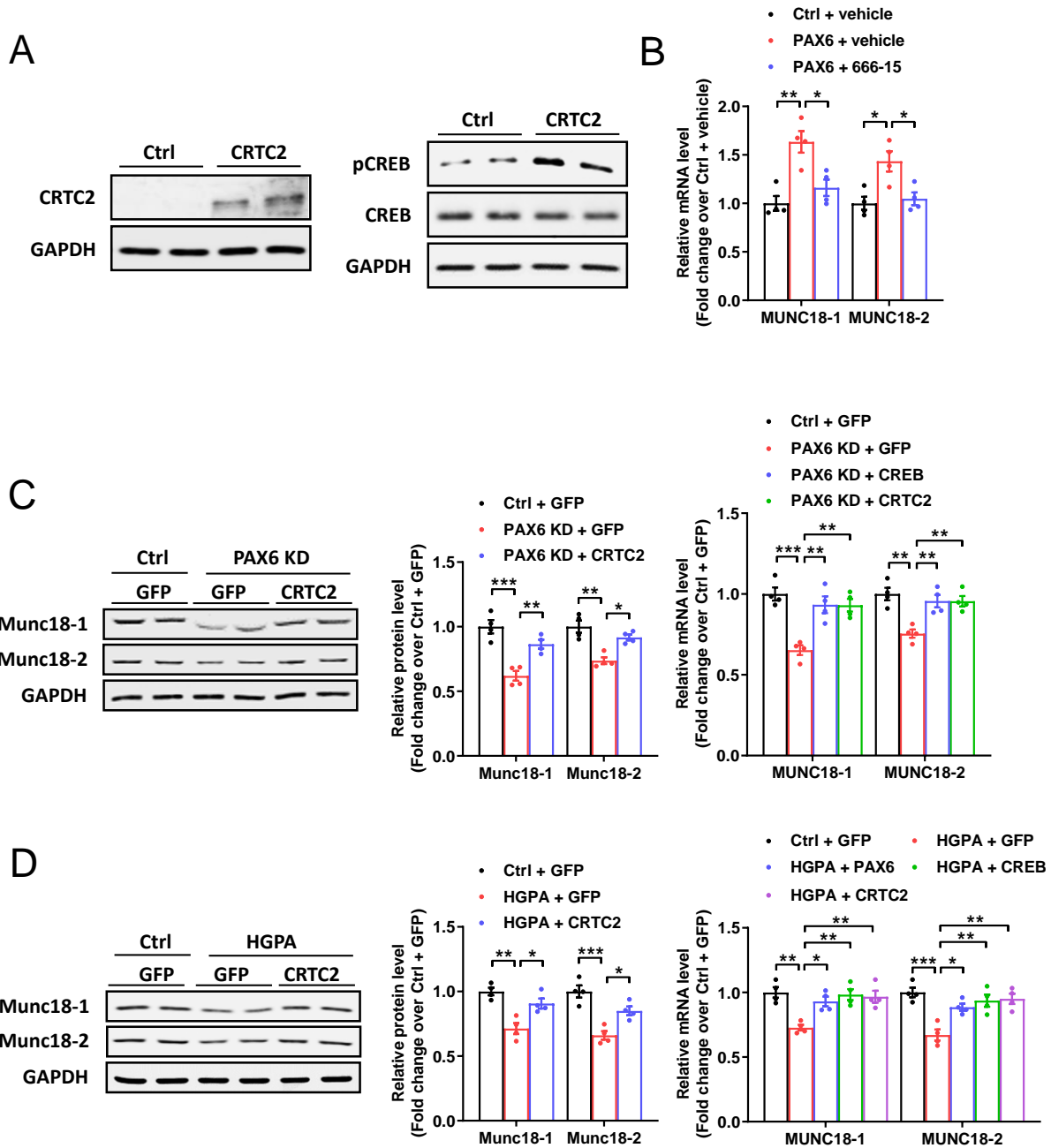


1

2 **Fig. S6.** (A) mRNA levels of *CREB*, *CREM* and *ATF1* in EndoC-βH1 cells with PAX6

3 knockdown and overexpression ( $n=5$ ). (B) Attenuated glucose-induced expression of *cFOS* and

1 *NR4A1* mRNA levels in PAX6 knockdown cells. EndoC-βH1 cells were stimulated with low (2.5  
2 mmol/l) or high (20 mmol/l) glucose for 1 h (*n*=4). Measurement of Ca<sup>2+</sup> influx by live-cell  
3 imaging of EndoC-βH1 cells under (C) control and PAX6 knockdown conditions and under (D)  
4 control and HGPA conditions with or without PAX6 overexpression. Representative images of  
5 cells after 3- and 6-min glucose (20 mmol/l) stimulation are shown. Scale bar = 20 μm. \**P*<0.05;  
6 \*\**P*<0.01; \*\*\**P*<0.001; ns: non-significant. Data are means ± SEMs.



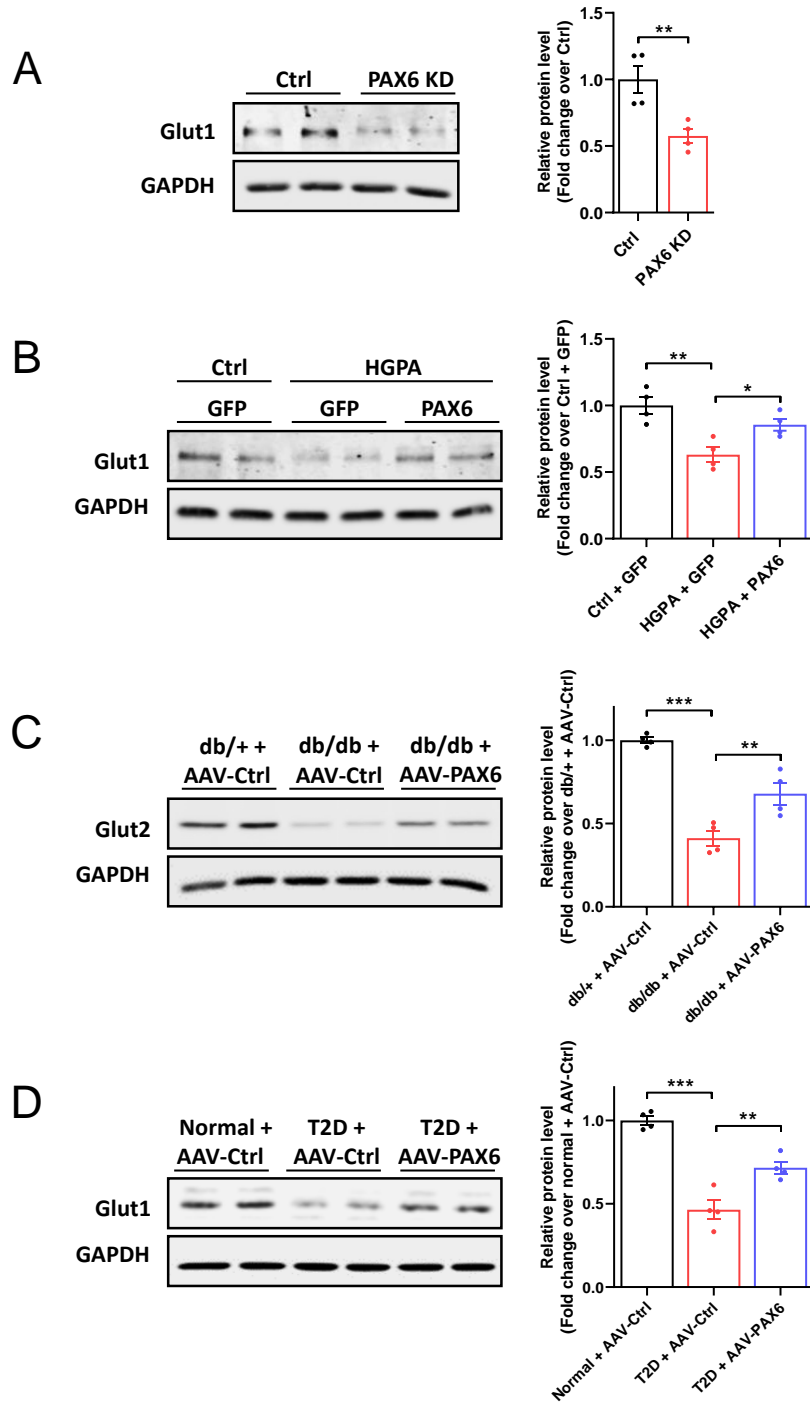
1

2 **Fig. S7.** (A) Western blotting showing overexpression of CRTC2 and increased phosphorylated

3 CREB in EndoC-βH1 cells. (B) mRNA levels of *MUNC18-1* and *MUNC18-2* in EndoC-βH1

1 cells with PAX6 overexpression and treatment of CREB inhibitor 666-15 (40 nM) for 72 h ( $n=4$ ).  
2 (C) Protein and mRNA expression of Munc18-1/2 in EndoC- $\beta$ H1 cells with PAX6 knockdown  
3 and CREB/ CRTC2 overexpression ( $n=4$ ). (D) Protein and mRNA expression of Munc18-1/2 in  
4 EndoC- $\beta$ H1 cells with CREB/ CRTC2 overexpression after 72-h exposure to normal or HGPA  
5 condition ( $n=4$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ . Data are means  $\pm$  SEMs.

6



1

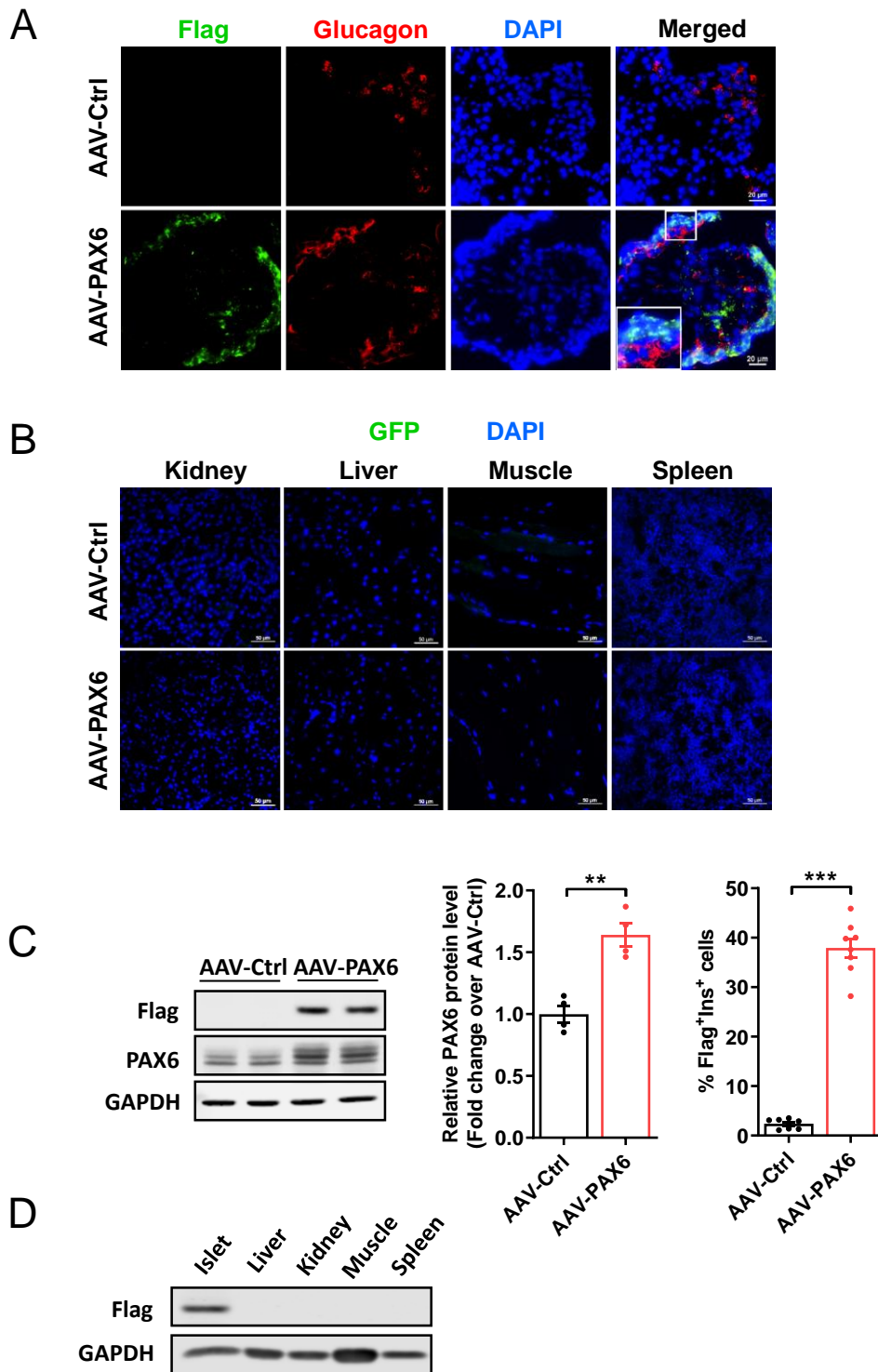
2 **Fig. S8.** Western blotting showing Glut1 and Glut2 protein expression. (A) Protein expression of  
 3 Glut1 in EndoC-βH1 cells with PAX6 knockdown. (B) Protein expression of Glut1 in EndoC-

- 1  $\beta$ H1 cells with PAX6 overexpression after 72-h exposure to normal or HGPA condition. (C)
- 2 Protein expression of Glut2 in isolated islets of *db/+* and *db/db* mice 6-week post AAV-Ctrl or
- 3 AAV-PAX6 injection. (D) Protein expression of Glut1 in human islets with AAV-Ctrl or AAV-
- 4 PAX6 transduction ( $n=4$ ). \* $P<0.05$ ; \*\* $P<0.01$ ; \*\*\* $P<0.001$ . Data are means  $\pm$  SEMs.



Mouse islets

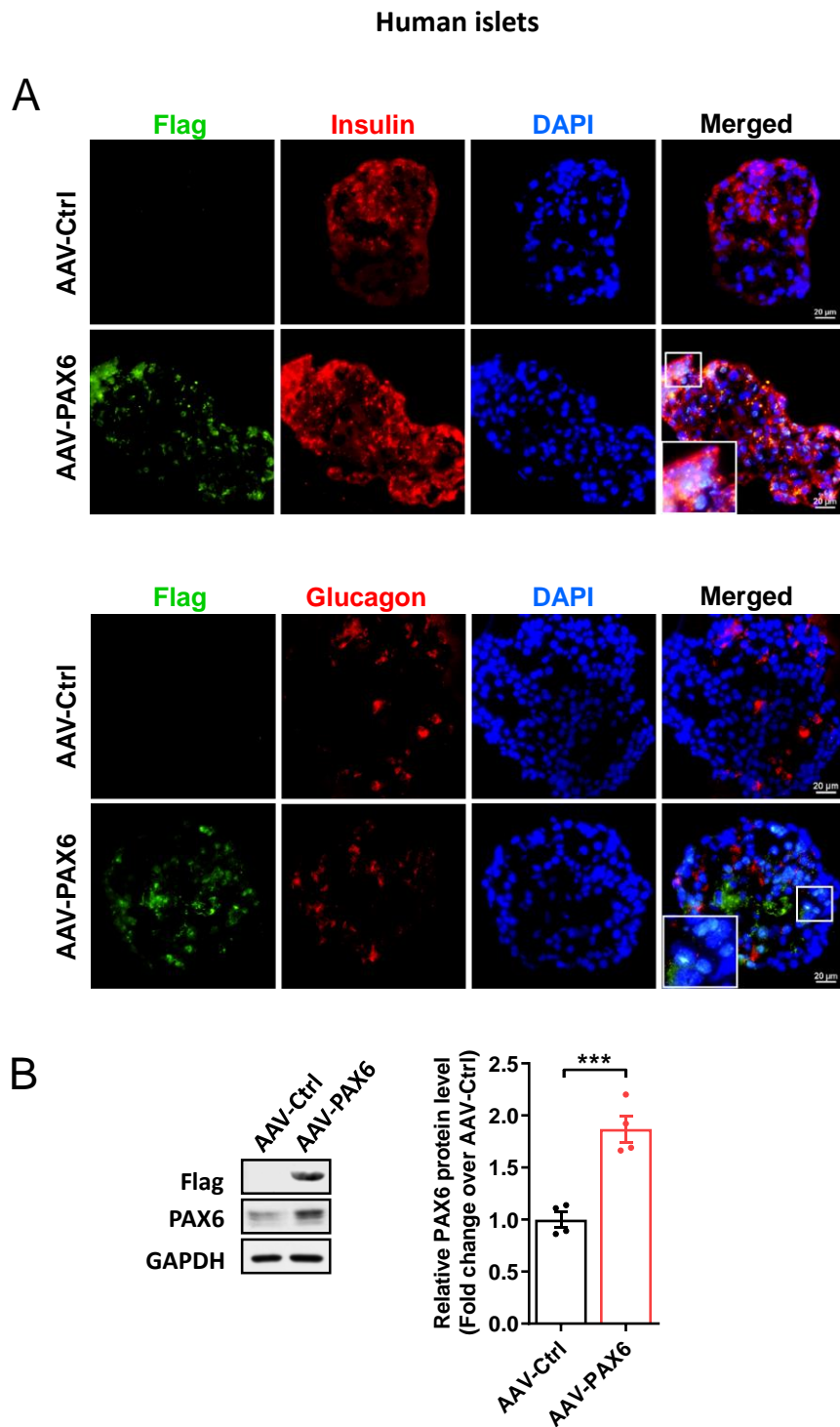
Figure S9



1

2

1 **Fig. S9.** (A) Representative immunostaining of *db/db* mouse islets labeled for flag (green),  
2 glucagon (red) and DAPI (blue). Scale bar = 20  $\mu$ m. (B) Representative immunostaining of  
3 *db/db* mouse kidney, liver, muscle and spleen labeled for GFP (green) and DAPI (blue). Scale  
4 bar = 50  $\mu$ m. Tissues were harvested in *db/db* mice 6-week post AAV injection. (C) Western  
5 blotting and quantification showing the expression of exogenous PAX6 in *db/db* mouse islets  
6 with AAV-PAX6 injection. (D) Western blotting showing the absence of flag tag in other *db/db*  
7 mouse organs. \*\* $P < 0.01$ ; \*\*\* $P < 0.001$ . Data are means  $\pm$  SEMs.



1

2 **Fig. S10.** (A) Representative immunostaining of human T2D islets labeled for flag (green),  
 3 insulin or glucagon (red) and DAPI (blue) with AAV transduction. Scale bar = 20  $\mu$ m. (B)

1 Western blotting and quantification showing the expression of exogenous PAX6 in T2D islets  
2 with AAV-PAX6 transduction. \*\*\* $P < 0.001$ . Data are means  $\pm$  SEMs.

3

**Table S1**

shRNA for gene knockdown	shRNA sequence	
<i>PAX6</i>	AATTCTGGGCAGGTATTACGA	
<i>MUNC18-1</i>	TATGATCTGCTGCCTATCGAA	
<i>MUNC18-2</i>	ATATCTTCTTCACCGACACCT	
<i>CREB1</i>	ACAGCACCCACTAGCACTATT	
Non-targeting control	CAACAAGATGAAGAGCACCAA	
Primers for qPCR	Forward primer (5'-3')	Reverse primer (5'-3')
<i>b-actin</i>	ACAGAGCCTCGCCTTTGCC	GATATCATCATCCATGGTGAGCTGG
<i>PAX6</i>	AACGATAACATACCAAGCGTGT	GGTCTGCCCGTTCAACATC
<i>MUNC18-1</i>	AAAGCTGTTGTCGGAGAGAAG	CACAATCGTTATGCCCTCGG
<i>MUNC18-2</i>	ATTCTGAGCGGAGTTATTCGGA	CCGCCGTTTGTGATGCTTTC
<i>MUNC18-3</i>	AGCAACAGTGTGGTACTGCTGC	TCTGACAGGTTACGGTCTTAA
<i>MUNC13-1</i>	CGATGATCGGGAACCTCTTG	AGCTCCTGGCTGCATTGAA
<i>SNAP25</i>	ATGGATGAAAACCTAGAGCAGG	ACACTTAACCACTTCCCAGC
<i>SYB2</i>	CTCAAGCGCAAATACTGGTGG	TGATGGCGCAAATCACTCCC
<i>STX1A</i>	TAAAGAGCATCGAGCAGTCCA	GACATGACCTCCAACTTTCT
<i>cFOS</i>	GGGGCAAGGTGGAACAGTTAT	AGGTTGGCAATCTCGGTCTG
<i>NR4A1</i>	AGGGCTGCAAGGGCTTCT	GGCAGATGTAAGGGCTTTT
<i>CREB1</i>	GTGACGGAGGAGCTTGTACC	GCTGCTGGCATAGATACTGG
<i>CREM</i>	ACAATGAGCAAATGTGCAAGGA	TCTGCAATTGCTGCTACCTGA
<i>ATF1</i>	GGTAGCTTAGGACAGTTGGC	CTGATAAAGATGATACCTGTTGAGC
Primers for luciferase vectors construction	Forward primer (5'-3')	Reverse primer (5'-3')
<i>MUNC18-1</i> (-1107 to +116)	ACTGACGCTAGCCTCCTATTAACAAAGAAA	ACTGACAAGCTTGGCGCTGCGGAGTCTCC
<i>MUNC18-1</i> (-1107 to -16)	ACTGACGCTAGCCTCCTATTAACAAAGAAA	ACTGACAAGCTTCGCGGACTGGGCGCGGGG
<i>MUNC18-1</i> (-1107 to -128)	ACTGACGCTAGCCTCCTATTAACAAAGAAA	ACTGACAAGCTTCCCCTGCTCCGCCAA
<i>MUNC18-2</i> (-2894 to -9)	ACTGACGCTAGCCACTGTGGCTTTCCTCA	ACTGACAAGCTTCTCCGGGTGTGTCCCAA
<i>MUNC18-2</i> (-2894 to -52)	ACTGACGCTAGCCACTGTGGCTTTCCTCA	ACTGACAAGCTTCCCACGTGGACCCGCC
<i>MUNC18-2</i> (-2894 to -739)	ACTGACGCTAGCCACTGTGGCTTTCCTCA	ACTGACAAGCTTCCCCTCAAATGCTTGCAA
Primers for ChIP assay	Forward primer (5'-3')	Reverse primer (5'-3')
<i>MUNC18-1</i>	GGGTTTCAGGCAGATGGA	TACCCAGGCCTCCTTTTC
<i>MUNC18-2</i>	CTCCCATCTGTGAGCGTGT	AGGCCAGGAAGTTGAGTC
<i>GAPDH</i>	TTGACTCACCTGCCCTCAAT	CCTCCCAAAGGCCTCCT

- 1
- 2 **Table S1.** Oligonucleotide sequences used in shRNA vectors construction, qPCR, luciferase
- 3 vectors construction and ChIP assay.

**Table S2**

<u>Species</u>	<u>Gene</u>	<u>Position</u>	<u>Sequence</u>
Human	<i>MUNC18-1</i>	-127 to -120	TGACGCCA
		-15 to -11	CGTCA
	<i>MUNC18-2</i>	-738 to -734	CGTCA
		-51 to -44	TGACGCGC
Mouse	<i>Munc18-1</i>	-195 to -191	TGACG
		-189 to -185	TGACG
		-83 to -79	CGTCA
	<i>Munc18-2</i>	-1296 to -1292	CGTCA
		-45 to -38	TGACGCGT

1

2 **Table S2.** Sequences and positions of putative CREs in promoters of Munc18-1 and Munc18-2

3 in both human and mouse.

**Table S3**

<u>Donor ID</u>	<u>Age</u>	<u>Gender</u>	<u>Weight (lbs)</u>	<u>BMI</u>	<u>HbA1c (%)</u>
Normal 1	23	F	157	24.5	4.9
Normal 2	41	M	218	35.12	5.8
Normal 3	51	M	195	31.45	5.2
Normal 4	55	M	160	23.63	4.7
T2D 1	55	F	164	29.9	7.4
T2D 2	63	M	234	31.69	6.9
T2D 3	52	M	158	24.75	7.6
T2D 4	59	M	168	25.54	7.3

4

5 **Table S3.** Human islet donor information.