

**Glucose-dependent downregulation of glucagon gene expression
mediated by selective interactions between ALX3 and PAX6 in mouse
alpha cells**

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Abstract

Aims/hypothesis The stimulation of glucagon secretion in response to decreased glucose levels has been studied extensively. In contrast, little is known about the regulation of glucagon gene expression in response to fluctuations in glucose concentration. PAX6 is a key transcription factor that regulates the glucagon promoter by binding to the G1 and G3 elements. Here, we investigated the role of the transcription factor ALX3 as a glucose-dependent modulator of PAX6 activity in alpha cells.

Methods Experiments were performed in wild type or *Alx3*-deficient islets and alphaTC1 cells. We used chromatin immunoprecipitations and electrophoretic mobility shift assays for DNA binding, immunoprecipitations and pull-down assays for protein interactions, transfected cells for promoter activity, and small interfering RNA and quantitative RT-PCR for gene expression.

Results Elevations in the concentration of glucose resulted in stimulated expression of *Alx3* and decreased glucagon gene expression in wild type islets. In ALX3-deficient islets, basal glucagon levels were non responsive to changes in glucose concentration. In basal conditions ALX3 bound to the glucagon promoter at G3, but not at G1. ALX3 could form heterodimers with PAX6 that are permissive for binding to G3 but not to G1. Thus, increasing the levels of ALX3 in response to glucose resulted in the sequestration of PAX6 by ALX3 for binding to G1, thus reducing glucagon promoter activation and glucagon gene expression.

Conclusions/Interpretation Glucose-stimulated expression of ALX3 in alpha cells provides a regulatory mechanism for the downregulation of glucagon gene expression by interfering with PAX6-mediated transactivation on the glucagon G1 promoter element.

Keywords Glucagon, Transcription, Homeodomain, Paired domain, Alpha cells, Glucose-dependent gene expression.

Abbreviations

EMSA Electrophoretic mobility shift assay

ChIP Chromatin immunoprecipitation

HBSS Hanks' balanced salt solution

GST Glutathione S-transferase

RNAi RNA interference

Introduction

Beta cell dysfunction, accompanied by reduced capacity of target tissues to respond to insulin, has been largely accepted as the main cause of chronic hyperglycaemia characteristic of diabetes [1, 2]. In recent years, however, dysfunctional glucagon hypersecretion contributing to high blood glucose has been recognized as an additional and important aetiopathogenic factor [3, 4]. Since hormonal stores depleted after secretion must be replenished by increasing glucagon gene expression and biosynthesis in a coordinated manner [5], the elucidation of the mechanisms that link glucagon gene transcription and glucagon secretion in response to fluctuations in glucose concentrations are of great importance for the understanding of the aetiopathogenic mechanisms of diabetes.

The stimulated secretion of glucagon from alpha cells in response to decreased blood glucose levels has been studied extensively [6-8]. In contrast, the study of glucose-regulated glucagon gene transcription has received relatively less attention and the mechanisms involved remain unclear [9-12]. Glucagon gene transcription in alpha cells is regulated primarily by at least four promoter elements, termed G1-G4 [13-15], that are recognized by several transcription factors including PAX6 [16-20]. PAX6 is particularly important because it is essential for the differentiation of alpha cells during development [21], and because it regulates the expression of the glucagon gene both directly [16, 22-24] and indirectly, acting on genes encoding transcription factors that in turn act on the glucagon promoter [25]. In addition, PAX6 regulates the expression of genes involved in prohormone processing [26] and glucagon secretion [27]. Therefore, PAX6 co-ordinately links glucagon production and secretion in alpha cells. However, it is unknown whether PAX6 or other G-element-binding transcription factors are involved in glucose-dependent regulation of glucagon gene expression.

In this study, we show that ALX3, an *aristaless*-type homeodomain transcription factor expressed in islets cells previously found to be important for glucose homeostasis [28, 29], dynamically regulates glucose-dependent glucagon gene expression by engaging in protein-protein interactions with PAX6 that result in reduced PAX6 accessibility to the glucagon promoter in the presence of increased glucose concentrations.

Methods

Mice. *Alx3* mutant mice [30] were maintained in our animal facilities and genotyped as described [31]. Experiments were performed with 12-16 week old male mice. Experimental protocols were approved by the institutional bioethics committee and meet the requirements of current Spanish and European Union legislation.

Islets. Mouse pancreatic islets were isolated as described [32]. Details are provided in ESM Methods.

Glucagon content. Batches of thirty islets were incubated overnight at 4°C in 50 ml lysis buffer (70% [vol/vol.] ethanol, 0.4% [vol/vol.] HCl at 30% [vol/vol.], 29.6% [vol/vol.] distilled water). After centrifugation the supernatant was used for glucagon detection using an ELISA kit (YK090; Gentaur, Brussels, Belgium).

Quantitative RT-PCR. Determinations were performed from total RNA samples as detailed in ESM Methods.

Cell lines. AlphaTC1 cells (alphaTC1-9, ATCC CRL-2350) were cultured as described [10] supplemented with 16 mmol/l glucose unless specified otherwise. BHK21 cells were cultured as described [28].

Electrophoretic mobility shift assays (EMSA). EMSA were performed as described [33]. The sequences of the oligonucleotides used are shown in ESM Table 2. When indicated, ALX3 [33] or PAX6 (Millipore, MAB5552) antibodies were added to the binding reaction. In Fig. 3e, the ab64985 ALX3 antibody (Abcam, Cambridge, UK) was used.

Plasmids. ALX3 and PAX6 expression plasmids have been described [22, 28]. Oligonucleotides used for plasmid constructions are indicated in ESM Table 3. Luciferase reporter plasmids, plasmids encoding glutathione S-transferase (GST) fusion proteins and those used for *in vitro* transcription/translation are described in ESM Methods.

Transfections. Cells (10^5 per well) were incubated with reporter plasmids (0.5 μ g) mixed with Transfectin (BioRad) in serum-free medium. Expression plasmids or the corresponding empty vectors were used when required, keeping the total amount of DNA constant. Luciferase activity was determined and corrected for transfection efficiency using the *Renilla* luciferase assay kit (Promega). The Rous sarcoma virus enhancer reporter plasmid RSV-Luc was used as a standard for normalization. All transfections were performed in duplicates.

GST Pull-down assays. [35 S]Met-labelled and GST fusion proteins were used as described [33]. Details are provided in ESM Methods.

Immunoprecipitation. Immunoprecipitations [28] were performed using extracts from alphaTC1 cells, and were followed by immunodetection of PAX6 or ALX3 by western blot as indicated in ESM Methods.

Chromatin immunoprecipitation (ChIP) assays. ChIP assays [29] were performed using chromatin extracted from mouse islets or alphaTC1 cells. Details are indicated in ESM methods.

RNA interference (RNAi). RNAi [32] was performed using double stranded RNA duplexes as described in ESM Methods.

Western blots. Nuclear extracts from alphaTC1 cells and primary antibodies for ALX3 [34] (1:4000 dilution), PAX6 (H-295, Santa Cruz Biotechnology) (1:1000 dilution) and actin (clone AC-15, Sigma; 1:10,000 dilution) were used [32].

Statistical analysis Results represent mean \pm SEM for the indicated number of experiments. Statistical significance was calculated using Student's *t* test.

Results

Decreased glucagon levels in Alx3-deficient islets Reduced levels of glucagon content and gene expression in islets of *Alx3*-deficient mice (Fig. 1) [29] could not be attributed to decreased expression of key transcriptional regulators, because PAX6 [16], FOXA2 [20] and ARX [35] did not change whereas MAFB [17] increased (Fig. 1b). Thus, we hypothesized that ALX3 could regulate the glucagon promoter directly, and therefore

fluctuations in the levels of ALX3 could contribute to the dynamic regulation of glucagon gene expression depending on the metabolic needs of the organism. To test this hypothesis we investigated whether the levels of ALX3 in alpha cells vary depending on fluctuations in the concentration of glucose.

Expression of Alx3 is stimulated by glucose and results in inhibition of glucagon gene expression The levels of *Alx3* mRNA and protein were elevated when alphaTC1 cells were cultured in the presence of high glucose as compared to those cultured in low glucose, indicating that *Alx3* is a glucose-responsive gene (Fig. 2a-c). We tested whether glucose-regulated *Alx3* affects glucagon gene expression using cultured mouse islets. In wild type islets, increasing the concentration of glucose in the medium resulted in a significant increase in *Alx3* expression (Fig. 2d), and a concomitant decrease in glucagon expression (Fig. 2e). In contrast, in *Alx3*-deficient islets, increasing the concentration of glucose did not result in a significant change in the expression of glucagon mRNA (Fig. 2e). A glucose-stimulated increase in insulin I expression was unaffected by lack of ALX3 (Fig. 2f).

Alx3 binds specific sites of the glucagon promoter EMSA with nuclear extracts from alphaTC1 cells and probes corresponding to the G3 or G1 sites showed that at least two of the bands detected on the G3 probe were disrupted by the addition of ALX3- or PAX6-specific antibodies (Fig. 3b). At the G1 site, the addition of the PAX6 antibody disrupted the formation of DNA-protein complexes, whereas the addition of the ALX3 antibody had no effect (Fig. 3c). The ALX3 antibody also inhibited the formation of a complex on the G1₅₀ site (Fig. 3d), a homeodomain-binding site located downstream from G1 (Fig. 3a) [36]. Mutations in the PAX6 paired domain-binding site of G3

indicated that lack of binding of PAX6 prevented recruitment of ALX3 (ESM Fig. 1). ChIP assays demonstrated that both ALX3 and PAX6 are physically bound to the same region of the native glucagon promoter *in vivo* (Fig. 3e). These experiments indicate that ALX3 and PAX6 are part of the protein complex assembled on G3, and that the G1 site is bound by PAX6 as previously determined [36] but not by ALX3, which in turn recognizes the downstream G1₅₀ site.

ALX3-PAX6 heterodimers bind G3 but not G1 To investigate whether ALX3 interacts directly with PAX6 we performed immunoprecipitations from alphaTC1 cells. PAX6 was detected in samples immunoprecipitated with an ALX3 antiserum (Fig. 4a), and *vice versa* (Fig. 4b), but not in those immunoprecipitated with control serum, indicating that ALX3 and PAX6 physically interact in the nuclei of alpha cells. The domains from each protein required for this interaction were studied using GST pull-down assays using full length or truncated proteins (Fig. 4c). We found that [³⁵S]Met-labelled full length PAX6 was able to interact specifically with GST-ALX3 or with GST-ALX3HD, containing only the ALX3 homeodomain, but not with control GST or with GST fusion proteins containing either the carboxyl or the amino terminus of ALX3 (GST-ALX3ΔN or GST-ALX3ΔC, respectively) (Fig. 4d). A similar pattern of interactions was found when the [³⁵S]Met-labelled *paired* domain of PAX6 (PAX6PD) was used (Fig. 4e). Finally, the [³⁵S]Met-labelled homeodomain of ALX3 (ALX3HD) could interact independently with either the *paired* domain or the homeodomain of PAX6 (GST-PAX6PD or GST-PAX6HD, respectively) (Fig. 4f). These results indicate that the homeodomain of ALX3 is sufficient for heterodimerization via direct interactions with the *paired* domain or the homeodomain of PAX6.

The DNA binding consequences of these interactions were investigated using EMSA. PAX6 synthesized *in vitro* using a reticulocyte lysate could bind to the G3 probe as expected (Fig. 4g). In contrast, GST-ALX3HD, which bound efficiently to a previously described control probe (ESM Fig. 2a), failed to bind to G3 on its own. However, when the homeodomain of ALX3 was incubated together with PAX6 an additional band was generated, indicating that PAX6-ALX3 heterodimers can recognize G3 (Fig. 4g). In the case of the G1 site, binding of *in vitro* synthesized PAX6 (Fig. 4h) or of PAX6 present in nuclear extracts of alphaTC1 cells (Fig. 4i) was confirmed by the addition of a specific antibody that resulted in the disruption of binding. In both cases, the complexes containing PAX6 disappeared by the addition to the binding reaction of GST-ALX3, but remained undisturbed by the addition of control GST (Fig. 4h and i). Lack of binding of ALX3 to the G1 site was confirmed using recombinant ALX3 (ESM Fig. 2c and d).

These experiments indicate that ALX3-PAX6 heterodimers can bind to G3 efficiently, but are not capable to recognize G1. Since G1 is a well-known target site for binding by PAX6, these results raised the possibility that increasing levels of ALX3 in alpha cells could result in inhibition of PAX6 binding to G1, thus resulting in decreased glucagon promoter activation. This notion was tested directly by transfections in alphaTC1 cells.

Functional interactions between ALX3 and PAX6 on the glucagon promoter

Overexpression of ALX3 resulted in decreased luciferase activity elicited by the glucagon promoter reporter plasmid Gcg370Luc (Fig. 5a). In marked contrast, overexpression of PAX6 resulted in increased luciferase activity (Fig. 5a). When overexpressed together, the transactivation activity of PAX6 was inhibited by ALX3 in

a concentration-dependent manner (Fig. 5b). In turn, increasing amounts of PAX6 were unable to transactivate the glucagon promoter in the presence of a fixed amount of ALX3 (Fig. 5b).

When using the Gcg370/205Luc reporter, including G3 but excluding G1, ALX3 stimulated luciferase activity to a similar degree than that elicited by PAX6, and overexpression of both transcription factors concomitantly did not further increase or inhibit reporter activity (Fig. 5c). In contrast, when using Gcg122Luc, including G1 but excluding G3, overexpression of PAX6 stimulated luciferase activity, but overexpression of ALX3 decreased basal reporter activity and prevented transactivation by PAX6 (Fig. 5d). A similar pattern was found when using the reporter plasmids 3xG3T81Luc, which carries three tandem copies of G3, or 3xG1T81Luc, which carries three tandem copies of G1, respectively (Fig. 5e and f). In BHK21 cells, which do not express ALX3 or PAX6 [33, 37], ALX3 did not increase Gcg370Luc activity on its own, but enhanced PAX6-dependent transactivation synergistically (ESM Fig. 3). However, this effect was reversed to an inhibition of luciferase activity with the highest amount of ALX3 used (ESM Fig. 3). These experiments support the notion that ALX3 contributes to glucagon promoter activity by acting on G3 co-ordinately with PAX6, and that in turn it is able to inhibit PAX6-dependent promoter activity on G1 when ALX3 levels increase.

Alx3 regulates glucose-dependent binding of PAX6 and glucagon expression in alpha cells Our data are consistent with a model according to which glucose-dependent decrease of glucagon gene expression could result from selective displacement of PAX6 from G1 as a consequence of the formation of new PAX6-ALX3 heterodimers generated in response to increased levels of ALX3 induced by glucose (Fig. 6). In

alphaTC1 cells, glucose-dependent increase in ALX3 levels without affecting the levels of PAX6 (Fig. 6c) was accompanied by increased ALX3-PAX6 heterodimerization demonstrated by immunoprecipitation (Fig. 6d). As predicted by our model, increased levels of ALX3 specifically induced by high glucose concentrations (Fig. 6e) were accompanied by decreased binding of PAX6 to G1 (Fig. 6f), whereas binding of PAX6 to G3 or of ALX3 to G1₅₀, respectively, were unaffected (Fig. 6g-h). Reduced binding to G1 was not observed when expression of ALX3 was silenced by siRNA (Fig. 6i-j).

We tested this model further in alphaTC1 cells with silenced expression of ALX3 (Fig. 7a-c). In the presence of 2.8 mmol/l glucose, glucagon expression was reduced in *Alx3*-knock down cells relative to control cells (Fig. 7d), likely reflecting loss of ALX3 binding to G3 and G1₅₀ (Fig. 7e). In contrast, in the presence of 16 mmol/l glucose, glucagon expression increased in *Alx3*-knock down cells relative to control cells (Fig. 7f), reflecting availability of unsequestered PAX6, a stronger transactivator, for binding to G1 (Fig. 7g). These experiments support the requirement of ALX3 for glucose-dependent downregulation of glucagon gene expression in alpha cells.

Discussion

Our work indicates that ALX3 participates in the regulation of glucagon expression by binding with PAX6 to G3, and independently to G1₅₀. PAX6 binds to G1 in the absence of ALX3, but PAX6-ALX3 heterodimers are unable to recognize G1. This has important functional implications because the levels of ALX3 are up-regulated by glucose, thus favouring the formation of ALX3-PAX6 heterodimers engaging G1-bound PAX6. In this manner, ALX3 sequesters PAX6 and prevents its binding to G1, leading to downregulation of transcriptional activity. The inhibitory effect of ALX3 on

G1-bound PAX6 is dominant because PAX6 is a stronger transactivator than ALX3. In agreement with this model, glucose-unregulated glucagon mRNA levels in ALX3-deficient islets are higher than those observed in control islets under high glucose conditions (see Fig. 2e and ESM Fig. 4). In ALX3-deficient islets, PAX6 would be bound to both G1 and G3 regardless of the concentration of glucose, whereas in wild type islets, PAX6 would only be bound to G3, and the contribution of ALX3-bound to G3 and G1₅₀ would be small because ALX3 is known to be a weak transactivator [28] (see Fig. 5c and e).

Transcription factor interactions resulting in inhibition of glucagon expression by interfering with PAX6 have been described before. Some of these interactions may have implications in the context of the pharmacological treatment of diabetes [24], or during embryonic development to prevent glucagon expression in non-alpha islet cells [23, 36, 39], but their physiological relevance has not been completely established [40].

Glucagon secretion in the presence of reduced concentrations of glucose is coupled to stimulated transcription of the glucagon gene triggered by an auto-regulatory loop ensuring continuous hormone availability [5]. Our data support the existence of an ALX3-dependent counter regulatory glucose-sensing mechanism acting at the transcriptional level, by virtue of which overproduction of glucagon would be prevented under conditions in which glucose concentration is elevated and secretion inhibited. Thus, downregulation of PAX6 function by glucose-induced modulation of the ALX3 to PAX6 ratio links transcriptional regulation of gene expression in alpha cells to fluctuations in the circulating concentrations of glucose to accommodate glucagon biosynthesis to the metabolic demands of the organism.

PAX6 links glucagon production and secretion by coordinating the expression of a gene network regulating these processes at different levels [16, 22-27]. Therefore, it is

possible that these genes are regulated in a glucose-dependent manner via a mechanism involving selective interactions between PAX6 and ALX3 on different promoters. For example, the levels of *MafB* were elevated in islets from *Alx3*-deficient mice suggesting that ALX3 inhibits *MafB* expression. Although MAFB can act as a repressor [41], it is known to regulate glucagon gene expression [16, 17], and the *MafB* promoter contains two potential PAX6 binding sites [25]. Thus, inhibition of *MafB* expression by glucose-induced ALX3-mediated downregulation of PAX6 transactivation may provide an additional mechanism to reduce glucagon promoter activity in the presence of high glucose levels.

Apart from genes encoding transcription factors regulating glucagon gene expression [25], PAX6 regulates the expression of additional genes in alpha cells including those encoding the prohormone convertase 2 and its chaperone 7B2, the fatty acid transporter GPR40, or the glucose-dependent insulinotropic peptide receptor (GIPR) [26, 27]. As these genes regulate glucagon production and secretion, the possible occurrence of glucose-dependent PAX6-ALX3 interactions on their promoters may be centrally important for the coordinated modulation of transcriptional mechanisms opposing glucagon overproduction under elevated glucose concentrations. This would place ALX3 as a key component of glucose-regulated glucagon production and secretion.

Both ALX3 and PAX6 contain *paired*-type homeodomains that can form dimers with selective DNA binding specificities [33, 42, 43]. Besides, PAX6, but not ALX3, contains a *paired* domain. Both the *paired* domain and the homeodomain of PAX proteins bind DNA, and in some cases they interact to promote cooperative binding to specific regulatory elements [44-46]. PAX6 heterodimerization and alternate utilization of different protein subdomains modulate sequence selection and target site specificity

of the binding partners [47]. In the glucagon promoter, the *paired* domain of PAX6 is sufficient for binding to G3, whereas the homeodomain is dispensable [48]. In contrast, both the *paired* domain and the homeodomain are essential for binding of PAX6 to G1 [48]. Therefore, PAX6 bound to G3 by the *paired* domain could tether ALX3 via interactions between their homeodomains, so that ALX3-PAX6 interactions involving homeodomain heterodimerization would not disturb the *paired* domain-mediated binding of PAX6 to G3, thus providing stable transactivation from this element. Indeed, the interaction between the ALX3 homeodomain and the PAX6 *paired* domain may even contribute to binding [44-46]. This type of interactions between PAX6 and other homeodomain transcription factors has been shown to be important for transcriptional regulation [49]. In contrast, since the *paired* domain is not sufficient for PAX6 binding to G1 [48], utilization of the G1-bound PAX6 homeodomain after recruitment of ALX3 for heterodimerization would preclude the use of PAX6 homeodomain for DNA binding, thus destabilizing the protein-DNA interaction at this site.

The mechanism by which *Alx3* expression is stimulated by glucose is unknown. In the case of intact islets, it cannot be excluded that insulin released from beta cells in response to glucose may contribute to downregulation of glucagon gene transcription. However, the results obtained from alphaTC1 cells, which retain important physiological features of intact alpha cells [5, 10, 50] strongly argue in favour of a direct effect of glucose on *Alx3* expression.

Our findings may have implications for the aetiopathogenic mechanisms of diabetes. Dysfunctional glucagon hypersecretion contributing to hyperglycaemia has been recognized as an important factor in this disease [3, 4], suggesting the existence of impaired glucose-dependent inhibitory mechanisms affecting glucagon gene expression and hormone secretion. As ALX3 is important for the maintenance of glucose

homeostasis in mice [29], it is possible that ALX3 loss-of-function mutations may contribute to dysregulated glucagon production in alpha cells.

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Contribution statement MM performed the experiments. MV designed the study and wrote the paper. Both authors analysed the data and revised the final version of the paper before submission, and are responsible for the integrity of this work.

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Figures

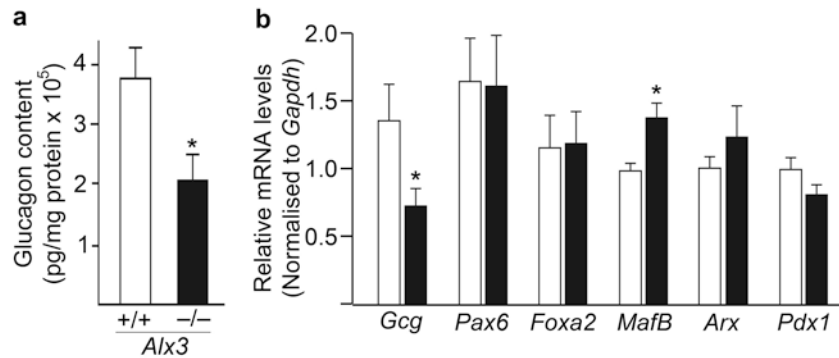


Figure 1. Reduced glucagon levels in *Alx3*-deficient mice. **a** Glucagon content in isolated islets from wild type (+/+) or *Alx3*-deficient (-/-) mice. n=6. **b** Relative mRNA levels of glucagon (*Gcg*) or alpha cell transcription factors and *Pdx1* in islets of wild-type (white bars) or *Alx3*-deficient (black bars) mice. n=5–15. **p*<0.05.

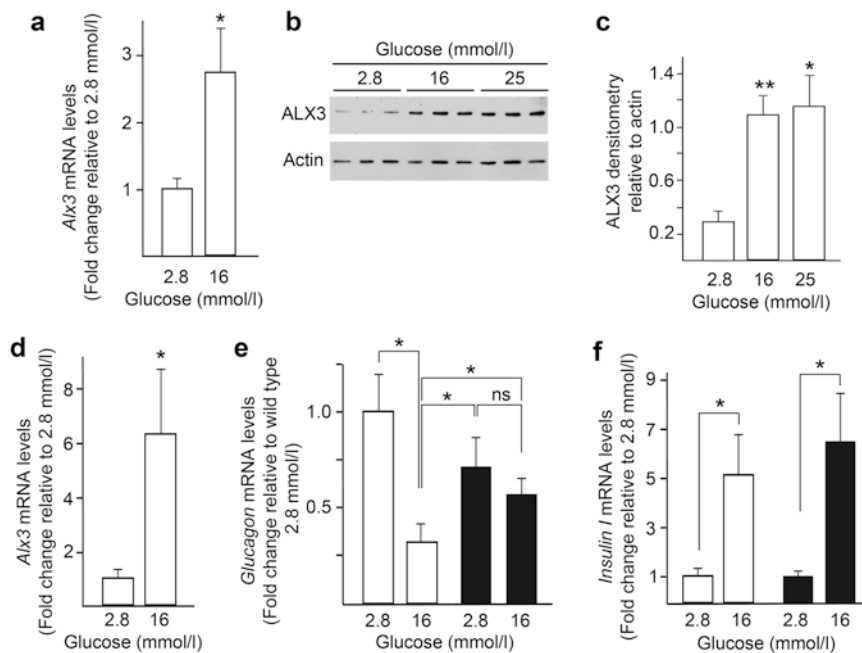


Figure 2. Glucose-stimulated expression of *Alx3* reduces glucagon gene expression. **a-b** Levels of *Alx3* mRNA (**a**) or protein (**b**) in alphaTC1 cells incubated in the presence of glucose at the indicated concentrations. In **b**, the results of three independent experiments for each condition are shown. **c** Densitometric quantification of the ALX3

bands relative to actin bands shown in **b**. **d-f** Levels of mRNA for *Alx3* (**d**), glucagon (**e**) or insulin I (**f**) in islets from wild-type (white bars) or *Alx3*-deficient (black bars) mice incubated at the indicated concentrations of glucose. n=9-10. * $p < 0.05$, ** $p < 0.01$.

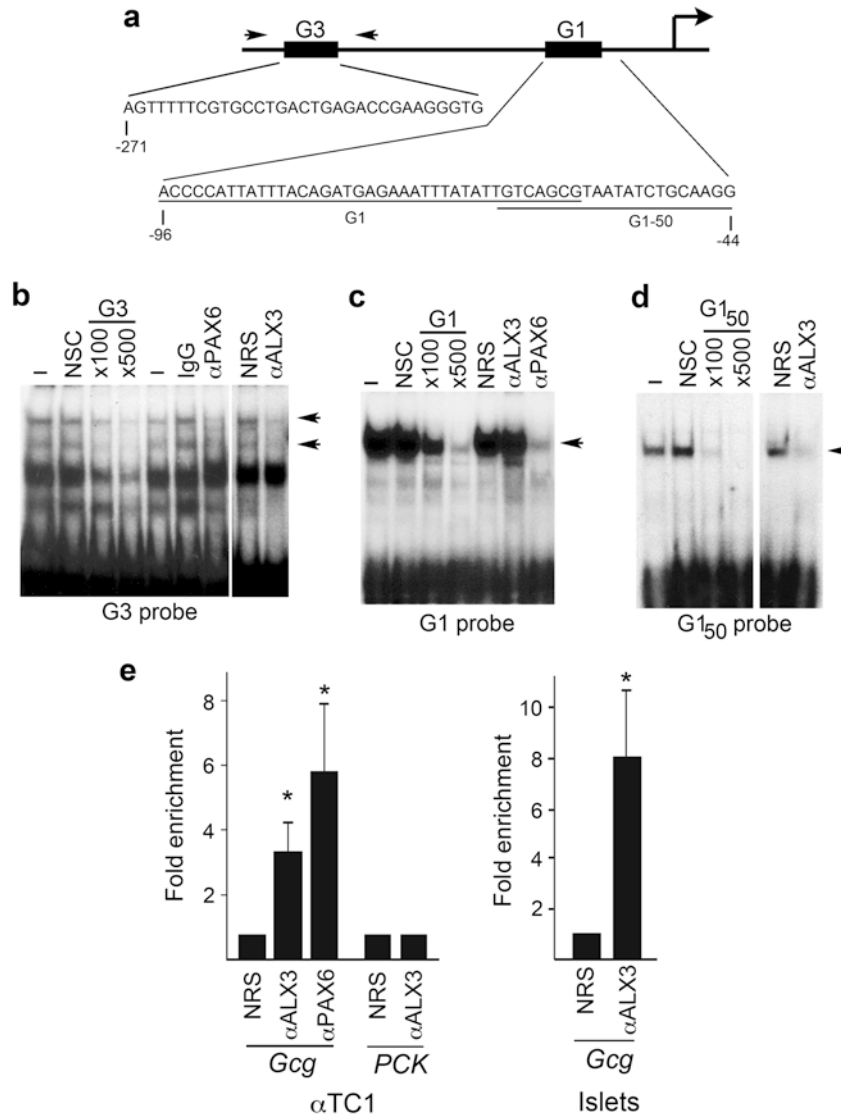


Figure 3. ALX3 binds to the glucagon gene promoter. **a** Relative location of the G3, G1 and G1₅₀ sites in the glucagon promoter. Arrows indicate the location of primers used for qPCR. **b-d** EMSAs showing the binding of nuclear proteins from alphaTC1 cells to G3 (**b**), G1 (**c**) or G1₅₀ (**d**). The absence (-) or presence of specific or non-specific (NSC) competing (500-fold molar excess) oligonucleotides, or the addition of PAX6 or

ALX3 antibodies, or control IgG or NRS is indicated. Arrowheads indicate complexes containing PAX6 and/or ALX3. **e** ChIP assays analysed by qPCR showing amplification of a region of the glucagon (Gcg) promoter containing G3 immunoprecipitated with ALX3 or PAX6 antibodies, or with control non-immune rabbit serum (NRS) from alphaTC1 cells or from mouse islets. The phosphoenolpyruvate carboxykinase (PCK) gene was used as negative control. * $p < 0.05$; n=3.

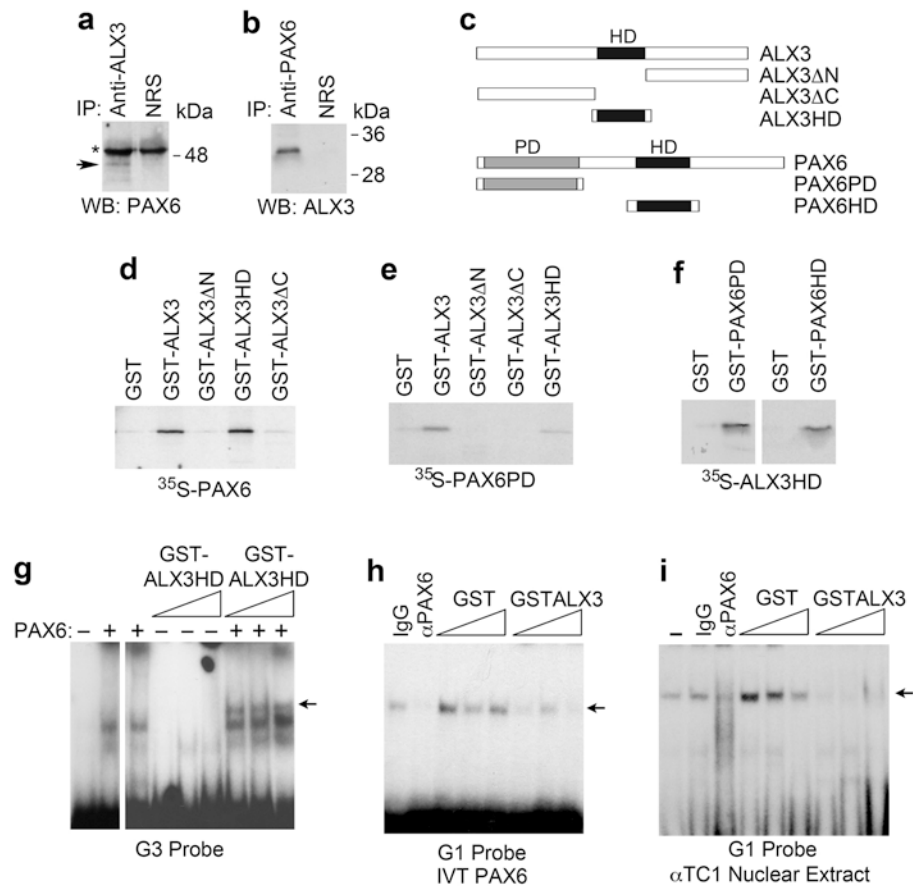


Figure 4. Physical interactions and DNA binding of ALX3 and PAX6. **a-b** Western blots (WB) for PAX6 (**a**) or ALX3 (**b**) on samples immunoprecipitated from alphaTC1 cells with an ALX3 or a PAX6 antibody, respectively. The band showing PAX6 interacting with ALX3 is indicated by arrow. *Non-specific bands. (NRS, non-immune rabbit serum). **c** Full length and truncated versions of ALX3 or PAX6 used in GST pull-

down experiments. HD, homeodomain; PD, *paired* domain. **d-f** GST pull-down experiments performed with ³⁵S-labeled PAX6 (**d**), PAX6 *paired* domain (**e**), or ALX3 homeodomain (**f**). **g** EMSA with *in vitro* transcribed/translated PAX6 (+) or control reticulocyte lysate (-) incubated with radiolabelled G3 in the absence or presence of GSTAlx3HD. Arrow indicates a PAX6-GSTAlx3HD heterodimer. **h-i** Binding of *in vitro* transcribed/translated (IVT) PAX6 (**h**) or alphaTC1 nuclear extracts (**i**) to radiolabelled G1. The complex containing PAX6 (arrow), identified by addition of a specific antibody, was inhibited by GSTAlx3 but not by control GST.

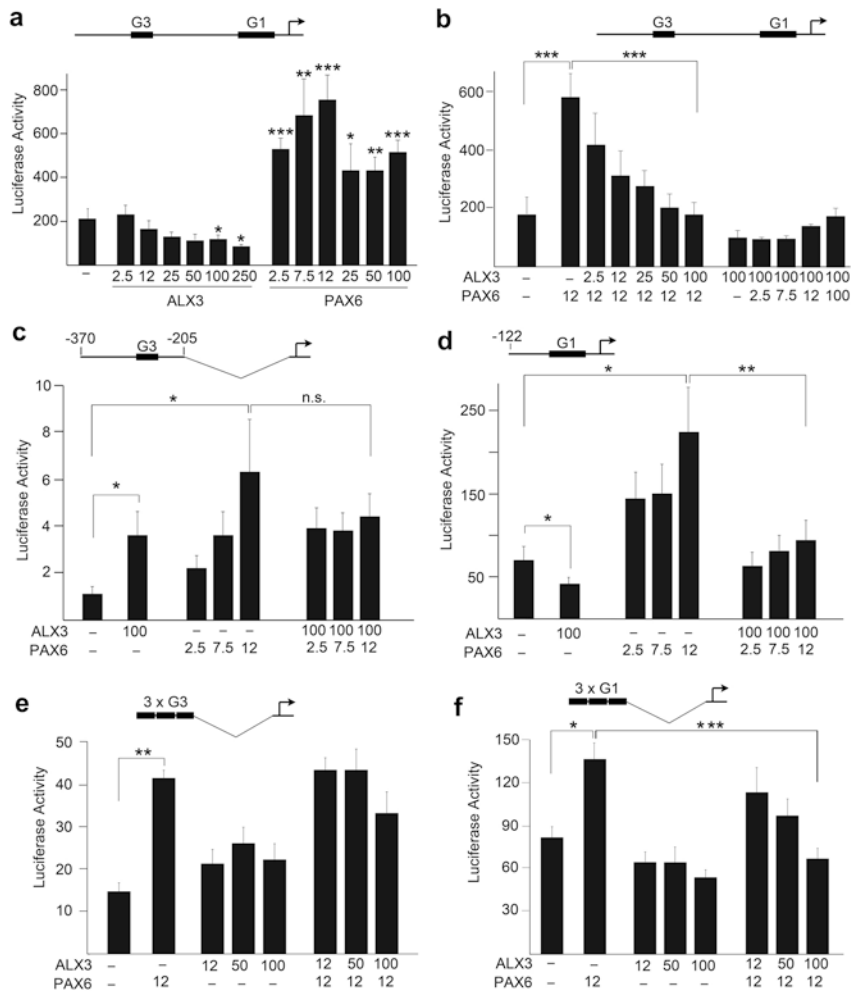


Figure 5. Functional interactions between ALX3 and PAX6 on the glucagon promoter.

Relative luciferase activities elicited in alphaTC1 cells cotransfected with Gcg370Luc

(a and b), Gcg370/205Luc (c), Gcg122Luc (d), 3xG3T81Luc (e) or 3xG1T81Luc (f) reporter plasmids and the indicated amounts (ng) of expression plasmids encoding either ALX3 or PAX6. The relative positions of the G3 and/or G1 elements in the glucagon promoter is depicted on top of the histograms. n=4-6. * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$.

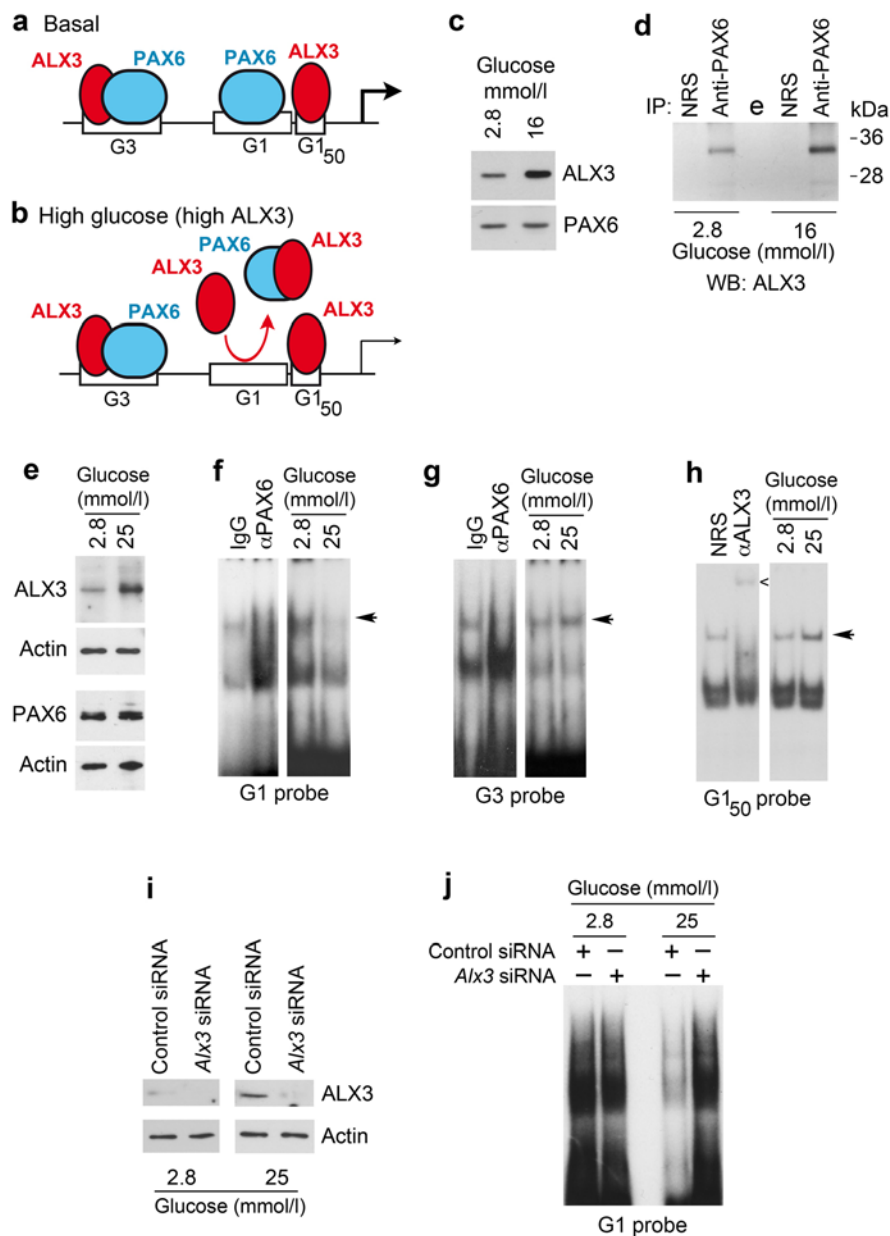


Figure 6. ALX3 prevents binding of PAX6 to G1 in a glucose-dependent manner. **a-b** Hypothetical model of the glucose-dependent regulation of glucagon gene expression in

alpha cells by ALX3 and PAX6. **c** Western blot showing glucose-dependent induction of ALX3 but not PAX6 in extracts of alphaTC1 cells used for immunoprecipitations (**d**). **d** Glucose-enhanced ALX3-PAX6 interaction shown by western blot (WB) for ALX3 on samples from alphaTC1 cells immunoprecipitated (IP) with a PAX6 antibody (e, empty lane). **e** Western blot showing glucose-dependent induction of ALX3 but not PAX6 in extracts of alphaTC1 cells used in EMSA. **f-h** EMSA showing binding of nuclear extracts from alphaTC1 cells cultured at the indicated concentrations of glucose to G1 (**f**), G3 (**g**) or G1₅₀ (**h**). Arrows indicate the presence of PAX6 (**f** and **g**) or ALX3 (**h**) identified by the addition of specific antibodies. Arrowhead, supershifted ALX3 (**h**). **i** Western blot showing levels of ALX3 in alphaTC1 cells transfected with control or *Alx3* siRNA. **j** EMSA showing binding to G1 of nuclear extracts from alphaTC1 cells transfected with control or *Alx3* siRNA. In **i** and **j**, glucose concentrations for cultured cells are indicated. NRS, non-immune rabbit serum.

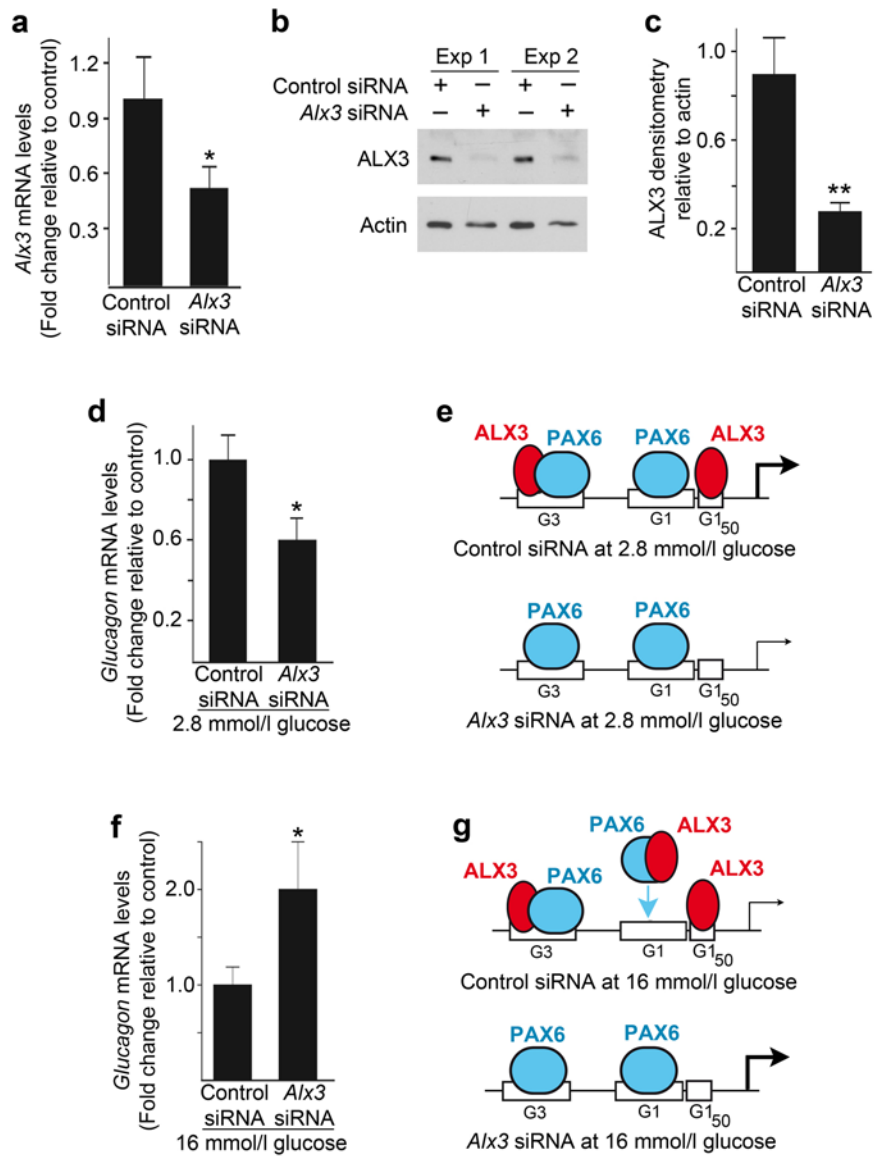


Figure 7. Glucose-dependent regulation of glucagon expression by ALX3. **a** Levels of *Alx3* mRNA in alphaTC1 cells transfected with control or *Alx3* siRNA (n=4). **b** Western blot showing levels of ALX3 in alphaTC1 cells transfected with control or *Alx3* siRNA. Results from two representative experiments are shown. **c** Densitometric quantification of ALX3 bands relative to actin bands obtained from western blots similar to those shown in **b** (n=6). **d-g** Relative levels of glucagon mRNA in alphaTC1 cells cultured at the indicated concentrations of glucose and transfected with control or *Alx3* siRNA (n=4) (**d** and **f**) and the respective schematic interpretation of the results in each condition (**e** and **g**). * $p < 0.05$; ** $p < 0.01$.