

Regulation of forkhead box O1 (FOXO1) by protein kinase B and glucocorticoids: different mechanisms of induction of beta cell death in vitro

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Abstract

Aims/hypothesis In steroid diabetes insulin secretion does not adequately compensate for enhanced hepatic gluconeogenesis and peripheral insulin resistance. Previous studies suggest that activation of the transcription factor forkhead box O1 (FOXO1) contributes to glucocorticoid-induced beta cell death. This study examines the role and regulation of FOXO1 in insulin-secreting cells.

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Methods INS-1E cells and mouse islet cells were cultured in the presence of dexamethasone. Signalling pathways were modified pharmacologically or by small interfering (si) RNA-mediated inhibition of protein synthesis. Changes in protein abundance and phosphorylation were analysed by western blotting, and subcellular localisation was assessed using confocal microscopy. Transcript levels were examined by RT-PCR.

Results Surprisingly, downregulation of FOXO1 by siRNA did not affect dexamethasone-induced apoptosis or *Bim* expression, but it prevented the effects of the pan protein kinase B (AKT) inhibitor (Akti-1/2). Indeed, dexamethasone and Akti-1/2 synergistically increased beta cell death and *Bim* expression. Akti-1/2 triggered dephosphorylation and nuclear translocation of FOXO1. Glucocorticoid-receptor activation stimulated *Foxo1* transcription, but FOXO1 phosphorylation was unchanged and the cytosolic concentration of FOXO1 remained high in relation to its nuclear concentration. However, subcellular fractionation revealed a significant increase in both cytosolic and nuclear FOXO1 compared with untreated cells. Dexamethasone diminished *Pdx1* mRNA level, an effect which was not reversed by siRNA against *Foxo1*. Downregulation of AKT isoforms and serum/glucocorticoid-regulated kinase 1 (SGK1) suggests that only sustained suppression of all three AKT isoforms caused dephosphorylation and nuclear accumulation of FOXO1.

Conclusions/interpretation This study reveals that FOXO1 is not the main mediator of glucocorticoid-receptor-induced beta cell apoptosis, but rather that it escalates beta cell death when AKT activity is inhibited by distinct pathways.

Keywords AKT isoforms · Apoptosis · Dexamethasone · FOXO1 · Glucocorticoids · Insulin-secreting cells · SGK1

Abbreviations

AKT	Protein kinase B
Akti-1/2	AKT inhibitor-1/2
BAD	BCL2-associated agonist of cell death
FOXO1	Forkhead box O1
p-	Phosphorylated
PARP	Poly(ADP-ribose) polymerase family, member 1
SGKi	SGK1 inhibitor GSK650394
SGK1	Serum/glucocorticoid-regulated kinase 1
si	Small interfering

Introduction

In clinical therapy, glucocorticoids such as the synthetic agent dexamethasone are commonly used in autoimmune and anti-inflammatory disease management. A frequent side effect of long-term and high-dose glucocorticoid treatment is the development of steroid diabetes [1]. In addition to an increase in hepatic glucose production [2] and peripheral insulin resistance [3], glucocorticoids impair beta cell function [4]. The underlying mechanism of glucocorticoid-induced beta cell dysfunction is not well understood. Many observations point to a crucial role of the transcription factor forkhead box O1 (FOXO1) in beta cell failure [5]. However, the role of FOXO1 in the maintenance of beta cell function and appropriate beta cell mass is controversial [6]. On one hand, FOXO1-mediated inhibition of the beta cell-specific transcription factor *Pdx1* impairs beta cell neogenesis and contributes to a reduction of beta cell mass [7]. On the other hand, FOXO1 induces *NeuroD* and *MafA* expression, which encode two transcription factors essential for insulin gene transcription [8]. A reduction in FOXO1 protein protects against beta cell death induced by the synthetic glucocorticoid dexamethasone, suggesting that FOXO1 activation might mediate the pro-apoptotic effects of glucocorticoids [9]. In insulin-secreting cells FOXO1 stays under the control of insulin/IGF-1 signalling through IRS-2 and protein kinase B (AKT) [10]. Previously, we described that dexamethasone reduces phosphorylation of AKT and of the pro-apoptotic protein BCL2-associated agonist of cell death (BAD), triggering the intrinsic apoptotic pathway [11].

Further effects of glucocorticoids on beta cell function comprise inhibition of insulin biosynthesis and insulin secretion [11–16]. Inhibition of insulin secretion is at least partly mediated by increased potassium channel activity [12]. The mechanism involves α_2 -adrenoceptor activation and serum/glucocorticoid-regulated kinase 1 (SGK1) induction [17]. SGK1, a serum- and glucocorticoid-inducible kinase with 75% structural homology with AKT, increases the activity of voltage-dependent potassium channels [18, 19]. Interestingly, a genetic polymorphism downstream of *SGK1* 3'-untranslated region (UTR) associates with increased insulin

secretion and reduced risk of developing diabetes mellitus, which is indicative of a regulatory role of SGK1 in human beta cell function [20]. SGK1 and the other two isoforms, SGK2 and SGK3, are similar to AKT in that they are stimulated by insulin/IGF-1 receptor signalling and share substrates. A common substrate of SGK and AKT isoforms is the transcription factor FOXO1. AKT phosphorylates FOXO1 at Thr24, Ser256 and Ser319, while SGK1 preferentially phosphorylates its Ser256 and Ser319 residues [21–23]. Phosphorylation of FOXO1 is believed to be one of the main anti-apoptotic signals downstream of AKT [24]. Phosphorylated FOXO1 is confined into the cytosol, while inhibition of AKT allows the translocation of the transcription factor into nuclei and activation of FOXO1-dependent pro-apoptotic genes.

This study questions the role of AKT isoforms and SGK1 on FOXO1 regulation and its impact on glucocorticoid-induced beta cell death.

Methods

Reagents Antibodies against FOXO1 (cat no. 2880), phosphorylated (p)-FOXO1 (Ser256, cat no. 9461), AKT (cat no. 9272), p-AKT (Ser473, cat no. 9271), AKT1 (cat no. 2967), poly(ADP-ribose) polymerase family, member 1 (PARP) (cat no. 9542), SGK1 (cat no. 3272) and α/β -tubulin (cat no. 2148) were from Cell Signaling Technology (Beverly, MA, USA). Antibodies against p-FOXO1 (Ser319, cat no. sc-101682), glyceraldehyde 3-phosphate dehydrogenase (GAPDH)-horse-radish peroxidase (HRP) (cat no. 20357-HRP) and Histone H1 (cat no. sc-10806) were purchased from Santa Cruz Biotechnology (Santa Cruz, CA, USA). Peroxidase-linked donkey anti-rabbit IgG was from GE Healthcare (GE Healthcare Europe, Freiburg, Germany). Alexa Fluor 488 anti-rabbit IgG was obtained from Invitrogen (Karlsruhe, Germany). The antibody for SGK1 used for immunohistochemical staining was made by Pineda (Berlin, Germany).

Cell culture media were obtained from Gibco (Invitrogen) and FCS from Biochrom (Berlin, Germany). The protein assay was purchased from BioRad Laboratories (München, Germany), the protein ladder from Invitrogen. AKT inhibitor-1/2 (Akti-1/2) was purchased from Merck Chemicals (Nottingham, UK), GSK650394 from Tocris Bioscience (R&D Systems Wiesbaden-Nordenstadt, Germany) and nuclear dye TOPRO3 from Invitrogen. All other chemicals were from Sigma-Aldrich (Deisenhofen, Germany) and of analytical grade unless otherwise stated.

INS-1E cell culture and transfection Cells of the rat insulinoma cell line INS-1E (kindly provided by C. B. Wollheim, University of Geneva, Switzerland) were cultured in HEPES-buffered RPMI 1640 supplemented with 10% (vol./vol.) FCS, 10 mmol/l HEPES, 2 mmol/l L-glutamine,

1 mmol/l sodium pyruvate and 10 μ mol/l β -mercaptoethanol, as described previously [18]. The culture medium contained 11 mmol/l glucose.

Cells were transfected with the plasmids hSgk1^{SD}/pIRES2-EGFP and hSgk1^{KN}/pIRES2-EGFP (Sgk1^{SD} encodes the constitutively active form and Sgk1^{KN} encodes the kinase dead form of SGK1, respectively [Clontech, Saint-Germain-en-Laye, France]) or small interfering (si)RNA directed against Sgk1 (Qiagen, Hilden, Germany), Akt1, Akt2, Akt3, Foxo1 (siGenome smart pool) or siGLO RISC-Free Control siRNA using DharmaFect Transfection Reagent 3 (Dharmacon, Chicago, USA). Cells were analysed 1 day after transfection with plasmid and 2 days after transfection with siRNA.

Mouse islet preparations Mouse islets from adult C57BL/6 mice were isolated using collagenase digestion (1 mg/ml collagenase, Serva, Heidelberg, Germany). Single islet cells were prepared by digestion with trypsin-EDTA in PBS for 4–8 min at 37°C and thereafter seeded on polyornithine-coated (0.001% (wt/vol.)) glass cover slips. Islets and islet cells were cultured overnight in the same medium as INS-1E cells without addition of β -mercaptoethanol. After medium change, cells and islets were cultured in the presence of dexamethasone, 100 nmol/l for 24 h. Akti-1/2, 3 μ mol/l, was added for the last 30 min.

Immunohistochemical staining and TUNEL assay INS-1E cells or mouse islet cells were cultured in the presence of test substances, as indicated. Cells were fixed with 4% paraformaldehyde in PBS for 1 h, permeabilised with 0.2% (vol./vol.) Triton X-100 for 2 min on ice, blocked with 10% (vol./vol.) FCS for 45 min at room temperature and incubated overnight with a primary antibody against FOXO1 (1:100) or SGK1 (1:100). After incubation with a secondary antibody (1:400 Alexa Fluor 488 anti-rabbit IgG) for 1 h, nuclei were stained with 1 μ mol/l TO-PRO3 in PBS for 1 h. TUNEL staining was performed according to the instructions of the manufacturer (Roche Diagnostics, Mannheim, Germany). Thereafter, nuclei were stained with DAPI, 0.1 μ g/ml in PBS, for 30 min at room temperature. Fluorescence was visualised using a laser scanning microscope (Leica Microsystems, Wetzlar, Germany).

Western blotting Cells were seeded and treated with test substances as indicated for each experiment. For experiments with the AKT inhibitor Akti-1/2 and the SGK1 inhibitor GSK650394 (SGKi), cells were pre-incubated for 1 h in KRB supplemented with 2.8 mmol/l glucose and 0.1% (wt/vol.) BSA (fraction V); dexamethasone was present throughout the preincubation. For western blotting, cells were lysed in buffer containing 125 mmol/l NaCl, 1% (vol./vol.) Triton X-100, 0.5% (wt/vol.) Na-deoxycholate, 0.1% (wt/vol.) SDS, 10 mmol/l EDTA, 25 mmol/l HEPES pH 7.3, 10 mmol/l

NaPP, 10 mmol/l NaF, 1 mmol/l Na-vanadate, 10 μ g/ml pepstatin A, 10 μ g/ml aprotinin and 0.1 mmol/l phenylmethylsulfonylfluoride (PMSF) as described previously [11]. Proteins (30 μ g) of a 10,000 g supernatant fraction were subjected to SDS-PAGE and blotted on nitrocellulose membranes (Schleicher and Schuell, Dassel, Germany). Antibodies were diluted 1:1000 in Tris-buffered saline (TBS) supplemented with 0.15% (vol./vol.) Tween and 5% (wt/vol.) BSA or 5% (wt/vol.) milk. Quantification of western blots was performed by measuring the optical densities of the respective bands on the film using ImageJ (NIH, Bethesda, MD, USA). For nuclear and cytosolic cell fractionation a commercial kit (Thermo Scientific; Rockford, IL, USA) was used and 30 μ g protein of the cytosolic fraction and 15 μ g protein of the nuclear fraction were loaded onto western blots.

Quantitative real-time RT-PCR analysis INS-1E cells were cultured under standard conditions and treated with substances as indicated. Thereafter, cells were lysed and total RNA isolated using a commercial kit (Qiagen). After cDNA synthesis using Oligo(dT)12-18 as primer (Roche Diagnostics), specific PCR was performed with the Light Cycler Fast Start Master SybrGreen method (Roche Diagnostics) and the Light Cycler Primer Set for Murine/Rat GAPDH (Roche Diagnostics) as external standard. The primers are listed in electronic supplementary material (ESM) Table 1.

Quantification of mRNA levels of mouse islets was performed with the Light Cycler System LC480 and the LC480 Probes Master (Roche Diagnostics). Gene expression relative to the housekeeping gene *Rps13* was quantified by the $2^{-\Delta\Delta C_t}$ method. The respective primers are listed in ESM Table 2.

Statistics Data are provided as mean \pm SEM; *n* represents the number of observations of at least three independent experiments. All data were tested for significance using Student's *t* test, ANOVA and Newman–Keuls test as post hoc test where appropriate. Results with *p*<0.05 were considered statistically significant.

Results

Activation of FOXO1 is not required for dexamethasone-induced cell death To examine whether FOXO1 contributes to dexamethasone-induced cell death, FOXO1 synthesis was downregulated by up to 86% in INS-1E cells using the siRNA strategy (Fig. 1a, b). Treatment of cells with the pan AKT inhibitor Akti-1/2 was used as a positive control for AKT-dependent FOXO1 regulation [25]. Apoptosis was examined by two independent methods, by immunohistochemical staining of TUNEL-positive cells and by detection of cleaved PARP, which reflects the activation of

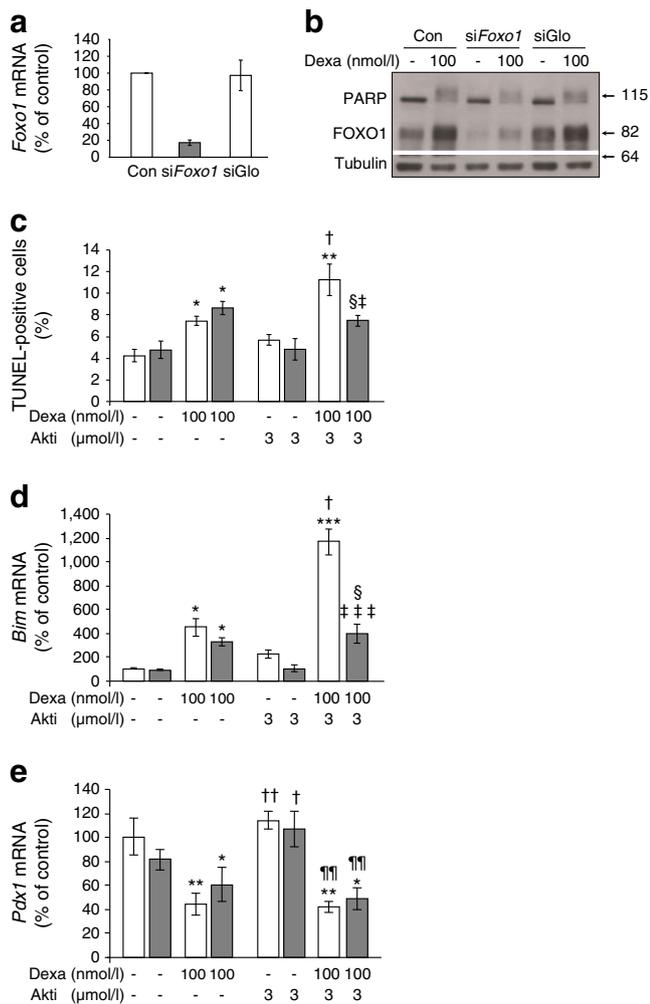


Fig. 1 Activation of FOXO1 is not required for dexamethasone-induced cell death. INS-1E cells were transfected with siRNA against *Foxo1* (grey bars) or siGLO (white bars) and treated with dexamethasone, 100 nmol/l for 1 day, and with Akti-1/2, 3 μmol/l, in the presence of dexamethasone for 2 h for TUNEL and for 8 h for mRNA analysis, as indicated and described in the **Methods**. **(a)** Relative amounts of *Foxo1* mRNA are expressed as mean ± SEM of $n=5$ independent experiments. **(b)** A representative western blot for PARP and FOXO1. Tubulin was used as loading control. **(c)** The percentage of TUNEL-positive cells is expressed as mean ± SEM of $n=5$ independent experiments. **(d)** *Bim* and **(e)** *Pdx1* mRNA levels are expressed as mean ± SEM of $n=4$ independent experiments. The content of mRNA under control was set to 100%. * $p<0.05$ and ** $p<0.01$ denote significance compared with siGlo untreated (first bar of each graph); † $p<0.05$ and †† $p<0.01$ denote significance compared with siGlo dexamethasone-treated; ‡ $p<0.05$ and ‡‡ $p<0.001$ denote significance compared with siGlo treated with dexamethasone and Akti-1/2; § $p<0.05$ denotes significance compared with siRNA *Foxo1* treated with Akti-1/2; ¶ $p<0.01$ denotes significance compared with the respective Akti-1/2 treated cells; Akti, Akti-1/2; Con, control; Dexa, dexamethasone; siFoxo1, cells transfected with siRNA against *Foxo1*

caspace-3 [26]. In FOXO1-deficient cells *Foxo3* and *Foxo4* mRNA remained unchanged (data not shown). The successful inhibition of FOXO1 synthesis did not prevent either the dexamethasone-induced degradation of PARP or the increase in the number of TUNEL-positive nuclei (Fig. 1b, c). The

significant augmentation of dexamethasone-induced apoptosis by the AKT inhibitor Akti-1/2 (3 μmol/l for 2 h), however, was completely abrogated by siRNA against *Foxo1* (Fig. 1c). In accordance, dexamethasone increased *Bim* mRNA levels in a FOXO1-independent manner, while the effect of Akti-1/2 on *Bim* mRNA was abrogated in FOXO1-deficient cells (Fig. 1d). Furthermore, dexamethasone reduced *Pdx1* mRNA levels independently of FOXO1, while Akti-1/2 did not alter mRNA levels of *Pdx1* (Fig. 1e). These results suggest that glucocorticoid-receptor activation triggers apoptotic beta cell death in a FOXO1-independent manner, which may further sensitise the cells toward FOXO1-mediated apoptotic signals.

Akti-1/2 but not dexamethasone inhibited phosphorylation and promoted nuclear accumulation of FOXO1 As AKT activity determines phosphorylation and subcellular localisation of FOXO1 and previous observations show that dexamethasone inhibits phosphorylation of AKT, phosphorylation of FOXO1 and its subcellular distribution were examined in dexamethasone-treated INS-1E cells. Surprisingly, phosphorylation of FOXO1 at Ser256 and Ser319 was significantly higher in dexamethasone-treated (100 nmol/l for 24 h) compared with control whole-INS-1E-cell lysates (Fig. 2a, b). The increase in FOXO1 protein occurred in parallel with its phosphorylation. Consequently, dexamethasone did not change the degree of phosphorylation of FOXO1 (Fig. 2c–e). As against this, Akti-1/2 (3 μmol/l for 30 min) strongly inhibited phosphorylation of FOXO1 (Fig. 2f). In control as well as in dexamethasone-treated cells (100 nmol/l for 4, 24 and 48 h) the concentration of FOXO1 was higher in the cytosolic compartments than in the nuclei, as confirmed by confocal microscopy (Fig. 3a). In contrast, exposure to Akti-1/2 for as short a period as 30 min induced nuclear accumulation of FOXO1 (Fig. 3b). These observations confirm that Akti-1/2 activates FOXO1 while dexamethasone does not.

Intriguingly, subcellular fractionation suggests that the amount of FOXO1 increased in the cytosolic fraction and also in the nuclear fraction on dexamethasone treatment (ESM Fig. 1a). Treatment with Akti-1/2 (3 μmol/l for 30 min) resulted in translocation of FOXO1 as the cytosolic amount of FOXO1 decreased while the nuclear amount increased. Additionally, in dexamethasone-exposed INS-1E cells AKT phosphorylation was significantly reduced, in line with previous observations (ESM Fig. 1b).

To confirm that regulation of FOXO1 by dexamethasone and Akti-1/2 in INS-1E cells translates to primary islet cells, comparable experiments were performed with islets isolated from adult mice (Fig. 4). In mouse islet cells cultured in the presence of dexamethasone (100 nmol/l for 24 h), FOXO1 immunoreactivity was more prominent in the cytosol than in nuclei, while after exposure to Akti-1/2 (3 μmol/l for 30 min) FOXO1 accumulated in nuclei (Fig. 4a). Dexamethasone treatment of mouse islets (100 nmol/l for 24 h) significantly

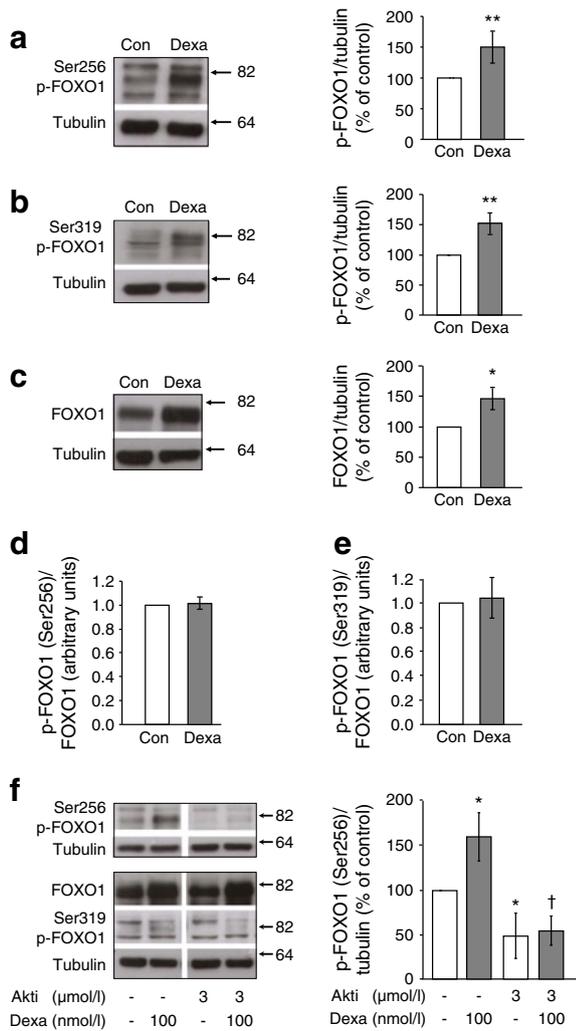


Fig. 2 Akti-1/2 but not dexamethasone inhibited phosphorylation of FOXO1. Cells were treated with dexamethasone, 100 nmol/l for 24 h, and Akti-1/2, 3 μmol/l for 30 min, as indicated and described in the Methods. (a–c, f) Representative western blots and mean ± SEM of three to five independent experiments for p-FOXO1 (Ser256) (a), p-FOXO1 (Ser319) (b), FOXO1 (c) and p-FOXO1 (Ser256), FOXO1 and p-FOXO1 (Ser319) (f). Tubulin was used as loading control. The ratio of p-FOXO1 (Ser256) (d) and p-FOXO1 (Ser319) (e) to FOXO1 protein is presented as mean ± SEM of $n=6$ independent experiments. * $p<0.05$ and ** $p<0.01$ denote significance compared with control; † $p<0.05$ denotes significance compared with dexamethasone. Akti, Akti-1/2; Con, control; Dexa, dexamethasone

increased *Foxo1* (Fig. 4b), *Foxo3* (Fig. 4c) and *Bim* (Fig. 4d) mRNA levels but reduced *Pdx1* (Fig. 4e) mRNA levels as in INS-1E cells. The mRNA concentration of *Foxo1* was notably higher (200-fold) than that of *Foxo3*. Thus, dexamethasone exerts similar effects in INS-1E cells and mouse islets.

Specific effects of glucocorticoid-receptor activation on *Foxo*, *Akt* and *Sgk* isoforms To examine the mechanism underlying increased FOXO1 protein in cells exposed to an excess of glucocorticoids, the effect of dexamethasone

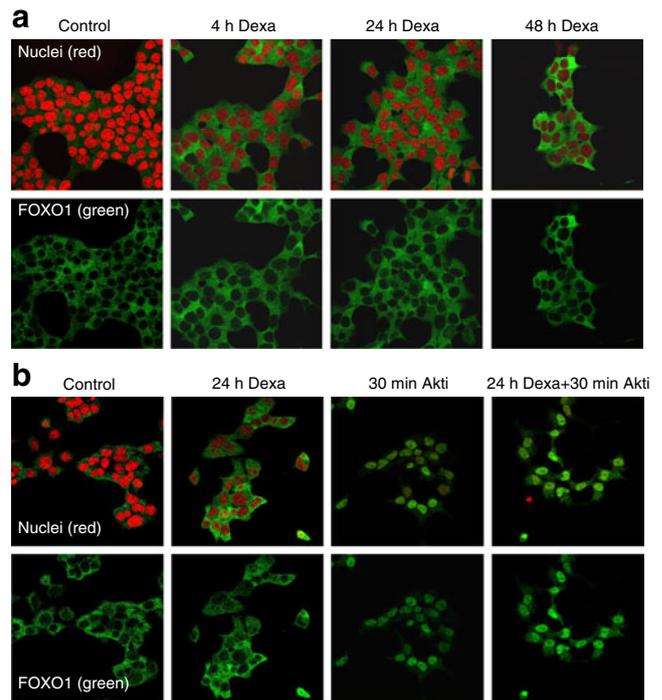
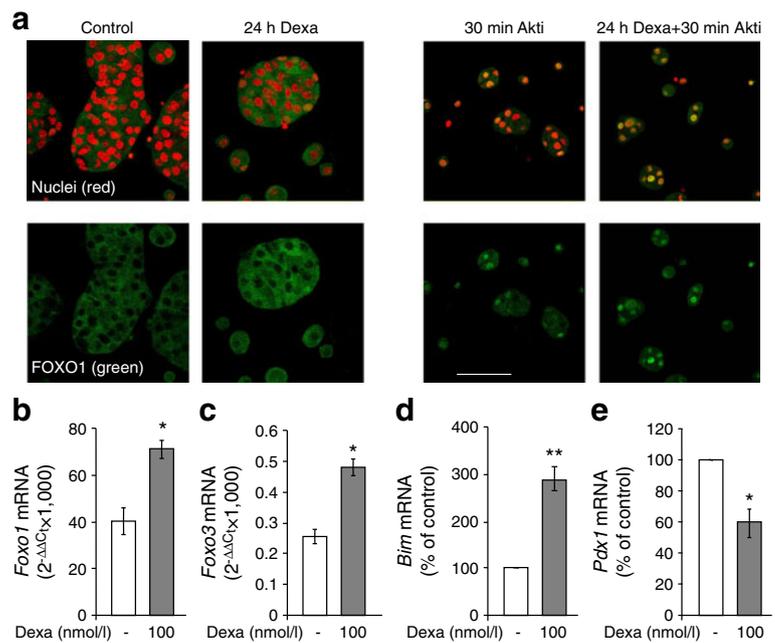


Fig. 3 The inhibitor Akti-1/2, but not dexamethasone, promotes nuclear accumulation of FOXO1. (a) INS-1E cells were cultured under standard conditions and treated with dexamethasone for the indicated time. (b) Cells were treated as described in Fig. 2. Representative pictures of immunostaining of FOXO1 (green) using confocal microscopy are shown. Nuclei are stained in red. Akti, Akti-1/2; Dexa, dexamethasone

on *Foxo1* transcript levels was evaluated by semiquantitative analysis of mRNA levels in the absence and presence of the glucocorticoid-receptor inhibitor RU486 (ESM Figs 2, 3). In INS-1E cells, *Foxo1* mRNA was as abundant as *Foxo3* mRNA, whereas the level of *Foxo4* mRNA was 30-fold lower (ESM Fig. 2a). Glucocorticoid treatment specifically increased *Foxo1* mRNA within 4 h (ESM Fig. 2b). A 2.5-fold increase observed throughout the treatment (24 and 48 h) was abrogated on exposure to the glucocorticoid-receptor antagonist RU486 (mifepristone, 10 μmol/l [ESM Fig. 3a]). *Foxo3* and *Foxo4* mRNA levels increased significantly only after prolonged exposure (24–48 h) to the glucocorticoid analogue (ESM Fig. 2c, d). These effects were again sensitive to mifepristone (ESM Fig. 3b, c).

In the next series of experiments mRNA levels of *Akt* and *Sgk* isoforms were analysed to identify the enzymes responsible for phosphorylation of FOXO1 (ESM Fig. 2e–l). Control INS-1E cells expressed comparable levels of *Akt1* and *Sgk1* mRNA (ESM Fig. 2e, i). *Akt2* and *Akt3* mRNA levels were tenfold and fivefold lower, respectively, than *Akt1* mRNA levels (ESM Fig. 2e); *Sgk2* and *Sgk3* mRNA levels were 100-fold and tenfold lower, respectively, than *Sgk1* mRNA levels (ESM Fig. 2i). Dexamethasone treatment resulted in a unique 20-fold increase in *Sgk1* mRNA (ESM Fig. 2j). Accordingly, SGK1 protein was detected by

Fig. 4 Effects of dexamethasone on *Foxo1*, *Foxo3*, *Bim* and *Pdx1* expression in mouse islets. Mouse islet cells were isolated and cultured as described in **Methods**. **(a)** Representative pictures of immunostaining of FOXO1 (green) and nuclei (red) using confocal microscopy; scale bar 50 μ m. **(b–e)** Relative gene expressions of *Foxo1* **(b)**, *Foxo3* **(c)**, *Bim* **(d)** and *Pdx1* **(e)** were determined by quantitative RT-PCR and are expressed as mean \pm SEM of $n=4$ independent experiments. *Rps13* was used as housekeeping gene. For *Bim* and *Pdx1*, mRNA content of untreated cells was set to 100%. * $p<0.05$ and ** $p<0.01$ denote significance to untreated cells. Akti, Akti-1/2; Dexa, dexamethasone



immunostaining in dexamethasone-treated, but not in control, INS-1E cells (ESM Fig. 3d). The relative mRNA concentrations of *Sgk2* and *Sgk3* (ESM Fig. 2k, l) as well as of all three *Akt* isoforms (ESM Fig. 2f–h) were not significantly altered by dexamethasone.

These findings suggest that increased FOXO1 protein is a result of specific glucocorticoid-receptor-mediated activation of transcription. FOXO1 phosphorylation and its cytosolic retention could be regulated by the action of multiple AKT and SGK isoforms, of which the most abundant are SGK1 and AKT1.

Phosphorylation of FOXO1 depended on AKT1, AKT2 and AKT3 To examine whether the discrepancy between dexamethasone-induced inhibition of AKT phosphorylation and maintenance of FOXO1 phosphorylation could be explained by effects of distinct AKT isoforms, isoform-specific siRNAs were used to inhibit selectively *Akt1*, *Akt2* and *Akt3*. Inhibition of the synthesis of individual AKT isoforms did not alter phosphorylation of FOXO1 (ESM Fig. 4a). Phosphorylation of FOXO1 persisted even when the synthesis of two AKT isoforms (AKT1 and AKT2 or AKT1 and AKT3 or AKT2 and AKT3) was simultaneously inhibited (ESM Fig. 4b). The individual mRNA levels were reduced to $16.0 \pm 1.1\%$ for *Akt1*, $21.1 \pm 2.2\%$ for *Akt2* and $18.4 \pm 1.5\%$ for *Akt3* (ESM Table 3). Immunostaining of FOXO1 remained prominent in the cytosolic compartment, confirming phosphorylation of FOXO1 (Fig. 5a). Finally, the treatment of INS-1E cells with siRNA against all three *Akt* isoforms, which resulted in an 80–90% reduction of their transcript levels, significantly inhibited phosphorylation of FOXO1 (Fig. 5b) and increased its nuclear staining in control INS-1E cells (Fig. 5e). The

dexamethasone-mediated increase in FOXO1 level was not affected by siRNAs against all *Akt* isoforms (Fig. 5c), whereas phosphorylation of FOXO1 was significantly reduced but not abrogated (Fig. 5d). In conclusion, all three isoforms, AKT1, AKT2 and AKT3, phosphorylate and facilitate nuclear extrusion of FOXO1 and, via this, mediate the inhibition of its transcriptional activity in insulin-secreting cells.

FOXO1 phosphorylation was not sensitive to SGK1 abundance or activity Dexamethasone treatment induces *Sgk1* expression and this kinase could contribute to FOXO1 inhibition in cells exposed to glucocorticoid excess. The role of SGK1 in the regulation of FOXO1 was clarified by specifically changing the activity and production of SGK1. Neither the inhibition of SGK1 activity by SGK1i, 10 μ mol/l, nor the prevention of dexamethasone-induced upregulation of SGK1 with siRNA reduced phosphorylation of FOXO1 at Ser256 and Ser319 (ESM Fig. 5a, b). In accordance, neither the overexpression of kinase dead *Sgk1* (*Sgk1*KN) nor the overexpression of constitutively active *Sgk1* (*Sgk1*SD) affected phosphorylation of FOXO1 at Ser256 (ESM Fig. 5c). The analysis of the cellular distribution of FOXO1 by confocal microscopy confirmed that in cells treated with dexamethasone and SGK1i, FOXO1 remained largely cytosolic (ESM Fig. 5d). These observations suggest that SGK1 does not account for cytosolic retention of FOXO1 in insulin-secreting cells under conditions of glucocorticoid excess.

In summary, beta cell death induced by glucocorticoids occurs in the absence of substantial activation of FOXO1, but is amplified when the overall activity of AKT is sufficiently reduced, resulting in the nuclear translocation and activation of FOXO1.

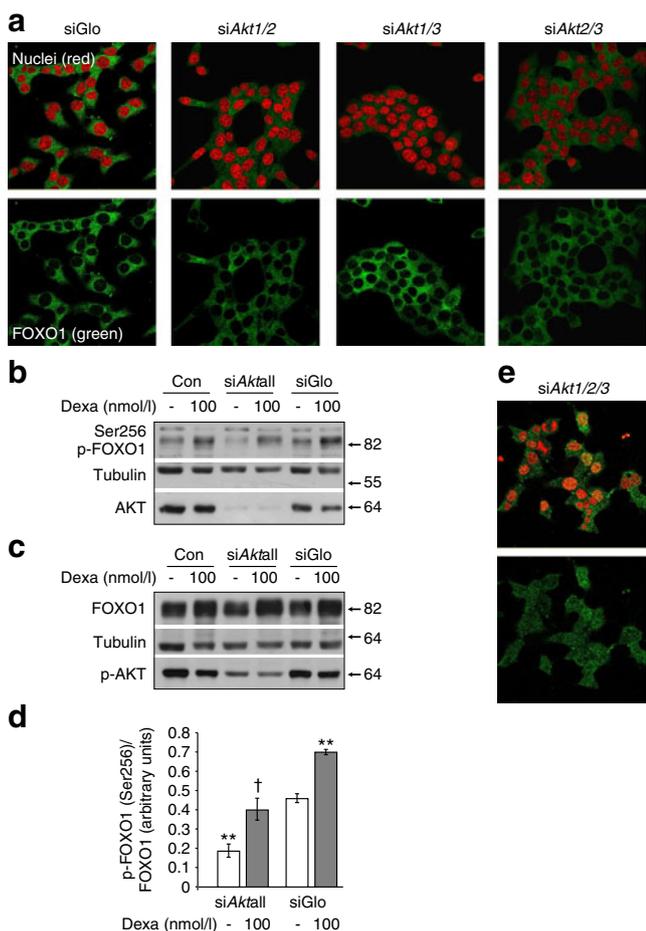


Fig. 5 Phosphorylation and nuclear accumulation of FOXO1 is dependent on AKT1, AKT2 and AKT3. (a–e) INS-1E cells were treated with siRNA as indicated and cultured under standard culture conditions (a, e) and in the presence of dexamethasone, 100 nmol/l for 1 day. (b–d) Representative pictures of immunostaining of FOXO1 (green) using confocal microscopy. Nuclei are stained in red. (a, e) Representative western blots of three independent experiments for p-FOXO1 (Ser256) (b) and FOXO1 (c) protein are shown. Tubulin was used as loading control. (d) The ratio of p-FOXO1 (Ser256) to FOXO1 protein is presented as mean ± SEM of n=3 independent experiments. **p<0.01 denotes significance compared with siGlo under control conditions; †p<0.05 denotes significance compared with siGlo treated with dexamethasone. Dexa, dexamethasone; siAktall, cells transfected with siRNA against *Akt1/Akt2/Akt3* isoforms

Discussion

This study shows that glucocorticoids induce beta cell death independently of FOXO1, as downregulation of FOXO1 using the siRNA strategy did not reduce the rate of cell death induced by dexamethasone. The detailed analysis of post-translational phosphorylation and subcellular distribution of FOXO1 suggests that FOXO1 remained phosphorylated and was largely localised in the cytosol even when apoptosis was significantly increased i.e. after 24 h exposure of INS-1E cells to dexamethasone. This observation is even more surprising

as dexamethasone increased the expression of *Foxo1*, confirming a previous study with INS-1 cells showing that dexamethasone increases, while prolactin decreases, the transcript levels of *Foxo1* [27]. Increased protein levels in combination with reduced phosphorylation and activation of FOXO1 after dexamethasone exposure has been described in a variety of cells, including hepatocytes, cardiomyocytes and tenocytes [28–30]. Interestingly, in tenocytes, dexamethasone-mediated inhibition of FOXO1 phosphorylation was counteracted by insulin. Previously, we described that IGF-1 reduces dexamethasone-induced cell death [16]. Thus, it is reasonable to speculate that insulin, which is secreted by INS-1E cells into the culture medium, may be responsible for sufficient activation of AKT and the subsequent phosphorylation and cytosolic localisation of FOXO1. Subcellular fractionation confirmed that the amount of FOXO1 increased in the cytosol after dexamethasone treatment. Intriguingly, the amount of FOXO1 in the nucleus was also increased when compared with untreated cells. There is a possibility that nuclear FOXO1 is not pro-apoptotic, as our results suggest that induction of apoptosis by dexamethasone occurs independently of FOXO1 [6].

Dexamethasone-induced cell death was significant even in the presence of secreted insulin, indicating that pathways other than FOXO1 signalling trigger apoptosis. Indeed, increased *Bim* expression induced by dexamethasone was not dependent on FOXO1 (Fig. 6). In fact, in lymphocytes, glucocorticoid-induced cell death was reversed by knock-down of BCL2-like 11 (apoptosis facilitator) (BIM) [31].

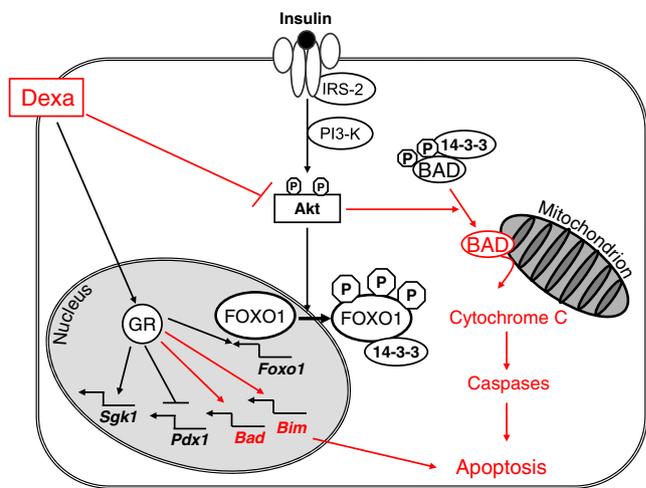


Fig. 6 Dexamethasone-induced beta cell death is not initiated by activation of FOXO1. Dexamethasone increases *Sgk1*, *Foxo1*, *Bim* and *Bad* expression through glucocorticoid-receptor activation and concomitantly reduces *Pdx1* mRNA. The reduced phosphorylation of AKT under excess of glucocorticoids does not activate FOXO1-dependent apoptosis but reduces BAD phosphorylation [11], indicating that dexamethasone promotes beta cell death directly through changes of mitochondrial BCL-2 proteins. Dexa, dexamethasone; GR, glucocorticoid receptor; PI3K, phosphatidylinositide 3-kinase

Our previous study revealed dexamethasone-dependent dephosphorylation of the pro-apoptotic BCL-2 protein BAD, which is a substrate of AKT [11]. On dephosphorylation BAD translocates to mitochondria, accomplishing an essential step of mitochondrial-linked apoptotic cell death [32]. In view of reduced AKT phosphorylation in dexamethasone-treated INS-1E cells, BAD seems to be more sensitive towards inhibition of AKT than FOXO1 (Fig. 6). An 80% reduction in a single or two *Akt* isoforms by siRNA did not impair FOXO1 phosphorylation. Quantification of relative *Akt* mRNA levels indicates that 20% of total AKT activity is sufficient to maintain FOXO1 in a phosphorylated state. The concomitant knockdown of all three AKT isoforms, AKT1, AKT2 and AKT3, finally suggests that each AKT isoform is able to compensate for the loss of AKT activity. These results were confirmed by the pan AKT inhibitor Akti-1/2, which effectively dephosphorylated FOXO1, leading to its nuclear accumulation. The addition of Akti-1/2 for 2 h to cells which were pre-exposed to dexamethasone for 22 h potentiated beta cell death, indicating additive effects through distinct pathways. A synergistic action of glucocorticoid receptor and FOXO1 has been described in muscle cells, pointing towards an independent activation of these pathways [33]. Here, we found that interaction of glucocorticoid receptor and FOXO1 leads to a potentiation of *Bim* expression, promoting mitochondrial-dependent apoptosis. During the development of diabetes such a synergistic effect could be induced by NEFA as they stimulate nuclear translocation of FOXO1 [34, 35].

The second surprising finding is that SGK1 does not contribute to phosphorylation of FOXO1 under glucocorticoid excess. This is contradictory to previous studies, in which SGK1 was found to phosphorylate FOXO1 at Ser256 and Ser319 [36, 37]. The expression of *Sgk1* is highly regulated in insulin-secreting cells and glucocorticoids augment mRNA and protein of SGK1 up to 20–50-fold [18]. However, downregulation of SGK1 by siRNA along with dexamethasone treatment did not reduce the phosphorylation of FOXO1. Moreover, overexpression of a constitutively active or inactive SGK1 (SGK1SD and SGK1KN) did not affect FOXO1 phosphorylation. The commercially available inhibitor was also ineffective. These observations underline that AKT isoforms are the main regulators of FOXO1 in insulin-secreting cells.

The dissociation between cell death and FOXO1 activation in insulin-secreting cells became obvious when phosphorylation and subcellular distribution of the transcription factor were simultaneously analysed. Our study unfolds an important molecular mechanism which determines subcellular localisation of FOXO1. Nuclear activity of FOXO1 depends on additional modifications such as acetylation [38]. Further studies are needed to understand the regulation of nuclear FOXO1 activity. Concerning the role of FOXO1 in beta cells, a significant study has recently been performed with beta cell-specific

deletion of FOXO1 in mice [39]. This study, published by Talchai and coworkers, shows that FOXO1 regulates differentiation of pancreatic endocrine cells rather than apoptosis. In addition, our results demonstrate the unique and promiscuous role of AKT isoforms for FOXO1 phosphorylation and reveal that AKT activity is decisive for beta cell survival.

Glucocorticoids regulate physiological blood glucose concentration by mobilising glucose from hepatocytes [40]. An excess of glucocorticoids, however, as during immunosuppressive drug treatment, leads frequently to the development of hyperglycaemia and steroid diabetes [1]. Apart from the induction of insulin resistance, glucocorticoids compromise beta cell function and induce beta cell death. The present study sheds light on the fact that glucocorticoids prime the cells for the cytotoxic effects seen on impairment of AKT signalling while denying the involvement of FOXO1 activation in glucocorticoid-induced beta cell apoptosis.

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Duality of interest The authors declare that there is no duality of interest associated with this manuscript.

Contribution statement GK and FG performed experiments, analysed data, wrote and revised the manuscript. DM, SB and BF performed experiments and analysed data and revised the manuscript; NS-S generated the SGK1 plasmids and designed experiments and revised the manuscript. FL and H-UH contributed to the design of experiments, interpretation of results and revised the manuscript. SU designed the study, analysed data and drafted, revised and edited the manuscript. All authors approved the final version of the manuscript.

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