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Direct regulation of insulin secretion by angiotensin II in human islets of Langerhans

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Abstract *Aims/hypothesis:* This study aimed to identify the expression of angiotensin II receptors in isolated human islets and beta cells and to examine the functional consequences of their activation. *Materials and methods:* Single-cell RT-PCR was used to identify whether human islet cells express mRNA for type 1 angiotensin II receptors (AT₁), and western blotting was used to determine AT₁ protein expression by human islets and MIN6 beta cells. We measured changes in intracellular calcium by microfluorimetry using Fura 2-loaded MIN6 cells and human islet cells. Dynamic insulin secretory responses were determined by RIA following perfusion of human islets and MIN6 cells. *Results:* Human islets expressed mRNAs for both the angiotensin precursor, angiotensinogen, and for angiotensin-converting enzyme. In addition, human and mouse beta cells expressed AT₁. These were functionally coupled to increases in intracellular calcium, which occurred at least in part through phospholipase-C-sensitive mechanisms and calcium influx through voltage-operated calcium channels.

Short-term exposure of human islets and MIN6 cells to angiotensin II caused a rapid, short-lived initiation of insulin secretion at 2 mmol/l glucose and potentiation of insulin secretion induced by glucose (at 8 and 16.7 mmol/l). *Conclusions/interpretation:* These data demonstrate that the AT₁ is expressed by beta cells and that angiotensin II effects a short-lived and direct stimulation of human and mouse beta cells to promote insulin secretion, most probably through elevations in intracellular calcium. Locally produced angiotensin II may be important in regulating a coordinated insulin secretory response from beta cells.

Keywords Angiotensin II · Calcium microfluorimetry · Human islets · Insulin secretion · MIN6 cells

Abbreviations AGT: Angiotensinogen · Ang II: Angiotensin II · AT₁: Type 1 Ang II receptor · AT₂: Type 2 Ang II receptor · [Ca²⁺]_i: Intracellular calcium concentration · HT: Hypertension · IP₃: Inositol trisphosphate · PLC: Phospholipase C · RAS: Renin–angiotensin system

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Introduction

Due to its vasoconstrictive effects, the systemic renin–angiotensin system (RAS) is classically linked to the regulation of blood pressure and fluid and electrolyte balance [1]. The systemic RAS is composed of several components including the precursor angiotensinogen (AGT), angiotensin II (Ang II) and multiple receptor subtypes [2], linked by alternative, non-exclusive components such as the enzymes renin and ACE [3, 4]. Ang II induces its effects via at least two G-protein-coupled transmembrane receptors, Ang II receptor 1 (AT₁) and Ang II receptor 2 (AT₂), of which the former is considered to be more physiologically important. Thus, AT₁ receptors are expressed abundantly in many adult tissues, but expression is limited in neonatal tissues [1], whereas AT₂ receptors are widely expressed in embryonic and neonatal tissues, but only in a few adult tissues [5, 6].

There is evidence of local angiotensin-generating systems in several organs, including the brain, pituitary, heart, kidney, gonads and pancreas (see [7]). An intrinsic pancreatic angiotensin-generating system was demonstrated in the canine pancreas [8, 9] by localisation of mRNA and protein for AGT and Ang II receptors, and subsequent studies have identified RAS components in rat, mouse and primate pancreas [7, 10, 11]. The highest densities of pancreatic Ang II receptors are reported to be localised to pancreatic ducts, vascular endothelium and acinar cells [10]. Although it has recently been reported that AT₁ receptors are localised to mouse pancreatic beta cells [11], there is currently no direct evidence of AT₁ receptor expression by human islet beta cells, nor of the effect of exogenous Ang II on human islet function.

The existence of an intrinsic pancreatic RAS may have pathophysiological implications, since type 2 diabetes is often associated with essential hypertension [12, 13], and hypertension is a significant risk factor for the development of type 2 diabetes [13]. Moreover, there is some evidence that ACE inhibition or Ang II receptor blockade protects from the development of glucose intolerance in hypertensive patients [14–16], and experimental manipulation of systemic RAS in humans can influence insulin secretory responses [14–17]. However, it is difficult for *in vivo* studies or experiments using the whole pancreas to differentiate between effects on the pancreatic vasculature and direct effects on islet endocrine function, and the inhibitory effects of Ang II on insulin release *in vivo* or in the perfused pancreas have been attributed to its vasoconstrictive effects on islet blood flow [18, 19].

Studies using isolated islets or beta cells avoid the complications of vascular effects of RAS activation, but have produced conflicting results in rodent models, in which inhibitory effects [11] and the absence of any effect [20] on hormone secretion have been reported. In the present study we therefore used human islets, isolated human beta cells and a mouse beta cell line to investigate the expression of Ang II receptors and the functional consequences of their activation.

Subjects, materials and methods

Isolation and maintenance of human islets of Langerhans

Human islets were obtained from the King's College Hospital Islet Transplantation Unit (King's College Hospital, London, UK). Briefly, islets were isolated from multi-organ donors using novel density gradient media [21] after donors' relatives had specifically consented to the use of their relatives' pancreases for islet research, in the event that it was not used for clinical transplantation. This study was approved by the Ethics Committee of King's College Hospital NHS Trust. Over 90% of islets isolated by this procedure are viable and show purities of between 80 and 95% [21]. Islets were maintained (5% CO₂) in CMRL culture medium supplemented with 10% FCS, and

100 U/ml penicillin/0.1 mg/ml streptomycin for up to 3 days prior to experimental use, and they were then further purified under a dissecting microscope using a finely drawn capillary pipette.

Maintenance of MIN6 cells

MIN6 insulin-secreting cells were maintained at 37°C (5% CO₂) in DMEM supplemented with 10% FCS, 2 mmol/l glutamine and 100 U/ml penicillin with 0.1 mg/ml streptomycin. The medium was changed every 3 days and the monolayers were trypsinised (0.1% trypsin, 0.02% EDTA) for experiments when approximately 80% confluent. For measurements of insulin secretion, MIN6 cell suspensions were maintained for 7–8 days in tissue culture flasks pre-coated with 1% (w/v) bovine gelatin to allow the formation of islet-like clusters, since configuration of MIN6 cells in these three-dimensional structures markedly improves their secretory responsiveness [22]. All tissue culture reagents were purchased from Sigma Aldrich (Poole, Dorset, UK).

RNA isolation and RT-PCR

Isolated rat and human islets, HepG2 human liver carcinoma cells, MIN6 beta cells, and small pieces of mouse heart were lysed in 200 µl ice-cold lysis buffer (100 mmol/l Tris-HCl, pH 8.0, 500 mmol/l LiCl, 10 mmol/l EDTA, 0.1% [w/v] lithium dodecylsulphate, 5 mmol/l dithiothreitol) and mRNAs were isolated using a Dynabeads oligo (dT)₂₅ kit (Dyna, Oslo, Norway) according to the manufacturer's instructions. cDNAs were synthesised from each mRNA sample immediately, using Superscript II reverse transcriptase (200 U), oligo(dT)₁₈ (1 µg) and random 10-mers (1 µg) as previously described [23], and cDNAs were diluted ten-fold in tRNA (10 µg/ml) and stored at –85°C. Human kidney mesangial cells, which were used as positive control tissue for human AT₁ receptor expression, were obtained by collagenase digestion from donor kidneys that were unsuitable for transplantation [24] and mesangial cell cDNA was prepared as described previously [25]. Hot-start PCR amplification was performed using synthetic oligonucleotide primers specific for human AGT and ACE, and for human and rodent AT₁ receptors (Molecular Biology Unit, King's College, London, UK) in 40-cycle reactions. In all PCR experiments, the presence of possible contaminants was checked by control reactions in which amplification was performed in reaction mixtures without a DNA template. No products were detected under these circumstances.

RT-PCR amplification of mRNA isolated from single human islet cells was conducted using a modification of a previously described protocol [26]. Hand-picked human islets (100) were dispersed into cell suspensions by trypsinisation, pelleted by centrifugation (1000×g, 5 min), washed twice in PBS containing 1 mmol/l glucose and resuspended in 500 µl CMRL medium supplemented with 10% FCS. Extracellular mRNA released from damaged cells was digested by incubation (15 min, 37°C) in the

presence of 50 µg/ml RNase A, and individual islet cells were selected under an inverted phase-contrast microscope: each single cell was drawn into a microcapillary filled with PBS and washed by serial transfer through four separate drops of PBS. Each cell was further washed twice using a second microcapillary, then transferred to a 0.2-ml PCR tube containing 5 µl lysis buffer composed of 0.8% Nonidet P-40, 100 µg/ml yeast tRNA, 40 U rRNasin and 10 mmol/l dithiothreitol. To ensure that RT-PCR products were not derived from nucleic acids released from damaged cells, 0.5 µl of the last wash drop of PBS was also used as a template for reverse transcription. Reverse transcription was performed (50 min, 42°C) using 50 pmol oligo(dT)₁₈, 100 pmol of the outer downstream primers, 200 U MMLV reverse transcriptase, 500 µmol/l of each dNTP and 10 mmol/l dithiothreitol in a final volume of 20 µl. AT₁ receptor and preproinsulin, proglucagon and preprosomatostatin cDNAs were amplified separately from 5 µl of the products of the RT reaction by addition of 40 µl of PCR reaction mixture containing 0.5 µmol/l of the specific outer upstream primers and 2.5 U Taq polymerase. The first-round PCR was performed for 45 cycles, after which 2 µl of the PCR product were added to 28 µl of a second PCR reaction mixture containing 125 µmol/l of each dNTP, 1 µmol/l of each inner primer and 1.5 U of Taq polymerase, and the second amplification was performed for 40 cycles. In single-cell PCR experiments, control reactions, in which mRNA was used instead of a DNA template, were used to check that genomic contamination could not contribute to product synthesis.

Primer sequences, MgCl₂ concentrations and PCR parameters for AGT, ACE, AT₁ receptor, proglucagon, preproinsulin and preprosomatostatin amplification are shown in Table 1. The PCR products were separated on 2% agarose gels containing ethidium bromide (10 µg/ml).

The products were excised from the gels using the Qiagen Gel Extraction kit (Qiagen, West Sussex, UK), according to the manufacturer's protocol, and their identities were confirmed by sequencing using fluorescent di-deoxy termination methods (Molecular Biology Unit, King's College London).

Protein extraction and western blotting

MIN6 cells were detached from tissue culture flasks using 0.02% (w/v) EDTA solution to avoid proteolytic damage to cell surface proteins. Extracts of MIN6 cells and of mouse heart and kidney and of human islets and kidney mesangial cells were prepared in lysis buffer, protein contents were determined by the Bradford assay [27] and proteins were fractionated by electrophoresis on 10% polyacrylamide Bis-Tris gels (Invitrogen, Paisley, UK). Proteins were transferred to polyvinylidene fluoride membranes and AT₁ receptor expression was identified by immunoprobings with 0.08 µg/ml of a rabbit polyclonal antibody raised against a peptide mapping near the amino terminus of the AT₁ receptor (Autogen Bioclear, Calne, UK) and a 1:5,000 dilution of horseradish-peroxidase-linked goat anti-rabbit secondary antibody (DakoCytomation, Cambs, UK). Blots were developed using the ECL Western Blotting Detection Kit (GE Healthcare, Bucks, UK) and standard curves were constructed of distances migrated by Rainbow Molecular Mass Markers (GE Healthcare).

Single-cell calcium microfluorimetry

Hand-picked human islets isolated with Liberase (Roche Diagnostics, East Sussex, UK) were gently dissociated by

Table 1 Primer sequences, annealing temperatures and MgCl₂ concentrations for amplification of rodent and human angiotensin type 1 receptor (AT₁) cDNAs, and human angiotensin converting enzyme (ACE), angiotensinogen (AGT), proglucagon (PPG), preproinsulin (PPI) and preprosomatostatin (PPS) cDNAs

cDNA amplified	Primer sequence	Annealing temp. (°C)	MgCl ₂ (mmol/l)
Rodent AT ₁ receptor	(F): 5'-cagcttggtggtgattgctc-3' (R): 5'-gccatcggtattccatagc-3'	52	1.0
Human ACE	(F): 5'-caccacagagaccagcaaga-3' (R): 5'-tcgggtaaaactggaggatg-3'	60	1.5
Human AGT	(F): 5'-gaactggatgttgctgctga-3' (R): 5'-agaggcatagtgaggctgga-3'	60	1.5
Human AT ₁ receptor			
Outer	(F): 5'-ccttcgacgcacaatgcttg-3' (R): 5'-agccctatcggagggttga-3'	54	2.0
Inner	(F): 5'-ccaaagtcacctgcatcac-3' (R): 5'-gggttgaatttgggactca-3'	60	1.5
Human PPI			
Outer	(F): 5'-ccctctggggactgacc-3' (R): 5'-acaatgccacgettctgc-3'	56	1.5
Inner	(F): 5'-aacgagcttctctacacac-3' (R): 5'-ggtacagcattgtccaca-3'	56	1.5
Human PPG			
Outer	(F): 5'-ccaggcagaccactcag-3' (R): 5'-tcaacaatggcgacctc-3'	56	1.5
Inner	(F): 5'-cattcacaggcacattcac-3' (R): 5'-gcttggcctccaataag-3'	56	1.5
Human PPS			
Outer	(F): 5'-cccagactccgtcagtttc-3' (R): 5'-gcctcatttcctctgctc-3'	56	1.5
Inner	(F): 5'-gactccgtcagtttctgca-3' (R): 5'-gcatcatttcctctggtt-3'	56	1.5

trituration in a 0.02% EDTA (w/v) solution and cells were seeded onto poly-D-lysine-coated glass coverslips, and MIN6 cells were trypsinised and seeded onto 3-aminopropyltriethoxysilane-coated glass coverslips. After maintenance overnight under standard tissue culture conditions, human islet cells and MIN6 cells were loaded for 30 min at 37°C with 2.5 $\mu\text{mol/l}$ of the Ca^{2+} -fluorophore Fura-2/AM. The coverslips were placed in a steel chamber that was mounted into a heating platform on the stage of an Axiovert 135 Research Inverted microscope (Carl Zeiss, Herts, UK) and changes in intracellular Ca^{2+} were determined as described previously [21, 28].

Dynamic measurement of insulin secretion

Groups of 100 Liberase-isolated human islets or approximately 1,000 MIN6 cell clusters were transferred to Swinnex chambers (Millipore UK, Watford, UK) containing 1- μm pore-size nylon filters and perfused at a flow rate of 0.5 ml/min at 37°C in a temperature-controlled environment. Perfusate samples were collected every 2 min for the duration of the experiments and insulin secretion was determined by RIA [29].

Data analysis

Mean changes in cytosolic calcium are expressed as 340/380 nm ratiometric data \pm SEM or as basal-to-peak fluorescence \pm SEM. Insulin secretion data are expressed as a percentage of basal insulin secretion at 2 mmol/l glucose. Student's *t*-tests were used to assess statistical significance between treatments: differences were considered statistically significant at $p < 0.05$.

Results

AGT, ACE and AT₁ receptor mRNA expression

PCR amplifications using primers for human *AGT*, the precursor of angiotensin I, produced a single product of 327 bp from human islet cDNA, as well as from the HepG2 human liver cell line, which was used as a positive control (Fig. 1a). In addition, human islets also expressed mRNA for ACE, the enzyme responsible for conversion of angiotensin I to angiotensin II (Fig. 1b). Reactions using rodent and human AT₁ receptor primers produced a single product of 146 bp from mouse islet cDNA, as well as from

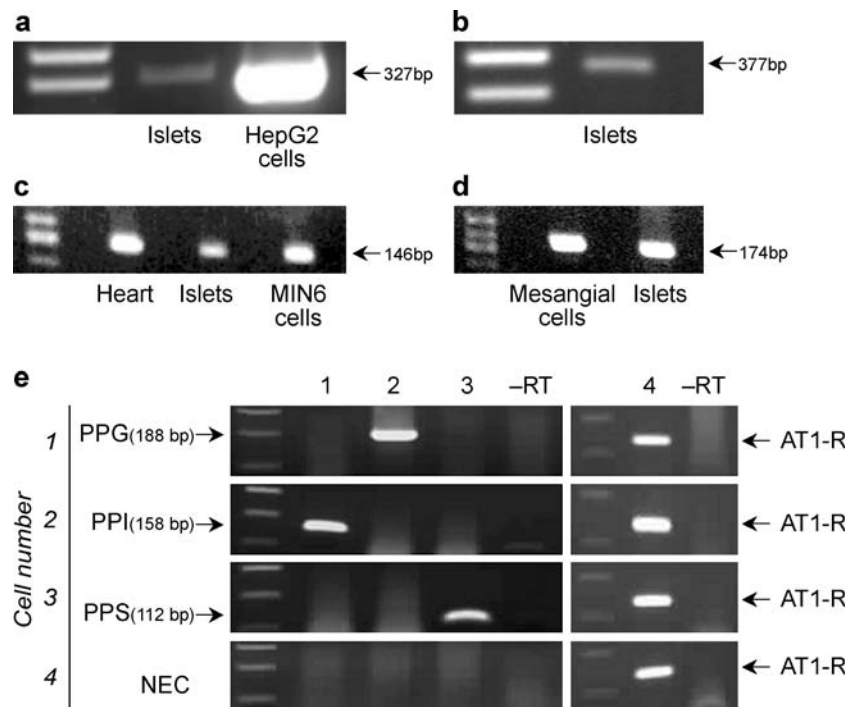


Fig. 1 Angiotensinogen, angiotensin-converting enzyme and AT₁ receptor mRNA expression. Standard RT-PCR was performed for human angiotensinogen (a), human angiotensin-converting enzyme (b), rodent AT₁ receptor (c) and human AT₁ receptor (d) using RNA extracted from human islets (a, b, d), HepG2 human liver cells (a), mouse heart, mouse islets and MIN6 cells (c) and from human kidney mesangial cells (d). e PCR amplifications from single dispersed human islet alpha, beta or delta cells or from non-

endocrine cells (NEC). The islet endocrine cells were identified by reactions using primers for preproinsulin (PPI, lane 1), preproglucagon (PPG, lane 2) and preprosomatostatin (PPS, lane 3). Cells that did not express PPG, PPI or PPS were designated NECs (row 4). Lane 4 shows AT₁ receptor (*AT1-R*) mRNA expression. -RT indicates the absence of product formation when using an mRNA template in the absence of reverse transcriptase

mouse heart cDNA, which is known to express the AT₁ receptor subtype (Fig. 1c). A similar product was also amplified from MIN6 cell cDNA (Fig. 1c), demonstrating the expression of AT₁ receptor mRNA in a pure beta cell population. A single PCR product of 174 bp was amplified from human islet cDNA and also from human kidney mesangial cell cDNA, which is known to express this receptor subtype (Fig. 1d). Single-cell PCR reactions identified AT₁ receptor mRNA expression in human beta cells, and also in human alpha and delta cells, as shown by the co-expression of mRNAs for preproinsulin, preproglucagon and preprosomatostatin and the AT₁ receptor (Fig. 1e). Human islet cells that did not express mRNAs for preproinsulin, preproglucagon or preprosomatostatin were designated 'non-endocrine cells', and AT₁ receptor mRNA was also amplified from these cells (Fig. 1e). In all, AT₁ receptor mRNA was amplified from one of five alpha cells, four of six beta cells, two of five delta cells and two of seven non-endocrine cells. Purification and sequencing of rodent and human PCR reaction products indicated that they showed 100% homology with the expected AT₁ receptor cDNAs.

AT₁ receptor protein expression

Expression of AT₁ receptors at the protein level was identified by western blotting using a polyclonal anti-AT₁ receptor antibody. A major immunoreactive protein of ~42.5 kDa was detected in mouse kidney and heart samples, as expected (Fig. 2a). A protein of the appropriate size was also detected in MIN6 cell extracts, indicating that the AT₁ receptor protein is localised to mouse pancreatic beta cells. Western blot analysis with the human mesangial and islet cell extracts revealed a major immunoreactive protein of ~42.5 kDa, corresponding to the AT₁ receptor (Fig. 2b).

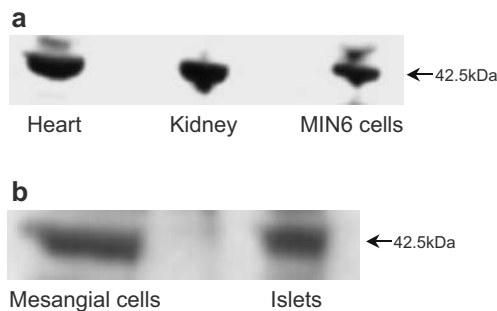


Fig. 2 AT₁ receptor protein expression. Western blotting was performed on fractionated samples of mouse heart, kidney and MIN6 cells (a) and human kidney mesangial cells and human islets (b) using a polyclonal antibody directed against the AT₁ receptor and a horseradish-peroxidase-labelled secondary antibody

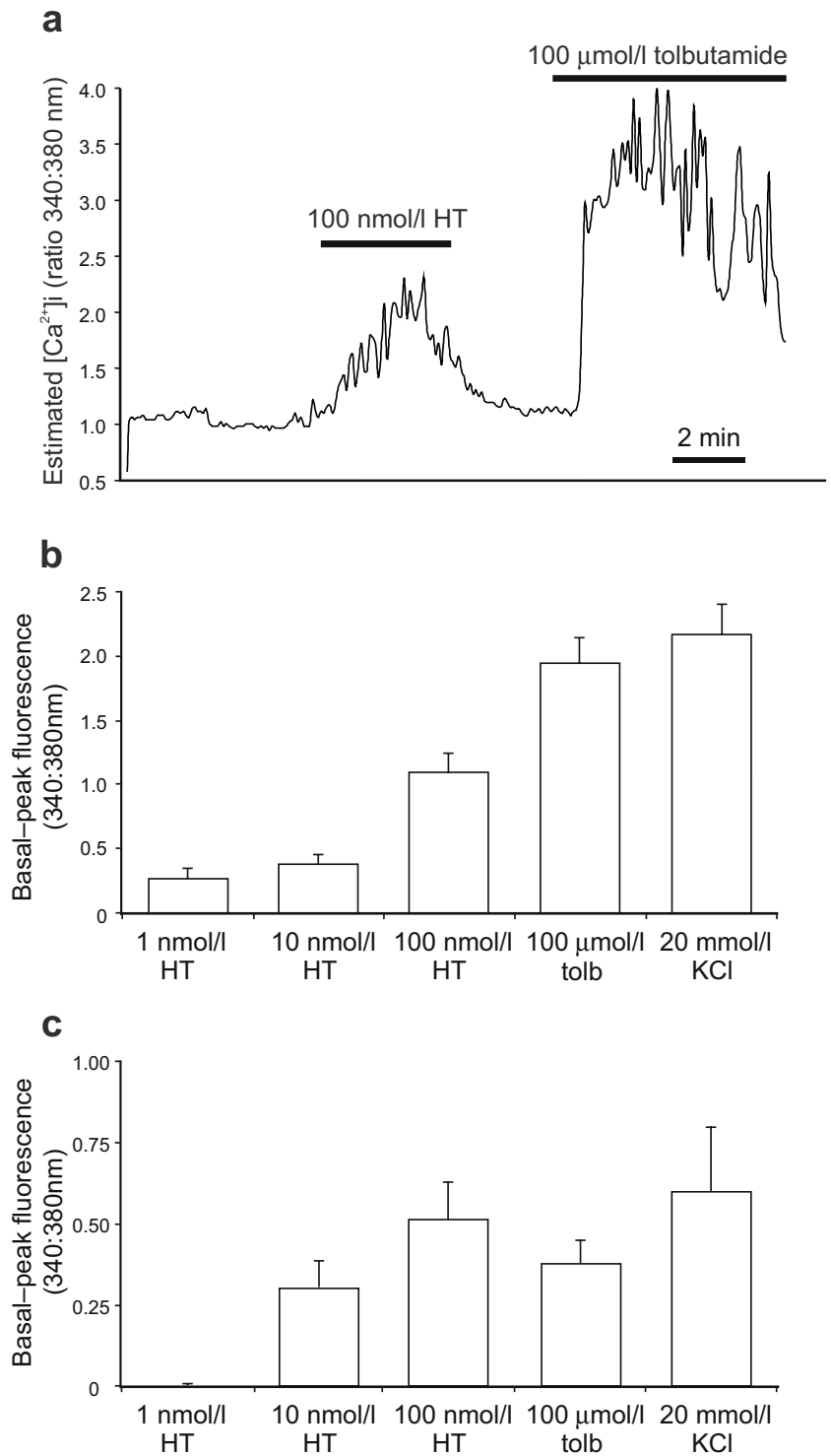
Effects of Ang II receptor activation on intracellular calcium concentration

The functional consequences of Ang II receptor activation were initially investigated by examining the effects of the stable Ang II analogue hypertensin (HT) on changes in intracellular calcium concentrations ($[Ca^{2+}]_i$) in MIN6 cells and dispersed human islet cells. As shown in Fig. 3a, perfusion of Fura-2-loaded MIN6 cells with the Ang II amide HT (100 nmol/l) evoked a rapid elevation in $[Ca^{2+}]_i$, with a mean basal-to-peak amplitude that was ~50% of the response evoked by 100 μ mol/l tolbutamide, a blocker of ATP-sensitive K⁺ channels. Figure 3b shows the cumulative results of several experiments, demonstrating that the effects of HT were concentration dependent, with 22% of MIN6 cells responding to 1 nmol/l HT, increasing to 44% at 10 nmol/l HT. A further tenfold increase in HT (100 nmol/l), produced a small increase in the proportion of cells showing a HT-evoked change in $[Ca^{2+}]_i$ (62%; 55 of 89 cells from five separate experiments), and caused a significant ($p < 0.001$) increase in the amplitude of the response. As expected, MIN6 cells showed robust increases in $[Ca^{2+}]_i$ in response to either tolbutamide (100 μ mol/l) or direct depolarisation by KCl (20 mmol/l, Fig. 3b). Similar effects were observed in experiments using human beta cells, as shown in Fig. 3c. Tolbutamide (100 μ mol/l) and KCl (20 mmol/l) increased $[Ca^{2+}]_i$ in human islet cells, and the tolbutamide-responsive cells were considered to be beta cells and were subsequently challenged with HT (1, 10 and 100 nmol/l). In all, 67% of tolbutamide-responsive cells showed an increase in cytosolic Ca^{2+} in response to 10 nmol/l HT, while all tolbutamide-responsive cells showed an increase in $[Ca^{2+}]_i$ at 100 nmol/l HT (Fig. 3c).

A non-selective Ang II receptor antagonist (saralasin) and an AT₂ receptor-specific antagonist (PD123319) were used to identify the Ang II receptor subtype involved in the effects of HT on $[Ca^{2+}]_i$ in beta cells. As shown in Fig. 4a, the addition of 100 nmol/l HT in the presence of 1 μ mol/l saralasin resulted in only a small increase in $[Ca^{2+}]_i$. However, upon removal of saralasin, HT elicited a rapid and reversible increase in $[Ca^{2+}]_i$. Similar effects were seen in experiments using tolbutamide-sensitive dispersed human islet cells, as shown in Fig. 4b, demonstrating AT-receptor-mediated effects of HT on $[Ca^{2+}]_i$ in both primary human beta cells and MIN6 cells. In contrast, selective blockade of the AT₂ receptor in tolbutamide-sensitive human beta cells by the presence of PD123319 (1 μ mol/l) did not significantly reduce HT-stimulated increases in $[Ca^{2+}]_i$ (Fig. 4c).

HT-induced increases in $[Ca^{2+}]_i$ in MIN6 cells were partially reduced by the L-type Ca^{2+} channel blocker, nifedipine. Figure 5 shows that exposure of cells to nifedipine (10 μ mol/l) alone did not affect basal $[Ca^{2+}]_i$ and that HT (100 nmol/l) caused a marked elevation in $[Ca^{2+}]_i$ in the absence of nifedipine. However, the concomitant application of nifedipine (10 μ mol/l) significantly inhibited the

Fig. 3 Changes in intracellular calcium in Fura-2-loaded beta cells. **a** Trace from a single Fura-2-loaded MIN6 cell. Mean+SEM basal-peak fluorescence data for **b** MIN6 cells ($n=16$) and **c** human islet beta cells ($n=6$)



effects of HT on $[Ca^{2+}]_i$, although there was a sustained plateau above basal fluorescence. Subsequent removal of nifedipine resulted in an immediate and rapid increase in $[Ca^{2+}]_i$, with a more enhanced Ca^{2+} peak than had been obtained on the first exposure to HT. HT-induced increases in $[Ca^{2+}]_i$ in MIN6 cells were also partially inhibited by the

presence of the phospholipase C (PLC) inhibitor, U73122. Thus the HT-induced increase in Fura-2 fluorescence (100 nmol/l, basal-to-peak fluorescence ratio 0.90 ± 0.07) was significantly reduced in the same cells by the presence of U73122 (10 $\mu\text{mol/l}$, 0.50 ± 0.03 , $p < 0.01$, $n = 15$).

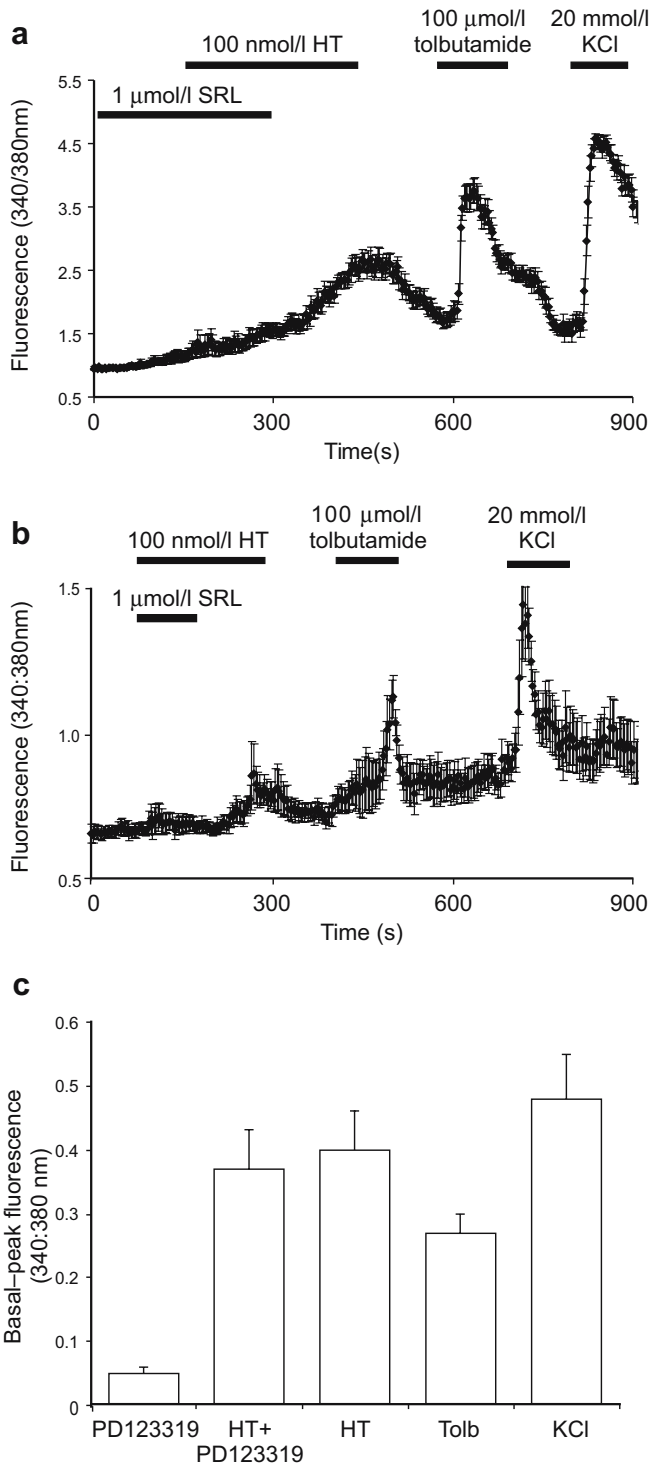


Fig. 4 Effect of AT receptor antagonists on intracellular calcium. Mean±SEM fluorescence traces from Fura-2-loaded MIN6 cells ($n=16$) (**a**) and from **b** human islet beta cells ($n=4$) showing the effects of hypertensin (HT) in the presence of saralasin (SRL). **c** Mean basal-peak fluorescence data (+SEM, $n=17-22$) of the effects of PD123319 in human islet beta cells

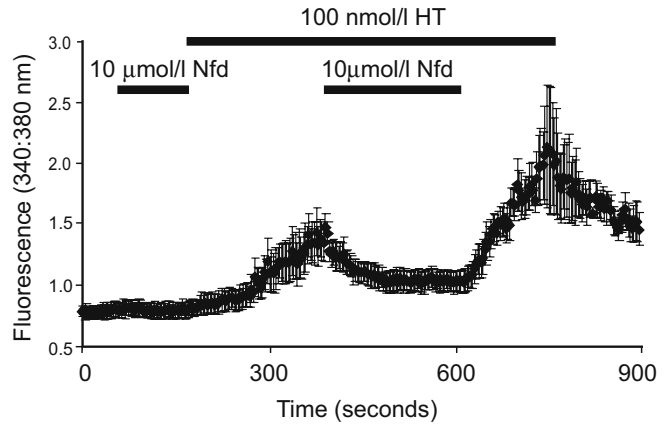
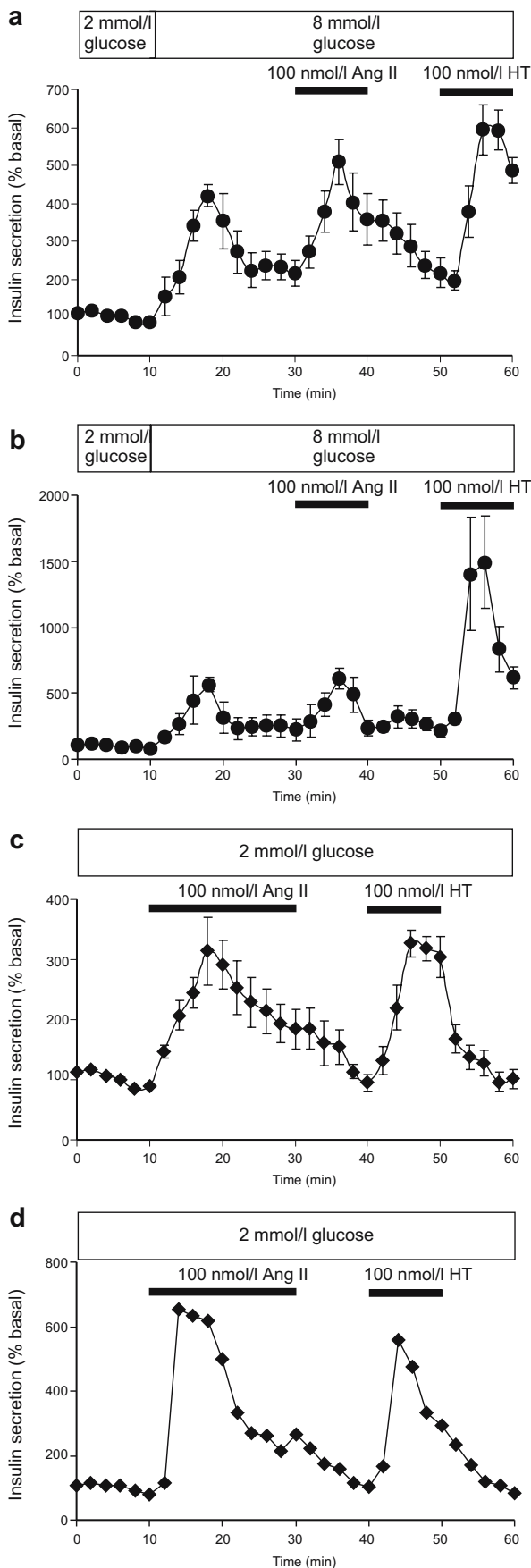


Fig. 5 Effect of nifedipine (Nfd) on intracellular calcium. Mean±SEM fluorescence traces from Fura-2-loaded MIN6 cells showing the effects of Nfd on hypertensin-stimulated increases in calcium ($n=4$)

Effects of Ang II receptor activation on insulin secretion

Perfusion experiments demonstrated that insulin secretion, as expected, was significantly ($p<0.001$) stimulated upon raising the glucose concentration from 2 to 8 mmol/l in isolated human islets (Fig. 6a) and MIN6 cell clusters (Fig. 6b). The insulin secretory response to glucose (8 mmol/l) was significantly ($p<0.05$) potentiated by the addition of Ang II (100 nmol/l) during the plateau phase of the glucose-induced response (Fig. 6a,b). The responses to Ang II were rapidly reversible on removal of Ang II from the perfusion medium, resulting in a restoration of insulin output to levels equivalent to the sustained secondary phase plateau of glucose-induced secretion. Both human islets and MIN6 cell clusters also responded to a subsequent exposure to HT (100 nmol/l) with a rapid increase in insulin secretion. Human islets showed similar responses to Ang II and HT (Fig. 6a), whereas glucose-induced secretion was enhanced to a greater extent by HT in the MIN6 clusters (Fig. 6b). In similar perfusion experiments an elevation in the glucose concentration from 2 to 16.7 mmol/l resulted in a maximal 10.6-fold increase in insulin secretion, followed by a secondary plateau phase of $187\pm64\%$ of basal. Ang II also significantly potentiated insulin secretion from human islets in the presence of 16.7 mmol/l glucose, further increasing secretion to $406\pm110\%$ of basal (means±SEM, $n=4$, $p<0.05$).

Activating the AT₁ receptor also exerted stimulatory effects on insulin secretion at a sub-stimulatory concentration of glucose (2 mmol/l). Thus, Ang II (100 nmol/l) provoked a rapid increase in insulin release from human islets (Fig. 6c) and from MIN6 cell clusters (Fig. 6d) and the secretory response was followed by a steady decline in secretion, despite the continued presence of Ang II. Secretion levels were restored to basal when islets and MIN6



◀ **Fig. 6** Insulin secretion from perfused human islets and MIN6 cell clusters. Isolated human islets (**a, c**) and MIN6 cell clusters (**b, d**) were perfused with media supplemented with the agents shown by the horizontal bars (*Ang II* angiotensin II; *HT* hypertensin) and insulin secretion was measured every 2 min by RIA. Data are from two to four separate perfusion chambers of islets from a single isolation and are expressed **a–c** as means \pm SEM ($n=4$) and **d** as means ($n=2$)

cell clusters were perfused with 2 mmol/l glucose in the absence of Ang II (time=30–40 min). Subsequent exposure to HT (100 nmol/l) also initiated a rapid and transient increase in basal insulin secretion, of similar amplitude to that obtained with 100 nmol/l Ang II (Fig. 6c,d). The effects of Ang II and HT were not additive, since simultaneous exposure to both agonists did not cause significant enhancement of the secretory response to the individual agonists (secretion, % 2 mmol/l glucose: 100 nmol/l Ang II: 195 ± 43 ; 100 nmol/l HT: 200 ± 59 ; 100 nmol/l Ang II+100 nmol/l HT: 224 ± 75 , means \pm SEM, $n=4$, NS). The presence of nifedipine (10 μ mol/l) reduced Ang II (100 nmol/l)-stimulated insulin secretion at 2 mmol/l glucose by $38.8\pm 8\%$ ($p<0.01$), indicating that the effects of Ang II were only partially dependent on Ca^{2+} influx through voltage-operated Ca^{2+} channels.

Discussion

In the systemic RAS, circulating renin cleaves Ang I from AGT, which is synthesised by the liver, and endothelial cells of the lung produce ACE to convert Ang I to Ang II. There is convincing evidence of an intrinsic pancreatic RAS, through which Ang II is produced locally [7–11, 30–32], but the physiological roles of this system remain unclear. The present results demonstrate unambiguously that human islets of Langerhans express both AGT and ACE, suggesting that the islets themselves may produce Ang II locally. Furthermore, receptors for Ang II were detected in human islets, and our use of single-cell RT-PCR to amplify mRNA from individual human islet cells demonstrates for the first time that primary human islet beta cells express the AT_1 receptor mRNA. Our identification of AT_1 receptor expression in human and mouse beta cells is consistent with previous reports suggesting expression of AT_1 receptors in dog [8], mouse [30], rat [31] and human [32] pancreas and in isolated mouse islets [11], and our study further indicates that alpha, delta and human islet non-endocrine cells also express AT_1 receptors. Our detection of AT_1 receptor mRNA and protein in human beta cells and mouse MIN6 cells is also consistent with our results using AT_1 receptor antagonists, which demonstrated the presence of a functional AT_1 receptor subtype coupled to elevations in beta cell $[Ca^{2+}]_i$, in agreement with previous pharmacological studies in which AT_1 receptor activation was associated with Ca^{2+} transients in a number of other tissues [33–37].

Plasma membrane Ca^{2+} channels, including L-type Ca^{2+} -channels, are known to play important roles in Ang II-induced cellular responses [38, 39], and our observation

that an L-type Ca^{2+} channel blocker reversibly inhibited Ang II-stimulated increases in beta cell $[\text{Ca}^{2+}]_i$ suggests that this mechanism is operational in beta cells. However, blockade of L-type Ca^{2+} channels was only partially effective in blocking AT_1 -induced increases in $[\text{Ca}^{2+}]_i$, suggesting the existence of another mechanism. Activation of the PLC–inositol trisphosphate (IP_3) signalling pathway is involved in the rapid Ang II-evoked changes in $[\text{Ca}^{2+}]_i$ in a number of other cell types [40–43]. The inhibition of HT-induced elevations in $[\text{Ca}^{2+}]_i$ in beta cells by U73122, a PLC inhibitor, suggests that the beta cell AT_1 receptor acts, at least in part, via IP_3 -mediated release of cytosolic Ca^{2+} from the endoplasmic reticulum. Our data therefore indicate that, in pancreatic beta cells, AT_1 receptor activation increases $[\text{Ca}^{2+}]_i$ both by promoting influx of extracellular Ca^{2+} and by the PLC-mediated IP_3 -sensitive release of Ca^{2+} release from intracellular stores.

Changes in $[\text{Ca}^{2+}]_i$ play a fundamental role in the regulation of insulin secretion, and the dual mode of action of AT_1 receptor activation on beta cell $[\text{Ca}^{2+}]_i$ is consistent with the concomitant stimulation of insulin secretion from human islets and MIN6 cells. Stimulus–response coupling in beta cells is complex, but it is generally agreed that the initiation of insulin secretion by nutrient stimuli, such as glucose, is normally dependent on an influx of extracellular Ca^{2+} through L-type Ca^{2+} channels in response to depolarisation. Our experiments suggest that AT_1 receptor-induced influx of extracellular Ca^{2+} is not the sole mechanism of action of Ang II on insulin secretion, since the increases in $[\text{Ca}^{2+}]_i$ and insulin secretion were not fully abolished by nifedipine. This is consistent with Ang II receptors being coupled to a PLC-mediated release of intracellular Ca^{2+} , and this mechanism could also explain the enhancement by Ang II of glucose-induced insulin secretion from human islets and MIN6 cells. Receptor agonists often potentiate rather than initiate insulin secretory responses, but we have recently reported that two other agonists (17beta-oestradiol and adrenocorticotrophin) stimulate Ca^{2+} -dependent insulin secretion from human islets at 2 mmol/l glucose [44, 45].

Irrespective of the coupling mechanism involved, our data clearly demonstrate that AT_1 receptor activation stimulates insulin secretion, in contrast to previous reports. It is difficult to dissect out direct effects on Ang II on pancreatic beta cells in *in vivo* studies, or in those using whole perfused pancreas, but previous studies using isolated rodent islets have reported either a lack of effect [20] or an inhibitory effect [11] on insulin secretion. An observed lack of effect of Ang II may be ascribed to the loss of functional cell-surface receptors during the collagenase digestion of rat pancreas during islet preparation [20], while the inhibitory effects in mouse islets were attributed to reduced (pro)insulin biosynthesis [11]. In common with other circulating bioactive peptides, Ang II has a short half-life and our experiments were specifically designed to measure immediate and rapid effects of AT_1 receptor activation on $[\text{Ca}^{2+}]_i$ and insulin secretion within a time-frame of minutes. Any inhibitory effects on insulin se-

cretion as a consequence of reduced (pro)insulin synthesis in response to prolonged AT_1 receptor activation would not be detected in our studies. Similarly, the present experiments were not designed to address the mechanisms underlying the reported protective effects of treatment with ACE inhibitors or Ang II receptor blockers on the incidence of type 2 diabetes [46–48], since these trials were based on many years of prolonged treatment, and the therapeutic effects may be secondary to blocking the pro-inflammatory effects of Ang II [49, 50]. Nonetheless, our study identifies the human beta cell as one potential target of pharmacological manipulation of the RAS.

The circulating concentration of Ang II is approximately 5 pmol/l in healthy individuals [51], considerably lower than the concentrations that stimulated insulin secretion in these studies. However, concentrations of locally produced Ang II would be expected to exceed systemic concentrations and may achieve the concentration range used in the current studies, although accurate quantification of tissue concentrations of Ang II is hampered by its short half-life [8, 11]. Our observations are therefore consistent with an intrinsic pancreatic RAS generating high, but localised, concentrations of Ang II, which has a short-lived and direct stimulatory effect on beta cells to promote insulin secretion. Cells within islets show coordinated responses [52], and it is an intriguing possibility that the local production of Ang II may be important in regulating a coordinated insulin secretory response from pancreatic beta cells.

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