Insulin acts via mitogen-activated protein kinase phosphorylation in rabbit blastocysts

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Abstract

The addition of insulin during *in vitro* culture has beneficial effects on rabbit preimplantation embryos leading to increased cell proliferation and reduced apoptosis. We have previously described the expression of the insulin receptor (IR) and the insulin-responsive glucose transporters (GLUT) 4 and 8 in rabbit preimplantation embryos. However, the effects of insulin on IR signaling and glucose metabolism have not been investigated in rabbit embryos. In the present study, the effects of 170 nM insulin on IR, GLUT4 and GLUT8 mRNA levels, Akt and Erk phosphorylation, GLUT4 translocation and methyl glucose transport were studied in cultured day 3 to day 6 rabbit embryos. Insulin stimulated phosphorylation of the mitogen-activated protein kinase (MAPK) Erk1/2 and levels of IR and GLUT4 mRNA, but not phosphorylation of the phosphatidylinositol 3-kinase-dependent protein kinase, Akt, GLUT8 mRNA levels, glucose uptake or GLUT4 translocation. Activation of the MAPK signaling pathway in the absence of GLUT4 translocation and of a glucose transport response suggest that in the rabbit preimplantation embryo insulin is acting as a growth factor rather than a component of glucose homeostatic control. *Reproduction* (2004) **128** 517–526

Introduction

Insulin and insulin-like growth factors (IGF)-I and -II mediate mitogenic, anti-apoptotic and anabolic effects in mammalian preimplantation embryos. Early embryos express all three receptor subtypes and the ligand for IGF-II. Whilst IGF-I expression has been demonstrated in mice, sheep and bovine embryos (Schultz et al. 1992, Kaye 1997), insulin is not expressed by any species studied to date. Insulin and IGF-I are found in the oviduct and uterine lumen during the pre- and peri-implantation period (Lighten et al. 1998, Chi et al. 2000). Thus, the embryo either expresses these growth factors itself or has access through genital tract secretions, implying that functional autocrine, paracrine and/or endocrine insulin/IGF circuits are operating during early development. The intracellular mechanisms by which the hormone signal is transduced following receptor binding and the induced cellular reactions, however, have not been studied in detail so far.

Whilst both the insulin receptor (IR) and IGF-I receptor (IGF-IR) are type 1 tyrosine kinase receptors, the type 2 IGF-II receptor (IGF-II-R) contains binding sites for a number of other ligands including mannose-6-phosphate and

retinoic acid, and has multiple functions depending on the binding domain that is activated. IGF-II binding to the type 2 receptor is thought to promote differentiative development in the early embryo (Pantaleon *et al.* 2003). Structural similarities between insulin and IGF-I allow each ligand to bind to both type 1 receptors but with a more than 100 times greater affinity for its homologous receptor. In rabbit embryos IR expression is first apparent in the early blastocyst (Navarrete Santos *et al.* 2004) whilst IGF-I binding is seen from the morula stage (Herrler *et al.* 1998). The rabbit IR sequence shows a high degree of identity to the human IR type A and to IR of various other mammalian species with conserved tyrosine phosphorylation sites (Navarrete Santos *et al.* 2004).

The IR/IGF-IR signaling involves two major pathways, the mitogenic-activated protein kinase (MAPK) and the phosphatidylinositol 3-kinase (PI3-K) pathway. The mitogenic response, mediated via the interaction of Grb2, SOS and RAS, leads to the activation of the extracellular signal-regulated kinase (Erk) which, in turn, regulates several transcriptional events involved in mitogenesis. Activated Erks, for example, mediate the growth-promoting effects of insulin by phosphorylating transcription factors such as ELK-1 (for review see Le Roith & Zick 2001).

Metabolic responses to insulin are primarily mediated via the PI3-K pathway. Following association of the p85/p110 complex of the PI3-K with insulin receptor substrate molecules, PI3-K activates phosphoinositol 3,4,5-phosphate (PIP₃), which binds to and activates the PI3-K-dependent kinase-1 (PDK-1) and Akt (protein serine/ threonine kinase B, also referred to as PKB). The stimulation of glucose transport, glycogenesis and protein synthesis are key metabolic effects of insulin mediated by the PI3-K/Akt pathway. Akt has been implicated in the translocation of glucose transporter (GLUT) 4 from cytoplasmic storage vesicles (Rea & James 1997) into the cell membrane in muscle and fat cells (Pessin *et al.* 1999, Patki *et al.* 2001).

Glucose is the main energy substrate during blastocyst formation and development in mammals. We have recently shown that two insulin-responsive GLUT isoforms, GLUT4 and GLUT8, are expressed in rabbit blastocysts in a developmentally regulated pattern (Navarrete Santos *et al.* 2004). These results prompted us to investigate insulin effects on IR signaling and glucose transport in preimplantation embryos of this species. We show that insulin acts on Erk phosphorylation and transcriptional regulation of GLUT4 and IR, but not on translocation of GLUT4 and stimulation of glucose transport in rabbit blastocysts.

Materials and Methods

Embryo recovery and in vitro culture

Embryos were collected from sexually mature rabbits stimulated with 100 I.U. follicle-stimulating hormone (Ovagen, Immuno-Chemical Products, Auckland, New Zealand, or Folligon, Intervet, Boxmeer, The Netherlands). Mating, embryo recovery (Navarrete Santos *et al.* 2000) and embryo culture (Kietz & Fischer 2003) were performed as described. On days 3, 4 and 6 post coitum (p.c.) embryos were flushed from oviducts or uteri, washed three times with PBS, pooled and randomly divided amongst the experimental groups.

To study the effects of insulin on IR and GLUT4 expression, day 6 blastocysts were cultured in groups of 10 in 500 µl BSM II medium at 37 °C in a saturated atmosphere of 5% O2, 5% CO2, 90% N2 (Lindenau & Fischer 1994) in a water-jacketed incubator (BB 6060, Heraeus, Hanau, Germany). Blastocysts were precultured for 2 h in serum- and insulin-free BSM II medium. Afterwards, 170 nM insulin (Invitrogen, Karlsruhe, Germany) was added to the culture medium and the culture continued for 1, 2, 3 or 4 h. Controls were cultured without insulin but otherwise were treated in an identical manner as the treated embryos. IR signaling was analyzed after culture of day 6 blastocysts in the presence or absence of insulin (170 nM, Invitrogen) for 10, 30, 60 or 120 min. For glucose transport studies, morulae (recovered at day 3 p.c.) and blastocysts (day 4) were cultured with or without 166 nM (morulae, blastocysts), 16 pM insulin (Humulin R, Lilly, North Ryde, Australia, diluted in M2 medium) or 1.3 nM IGF-I (Amersham Pharmacia Biotech, UK) (only day 4 blastocysts) for 1 h in 100 μ l droplets of glucose-containing M2 medium (Pantaleon *et al.* 1997) under oil. Three to nine blastocysts and 6 to 15 morulae were cultured per drop.

RNA extraction

Preparation of total RNA from embryos was performed using TRIzol reagent (Invitrogen) according to a previously described protocol (Koerber *et al.* 1998). RNA was treated with DNAse for 1 h. The amount of total RNA was determined spectrophotometrically at 260 nm.

Semiquantitative RT-PCR of IR, GLUT4 and GLUT8 in rabbit blastocysts

Semiquantitative RT-PCR was performed essentially as previously described (Navarrete Santos et al. 2000, Kietz & Fischer 2003) with oligonucleotide primers specific for rabbit IR, GLUT4, GLUT8, β-actin and GAPDH. The IR, GLUT4 and GLUT8 primer sequences were derived from published rabbit (rab) sequences (Navarrete Santos et al. 2004). Two micrograms total RNA were reverse transcribed in a volume of 20 µl containing 0.5 mM dNTPs, 10 mM dithiothreitol (DTT), 200 units superscript II, 20 units RNAse inhibitor (Roche Diagnostics, Mannheim, Germany), 1 µl random primer and 2 µl reverse transcriptase buffer at 42 °C for 1 h, followed by an incubation at 90 °C for 5 min. H_2O (20 µl) was added to each cDNA reaction. As a control for DNA contamination, 0.1 µg RNA was amplified without preceding reverse transcription. Amplification was carried out with 1 to 5 µl cDNA in a 50 µl volume containing 5 µl dNTP, 2.5 units Taq polymerase (Invitrogen), employing the primer combinations: rabIRp1 5'-GCTGGTGGTGATGGAGTTG-3[′]/rablRp2 5'-TCTCTCTGGACAGTTGTCGG-3'; rab-GLUT4p1 5'-GGCGGCATGATTTCCTCC-3'/rabGLUT4p2 5'-GAAGGGCAGCAGGATCAGCT-3'; and rabGLUT8p1 5'-CACGTCAAGGGTGTGGCT-3'/rabGLUT8p2 5'-CAGG-GACGCAGAACAAAGTG-3' for IR, GLUT4 and GLUT8 respectively. Primers for β-actin (actinp1 5'-CTACAATGA-GCTGCGTGTGG-3', actinp2 5'-TAGCTCTTCTCCAGGG-AGGA-3') and GAPDH (GAPDHp1 5'-CGGGAAGCTTGT-GATCAATGG-3', GAPDHp2 5'-GGCAGTGATGGCATGG-ACTG-3) were used to check DNA contamination and quantity of the reverse transcribed mRNA. For all PCR reactions the optimal amplification range was estimated by variation of cycle numbers and cDNA amount (IR: 1μ l cDNA, 35 cycles; GLUT4: 2 µl cDNA, 38 cycles; GAPDH: 1 µl cDNA, 35 cycles). The resulting PCR fragments for GLUT4 (398 bp), IR (497 bp) and GAPDH (507 bp) were separated by electrophoresis on a 1.8% agarose gel and visualized by ethidium bromide staining. Gels were photographed and the product bands were quantified by densitometric analysis employing the software BIO-Profile 1D (LTF-Labortechnik, Wasserburg, Germany). The relative amounts of IR and GLUT4 mRNA were calculated as a ratio of the specific product (for IR or GLUT4) and the housekeeping gene band volume (GAPDH) (Kietz & Fischer 2003). All PCR reactions were performed at least three times in two independent experiments.

Protein preparation

Blastocysts were washed three times with PBS after culture and transferred to a 1.5 ml Eppendorf tube. They were homogenized in 100 μ l cold RIPA buffer (PBS, 1% NP-40, 0.5% sodium deoxycholate, 0.1% sodium dodecyl sulfate (SDS)) with protease inhibitor cocktail (Sigma, St Louis, MO, USA). The samples were centrifuged at 5000 g for 10 min. The supernatant was stored at -80 °C until use. The total protein content was determined using the Bio-RAD Protein Assay (Bio-RAD, München, Germany).

Immunoblotting

Western blots were performed with 25 µg total protein from groups of 10 day 6 blastocysts cultured with or without insulin (170 nM) for 10, 30, 60 or 120 min. Western analysis was repeated twice in two independent experiments. Equal amounts of protein (25 µg) were heated at 100 °C for 5 min, solubilized in Laemmli buffer containing 200 mM DTT and electrophoresed on a 8% SDS-PAGE. Proteins were electrotransferred to nitrocellulose membranes. For detection of phospho-Erk1/2, Erk1/2, phospho-Akt and Akt, membranes were blocked in Trisbuffered saline containing 0.1% Triton (0.1% TBST) with 5% nonfat dry milk at room temperature for 1 h. Blots were incubated in 0.1% TBST containing 5% BSA or nonfat dry milk respectively, according to the manufacturers' protocols, with monoclonal antiphospho-Erk1/2 antibody (1:2000, phospho-p42/p44 MAPK Thr202/Tyr204 E10; Cell Signaling Technologies, Inc., Frankfurt, Germany), monoclonal anti-Erk1/2 antibody (1:2000, p42 MAPK 3A7, Cell Signaling Technologies, Inc.), rabbit-antiphospho-Akt (1:1000, phospho-Akt Ser473, Cell Signaling Technologies, Inc.) and monoclonal anti-Akt (1:1000, Akt 5G3, Cell Signaling Technologies Inc.) at 4°C overnight. The phospho-specific activity and the cross-reactivity with rabbit Erk1/2 and Akt were controlled for all antibodies on rabbit liver and heart muscle and on the insulin-treated and -untreated human cell line, MCF7. The antibodies were specific for rabbit tissue. The expected phosphorylation signal was only found in insulin-treated MCF7 cells (see Fig. 1). The rabbit antiphospho-Akt recognized phosphorylated and non-phosphorylated Akt (see Fig. 1A). The Ser473 phosphorylated Akt corresponds with the upper band in Fig. 1A. Only this band was used for calculation (see below). Blots were subjected to three 20-min washes in 0.1% TBST and incubated for 1 h at room temperature with anti-mouse (1:20000) or anti-rabbit (1:10000) IgG conjugated to horseradish peroxidase (Dianova, Hamburg, Germany) in 5% BSA/TBST. After three 20-min washes in 0.1% TBST, immunoreactive signals were visualized by enhanced chemiluminescence detection (ECL Plus, Amersham Biotech, Freiburg, Germany) and quantified by densitometry (BIO-Profile 1D, LTF-Labortechnik, Wasserburg, Germany). Apparent molecular weights were determined by comparison with standard molecular weight markers (high range, Promega Corp., Mannheim, Germany). The amounts of Akt and Erk proteins were evaluated by stripping the membranes and re-blotting with non-phospho antibodies. For this the blots were rinsed with 2% SDS, 60 mM Tris-HCl, pH 6-7, and 100 mM β-mercaptoethanol at 50 °C for 30 min and re-probed with total Akt and total Erk. As control for the stripping efficiency, the membranes were incubated without adding primary antibody to check that all antibodies had been removed. Protein phosphorylation was calculated as the ratio of band intensities (pAkt vs Akt, pErk1/2 vs Erk1/2) in the same blot in order to correct for differences in protein loading. The relative stimulation of Akt and Erk1/2 phosphorylation by insulin was calculated as the ratio of the phosphorylation signal of insulin-treated and non-treated embryos at each time point. Non-cultured blastocysts (0 min in Fig. 1A,B) represent the phosphorylation status of in vitro handled but otherwise untreated in vivo controls. Western blot analyses were performed at least twice in two independent experiments.

Immunohistochemistry (IHC) and translocation of GLUT4

The GLUT4 antigen was localized on embryonic sections and whole blastocysts. Bouin-fixed, paraffin-embedded day 6 rabbit blastocysts were sectioned at 5 µm. At least 5 embryos were examined per group. Sections were mounted on silanized slides, deparaffinized in xylene and rehydrated through a series of graded alcohols. For whole mount IHC the paraformaldehyde-fixed blastocysts were rehydrated through a series of graded alcohols. The neozona was removed mechanically before peroxidase blocking. Endogenous peroxidase was quenched by treatment with 3% hydrogen peroxide in methanol for 30 min. Nonspecific antibody binding was blocked with 10% normal goat or donkey serum in PBS at room temperature for 1 h and incubated with the primary antiserum overnight at 4°C in a humidified chamber. GLUT4 antibody (mouse anti-GLUT4 antibody 1:2500, DPC Biermann, Bad Nauenheim, Germany) was diluted in PBS with 1% BSA. Sections and whole blastocysts were rinsed with PBS/0.1% Tween-20 (PBST) and incubated with the peroxidaselabeled secondary antiserum (DAKO EnVision + /HRPgoat-anti mouse IgG, DAKO, Hamburg, Germany) for conventional light microscopy. The antigen was visualized with the diamino-benzidine (DAB, WAK-Chemie Medikal, Bad Soden, Germany) substrate. The development of DAB was stopped in water after 5 min. Sections were A)



B)



counterstained with hematoxylin, dehydrated and cleared in xylene. The slides were mounted in DPX and examined under bright-field microscopy with an AH-3 microscope (Olympus, Hamburg, Germany). For whole mount confocal microscopy the GLUT4 protein was localized by immunofluorescence detection with the secondary antibody fluorescein (FITC)-conjugated AffinPure donkeyanti-mouse IgG (1:300, Jackson ImmunoResearch Lab., Cambridgeshire, UK). The nuclei were counterstained with 7-amino-actinomycin (7-AAD, DAKO). Whole blastocysts were examined by fluorescence microscopy with Zeiss Axioplan (Zeiss, Oberkochen, Germany). The specificity of immunostaining was demonstrated by the absence of signals in sections incubated with control mouse IgG Figure 1 Akt and Erk1/2 phosphorylation assays in day 6 rabbit blastocysts. Day 6 (d6) rabbit blastocysts were cultured, in groups of 10, with or without 170 nM insulin for 10, 30, 60 and 120 min. The phosphorylation of Akt and Erk1/2 was analyzed by Western blotting with antiphospho Akt and anti-phospho Erk1/2 antibodies, respectively, in relation to non-phospho anti-Akt and anti-Erk1/2 antibody staining (see Materials and Methods). Representative Western blots for (A) Akt and (B) Erk1/2 are shown. In C (Akt) and D (Erk) the relative stimulation of Akt1 and Erk1/2 phosphorylation by insulin is illustrated (data from two independent experiments; protein phosphorylation in non-treated embryos = 1). Asterisks (*) indicate significant differences (P < 0.05). As references, rabbit liver (L) and heart muscle (HM) and insulin-treated human MCF7 cells were added to the blot and analyzed. Insulin treatment increased Erk1/2 but not Akt phosphorylation.

(DAKO) or in sections processed after omission of the primary antibody. Only reactions with negative controls were included in the study.

Glucose transport studies

Glucose uptake was measured as described (Pantaleon & Kaye 1996, Pantaleon *et al.* 1997). In brief, embryos were washed twice in glucose-free M2 after 1 h exposure to +/- insulin/IGF-I (16 pM or 166 nM insulin, 1.3 nM IGF-I) and transferred into 100 µl pulse droplets, kept strictly at 37 °C, for 3 min. The glucose-free pulse medium contained 0.3 mM 3-*O*-methyl-D-[1-³H]glucose ([³H]OMG, 37 GBq/l, Amersham) and 25 mM 3-OMG (Sigma Chemical Co.).

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Uptake was stopped after 3 min by transferring the embryos through four washes of ice-cold glucose-free M2 medium. The diameters of the embryos were recorded using a calibrated ocular micrometer. Radioactivity of individual embryos was determined in a Packard 1600TR liquid scintillation analyzer. In 8 experimental replicates, 56 morulae, 9 early blastocysts and 91 blastocysts were studied. Uptake of 3-OMG is expressed in nmol per 3 min per surface area (cm²). The surface area of the spherical blastocysts ($4\pi r^2$) was calculated as previously described (Robinson *et al.* 1990), and used to standardize for the differences in size between 4-day-old rabbit blastocysts. When corrected for the different time of [³H]3-OMG measurement (3 vs 10 min), these data are in close agreement with those reported by Robinson *et al.* (1990).

To determine whether the extracellular coverings of rabbit embryos (see Fischer *et al.* 1991) exerted an influence on glucose transport, the coats of morulae and blastocysts were removed with 0.05% pronase E (Herrler *et al.* 1998) prior to (170 nM) insulin treatment. In total, 30 blastocysts and 6 morulae were studied in two independent experiments. Uptake of 3-OMG was unaffected by insulin treatment in coat-free embryos of both stages (P > 0.05; data not shown).

Statistics

Uptake of 3-OMG was analyzed by two-way ANOVA and multiple means range tests using Fisher's protected procedures. Statistical analysis of the GLUT4, GLUT8 and IR RNA levels and the relative stimulation of Akt and Erk1/2 phosphorylation were performed with the paired *t*-test between + and - insulin at the various time points studied (SigmaPlot 4.0, Jandel Corporation (San Rafael, CA, USA) mathematical and statistical analysis). The data are expressed as means \pm S.E.M.

Results

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Insulin does not stimulate the phosphorylation of Akt in day 6 blastocysts

Active Akt is phosphorylated at Ser473. Western blot analysis with phosphospecific anti-Akt antibody showed that addition of 170 nM insulin for 10 to 120 min did not alter the level of Akt phosphorylation in cultured blastocysts (Fig. 1A). The antibody used (phospho-Akt-Ser473) detects Akt when phosphorylated at Ser473 (upper band in Fig. 1A). The amount of phosphorylated Akt relative to total Akt was measured in each sample by image analysis. Protein phosphorylation was calculated as the ratio of band intensities (see Materials and Methods). The relative increase in phosphorylation by insulin treatment in comparison with untreated controls was analyzed at each time point (Fig. 1C). Akt phosphorylation was not stimulated by insulin treatment for up to 2 h (P > 0.05).

Insulin stimulates the rapid phosphorylation of Erk in day 6 blastocysts

Erk-1 and -2 are activated by various cell surface receptors, including tyrosine kinase receptors, receptors coupled to cytoplasmic tyrosine kinase and G protein-coupled receptors. Erk1/2 Western blots from insulin-treated and non-treated blastocysts are shown in Fig. 1B. The amount of phosphorylated Erk-1 and -2, calculated in relation to total Erk, was increased after insulin treatment (Fig. 1D). Addition of 170 nM insulin to cultured day 6 blastocysts caused a rapid increase in the phosphorylation level of both Erk-1 and -2, reaching a maximum increase of about 100% at 10min after insulin exposure and returning to control levels by 60min (10 and 30min vs 60min: P < 0.05).

Insulin receptor and GLUT4 but not GLUT8 mRNA levels are increased by insulin in day 6 rabbit blastocysts

Insulin treatment almost doubled the level of GLUT4 mRNA but was without effect on GLUT8 mRNA levels (Fig. 2A,B). Culture for 1 h increased the levels of IR mRNA by about 50% and insulin treatment for 1 h led to a further increase of 20%. In the groups exposed to insulin for longer than 2 h the IR transcript levels declined to control levels (Fig. 2C).

Insulin does not stimulate GLUT4 translocation in day 6 rabbit blastocysts

GLUT4 immunoreactivity in trophoblast and embryoblast cells following 1 h insulin stimulation was examined by immunofluorescence. GLUT4 was localized in the perinuclear cytoplasm (Fig. 3B,C). This localization was unchanged by insulin treatment (Fig. 3). It did, however, cause a slight increase in the GLUT4 fluorescence signal (Fig. 3C).

Insulin does not stimulate glucose transport in day 3 rabbit morulae and day 4 blastocysts

The rate of 3-OMG uptake was higher in blastocysts than in early blastocysts and morulae (blastocysts: 34.7 ± 4.3 (mean±s.E.M.) nmol 3-OMG/3 min/cm² surface area, n = 91; early blastocysts: 22.5 ± 12.9, n = 9; morulae: 16.8 ± 6.2 , n = 56; blastocysts vs morulae: P < 0.05). An approximate 50% increase in glucose uptake rate induced by 166 nM insulin was not significant whilst 16 pM insulin or 1.3 nM IGF-I were clearly ineffective (P > 0.05; Fig. 4).

Discussion

The massive stimulation of glucose uptake by insulin in adipose and muscle tissues occurs through a complex receptor signal transduction process promoting the movement of GLUT4 protein from intracellular storage sites to the plasma membrane. In adipocytes and myocytes



GLUT4 translocation is the rate-limiting step of insulinstimulated glucose uptake (Saltiel 2001). In preimplantation embryos GLUT4 expression has been shown for bovine (Navarrete Santos *et al.* 2000, Augustin *et al.* 2001), rat (Korgun *et al.* 2001) and rabbit (Navarrete Santos *et al.* 2004) blastocysts. The expression data from mouse preimplantation embryos are controversial. Whereas GLUT4 was not found by Hogan and Aghayan and co-authors (Hogan *et al.* 1991, Aghayan *et al.* 1992) Figure 2 Transcriptional regulation of glucose transporter 4 (GLUT4) and 8 (GLUT8) and insulin receptor (IR) by insulin in day 6 rabbit blastocysts. Day 6 rabbit blastocysts were cultured, in groups of 10, with or without 170 nM insulin for 1, 2, 3 and 4 h. The relative amounts of GLUT4 and IR RNA were quantified by semiquantitative RT-PCR in relation to the housekeeping gene, GAPDH, as described in Materials and Methods. The relative amounts of two experiments are shown. Data from all time points studied (1, 2, 3 and 4 h) were combined (N = 2 replicates, n = 4 blastocysts for each experimental group; means±s.E.M.). Insulin effects on (A) GLUT4 and (B) GLUT8 transcription reveal a difference between the GLUT isoforms. While transcript numbers of GLUT4 were significantly increased, GLUT8 mRNA levels were unaffected (hatched bars = cultured blastocysts without insulin, solid bars = cultured blastocysts with insulin, control = non-cultured blastocysts set to 100 %). Asterisks (*) indicate significant differences compared with controls (P < 0.05). For IR (C) the means of two individual experiments are shown for each time point studied (n = 4 blastocysts for each experimental group in each replicate; means \pm s.E.M.; open bars = cultured blastocysts without insulin, hatched bars = cultured blastocysts with insulin).

and insulin stimulation of glucose uptake was only about 40% suggesting the functional absence of GLUT4 (Pantaleon & Kaye 1996), we have recently detected GLUT4 mRNA in mouse blastocysts and embryonic stem cells (Tonack *et al.* 2004). In rabbit blastocysts, GLUT4 is expressed in trophoblast and embryoblast cells. In trophoblast cells of embryos grown *in vivo*, GLUT4 was localized in the perinuclear compartment and in the cell membrane (Navarrete Santos *et al.* 2004). This was

– Insulin 1h



+ Insulin 1h



Figure 3 Localization of GLUT4 in day 6 rabbit blastocysts cultured in the presence or absence of insulin. GLUT4 protein was localized perinuclearly in the cytoplasm of trophoblast and embryoblast cells. Detection was performed with a fluorescence-labeled mouse anti-GLUT4 (FITC, green color) in day 6 rabbit blastocysts cultured without (B) or with (C) 170 nM insulin for 1 h. A representative staining of a blastocyst from each group in the embryoblast (Em) and trophoblast (Tr) area is shown. The rim of the embryoblast is marked by white arrowheads. Nuclei were counterstained with 7-AAD (red color). Inserts A and D show control reactions for B and C respectively. Bars = $20 \,\mu$ m.

regarded as good evidence for a potentially functional role in the insulin-stimulated uptake of maternal glucose *in utero*. In order to better understand the cellular and molecular mechanisms of glucose transport and insulin action in preimplantation embryos, *in vitro* culture of rabbit blastocysts was employed for further studies. We found (i) activation of the IR signaling pathway with (ii) stimulation of the MAPK but not the Akt pathway, (iii) no GLUT4 translocation and no stimulation of glucose uptake, and (iv) an increased GLUT4 and IR transcription in the presence of insulin. These data suggest that insulin stimulates mitogenesis but not glucose transport in rabbit blastocysts.



Figure 4 Uptake of 3-OMG in insulin/IGF-I-treated and untreated day 4 rabbit blastocysts. Four-day-old blastocysts were cultured with or without insulin (166 nM, 16 pM) or IGF-I (1.3 nM) in glucose-supplemented M2 medium for 1 h. Glucose transport was measured by 3-OMG uptake during a 3-min pulse period. Data are means \pm s.E.M. and are expressed as nmol 3-OMG/3 min/cm² (see Materials and Methods). The number of embryos is given in each column. Insulin/IGF-I treatment did not increase OMG uptake (P > 0.05).

Activation of insulin signaling pathways in rabbit blastocysts by insulin

It is well established that the insulin-stimulated translocation of GLUT4 to the plasma membrane depends on the production of PIP₃ by PI3-K (Fruman et al. 1998). The phosphorylation of the 3' position recruits and activates proteins containing pleckstrin homology domains, such as 3' phosphoinositide-dependent kinase-1 (PDK-1) and Akt (Rameh & Cantley 1999). PDK-1, in turn, phosphorylates and activates downstream effectors including Akt. A role for this pathway in insulin-stimulated GLUT4 translocation has been confirmed. Inhibition of PI3-K using wortmannin and LY294002, and blockade of the PI3-K signaling pathway by dominant-negative mutations inhibited insulin-stimulated GLUT4 translocation and glucose uptake (Cheatham et al. 1994, Okada et al. 1994, Haruta et al. 1995, Kotani et al. 1995). These findings demonstrate that activation of the PI3-K and formation of PIP₃ are necessary for GLUT4 translocation. However, some controversial findings exist regarding the role of PIP₃ downstream effectors, mainly Akt, and their contribution to insulin-stimulated GLUT4 translocation. Akt is clearly activated in response to insulin in a variety of cell types (Kohn et al. 1996). However, overexpression of dominant-negative Akt mutants has produced divergent results. Whereas expression of the phosphorylation site-deficient Akt did not affect insulin-stimulated GLUT4 translocation, both a kinase dead- and a dual kinase dead-mutant and phosphorylation-deficient construct inhibited translocation (Cong et al. 1997, Kitamura et al. 1998, Wang et al. 1999). The expression of a constitutively active Akt with a membrane targeting sequence promoted GLUT4 translocation in a hormone-independent manner but in a substantially longer time interval than that observed after

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insulin stimulation (Kohn *et al.* 1996, 1998, Tanti *et al.* 1997, Foran *et al.* 1999).

In rabbit blastocysts we did not detect a significant increase in Akt phosphorylation after insulin treatment for up to 2 h, indicating that this pathway was not activated by 170 nM insulin. The antibody used for pAkt detection in the present study would have recognized all three phosphorylated Akt isoforms. The level of Akt phosphorylation was almost identical in insulin-stimulated and control blastocysts at the time points studied after insulin treatment (10 to 120 min), suggesting a basic Akt activation by exogenous (culture conditions, handling) or endogenous factors. For placental JAr cells it has been shown that despite a fourfold Akt activation, GLUT4 translocation, glucose uptake and glycogen synthesis were not stimulated by insulin. Mitogenesis was increased 1.9-fold and accompanied by phosphorylation of the MAPK cascade via Erk1/2 (Boileau et al. 2001). A similar activation of Erk1 and 2 was found in the present study in rabbit blastocysts 10 min after treatment with 170 nM insulin. Restoration of normal Erk1/2 levels occurred after 1 h in culture. The activated Erk1/2 cascade preferentially regulates cell growth, differentiation and development (for review see Schaeffer & Weber 1999). Herrler and co-authors (1998) have previously shown mitogenic and anti-apoptotic effects of insulin in rabbit embryos. Supplementation with IGF-I or insulin improved blastocyst formation producing larger day 3 blastocysts. In day 4 blastocysts addition of increasing amounts of insulin decreased apoptosis and increased cell proliferation in a dose-dependent manner. In cultured bovine embryos, insulin exerted comparable mitogenic and anti-apoptotic activities (Matsui et al. 1995a, b, Byrne et al. 2002, Augustin et al. 2003). Insulin increased blastocyst formation rate and total blastocyst cell number by 33% and 18% respectively (Augustin et al. 2003). In cultured mouse embryos insulin increased cell number, mitotic index (Gardner & Kaye 1991) and blastocyst cell number by specifically increasing inner cell mass (ICM) cell numbers (Harvey & Kaye 1990, Smith et al. 1993).

Transcriptional regulation of GLUT4 and IR by insulin

Because glucose transport is a rate-limiting step in glucose metabolism, GLUT4 expression in muscle and adipose is tightly regulated at both the mRNA and protein levels. Both are influenced by numerous hormonal and metabolic signals and physiological states. A major form of regulation involves the translocation of GLUT4 protein from the cytoplasmic vesicles to the plasma membrane in response to insulin (reviewed in Shepherd & Kahn 1999). Studies in adipose and muscle tissues showed that expression of the GLUT4 glucose transporter is controlled at the level of transcription (Gerrits *et al.* 1993, Santalucia *et al.* 1999). It has been shown in adipose, cardiac, and skeletal muscle that insulin-deficient states such as fasting

or streptozotocin-induced diabetes resulted in a severe downregulation of GLUT4 messenger RNA and protein (Bourey *et al.* 1990, Camps *et al.* 1992, Neufer *et al.* 1993, Olson *et al.* 1993, Dombrowski & Marette 1995, Depre *et al.* 2000). In our study, insulin treatment increased GLUT4 transcript numbers whereas GLUT4 translocation and glucose uptake were not affected. This finding indicates an involvement of other, so far not clearly known biological processes in GLUT regulation. It is known, however, that GLUT4 expression can be regulated by several transcription factors such as p53 (Schwartzenberg-Bar-Yoseph *et al.* 2004), myocyte enhancer factor 2A (Knight *et al.* 2003) and PAX3/forkhead homolog (Armoni *et al.* 2002).

GLUT4 translocation and stimulation of glucose uptake

The findings from our glucose uptake study and from GLUT4 immunohistochemistry after insulin treatment point to the same conclusion. GLUT4 localization was not different in embryos cultured for 1 h in an insulin-free or insulin-supplemented medium. It was localized intracellularly in trophoblast and embryoblast cells. An increase in membrane-associated GLUT4 was not found in the trophoblast cells of insulin-treated rabbit blastocysts and neither insulin nor IGF-I stimulated glucose transport. These and other results underline the significance of ligand concentrations and/or species when comparing studies on IR/IGF-IR effects in preimplantation embryos. For example, both insulin (700 nM) and IGF-I (130 nM) in high concentrations increased apoptosis through the downregulation of the IGF-I receptor in a recent study in mouse blastocysts (Chi et al. 2000). A significant increase (40%) in 3-OMG uptake in mouse blastocysts had been achieved with considerably lower IGF-I concentrations (0.17-1.7 pM; Pantaleon & Kaye 1996) than those employed in the present study, suggesting that in the mouse embryo IGF-I is the key hormone for stimulation of glucose uptake (Pantaleon & Kaye 1996). The increase in glucose uptake, however, was only moderate compared with insulin effects in myocytes and adipocytes. Insulin action in preimplantation embryos is aimed towards anabolic, mitogenic and anti-apoptotic effects.

In some species the mitogenic and anti-apoptotic effects of insulin and IGF-I were specifically directed towards ICM cells. For example, supplementation with physiological concentrations of IGF-I caused a specific increase in the number of the ICM cells (1.7 nM, human embryos, Lighten *et al.* 1998; 6.5 nM, bovine blastocysts, Sirisathien *et al.* 2003; 5.2 nM, mouse blastocysts, Smith *et al.* 1993), thus complementing earlier studies with insulin in mice (Harvey & Kaye 1990). IGF-I and insulin acted as ICMspecific growth factors. The reasons for this phenomenon are unknown. Implications for long-term development have been investigated by culturing two-cell embryos for 2 days *in vitro* with 170 nM insulin and subsequent embryo transfer into foster mothers. Insulin-treated embryos showed an increased fetal growth at day 19 and 20 by 4 to 6% (Kaye & Gardner 1999).

In conclusion, despite the stage-specific expression of IR and the insulin-responsive isoforms GLUT4 and GLUT 8, insulin did not stimulate glucose transport in rabbit blastocysts. Analysis of IR signaling indicates that insulin action is directed towards mitogenic rather than metabolic effects during this ontogenetic stage of development in the rabbit. Stimulation of embryonic glucose transport by hormones of the IGF family might not be necessary during this developmental period, perhaps due to an efficient uptake by glucose transporters under the physiological conditions *in utero*.

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