Adiponectin stimulates glucose uptake in mouse blastocysts and embryonic carcinoma cells

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Abstract

Preimplantation embryos are sensitive to maternal hormones affecting embryonic signal transduction and metabolic functions. We examined whether adiponectin, the most abundantly secreted adipokine, can influence glucose transport in mouse embryonic cells. In mouse blastocysts full-length adiponectin stimulated glucose uptake, while no effect of globular adiponectin was found. Full-length adiponectin stimulated translocation of GLUT8 glucose transporter to the cell membrane; we did not detect significant changes in the intracellular localization of GLUT4 glucose transporter in adiponectin-treated blastocysts. To study adiponectin signaling in detail, we used embryoid bodies formed from mouse embryonic carcinoma cell (ECC) line P19. We confirmed the expression of adiponectin receptors in these cells. Similar to mouse blastocysts, full-length adiponectin, but not globular adiponectin, stimulated glucose uptake in ECC P19 embryoid bodies. Moreover, full-length adiponectin stimulated AMPK and p38 MAPK phosphorylation. These results indicate that besides AMPK, p38 MAPK is a potential target of adiponectin in mouse embryonic cells. AMPK inhibitor did not influence the adiponectin-stimulated p38 MAPK phosphorylation, indicating independent action of these two signaling pathways. In mouse embryos adiponectin acts as a hormonal regulator of glucose uptake, which becomes especially important in phases with reduced levels of circulating insulin. Our results suggest that adiponectin maintains the glucose supply for early embryos under hypoinsulinaemic conditions, for example, in mothers suffering from type 1 diabetes mellitus.

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Introduction

The preimplantation embryo requires adequate energy production and distribution. Glucose is the main energy source from the blastocyst stage onwards (Pantaleon & Kaye 1998), and preimplantation embryos of various mammalian species express several facilitative glucose transporters (SLC2A/GLUT family, for review see Purcell & Moley 2009). Experiments using rabbits have shown that adiponectin, the most abundantly secreted adipokine, can stimulate glucose uptake in blastocysts, and the increase in glucose transport was connected with translocation of glucose transporter GLUT4 into the cell membrane (Fischer *et al.* 2010). Interestingly, insulin stimulus neither induced GLUT4 translocation nor increased glucose uptake in rabbit blastocysts (Navarrete Santos *et al.* 2004a). This is in contrast with the mouse, where insulin has been shown to stimulate blastocyst glucose uptake via IGF1 receptor-mediated translocation of GLUT8 glucose transporter (Pantaleon & Kaye 1996, Carayannopoulos *et al.* 2000, Pinto *et al.* 2002). These results suggest possible interspecies differences in the

regulation of blastocyst glucose uptake, which seems to be insulin- and IGF1-dependent in the mouse, and insulin-independent/adiponectin-stimulated in the rabbit. On the other hand, since adiponectin receptors have been detected in blastocysts of several mammalian species (for review see Čikoš 2012), adiponectin-regulated glucose transport could exist in blastocysts of other species as well.

Two insulin- or adiponectin-responsive facilitative glucose transporters have been identified in preimplantation embryos. Both GLUT4 (SLC2A4) and GLUT8 (SLC2A8) were found in bovine and rabbit blastocysts (Navarrete Santos *et al.* 2000, 2004b, Augustin *et al.* 2001). GLUT4 was also detected in rat blastocysts (Korgun *et al.* 2001) and GLUT8 was identified in mouse and hamster blastocysts (Carayannopoulos *et al.* 2000, Pinto *et al.* 2002, Tonack *et al.* 2009). A controversy exists on the expression of GLUT4 in mouse blastocysts (Tonack *et al.* 2004 vs Hogan *et al.* 1991, Aghayan *et al.* 1992).

The aim of this study was to examine whether adiponectin can stimulate glucose uptake in mouse

blastocysts, and whether GLUT4 and GLUT8 are involved in a possible adiponectin-regulated glucose transport. Embryoid bodies (EBs) developed from mouse embryonic carcinoma P19 (ECC P19) cells were used as a model for studying adiponectin signaling in embryonic cells. To validate this model, we verified the expression of adiponectin receptors in ECC P19 EBs. Finally, we analysed glucose uptake and activation of the adiponectin signaling pathway in the ECC P19 EBs.

Materials and methods

Embryo recovery

Female mice (CD-1 strain, 4 to 7 weeks old, Velaz, Prague, Czech Republic and Charles River, Freiburg, Germany) underwent superovulation treatment with intraperitoneal injection of 4 IU of serum gonadotropin (Folligon, Intervet International Bv. Boxmeer, Holland), followed 47 h later by administration of 4 IU of human chorionic gonadotropin (hCG, Organon, Oss, Holland). Twelve to fourteen hours after hCG administration, the mice were killed by cervical dislocation, and unfertilized oocytes were isolated by flushing from the oviduct. To obtain preimplantation embryos, females were mated with males of the same strain overnight (mating was confirmed by identification of a vaginal plug) and killed by cervical dislocation (57-, 72-, 86- or 98 h after hCG administration), and the embryos (4-cell, 8- to 16-cell embryos, morulas and blastocysts) were isolated by flushing from the oviduct and/or uterus. Oocytes and embryos were washed in several drops of FHM medium (Lawitts & Biggers 1993) containing 1% BSA (Sigma–Aldrich) and pooled according to their morphology. Cumulus cells were removed with 0.1% hyaluronidase (Sevac, Prague, Czech Republic). All animal experiments were performed in accordance with the principles of the Ethical Committee for Animal Experimentation at the Institute of Animal Physiology, and approved through the State Veterinary and Food Administration of the Slovak Republic (Ro 3155/14-221) and Germany (MLU I11M21) in strict accordance with legislation based on EU Directive 2010/63/EU on the protection of animals used for experimental and other scientific purposes. All efforts were made to reduce animal suffering and distress and to minimize the number of animals necessary to produce reliable results.

Mouse embryonic carcinoma P19 cell culture

Mouse embryonic carcinoma cells (ECC) line P19 (McBurney *et al.* 1982) were cultured under standard conditions (5% CO₂ and 37°C) on gelatin (0.1%)-coated tissue culture dishes in DMEM (Invitrogen) supplemented with 15% heat inactivated fetal calf serum (selected batches of Biochrom KG, Berlin, Germany), L-glutamine (2 mm; Life Technologies, Inc.), β-mercaptoethanol (final concentration 50 mm; Serva, Heidelberg, Germany), nonessential amino acids (stock solution diluted 1:100; Invitrogen), and streptomycin/ampicillin (5000 U/mL ampicillin, 5000 µg/mL streptomycin; Invitrogen) (Tonack *et al.* 2007). Drops of differentiation medium (20 µL) with 400 cells were placed onto the lids

of Petri dishes filled with PBS and cultured for 2 days. Cell aggregates formed in the hanging drops were transferred into non-adhesive bacteriologic-grade Petri dishes (6 cm) containing 5 mL culture medium to develop older (4–6 days old) embryoid bodies (EBs; Wobus *et al.* 1994).

RNA isolation and cDNA synthesis

To analyse expression of glucose transporter (GLUT4, GLUT8) and adiponectin receptor (AdipoR1, AdipoR2, T-cadherin) transcripts, total RNA was extracted from batches of 80–100 mouse blastocysts, early stage embryos and oocytes, from 15 ECC P19 embryoid bodies, and also from mouse heart and brain (positive controls). TRIzol Reagent (Invitrogen Life Technologies) was used for the extraction according to the manufacturer's instructions. Contaminating DNA in RNA preparations was digested with amplification-grade DNase I (Invitrogen Life Technologies). Three independent experiments were performed, using separate batches of embryos or embryoid bodies for the RNA isolation.

The RNA (from 80 to 100 embryos or 0.5 µg RNA from ECC P19 embryoid bodies, heart or brain) was reverse transcribed with Superscript™ III RNase H[−] Reverse Transcriptase (Invitrogen Life Technologies) using 4 µM anchored oligo dT primers and 1.5 µM random hexamer primers (Thermo Fisher Scientific ABgene). To check for the presence of genomic DNA contamination in the RNA preparations, reverse transcriptase negative controls (no reverse transcriptase in the reaction) were carried out in parallel, using part of each RNA sample. The cDNA preparations were then cleaned by ethanol precipitation, and the cDNA pellets from blastocysts were diluted in an appropriate amount of 10 mM Tris (pH 8.3) so that 1 µL of the cDNA corresponded to 2.5 embryo equivalents or to 25 ng of ECC P19 total RNA.

PCR and PCR product analysis

PCR amplifications were carried out in 25 µL volumes containing 1 µL of cDNA, 0.5 µM of each oligonucleotide primer, 50 mM KCl, 10 mM Tris–HCl pH 8.3, 2 mM MgCl₂, 0.2 mM dNTPs (dATP, dTTP, dCTP, dGTP) and 0.04 units/µL Platinum Taq DNA polymerase (Invitrogen Life Technologies). For detection of GLUT4 and GLUT8 transcripts in blastocysts we used primers designed by Tonack *et al.* (2004), for detection of AdipoR1 and AdipoR2 transcripts in ECC P19 embryoid bodies we used primers designed by Schmidt *et al.* (2008), and for detection of T-cadherin transcripts in blastocysts and ECC P19 embryoid bodies we used primers designed by Zhou *et al.* (2002). An initial denaturation step at 95°C for 2 min was followed by 35 (ECC P19, heart and brain cDNAs) or 40 (blastocyst cDNAs) cycles at 94°C for 30 s, 60°C (GLUT4, GLUT8, AdipoR1 and AdipoR2) or 64°C (T-cadherin) for 30 s and 72°C for 60 s. To check for the presence of cross contamination, the reaction with water instead of cDNA was performed concurrently (blank reaction). PCR products were analysed using electrophoresis on a 2% agarose gel stained with GelGreen (Biotium, Hayward, USA), and 100-bp DNA ladder (Jena Bioscience, Jena, Germany) was used as a marker to determine the size of PCR products. PCR products were

sequenced (GATC Biotech AG, Konstanz, Germany), and the obtained sequences were compared (using Clustal W and Blast 2 algorithms) with known mRNA sequences of mouse GLUT4, GLUT8, AdipoR1, AdipoR2 and T-cadherin (GenBank accession numbers NM_009204, NM_019488, NM_028320, NM_197985 and NM_019707 respectively).

Real-time PCR relative quantification of mRNA

To compare relative quantities of AdipoR1 and AdipoR2 transcripts as well as GLUT4 and GLUT8 transcripts in mouse blastocysts and ECC P19 EBs, we amplified these sequences using real-time PCR with SYBRGreen fluorescent dye. Amplification efficiency, threshold fluorescence and Ct values were measured in each amplification, and starting fluorescences were then calculated. The theoretical starting fluorescence is proportional to the amount of PCR template in the sample, and represents the quantity of target nucleic acid sequence expressed in arbitrary fluorescence units (Liu & Saint 2002, Peirson *et al.* 2003, Ramakers *et al.* 2003, Čikoš & Koppel 2009).

PCR amplifications for the quantitative analysis were performed in the Mx 3000P real-time PCR system (Stratagene, La Jolla, CA, USA). The reactions were carried out in 25 µL volumes containing 1 µL of cDNA, 1× PCR Master Mix with SYBR Green (Qiagen), and 0.4 µM of each specific forward and reverse primer. For adiponectin receptors (AdipoR1 and AdipoR2) we used primers designed by Schmidt *et al.* (2008), and for glucose transporters (GLUT4 and GLUT8) and beta actin (ActB) we used primer assays from Qiagen (GLUT4: Slc2a4 assay #PPM04166F, GLUT8: Slc2a8 assay #PPM041165A, ActB: ActB assay #PPM02945B). An initial step at 95°C for 10 min was followed by 40 cycles (or 36 cycles for ActB and 45 cycles for GLUT4) at 95°C for 20 s, 64°C (AdipoR1), 66°C (AdipoR2) or 65°C (GLUT4, GLUT8, ActB) for 30 s, 72°C for 20 s and 78°C (or 83°C for ActB) for 20 s (acquiring of fluorescence). Fluorescence was measured after the elongation step. Amplification specificity was checked using melting curve analysis.

Amplification efficiency was measured with the use of a relative standard curve and LinRegPCR software (Ramakers *et al.* 2003). Theoretical starting fluorescence for each analyzed sequence (AdipoR1, AdipoR2, GLUT4, GLUT8) and for reference gene (ActB) in each sample was calculated using the formula:

$$R_0 = R_{Ct} / (E + 1)^{Ct}$$

where R_0 is the starting fluorescence (proportional to the amount of PCR template), R_{Ct} is the threshold fluorescence value, E is the amplification efficiency and Ct is the fractional cycle number necessary for reaching the threshold fluorescence (Čikoš & Koppel 2009). The calculated starting fluorescences were corrected for the PCR product length (R_{0bp} values, obtained by dividing the R_0 value with the PCR product length in bps). Validation of this procedure was done in a preliminary experiment (Supplementary Table 1, see section on supplementary

materials given at the end of this article). To normalize target gene quantities, AdipoR1, AdipoR2, GLUT4 and GLUT8 starting fluorescences were divided by ActB starting fluorescences.

Immunostaining

We used the immunofluorescence technique to localise expression of GLUT4 and GLUT8 proteins in mouse blastocysts, and expression of AdipoR1 and AdipoR2 proteins in ECC P19 embryoid bodies. To examine the effect of adiponectin on GLUT4 and GLUT8 intracellular localization we stimulated mouse blastocysts with adiponectin. The blastocysts were isolated as described above and cultured under standard conditions (5% CO₂ and 37°C; KSOM culture medium, Specialty Media Group, Phillipsburg, NJ, USA) for 45 min in the presence or absence of adiponectin. The culture drops of adiponectin-stimulated blastocysts were supplemented with full-length adiponectin (mouse Acrp30, recombinant protein, produced in HEK 293 cells, BioVendor) at a final concentration of 10 µg/mL. No supplement was added in the culture drops of control blastocysts. Blastocysts were fixed in 4% paraformaldehyde, then they were washed in PBS/BSA and permeabilized for 1 h in PBS with 0.5% Triton X-100 (Sigma-Aldrich). Free aldehyde groups were blocked with 0.1 M glycine (Merck), and washed in PBS/BSA. Non-specific immunoreactions were blocked with 10% normal goat serum (Santa Cruz Biotechnology) for 2 h at room temperature. The blastocysts were then incubated with primary antibodies raised against mouse GLUT4 or GLUT8 (purchased from LifeSpan BioSciences, Seattle, WA, USA; for detailed information see Table 1) in 10% goat serum at 4°C overnight. Specific secondary antibody coupled with Alexa Fluor 488 (Alexa Fluor 488 goat anti-rabbit IgG, Invitrogen Life Technologies) was used to visualize primary antibodies (30 min at room temperature). Cell nuclei were stained with Hoechst 33342 in PBS/BSA (20 µg/mL; Sigma-Aldrich). Afterwards, the blastocysts were mounted in ProLong Gold antifade reagent (Molecular Probes, Invitrogen) on glass slides, sealed with coverslips and observed using a confocal microscope (Leica TCS SP5, Leica Microsystems). Negative control groups of embryos were incubated without the primary antibody or without the primary and secondary antibody.

ECC P19 embryoid bodies (EBs) were prepared as described above. EBs were fixed, dehydrated for paraffin embedding, sectioned and attached to slides as described previously (Tonack *et al.* 2006). The slides were incubated in a heating cabinet (60°C) over night, then dewaxed in xylene and rehydrated through a series of graded alcohols. For AdipoR1 detection, a heat-induced antigen retrieval step in 20 mM at pH9 was included. After washing in PBST (PBS with 0.1% Tween), non-specific antibody binding was blocked with 10% normal goat serum in PBS for 1 h at room temperature. The sections were incubated with primary antibodies raised against adiponectin receptor 1 or adiponectin receptor 2 (purchased from LifeSpan BioSciences, Inc. and Alpha Diagnostic International, San Antonio, TX, USA; for detailed information see Table 1) in 10% goat serum at 4°C overnight. Afterwards, the sections were incubated with secondary antibody coupled with Alexa Fluor

Table 1 Primary antibodies used in the study.

Antibody name	Host species and antibody clonality	Application	Dilution used	Source, catalog #
SLC2A4/GLUT4 Antibody (aa480-509)	Rabbit polyclonal	IHC	1:200	LifeSpan BioSciences # LS-C143467
SLC2A8/GLUT8 Antibody (aa461-477)	Rabbit polyclonal	IHC	1:200	LifeSpan BioSciences #LS-C313271
Adiponectin Receptor 1 Antibody (C-Terminus)	Rabbit Polyclonal	IHC	1:100	LifeSpan BioSciences # LS-C151518
Adiponectin Receptor 2 (ADIPOR2) Antibody	Rabbit polyclonal	IHC	1:100	Alpha Diagnostic International # ADIPOR22-A
AMPK α Antibody	Rabbit polyclonal	Western	1:1000	Cell Signaling Technology, # 2532
Phospho-AMPK α (Thr172) Antibody	Rabbit polyclonal	Western	1:1000	Cell Signaling Technology, # 2531
Phospho-Acetyl-CoA Carboxylase (Ser79) Antibody	Rabbit polyclonal	Western	1:1000	Cell Signaling Technology, # 3661
Phospho-p38 MAPK (Thr180/Tyr182) (12F8) Antibody	Rabbit monoclonal	Western	1:1000	Cell Signaling Technology, # 4631
Monoclonal Anti- β -Actin Antibody Clone AC-15	Mouse monoclonal	Western	1:40000	Sigma-Aldrich #A5441

488 (1–2 h at room temperature), counterstained with Hoechst 33342 (as described above for the blastocysts), mounted in Vectashield (Vector Laboratories) and observed using the confocal microscope (Leica TCS SP5, Leica Microsystems). To visualize the plasma membrane, F-actin staining with Phalloidin-Tetramethylrhodamine B isothiocyanate (5 min incubation at room temperature; Santa Cruz Biotechnology) was performed in some samples. Negative control groups of EBs were incubated without the primary antibody or without both the primary and secondary antibody.

Glucose uptake studies

Blastocysts

The blastocysts were isolated as described above, washed in several drops of FHM medium containing 0.1% BSA, and then pooled, randomly divided into experimental and control groups, washed in KSOM culture medium containing 5 mM glucose, and transferred to culture drops (1 embryo/1 μ L KSOM, covered with mineral oil). The culture drops of experimental groups were supplemented with adiponectin at a final concentration of 10 μ g/mL (adiponectin water stock solution (600 μ g/mL) was appropriately diluted with the KSOM culture medium). No supplement was added in the culture drops of the control (CTRL) group. The embryos were cultured in standard conditions (5% CO₂ and 37°C) for 2 h. Two isoforms of adiponectin were used in the experiments: full-length adiponectin (mouse Acrp30, recombinant protein, produced in HEK 293 cells, BioVendor; FLAD group) and the truncated isoform - globular adiponectin (mouse Acrp30, globular recombinant protein, produced in *Escherichia coli*, Abcam; GLAD group). Two experiments were carried out with each adiponectin isoform and the results were pooled.

Glucose uptake was measured as previously described (Fischer *et al.* 2010). In brief, after the 2-h culture with or without adiponectin, the embryos were washed three times in glucose-free FHM medium, and then transferred into 400- μ L 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 (3-OMG; 37 GBq/L; Amersham) and 25 mM 2,3,4,6-Tetra-O-methyl-D-glucose (Santa Cruz Biotechnology). The uptake was stopped after 3 min by transferring the embryos

through four wash steps with ice-cold glucose-free FHM medium. Radioactivity of embryos was determined in a liquid scintillation counter (LSC) Wallac 1409 (PerkinElmer). We put five blastocysts to each sample and calculated the 3-OMG uptake in picomoles per minute per blastocyst.

P19 cells

Mouse embryonic carcinoma P19 cells were cultured as described above. Embryoid bodies were then randomly divided into several groups to measure glucose uptake. The two isoforms of adiponectin were used in the experiments and the final concentration of adiponectin in the culture medium was 10 μ g/mL. Groups containing 15–25 EBs were cultured in 96-well sterile cell culture plates in a volume of 180 μ L for 1 h under standard conditions (5% CO₂ and 37°C) in the presence or absence (controls) of adiponectin.

In the first experiment, three groups were used: the CTRL group was cultured in the differentiation medium, the FLAD group was cultured in the differentiation medium supplemented with full-length adiponectin, and the GLAD group was cultured in the differentiation medium supplemented with globular adiponectin. To examine the effect of compound C on the full-length adiponectin-stimulated glucose uptake, a second experiment was performed. Three groups were used: the Control (DMSO) group was cultured in the differentiation medium supplemented with 0.1% DMSO (as compound C was dissolved in DMSO), the FLAD group was cultured in the differentiation medium containing 0.1% DMSO supplemented with full-length adiponectin, and the FLAD+CC group was cultured in the differentiation medium containing 0.1% DMSO supplemented with full-length adiponectin and 10 μ M compound C (these cells were preincubated in medium with 10 μ M compound C for 30 min before the stimulation with full-length adiponectin; compound C was purchased from Sigma-Aldrich).

After 1 h culture with or without adiponectin, the EBs were washed three times in 400 μ L of glucose-free DMEM and transferred into 400 μ L pulse droplets, kept 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/liter; Amersham Biosciences, Freiburg, Germany) and 25 mM 3-OMG (Sigma-Aldrich). The uptake was stopped after 3 min

by transferring the EBs through four wash steps with 400 μ L ice-cold glucose-free DMEM. The diameter of the EBs was recorded using a calibrated ocular micrometer. Radioactivity of individual EBs was determined in the LSC Walac 1409. The uptake of 3-OMG was expressed in nmol/vol (cm^3) $\times$ 3 min. The volume of the spherical EBs was calculated as $1/6 \mu\text{d}^3$ and used to standardise the differences in size between EBs.

Analysis of AMP and p38 MAP kinases in adiponectin-stimulated ECC P19 embryoid bodies

The EBs were stimulated with full-length adiponectin for 5, 10, 15, 60 and 120 min. Specific AMPK inhibitor, Compound C (CC) was added before stimulation with adiponectin to check the AMPK dependency of downstream targets. For Western blot analysis, 25 μ g total protein lysates were subjected to SDS-polyacrylamide electrophoresis and electrotransferred to nitrocellulose membranes. For detection of p-p38 MAPK and β -Actin, the membranes were blocked in Tris-buffered saline containing 0.1% Tween with 3% (wt/vol) non-fat dry milk at room temperature for at least 1 h. For pACC, pAMPK and AMPK detection, the membranes were blocked in Tris-buffered saline containing 0.1% Tween with 3% (wt/vol) BSA for at least 1 h. The primary antibody was incubated at 4°C overnight. Primary antibodies against the following targets were used: alpha subunit of AMP-activated protein kinase (AMPK), AMPK phosphorylated at threonine 172 (pAMPK), Acetyl-CoA Carboxylase phosphorylated at serine 79 (pACC), p38 MAP kinase phosphorylated at threonine 180 and tyrosine 182 (p-p38MAPK) and β -actin. The antibodies were purchased from Cell Signaling Technology and Sigma-Aldrich (for detailed information see Table 1). Secondary antibodies conjugated to HRP (anti-rabbit IgG – HRP, DAKO Cytomation, Germany, 1:6000 and anti-mouse IgG – HRP, Dianova, Germany, 1:20,000) were used to detect specific protein bands. The protein amount was calculated as the ratio of band intensities (pACC, p-p38 MAPK, AMPK, pAMPK protein vs β -actin protein and pAMPK protein vs AMPK protein respectively) in the same blot to correct for differences in protein loading.

Statistical analysis

Statistical analysis was performed using GraphPad Prism software (GraphPad Software, Inc.). Student's *t*-test was used (after checking for normal distribution) to compare the relative quantity of two adiponectin receptor transcripts (AdipoR1 and AdipoR2) and two glucose transporter transcripts (GLUT4 and GLUT8). Student's *t*-test was used to compare the glucose uptake between control and adiponectin-stimulated blastocysts; one-way ANOVA and Tukey's *post hoc* test were used to compare the glucose uptake between the three groups of embryoid bodies. Relative protein amounts were evaluated using SimaPlot v.12.0 (Systat Software GmbH, Erkrath, Germany). Levels of significance between groups were calculated using Student's *t*-test after checking for normal distribution. Multiple comparisons were made by factorial variance analysis (ANOVA) adjusted according to Bonferroni. All experiments were repeated at least three times. Differences of $P < 0.05$ were considered significant.

Figure 1 RT-PCR analysis of GLUT4 and GLUT8 transcripts in mouse blastocysts of CD-1 strain. Agarose gels with separated PCR products are shown. Lanes: MW, molecular weight markers; RT+, cDNA synthesised with reverse transcriptase; RT-, reverse transcriptase negative controls; Blank, reaction with water instead cDNA; PC, positive control tissue cDNA (heart for GLUT4, brain for GLUT8). The sizes of the MW and the predicted sizes of the PCR products in base pairs (bp) are indicated to the left and the right of the panels, respectively.

Results

Expression of GLUT4 and GLUT8 transcripts in mouse blastocysts

Controversial results have been published on the expression of GLUT4 in mouse blastocysts. We examined the expression of two hormone-responsive glucose transporters GLUT4 (*Slc2a4*) and GLUT8 (*Slc2a8*) in mouse blastocysts of CD-1 strain. We detected PCR products corresponding to both examined glucose transporters GLUT4 (410 bp) and GLUT8 (464 bp) in mouse blastocysts of CD-1 strain. No PCR products were detected in the reactions where reverse transcriptase (RT-) or cDNA (Blank) were omitted (Fig. 1). Sequencing the PCR products confirmed the GLUT4 and GLUT8 sequence: 100% sequence identity for GLUT4 and 99.8% sequence identity for GLUT8 was found between the PCR products and corresponding GenBank reference sequences (NM_009204 and NM_019488 respectively).

Expression of GLUT4 and GLUT8 proteins in mouse blastocysts and the effect of adiponectin on their intracellular localization

Our immunohistochemical study showed the presence of GLUT4 and GLUT8 proteins in CD-1 mouse blastocysts. The fluorescence signal produced by GLUT4 was evenly distributed in TE and ICM cells, the signal for GLUT8 was slightly stronger in TE cells (Fig. 2A and C). No signal was detected in controls incubated without the primary antibody (Fig. 2E) or without the primary and the secondary antibody (data not shown). After stimulation with adiponectin (10 μ g/mL for 45 min), GLUT8 relocated to the periphery of the cell (Fig. 2B vs A). On the other hand, we did not detect significant changes in GLUT4 localization after the adiponectin treatment (Fig. 2D vs C).

Figure 2 Immunohistochemical detection of GLUT4 and GLUT8 in mouse blastocysts of CD-1 strain. (A and B) Blastocysts incubated with the antibody against GLUT8 (green fluorescence signal); (C and D) blastocysts incubated with the antibody against GLUT4 (green fluorescence signal); (E) negative control with omitted primary antibody; (F, G, H, I and J) cell nuclei were counter-stained with Hoechst 33342 (blue fluorescence signal; identical samples as in images A, B, C, D and E); (K, L, M and N) merged images (K=A+F, L=B+G, M=C+H, N=D+I). Embryos presented in images B, (G, L) and D (I, N) were treated with adiponectin (10 µg/mL for 45 min). Upon the adiponectin treatment, GLUT8 fluorescence signal was concentrated at the cell periphery (B vs. A). Only a slight increase in the GLUT4 signal intensity at the cell periphery was detected after the adiponectin treatment (D vs C). A representative staining of a blastocyst from each group is shown. Scale bars: 30 µm.

Expression of AdipoR1 and AdipoR2 receptors in ECC P19 embryoid bodies

Since no information on expression of AdipoR1 and AdipoR2 receptors in mouse embryonic carcinoma cells is currently available, we examined the expression of AdipoR1 and AdipoR2 receptors in embryoid bodies (EBs) formed from embryonic carcinoma P19 (ECC P19) cells. We detected PCR products corresponding to both adiponectin receptors *AdipoR1* (132 bp) and *AdipoR2* (258 bp) in the EBs formed from ECC P19 cells. No PCR products were detected in the reactions where reverse transcriptase or cDNA were omitted (Fig. 3A). Sequencing the PCR products confirmed the *AdipoR1* and *AdipoR2* sequence: 99.2% sequence identity for *AdipoR1* and 99.6% sequence identity for *AdipoR2* was found between the PCR products and corresponding GenBank reference sequences (NM_028320 and NM_197985 respectively).

In our immunohistochemical study we examined the presence of AdipoR1 and AdipoR2 proteins in the ECC P19 EBs. A strong fluorescence signal was detected in

EBs incubated with the antibody against the AdipoR1 and AdipoR2 respectively (Fig. 3B). A weak background signal was detected in controls incubated without primary (Fig. 3B) or without primary and secondary antibodies (data not shown).

Expression of T-cadherin mRNA in mouse preimplantation embryos and ECC P19 embryoid bodies

T-cadherin can bind several molecules including hexameric and high-molecular-weight forms of adiponectin (Hug et al. 2004). T-cadherin is expressed in different tissues (for review see Philippova et al. 2009); however, no data on its expression in early embryonic cells are available. We examined T-cadherin transcript expression in ovulated oocytes and preimplantation embryos at four developmental stages, using several independent cDNA samples for each stage (eight samples for blastocysts and five samples for lower embryo stages and oocytes were used). In ovulated oocytes, T-cadherin mRNA was consistently detected

Figure 3 AdipoR1 and AdipoR2 expression in EBs formed from ECC P19 cells. (A) RT-PCR analysis of AdipoR1 and AdipoR2 receptor transcripts in ECC P19 EBs. Agarose gels with separated PCR products are shown. Lanes: MW, molecular weight markers; RT+, cDNA synthesised with reverse transcriptase; RT-, reverse transcriptase negative controls; Blank, reaction with water instead of cDNA; PC, positive control tissue cDNA (brain). The sizes of the MW and the predicted sizes of the PCR products in base pairs (bp) are indicated to the right and the left of the panels, respectively. (B) Immunohistochemical detection of AdipoR1 and AdipoR2 receptors in ECC P19 EBs (representative images are shown); (a and b) EBs incubated with the antibody against AdipoR1 and AdipoR2, respectively; (c) negative control with omitted primary antibody; (d, e and f) cell nuclei stained with Hoechst 33342 (identical samples as in images a, b and c). (g) Adiponectin receptor cellular localization; a merged image of F-actin staining (plasma membrane visualization, red), Hoechst 33342 staining (cell nuclei, blue) and the antibody against adiponectin receptor visualized with Alexa Fluor 488-coupled secondary antibody (green signal). Scale bars: 30 μ m.

(i.e., in 100% samples). In preimplantation embryos, T-cadherin transcript was detected inconsistently: it was present in 40% of four-cell embryo samples, in 20% of 8- to 16-cell embryo samples and in 37.5% of blastocyst samples; T-cadherin mRNA was not detected in any morula sample (Supplementary Fig. 1). In ECC P19 EBs we detected mRNA for T-cadherin in all four examined cDNA samples (Supplementary Fig. 2).

Relative quantity of adiponectin receptor and glucose transporter transcripts in mouse blastocysts and ECC P19 embryoid bodies

We compared the relative quantity of consistently expressed AdipoR1 and AdipoR2 transcripts, and GLUT4 and GLUT8 transcripts in both mouse blastocysts and ECC P19 EBs. Starting fluorescences (proportional to mRNA quantities) were calculated from AdipoR1, AdipoR2, GLUT4 and GLUT8 amplifications, and normalized with beta actin starting fluorescences. The normalized starting fluorescences are shown in Table 2. Our results showed that the amount of AdipoR1 transcript was about two times higher than the amount of AdipoR2 transcript in both cell types. The amount of GLUT8 transcript was much higher than the amount of GLUT 4 transcript in both cell types (about 15 times higher in blastocysts and about 10 times higher in embryoid bodies; Table 2).

Effect of adiponectin on glucose uptake in mouse blastocysts and ECC P19 embryoid bodies

We examined the effects of two adiponectin protein isoforms on glucose uptake in mouse blastocysts and ECC

P19 EBs. We found significantly higher uptake of 3-OMG in the blastocysts treated with full-length adiponectin (FLAD) compared to the control group (CTRL). On the other hand, we did not find any significant increase in glucose uptake after treatment with globular adiponectin (GLAD) compared to the control group (Fig. 4).

Similarly to the blastocysts, we found significantly higher uptake of 3-OMG in the ECC P19 EBs stimulated with full-length adiponectin compared to controls. No difference in glucose uptake between globular adiponectin-treated and control blastocysts was found (Fig. 5).

To study the involvement of AMP kinase, the effect of compound C (dorsomorphin, a potent AMPK inhibitor) on the full-length adiponectin-stimulated glucose uptake was examined in the next experiment. The increased

Table 2 Relative quantity of *AdipoR1* and *AdipoR2* transcripts, and *GLUT4* and *GLUT8* transcripts in mouse blastocysts and ECC P19 EBs.

Genes (ratios)	Blastocysts	ECC P19 Ebs
<i>AdipoR1</i>	4.48 $\pm$ 0.15	2.13 $\pm$ 0.10
<i>AdipoR2</i>	2.43 $\pm$ 0.09***	1.04 $\pm$ 0.06***
<i>AdipoR1/AdipoR2</i> ratio	1.85 $\pm$ 0.09	2.05 $\pm$ 0.08
<i>GLUT4</i>	7.77 $\pm$ 0.49	2.38 $\pm$ 0.09
<i>GLUT8</i>	114.9 $\pm$ 6.23***	24.29 $\pm$ 2.45***
<i>GLUT8/GLUT4</i> ratio	14.93 $\pm$ 0.82	10.24 $\pm$ 0.78

Normalized starting fluorescences were multiplied by a factor of 1×10^2 (*AdipoR1* and *AdipoR2* fluorescences) or by a factor of 1×10^4 (*GLUT4* and *GLUT8* fluorescences) to obtain numbers in a more presentable format. Values are arithmetical means $\pm$ s.e.m. ($n=6$). Statistical significance of differences between *AdipoR1* and *AdipoR2* values, and between *GLUT4* and *GLUT8* values was assessed with unpaired Student's *t*-test: *** $P < 0.001$.

Figure 4 Uptake of glucose (3-OMG) in mice blastocysts. In the first set of experiments, the blastocysts were incubated in the presence or absence of full-length adiponectin for 2 h; CTRL, control blastocysts, $n=50$; FLAD, full-length adiponectin-treated blastocysts $n=70$ (5 blastocysts per sample were used for radioactivity measurement; data from two experiments were pooled). In the second set of experiments, the blastocysts were incubated in the presence or absence of globular adiponectin for 2 h; CTRL, control blastocysts, $n=60$; GLAD, globular adiponectin-treated blastocysts, $n=70$ (5 blastocysts per sample were used for radioactivity measurement; data from two experiments were pooled). Statistical differences between the control and adiponectin-treated groups were assessed with unpaired Student's *t*-test: *** $P<0.001$. Means of the glucose uptake in control groups were set 1 and values of glucose uptake in adiponectin-treated groups are expressed relative to them. Values of relative glucose uptake in adiponectin-treated groups are arithmetical means $\pm$ S.E.M.

glucose uptake in adiponectin-stimulated EBs was only slightly reduced through compound C treatment (Fig. 6).

Activation of the adiponectin signaling pathway in ECC P19 embryoid bodies

To analyse the involvement of AMP and p38 MAP kinases in adiponectin signaling the ECC P19 embryoid body model was used. Adiponectin treatment stimulated AMPK phosphorylation. This effect was detectable after 10-min treatment and was most evident after the 2-h treatment (Fig. 7). The adiponectin-induced phosphorylation of AMPK was blocked by AMPK inhibitor Compound C (Fig. 8). In correlation with AMPK activation, phosphorylation of ACC was tendentially increased (1.2 fold with $P=0.06$), and this increase was reduced by Compound C (Figs 7 and 8).

Adiponectin stimulated p38 MAPK phosphorylation after 5 and 10 min of treatment. After 60-min treatment the amount of phosphorylated p38 MAPK decreased to the unstimulated level (Fig. 7). Compound C did not influence the adiponectin-induced phosphorylation of p38 MAPK, indicating that AMPK did not mediate this effect (Fig. 8).

Figure 5 Uptake of glucose (3-OMG) in single ECC P19 EBs. The EBs were incubated in the presence or absence of full-length adiponectin or globular adiponectin for 1 h. CTRL, control EBs ($n=35$), FLAD, full-length adiponectin-treated EBs ($n=20$); GLAD, globular adiponectin-treated EBs ($n=26$). Values are arithmetical means $\pm$ S.E.M. Statistical differences between the groups were assessed with ANOVA followed by Tukey's test, * $P<0.05$, ** $P<0.01$.

Discussion

Several studies have demonstrated that adiponectin is involved in the female fertility regulation and can influence oocyte maturation and preimplantation embryo development. Adiponectin receptors AdipoR1 and AdipoR2 have been detected in oocytes and

Figure 6 Effect of compound C on adiponectin-stimulated glucose uptake in ECC P19 EBs. The EBs were incubated in the presence or absence of full-length adiponectin or full-length adiponectin with 10 μ M compound C. DMSO, control EBs (0.1% DMSO in the medium, $n=20$); FLAD, full-length adiponectin-treated EBs ($n=20$); FLAD+CC full-length adiponectin and 10 μ M compound C-treated EBs ($n=19$). Values are arithmetical means $\pm$ S.E.M. Statistical differences between the groups were assessed with ANOVA followed by Tukey's test, * $P<0.05$.

Figure 7 Relative protein amounts of phospho-p38 MAPK (p-p38MAPK), phospho-AMPK (pAMPK), AMPK and phospho-ACC (pACC) in adiponectin-stimulated P19 ECC EBs. Relative protein amounts after 5, 10, 15, 60 and 120 min of adiponectin treatment. Protein amounts are given relative to untreated controls which were set 1 and are indicated with the dashed line. Values are arithmetical means $\pm$ S.E.M. ($n \geq 3$). Statistical differences between the control and adiponectin-treated groups were assessed with unpaired Student's *t*-test: $*P < 0.05$. Representative immunoblots are shown at the bottom right.

preimplantation embryos of several mammalian species (for review see Čikoš 2012, Palin *et al.* 2012, Dobrzym *et al.* 2018, Barbe *et al.* 2019). The results presented in this work show that T-cadherin, the third putative adiponectin receptor, is expressed in mouse oocytes. Our results suggest that T-cadherin maternal transcript is gradually degraded after fertilization, and the expression is resumed at the blastocyst stage. T-cadherin is a glycosylphosphatidylinositol-anchored extracellular protein that may act as a coreceptor in adiponectin binding and signaling (Hug *et al.* 2004); however, the exact molecular mechanisms of adiponectin action in early embryonic cells are not known. There are data

indicating that adiponectin (but not insulin) can stimulate glucose transport in rabbit blastocysts (Navarrete Santos *et al.* 2004a, Fischer *et al.* 2010). On the other hand, the stimulation of glucose uptake by insulin has been documented in mouse blastocysts (Pantaleon & Kaye 1996, Carayannopoulos *et al.* 2000, Pinto *et al.* 2002). Our results show that, in addition to insulin, adiponectin is another hormone capable of stimulating glucose uptake in mouse blastocysts. Moreover, our comparison of two adiponectin isoforms shows that full-length adiponectin, but not globular adiponectin, can stimulate glucose uptake in mouse blastocysts. This is in contrast with data obtained from muscle cells, where globular

Figure 8 Effect of Compound C (CC) on adiponectin-stimulated protein amounts. Protein amounts after 10 min of adiponectin treatment are given relative to untreated controls (ctrl). Values are arithmetical means $\pm$ S.E.M. ($n \geq 3$). Statistical differences between the groups were assessed with factorial variance analysis (ANOVA) adjusted according to Bonferroni. Statistical significances are indicated with different letters ($P < 0.05$). Ctrl, EBs without treatment; DMSO (dimethyl sulfoxide), solvent control for Compound C. Representative immunoblots are shown at the bottom right.

adiponectin was shown to stimulate glucose uptake. In other experiments using muscle cells, the globular but not the full-length isoform was effective (Ceddia *et al.* 2005, Fang *et al.* 2005, Palanivel *et al.* 2007, Vu *et al.* 2013).

In rabbit blastocysts, GLUT4 has been identified as the glucose transporter responsible for adiponectin-stimulated glucose transport (GLUT8 was not examined for this function, Fischer *et al.* 2010). In mouse blastocysts, the expression of GLUT4 was disputed (Purcell & Moley 2009), and GLUT8 was identified as the glucose transporter responsible for insulin-stimulated glucose transport (Carayannopoulos *et al.* 2000, Pinto *et al.* 2002). These data indicate that two insulin and/or adiponectin-sensitive facilitative glucose transporters are expressed and function in preimplantation embryos. To answer the question whether both hormone-sensitive transporters are present in mouse blastocysts, we examined GLUT4 and GLUT8 expression and detected both transporters (at mRNA as well as protein level) in mouse CD-1 blastocysts. Comparing relative quantity of transcripts, we found that the amount of GLUT8 transcript was about 15 times higher than the amount of GLUT 4 transcript. Moreover, our results showed that GLUT8 protein translocated to the cell membrane in blastocyst cells stimulated with full-length adiponectin, indicating that GLUT8 is involved in the adiponectin-stimulated glucose uptake in mouse blastocysts. On the other hand, we did not detect significant changes in GLUT4 protein localization after the adiponectin treatment, which could be caused by the lower amount of GLUT 4 than GLUT8 in mouse blastocyst cells.

Mouse blastocysts provide very limited amounts of biological material for biochemical analyses, so we used embryoid bodies (EBs) derived from mouse embryo carcinoma cell line P19 (ECC P19) as a model for studying adiponectin signaling in detail. To validate this model, we first examined the expression of adiponectin receptors in these cells. Our results show that adiponectin receptors are expressed in ECC P19 EBs and that the relative quantity of AdipoR1 and AdipoR2 transcripts is similar to that found in blastocysts. Several previous studies have demonstrated the expression of glucose transporters in ECC P19 EBs (Tonack *et al.* 2007, Knelangen *et al.* 2012), and the relative quantification of GLUT4 and GLUT8 mRNA showed a similar ratio of these transcripts in ECC P19 EBs and in mouse blastocysts. These results indicate that ECC P19 embryoid bodies represent a suitable model for the study of adiponectin signaling and glucose transport regulation in early embryonic cells.

Previous experimental data indicate that adiponectin regulates glucose and lipid metabolism via AMPK in rabbit blastocysts (Fischer *et al.* 2010, Schindler *et al.* 2017). We analysed the potential involvement of AMPK in the adiponectin-stimulated glucose uptake in mouse ECC P19 EBs. Administration of a potent

AMPK inhibitor, Compound C, only slightly reduced the adiponectin-stimulated glucose uptake which suggests that other signaling molecules participate in the regulation of glucose transport by adiponectin in these cells. Our next results supported this assumption. AMPK phosphorylation was moderately increased after 10 min of adiponectin treatment, and was most evident at the last checked time point in the 2-h treatment. A stronger response to adiponectin stimulation was observed in p38 MAPK, where the amount of phosphorylated protein doubled after 5 and 10 min of treatment. The AMPK inhibitor did not influence the adiponectin-stimulated p38 MAPK phosphorylation, indicating independent actions of the two signaling pathways. These results indicate that the p38 MAPK protein is a potential target of adiponectin in mouse embryonic cells. Activation of p38 MAPK by adiponectin has been demonstrated in several cell types. Yamauchi *et al.* (2003) showed that besides AMPK, p38 MAPK activation mediated adiponectin-stimulated glucose uptake and fatty acid oxidation in C2C12 myocytes. Although some reports suggest that p38 MAPK activation is downstream to AMPK activation (Yoon *et al.* 2006), other studies have demonstrated AMPK-independent activation of p38 MAPK by adiponectin (Charlton *et al.* 2010, Xin *et al.* 2011). The involvement of p38 MAPK in the regulation of glucose transport has been shown in rat cardiomyocytes in primary culture, hibernating swine myocardium and human gastric cancer cells (McFalls *et al.* 2004, Montessuit *et al.* 2004, Liu *et al.* 2015). Studying early embryos, Sozen *et al.* (2015) demonstrated reduced glucose uptake and decreased GLUT4 expression in mouse preimplantation embryos treated with the p38 MAPK inhibitor SB203580. Rabbit blastocysts developing in diabetic conditions showed increased adiponectin signaling (Schindler *et al.* 2013) accompanied by increased p-p38 MAPK levels (Schindler *et al.* 2017).

The stimulation of glucose transport by adiponectin can be important in the case of reduced levels of circulating insulin. Elevation of adiponectin concentration has been well documented in patients with type 1 diabetes mellitus (Imagawa *et al.* 2002, Lindström *et al.* 2006, Majewska *et al.* 2016), although a controversy exists regarding the molecular mechanisms involved (Fasshauer *et al.* 2002, Blümer *et al.* 2008). Increased expression of AdipoR1 and AdipoR2 mRNAs has been found in the muscle tissue of streptozotocin-induced diabetic mice, and insulin administration restored the expression level (Tsuchida *et al.* 2004, Inukai *et al.* 2005). Negative correlation between adiponectin receptor expression and insulin has been confirmed in several studies. Insulin downregulated AdipoR1 and AdipoR2 transcripts in rat skeletal muscle (L6) cells, but not in mouse 3T3-L1 adipocytes (Sattar & Sattar 2012). On the other hand, the inhibitory effect of insulin on AdipoR2 mRNA expression was shown in porcine adipocytes (Liu *et al.*

2008), and a similar effect of insulin on adiponectin receptor expression was demonstrated in mouse and human hepatocytes (Tsuchida *et al.* 2004, Shimizu *et al.* 2007) as well. These data indicate that a decrease in insulin can stimulate adiponectin action by elevating the adiponectin concentration in the circulation and by increasing adiponectin receptor expression in some tissues. Increased expression of AdipoR1 transcript found in rabbit blastocysts isolated from diabetic females suggests a compensatory role of adiponectin for lack of insulin in early embryos (Schindler *et al.* 2013).

In summary our results demonstrate that besides insulin, adiponectin can be another hormone capable of stimulating glucose uptake in mouse blastocysts. We show that both hormone-sensitive facilitative glucose transporters (GLUT4 and GLUT8) are expressed in mouse blastocysts and especially GLUT8 is involved in the adiponectin-stimulated glucose uptake. Using a mouse embryonic carcinoma cell model, we show that besides AMPK, p38 MAPK participates in adiponectin signaling in early embryonic cells. Our results suggest that adiponectin can help maintain the glucose supply for early embryos under hypoinsulinaemic conditions, for example, in mothers suffering from type 1 diabetes mellitus.

Supplementary materials

This is linked to the online version of the paper at <https://doi.org/10.1530/REP-19-0251>.

Declaration of interest

The authors declare that there is no conflict of interest that could be perceived as prejudicing the impartiality of the research reported.

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Author contribution statement

J Bu and J Ba designed and performed experiments, analysed and interpreted data. M S, J S J, A Š, M K and V K performed experiments and analysed results. D F was involved in designing experiments, analysed and interpreted data, critically revised the article. B F and J K interpreted data and critically revised the article. A N S contributed to the

conception of the work, designed and performed experiments and analysed results. Š Č contributed to the conception of the work, designed experiments, analysed and interpreted data and wrote the article.

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