FOSL1 Is Integral to Establishing the Maternal-Fetal Interface[∇]

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Remodeling of uterine spiral arteries by trophoblast cells is a requisite process for hemochorial placentation and successful pregnancy. The rat exhibits deep intrauterine trophoblast invasion and accompanying trophoblast-directed vascular modification. The involvement of phosphatidylinositol 3 kinase (PI3K), AKT, and Fos-like antigen 1 (FOSL1) in regulating invasive trophoblast and hemochorial placentation was investigated using Rcho-1 trophoblast stem cells and rat models. Disruption of PI3K/AKT with small-molecule inhibitors interfered with the differentiation-dependent elaboration of a signature invasive-vascular remodeling trophoblast gene expression profile and trophoblast invasion. AKT isoform-specific knockdown also affected the signature invasive-vascular remodeling trophoblast gene expression profile. Nuclear FOSL1 increased during trophoblast cell differentiation in a PI3K/AKT-dependent manner. Knockdown of FOSL1 disrupted the expression of a subset of genes associated with the invasive-vascular remodeling trophoblast phenotype, including the matrix metallopeptidase 9 gene (Mmp9). FOSL1 was shown to occupy regions of the Mmp9 promoter in trophoblast cells critical for the regulation of Mmp9 gene expression. Inhibition of FOSL1 expression also abrogated trophoblast invasion, as assessed in vitro and following in vivo trophoblast-specific lentivirally delivered FOSL1 short hairpin RNA (shRNA). In summary, FOSL1 is a key downstream effector of the PI3K/AKT signaling pathway responsible for development of trophoblast lineages integral to establishing the maternal-fetal interface.

Hemochorial placental development is characterized by close contact between maternal and fetal tissues and occurs in primates and rodents such as the rat and mouse. Trophoblast cells are the functional units of the placenta. One of the key activities of trophoblast cells is remodeling uterine spiral arteries. Specific lineages of trophoblast cells exit the placenta and enter into the uterine parenchyma, where they interact with the vasculature. Vascular remodeling transforms tightly coiled uterine spiral arteries into dilated vessels that are no longer under maternal control (45, 63). Restructuring maternal vasculature is essential for the optimal delivery of nutrients to the fetus. Poor trophoblast invasion is linked to miscarriage, preeclampsia, preterm birth, and fetal growth restriction (36, 45, 63). Hemochorial placentation and especially trophoblastdirected vascular remodeling in the human and the rat are remarkably similar (3, 14, 49, 63, 76). In both species, invasive trophoblast cells penetrate into the uterus through both endovascular and interstitial routes.

In the rat, invasive trophoblast cells arise from the junctional zone and move into the uterus in two waves. The first wave is initiated during midgestation and consists of trophoblast cell invasion into spiral arteries situated within the central area of the uterine mesometrial implantation site (3). These invasive endovascular trophoblast cells effectively replace the maternal endothelium (3, 68). The depth of entry into the uterus is generally restricted to the decidua basalis. The second wave of trophoblast cell invasion begins around gestation day 14.5, exhibiting both endovascular and interstitial courses of entry and characterized by deep penetration into the metrial gland (3, 14, 49, 76). Signaling pathways controlling trophoblast invasion and vascular remodeling are not well understood.

The phosphatidylinositol 3-kinase (PI3K)/AKT signaling pathway has been implicated as a potential regulator of trophoblast invasion. PI3K responds to various extracellular signals, leading to the activation of a serine/threonine kinase termed AKT (13, 15, 29). AKT has a wide range of substrates involved in many cellular processes, including metabolism, cell cycle, survival, protein synthesis, and differentiation (29). There are three AKT isoforms in mammals, AKT1, AKT2, and AKT3 (12). These enzymes possess shared and potentially unique substrate specificities (12, 22). Null mouse models for the three AKT isoforms exhibit a range of phenotypes, characterized by disruptions in placentation, fetal growth, and/or postnatal growth and metabolism (19, 20, 27, 84, 85). PI3K/AKT becomes constitutively activated upon differentiation of trophoblast stem (TS) cells (43) and regulates a set of genes associated with an invasive-vascular remodeling trophoblast phenotype (43, 46, 65, 66, 74). Among these PI3K/ AKT-regulated genes is the matrix metallopeptidase 9 gene (Mmp9). MMP9 is a proteolytic enzyme involved in the breakdown of extracellular matrix and tissue remodeling (47, 61). In trophoblast cells, MMP9 is proinvasive (11, 53, 55, 62, 66) and also participates in the regulation of cancer cell invasion and metastasis (23, 47). Downstream molecular mechanisms un-

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derlying PI3K/AKT regulation of trophoblast cell invasion have not been elucidated.

Fos-like antigen 1 (FOSL1) is a potential downstream target/effector of the PI3K/AKT signaling pathway leading to the activation of cell invasion (16, 18, 67). FOSL1 is a member of the basic region-leucine zipper transcription factor family and a contributor to the formation of the activator protein-1 (AP-1) transcription factor complex (28). In the human placenta, FOSL1 is expressed in several trophoblast cell types, including the extravillous/invasive trophoblast cell population (9, 57). *Fosl1* is also expressed in differentiating rat trophoblast cells (46). Mutant mice possessing a null mutation at the *Fosl1* locus die *in utero* due to placental defects (72). FOSL1 represents a candidate mediator of the PI3K/AKT signaling pathway in trophoblast cells.

In this study, we investigated the involvement of PI3K/AKT and FOSL1 signaling in the regulation of trophoblast cell differentiation, especially the acquisition of the invasive phenotype. *In vitro* and *in vivo* research strategies were performed utilizing Rcho-1 TS cells (30, 69) and trophoblast-specific lentivirally mediated gene manipulation (52), respectively. Collectively, the experimental findings demonstrate that PI3K/AKT and FOSL1 participate in a signal transduction pathway controlling a proinvasive/provascular remodeling trophoblast cell phenotype.

MATERIALS AND METHODS

Animals and tissue collection. Holtzman Sprague-Dawley rats were obtained from Harlan Laboratories (Indianapolis, IN). Animals were housed in an environmentally controlled facility with lights on from 0600 to 2000 h and were allowed free access to food and water. Timed pregnancies were generated by cohabitation of male and female rats. The presence of a seminal plug or sperm in the vaginal smear was designated day 0.5 of pregnancy. Rat placental tissues were collected on gestation days 7.5, 11.5, and 18.5. Tissues for histological analysis were frozen in dry ice-cooled heptane and stored at -80° C until processing. Tissue samples for RNA or protein extraction were frozen in liquid nitrogen and stored at -80° C until processing. Female rats were made pseudo-pregnant by mating with vasectomized males. The presence of seminal plugs was designated day 0.5 of pseudopregnancy. The University of Kansas Animal Care and Use Committee approved protocols for the care and use of animals.

Maintenance of the Rcho-1 TS cells. Rcho-1 TS cells were maintained at subconfluent conditions in stem cell medium (RPMI-1640 culture medium [Cellgro, Herndon, VA] supplemented with 20% fetal bovine serum [FBS; Atlanta Biologicals, Norcross, GA], 50 µM 2-mercaptoethanol [Sigma-Aldrich, St. Louis, MO], 1 mM sodium pyruvate [Cellgro], 100 µM penicillin, and 100 U/ml streptomycin [Cellgro]), as previously reported (69). Differentiation was induced by growing cells to near confluence and then replacing the stem cell medium with differentiation medium (NCTC-135 medium supplemented with 1% horse serum [HS; Atlanta Biologicals], 50 µM 2-mercaptoethanol, 1 mM sodium pyruvate, 10 mM HEPES, 4-(2-hydroxyethyl)-1-piperazineethanesulfonic acid [Fisher, Pittsburgh, PA], 38 mM sodium bicarbonate [Fisher], 100 µM penicillin, and 100 U/ml streptomycin [Cellgro]). High cell density and the absence of sufficient growth stimulatory factors (removal of FBS) facilitate trophoblast cell differentiation (69). Trypsin-EDTA (EDTA, in Hanks' balanced salt solution [Cellgro]) was used to passage the cells. Cells in the stem cell condition were grown in stem cell medium and collected 24 h after subculture to restrict the accumulation of spontaneously differentiating cells. Cells in the differentiation condition were grown for 8 days in differentiation medium prior to harvesting unless otherwise noted.

Inhibition of PI3K and AKT. LY294002 (LY; Calbiochem, La Jolla, CA) was used to inhibit PI3K (77). AKT inhibitor IX (Calbiochem) was used to inhibit AKT activities (42). Rcho-1 TS cells were grown to near confluence and then shifted to differentiation medium containing vehicle (0.1% final concentration of dimethyl sulfoxide [DMSO]), LY (10μ M), or AKT inhibitor IX (10μ M). Cells were harvested after 8 days of treatment. Culture medium and small-molecule inhibitors were replaced daily.

Western blotting. Whole-cell lysates were prepared in radioimmunoprecipitation buffer (10 mM Tris-HCl, pH 7.2, 1% Triton X-100 or 1% Nonidet P-40, 1% sodium deoxycholate, 0.1% SDS, 150 mM NaCl, 5 mM EDTA, 1 mM sodium orthovanadate, 1 mM phenylmethylsulfonyl fluoride [PMSF], 10 µg/ml aprotinin) or in cell lysate buffer (Cell Signaling Technology, Danvers, MA). Nuclear lysates were prepared as previously described (24) using an extraction buffer consisting of 25 mM HEPES, pH 7.5, 1 mM dithiothreitol (DTT), 10% glycerol, 0.5 M KCl, 0.5 mM PMSF, 10 mM β-glycerophosphate, 2 mM sodium orthovanadate, 1 mM sodium pyrophosphate, and 10 mM sodium fluoride. Whole-cell lysates or nuclear lysates were fractionated by SDS-PAGE. Separated proteins were electrophoretically transferred to nitrocellulose. Filters were probed with the designated antibodies overnight at 4°C. Antibodies specific for pan-AKT, phospho-Ser 473 AKT, AKT1, and AKT2 were obtained from Cell Signaling Technology. Antibodies to FOSL1 and TATA box binding protein (TBP) were obtained from Santa Cruz Biotechnology (Santa Cruz, CA). The antibody for ACTB (clone AC-15) was obtained from Sigma-Aldrich. The following antibodies were used at the dilutions indicated (in parentheses): pan-AKT (1:2,000), phospho-Ser 473 AKT (1:2,000), AKT1 (1:1,000), AKT2 (1:1,000), FOSL1 (1:1,000), TBP (1:2,000), and ACTB (1:8,000). Blots were incubated with horseradish peroxidase-conjugated antibodies to rabbit (Cell Signaling Technology) or mouse (Sigma-Aldrich) IgG for 1 h at room temperature. Reaction products were visualized by incubation with enhanced chemiluminescence according to the manufacturer's instructions (ECL, Amersham Pharmacia, Piscataway, NJ).

AKT kinase assay. The activity of AKT was measured in Rcho-1 TS cell lysates using an AKT kinase assay kit (Cell Signaling Technology). AKT activity was measured in whole-cell lysates from Rcho-1 TS cells treated with LY, AKT inhibitor IX, or vehicle control. The assay was performed according to the manufacturer's instructions. Briefly, phospho-AKT (Ser 473) was immunoprecipitated from 400 μ g of Rcho-1 TS cell lysates. Kinase activity was assessed from the immunoprecipitates in the presence of ATP (200 μ M) and glycogen synthase kinase $3\alpha/\beta$ (GSK $3\alpha/\beta$)–glutathione *S*-transferase fusion protein (1 μ g). Phosphorylation (P) of the GSK $3\alpha/\beta$ fusion protein substrate was identified by Western blotting and represented a measure of AKT kinase activity.

Quantitative reverse transcription-PCR (qRT-PCR). RNA samples were extracted using TRIzol (Invitrogen) according to the manufacturer's instructions. cDNAs were reverse transcribed from RNA by using reagents from Promega (Madison, WI) according to the manufacturer's instructions. SYBR green PCR master mix (Applied Biosystems, Foster City, CA) was used in the PCR. Reactions were processed using a 7500 real-time PCR system (Applied Biosystems). Conditions included an initial holding stage (50°C for 2 min and 95°C for 10 min) and 40 cycles (95°C for 15 s and 60°C for 1 min) followed by a dissociation stage (95°C for 15 s, 60°C for 1 min, and then 95°C for 15 s). Primers are provided in Table 1. The comparative cycle threshold method ($\Delta\Delta CT$) was used for relative quantification of the amount of mRNA for each sample normalized to 18S RNA. Values are presented relative to controls for each gene.

Matrigel invasion assay. The invasive ability of Rcho-1 TS cells were measured as previously described (62). Rcho-1 TS cells were differentiated for 6 days, trypsinized, and placed on Matrigel invasion chambers (BD Biosciences, San Jose, CA) at a density of 2.5×10^4 cells per chamber. Cells were initially plated in stem cell medium and then switched to differentiation medium and allowed to invade for 72 h. After 72 h, membranes were collected and stained with Diff-Quick (Dade Behring, Newark, DE). Invading cells were visualized by light microscopy and counted for each sample.

Gelatin zymography. MMP9 activity was measured in Rcho-1 TS cell-conditioned medium as previously described (62). Conditioned medium was normalized to cell number and run on 8% polyacrylamide gels containing 1 mg/ml porcine gelatin. Gels were washed (wash buffer: 50 mM Tris, pH 8.0, 5 mM calcium chloride, 1 μ M zinc chloride, and 0.5% Triton X-100) and incubated with wash buffer without Triton X-100. Proteins were stained using Coomassie staining solution (45% methanol, 10% acetic acid, and 0.2% Coomassie brilliant blue R-250) and incubated in destaining solution (12% methanol and 7% acetic acid).

Immunohistochemistry. Immunohistochemical analyses were used to localize proteins in placental tissues. Ten-micrometer cryosections of placentation sites were prepared and stored at -80°C until use. Sections were fixed in ice-cold 4% paraformaldehyde and blocked in 10% normal goat serum for 30 min at room temperature. Primary antibody incubation was for 1 h at room temperature with antibodies specific for FOSL1 (dilution of 1:50; Santa Cruz Biotechnology), prolactin family 3, subfamily D, member 1 (PRL3D1, also called placental lactogen-I [PL-I]; 1:50 dilution) (35), prolactin family 3, subfamily B, member 1 (PRL3B1, also called PL-II; 1:50 dilution) (25), or pan-cytokeratin (1:300; Sigma-Aldrich). Sections were incubated with secondary antibodies for 30

Gene	GenBank accession no.	Forward primer	Reverse primer
Akt1	NM 033230	GGCACCTTTATTGGCTACAAG	AGCACCTGAGTTGTCACTG
Akt2	NM_017093	TGGCAGGATGTGGTACAGAA	AGGAGAACTGGGGGAAGTGT
Akt3	NM_031575	TGGTTCGAGAGAAGGCAAGT	ACAGCTCTCCCCCATTAACA
Adm	NM_012715	ACGTCTCGGACTTTCTGCTT	GCTGCTGGACGCTTGTAGTT
Cgm4	NM_012525	TAGCCCGATACAGAACAGCAA	AGGGTCACAGCATGAGGAAA
Čtsd	NM ⁻ 134334	TACCTGAACGTCACCCGAAA	CAGGCTGGACACCTTCTCAC
Faslg	NM_012908	TCTGGTTGGAATGGGGTTAG	CTTGGCTTTTTGGTTTCAGAG
Fosl1	NM_012953	ATCCCCGACCTCTGACCTAT	CAAGGCGTTCCTTCTGCTT
Igf2	NM_031511	GGAAGTCGATGTTGGTGCTT	CTTGCCCACGGGGTATCT
ĨĨ17f	NM 001015011	CAAAACCAGGGCATTTCTGT	GACCAGGATTTCTTGCTGGA
Mmp9	NM_031055	AACTTCGACGCTGACAAGAA	TTTAGAGCCACGACCATACAGA
Prl4a1	NM_017036	GACCACCAGATGCCACACTT	CAGGAGCTTTATGTTTGATTCCTT
Sema6d	NM 001107768	GGCCAGTGATGCTGTCATTT	TATGTTCCACGGCGATTTCT
Serpine1	NM_012620	AGTCTTTCCGACCAAGAGCA	GTGCCGAACCACAAAGAGAA
18Ŝ	M11188	GCAATTATTCCCCATGAACG	GGCCTCACTAAACCATCCAA

TABLE 1. qRT-PCR primer sequences

min at room temperature. For bright-field immunohistochemical analysis, a secondary goat anti-rabbit antibody conjugated with biotin (Sigma-Aldrich) was used. Avidin-peroxidase was added for 30 min at room temperature followed by color development with an AEC kit (Zymed Laboratories, San Francisco, CA). Tissues were counterstained with Mayer's hematoxylin. For fluorescence detection, a secondary goat anti-rabbit antibody conjugated with cyanine 3 bis-*N*-hydroxysuccinimide (NHS) ester (Cy3; Jackson ImmunoResearch Laboratories, Inc., West Grove, PA) or alternatively a secondary goat anti-mouse antibody conjugated with fluorescein isothiocyanate (FITC; Sigma-Aldrich) was used. Nuclei were visualized with DAPI (4',6-diamidino-2-phenylindole; Molecular Probes, Carlsbad, CA). Bright-field and fluorescence images were captured using a Leica MZFLIII stereomicroscope or a DMI 4000 microscope equipped with charge-coupled-device (CCD) cameras (Leica Microsystems GmbH, Welzlar, Germany).

shRNA constructs and production of lentivirus. Akt short hairpin RNA (shRNA) constructs in the pLKO.1 vector were obtained from Open Biosystems (Huntsville, AL). Two Fosl1 shRNAs were designed and subcloned into the pLKO.1 vector by using AgeI and EcoRI restriction sites. Control shRNA that does not target any known mammalian gene, pLKO.1-shSCR (plasmid 1864), was obtained from Addgene (Cambridge, MA) (71). Sequences representing the sense target site for each of the shRNAs used in the analyses are as follows: Akt1, GCACATCAAGATAACGGACTT; Akt2, GATGGATCTTTCATTGGGTAT; Akt3, GCTCTTGATAAAGGATCCAAA; Fosl1 No. 1, GACAAGTTGGAGG ATGAGAAAT; Fosl1 No. 2, GTTCCTCAGCCCATCGAAAGAGTA; Control-N, CCTAAGGTTAAGTCGCCCTCG. Third-generation lentiviral packaging vectors were acquired from Addgene and included the following: pMDLg/ pRRE (plasmid 12251), pRSV-Rev (plasmid 12253), and pMD2.G (plasmid 12259). Lentiviral particles were produced as previously reported (52). To summarize, 293FT cells (Invitrogen, Carlsbad, CA) were transiently transfected using Lipofectamine 2000 (Invitrogen) with the following plasmids: shRNA containing transducing vector, third-generation packaging system plasmids (pMDLg/pRRE and pRSV-Rev) (26), and a vesicular stomatitis virus glycoprotein (VSVG) envelope plasmid (pMD2.G). After transfection, cells were maintained in 50% Dulbecco's modified Eagle's medium (DMEM) with high glucose (Cellgro), 45% Opti-MEM I (Invitrogen, Carlsbad, CA), and 5% FBS. Culture supernatants containing lentiviral particles were harvested every 24 h for 2 to 3 days. Supernatants were centrifuged to remove cell debris, filter sterilized, concentrated by ultracentrifugation, and stored at -80°C until used. Lentiviral vector titers were determined by measurement of p24 gag antigen by enzyme-linked immunosorbent assay (ELISA) (Advanced Bioscience Laboratories, Kensington, MD).

In vitro lentiviral transduction. Rcho-1 TS cells were exposed to lentiviral particles, selected with puromycin dihydrochloride (Sigma-Aldrich; $2 \mu g/\mu l$) for 2 to 4 days, and then maintained in a reduced concentration of the antibiotic (1 $\mu g/\mu l$). The puromycin selective pressure was removed during Rcho-1 TS cell differentiation.

In vivo lentiviral transduction. Rat embryos were transduced with lentiviral particles as previously described (52). Briefly, rat embryos were collected on gestation day 4.5. Recovered blastocysts were washed with KSOM medium (Millipore). Zonae pellucidae were removed with Pronase (Sigma-Aldrich; 10 mg/ml for 10 min) and incubated with concentrated lentiviral particles (750 ng of

p24/ml) for 4.5 h. Transduced blastocysts were transferred to uteri of day 3.5 pseudopregnant rats for subsequent evaluation of gene knockdown and placentation site phenotypes on gestation day 11.5.

Measurement of *in vivo* **FOSL1 expression.** Placentation sites derived from embryos treated with control or *Fosl1* shRNAs were processed for the detection of *Fosl1* mRNA by qRT-PCR and FOSL1 protein by Western blotting and immunofluorescence analysis. Trophoblast dissections were performed as previously described (5) and qRT-PCR and Western blotting conducted as described above. FOSL1 expression levels detected on tissue sections with immunofluorescence were measured by calculating integrated density using Photoshop CS5 extended edition (Adobe Systems Inc., San Jose, CA). All values were normalized to control samples. Placentation sites exhibiting a decrease in FOSL1 protein expression was defined as FOSL1 protein expression 1 standard deviation below the mean of control samples.

Measurement of *in vivo* invasion. Invasive trophoblast cells within placentation sites derived from embryos treated with control or *Fosl1* shRNAs were monitored by immunostaining with antibodies to pancytokeratin. An invasion index was calculated by measuring the ratio between the depth of endovascular trophoblast invasion into the decidua and the total depth of the decidua by use of National Institutes of Health ImageJ software (Bethesda, MD). All values were normalized to control samples.

ChIP. Chromatin immunoprecipitation (ChIP) analysis was performed according to a previously published procedure (50). Briefly, Rcho-1 TS cells stably transduced with control or Fosl1 shRNA-expressing lentivirus were grown to confluence in 150-mm dishes and differentiated for 8 days. Cells were then fixed with 1% formaldehyde, and purified nuclear lysates were sonicated on ice to prepare the DNA fragments at a size of approximately 500 bp. Lysates (300 µg of protein) were immunoprecipitated with 10 µg of FOSL1 antibodies (Santa Cruz Biotechnology). Rabbit IgG (BD Biosciences) was used as a nonspecific control. Immunoprecipitated chromatin fragments were eluted from agarose beads. DNA-protein interactions were reverse cross-linked and purified using a QiaQuick PCR purification kit (Qiagen). Purified DNA fragments were assessed by quantitative PCR (qPCR) using Mmp9 promoter-specific primers and SYBR green PCR master mix (Applied Biosystems). Five putative AP-1 response elements (TGAGTCA) were identified within 5 kbp upstream of the Mmp9 translation start site using rVista 2.0 (54) (http://rvista.dcode.org/). Two of the AP-1 response elements, located at -107 bp and -533 bp (upstream of the translation start site), were conserved in rat, mouse, and human and were examined for FOSL1 occupancy. Specific primers were used for qPCR measurements of AP-1 site A (-107 to -101) occupancy (forward, GAGTCAGCGTAAGCCTGGAG; reverse, AGCAGAATTTGCGGAGGTTT; 81-bp product spanning -105 to -26) and AP-1 site B (-533 to -527) occupancy (forward, AAAGAGCCTGC TCCCAGAG; reverse, CATTCCCACCCCTGTTAGT; 66-bp product spanning -567 to -502). Relative occupancy/enrichment was normalized to input samples by use of the $\Delta\Delta CT$ method.

Mmp9 promoter analysis. Rat *Mmp9* 5'-flanking DNA spanning -1107 to +120 (relative to the translation start site NC_005102.2) was PCR amplified using genomic DNA templates isolated from Rcho-1 TS cells (forward primer, GGAGGCTCAATCAGAACAGCTT; reverse primer, GAGGTTGGAGGTTT TCAGGTCT). PCR amplified 5'-flanking regions were initially cloned into the



FIG. 1. AKT expression and activity in trophoblast cells. (A) qRT-PCR analysis of *Akt1*, *Akt2* and *Akt3* in stem (Stem) and differentiated (Dif) Rcho-1 TS cells. Sample size: n = 4. Bars represent means \pm standard error of the mean (SEM). (B) Western blot analysis of total AKT (pan-AKT) and phosphorylated AKT (P-AKT) in stem and differentiated Rcho-1 TS cells. (C) Analysis of AKT phosphorylation state in differentiated Rcho-1 TS cells treated with vehicle, LY, or AI9. P-AKT, Western blot of phosphorylated AKT. pan-AKT, Western blot analysis of total AKT. ACTB was used as a loading control. (D) Analysis of AKT kinase activity in differentiated Rcho-1 TS cells treated with vehicle (0.01% DMSO), LY294002 (LY; 10 μ M), or AKT inhibitor IX (AI9; 10 μ M). P-GSK3 α/β , GSK3 α/β -glutathione *S*-transferase fusion protein was used as a substrate to monitor AKT kinase activity (P-GSK3 α/β , phosphorylated GSK3 α/β). ACTB was used as a loading control for the input Rcho-1 TS cell lysates used for immunoprecipitation of phospho-AKT (Ser 473) and the subsequent kinase assay (ACTB*).

TA cloning vector (Invitrogen) for sequence confirmation and then subcloned (-1024 to +47) into KpnI and BgIII sites (-1024 Mmp9 KpnI, GGAAGAGG GAAGGTACCGAGGCT; +47 Mmp9 BgIII, TGAGGGGCAGCAAAGATCT AGCCTA) of the pGL3 basic luciferase vector (Promega, Madison, WI) for in vitro assays using Rcho-1 TS cells. The two conserved AP-1 response elements, located at -107 bp and -533 bp (upstream of the translation start site) were modified (CAGGACA) by site-directed mutagenesis. Activities of wild-type and mutant promoter-reporter constructs were assessed as previously described (61). Briefly, cells grown in 6-well plates were cotransfected with 500 ng of reporter vector and 50 ng of control Renilla vector by use of Lipofectamine 2000 (Invitrogen). Twelve hours after transfection, the medium was changed, and incubations were continued for an additional 48 h before passive cell lysis and processing with a standard dual luciferase assay (Promega). Firefly luciferase activities were normalized to the cotransfected Renilla luciferase reporter activities. Results were expressed as the relative luciferase activity of the Mmp9 promoterreporter constructs compared to the luciferase activity of the empty (promoterless) luciferase reporter construct.

Statistical analyses. Statistical comparisons of two means were performed with Student's *t* test or Welch's *t* test. Comparisons of multiple groups were evaluated with analysis of variance. The source of variation from significant F-ratios was determined with Tukey's honestly significant difference (HSD) multiple comparison test. Statistical analyses were performed using the R statistical package (http://www.r-project.org/).

RESULTS

Trophoblast AKT expression and disruption with smallmolecule inhibitors. The PI3K/AKT pathway is activated during trophoblast differentiation (43). In the following experiments, we examined roles for the PI3K/AKT pathway in the regulation of trophoblast cell differentiation and the invasive trophoblast phenotype.

Each of the three isoforms of *Akt (Akt1, Akt2, and Akt3)* is expressed in Rcho-1 TS cells (Fig. 1A). AKT is activated upon phosphorylation of serine 473. In Rcho-1 TS cells, phosphorylation of AKT on serine 473 AKT was increased following differentiation (Fig. 1B) (43), indicating that AKT is activated in differentiated Rcho-1 TS cells.

AKT activity was blocked using a small-molecule inhibitor of PI3K (LY) (78) or using an inhibitor that blocks the activity of

all AKT isoforms (AKT inhibitor IX) (42). Treatment of Rcho-1 TS cells with LY decreased both AKT phosphorylation and kinase activity (Fig. 1C and D), while cells treated with AKT inhibitor IX retained AKT phosphorylation but lost AKT kinase activity (Fig. 1C and D). This strategy for using small-molecule inhibitors to disrupt PI3K and/or AKT was used in experiments described below to investigate the role of the PI3K/AKT signaling pathway in the regulation of the invasive trophoblast-vascular remodeling cell phenotype.

PI3K/AKT signaling: regulation of invasion-vascular remodeling genes. Previous gene profiling in trophoblast cells led to the identification of a subset of genes that are upregulated during differentiation and sensitive to PI3K inhibition (46). In this experiment, we focused on a group of 10 PI3Ksensitive genes that are potential regulators of trophoblast invasion and/or vascular remodeling (*Adm* [44, 81], *Cgm4* [82], *Ctsd* [59], *Faslg* [8, 37], *Igf2* [7, 40], *Il17f* [32], *Mmp9* [53, 62], *Prl4a1* [4, 6, 60], *Sema6d* [75], and *Serpine1* [33]) (Table 2). In Fig. 2A, we show that these 10 genes are sensitive to disruption of AKT activation. These observations are consistent with a role for the PI3K/AKT signaling pathway in the regulation of invasive and vascular remodeling trophoblast cell phenotypes.

PI3K/AKT signaling: MMP9 activity and trophoblast invasion. Of the 10 invasion-vascular remodeling genes, 1 gene in particular, *Mmp9*, which encodes matrix metallopeptidase 9 (MMP9), has been associated with invasive behavior of trophoblast (11, 53, 55, 62, 66) and cancer (23, 47) cells. *Mmp9* mRNA expression increases during trophoblast differentiation (Fig. 2B). MMP9 gelatinolytic activity is decreased in conditioned medium from cells treated with PI3K or AKT inhibitors (Fig. 2C). Disruption of PI3K and/or AKT activities impaired trophoblast cell invasion as assessed using Matrigel invasion chamber assays (Fig. 2D and E). These experimental results indicate that PI3K/AKT signaling promotes

Gene name	Abbreviation	Synonym(s)	Functional group	GenBank accession no.
Adrenomedullin	Adm		Hypotensive peptide	NM 012715
Carcinoembryonic antigen gene family 4	Cgm4	Psg16	Secretory protein, unknown function	NM_012525
Cathepsin D	Ctsd	CĎ, CatD	Lysosomal aspartic endopeptidase	NM ⁻ 134334
Fas ligand (TNF superfamily, member 6)	Faslg	Faslg, Tnfsf6	Ligand/membrane anchored	NM_012908
Insulin-like growth factor 2	Igf2	Igf-II	Ligand, growth factor	NM_031511
Interleukin 17F	IĨ17f	ML1	Ligand/cytokine	NM 001015011
Matrix metallopeptidase 9	Mmp9	Gelatinase B	Extracellular matrix remodeling	NM_031055
Prolactin family 4, subfamily a, member 1	Prl4a1	PLP-A	Ligand/cytokine	NM_017036
Sema domain, transmembrane domain (TM), and cytoplasmic domain, (semaphorin) 6D	Sema6d		Receptor	NM_001107768
Serine (or cysteine) peptidase inhibitor, clade E, member 1	Serpine1	Pai1, Planh	Blood coagulation, angiogenesis	NM_012620

TABLE 2. Potential regulators of trophoblast invasion and/or vascular remodeling

the trophoblast cell-invasive phenotype, potentially through modulation of the expression of *Mmp9* and other proinvasive genes.

AKT isoforms: modulation of the trophoblast phenotype. Cellular AKT activity is the net result of the activities of three AKT isoenzymes (AKT1, AKT2, AKT3) (12, 13). To investigate relationships between AKT isoforms and expression of the invasion-vascular remodeling genes, the expression of each AKT isoform was specifically disrupted using lentiviral delivery of isoform-specific shRNAs. Confirmation that isoform-specific shRNAs were effective in inhibiting expression at transcript and protein levels is shown in Fig. 3. Total AKT was affected by AKT1 knockdown but changed minimally following the knockdown of AKT2 or AKT3 (Fig. 3B), supporting earlier reports suggesting that AKT1 is the predominant isoform in the placenta (43, 85). Disruption of each individual AKT isoform yielded an isoform-specific effect on expression of the invasion-vascular remodeling genes. Knockdown of AKT1 resulted in a significant decrease in Cgm4, Faslg, Prl4a1, Sema6d, and Serpine1, while AKT2 disruption inhibited Cgm4, Igf2, Mmp9, Prl4a1, and Sema6d, and AKT3 contributed to the regulation of Cgm4, Igf2, Mmp9, and Sema6d (Fig. 3C). Expression of some invasion-vascular remodeling genes (Adm, Ctsd, and Il17f) was not affected by single AKT isoform disruption (Fig. 3C). Knockdown of individual AKT isoforms was not sufficient to disrupt the invasive abilities of trophoblast cells as assessed in Matrigel invasion chamber assays (Fig. 3D). Stable double-AKT-isoform knockdowns had effects on growth and survival and were problematic to establish. Trophoblast cell phenotypes resulting from the knockdown of individual AKT isoforms provide further support for the involvement of the PI3K/AKT signaling pathway in the regulation of trophoblast invasion and vascular remodeling. However, the results also demonstrate that there is a potentially complex interplay of each isoform in modulating the trophoblast cell phenotype.

FOSL1: expression during trophoblast differentiation and regulation by PI3K/AKT. We next sought to identify potential downstream mediators of the PI3K/AKT signaling pathway responsible for regulating the invasive trophoblast and vascular remodeling phenotype. In cancer cells, FOSL1 is regulated by PI3K/AKT signaling (16, 18, 67). FOSL1 has also been linked to placentation (72) and is expressed in a differentiation-dependent pattern in rat trophoblast cells (46). Expression of FOSL1 was examined in trophoblast cell populations maintained *in vitro*. FOSL1 transcripts and protein increase during Rcho-1 TS cell differentiation (Fig. 4A and B). To determine if the PI3K/AKT signaling pathway can regulate FOSL1, we examined the presence of FOSL1 protein in nuclear lysates from control and PI3K/AKT-disrupted Rcho-1 TS cells. FOSL1 expression was greatly diminished in nuclear lysates of cells treated with the PI3K inhibitor (Fig. 4B). These findings indicate that the PI3K/AKT signaling pathway regulates nuclear accumulation of FOSL1, the site of FOSL1 action.

FOSL1 expression at the placentation site. FOSL1 is expressed in the rat placenta on gestation days 11.5 and 18.5 as assessed by Western blotting. At day 18.5, FOSL1 protein was restricted to the junctional zone of the chorioallantoic placenta (Fig. 4C). FOSL1 was localized to the nuclei of trophoblast giant cells by immunohistochemistry at gestation days 7.5, 11.5, and 18.5. Trophoblast giant cells were identified morphologically and by their expression of PRL3D1 (PL-I) and PRL3B1 (PL-II) (Fig. 4D). Costaining with anti-FOSL1 and anti-cytokeratin (an epithelial cell marker routinely used to identify trophoblast cells) demonstrated that FOSL1 is also expressed in the nuclei of invasive endovascular trophoblast cells on gestation day 11.5 (Fig. 4E). Thus, FOSL1 is a candidate regulator of trophoblast cell invasion and/or uterine spiral remodeling.

In vitro and *in vivo* lentiviral knockdown of FOSL1. In the next series of experiments, we disrupted FOSL1 by using both *in vitro* and *in vivo* model systems for the purpose of investigating the biology of FOSL1 in trophoblast cells. *Fosl1* mRNA and FOSL1 protein were effectively decreased in Rcho-1 TS cells using lentiviral delivery of two independent shRNAs (Fig. 5A and B). *In vivo* FOSL1 knockdown was achieved using a trophoblast cell-specific lentiviral delivery of *Fosl1*-specific shRNAs (52). Gestation day 4.5 rat embryos were exposed to lentiviral particles expressing control shRNA or shRNA specific for *Fosl1* and then transferred to day 3.5 pseudopregnant female rats and allowed to develop until day 11.5. The *in vivo* lentiviral *Fosl1* shRNA delivery system significantly knocked down *Fosl1* mRNA (Fig. 5C) and FOSL1 protein (Fig. 5D and E) expression in gestation day 11.5 trophoblast tissue.

FOSL1: regulation of invasion-vascular remodeling genes. Knockdown of FOSL1 *in vitro* resulted in a decrease in expression of several invasion-vascular remodeling genes, including *Cgm4*, *Mmp9*, *Prl4a1*, and *Sema6d* (Fig. 6). Other



FIG. 2. PI3K and AKT regulate the expression of invasion-vascular remodeling genes and invasive abilities of trophoblast cells. (A) qRT-PCR analysis of invasion-vascular remodeling genes in differentiated Rcho-1 TS cells treated with vehicle (V; 0.01% DMSO), LY294002 (LY; 10μ M), or AKT inhibitor IX (AI9; 10μ M). Experiments were performed with a sample size ranging from 4 to 6. (B) qRT-PCR analysis of *Mmp9* expression during Rcho-1 TS cell differentiated Rcho-1 TS cells treated with V, LY, or AI9. (D) may activity by gelatin zymography in conditioned media from differentiated Rcho-1 TS cells treated with V, LY, or AI9. (D) may activity by gelatin removes a measured using the Matrigel chamber assay. (D) Representative filters showing trophoblast invasion through Matrigel in V, LY, or AI9 treated cultures are presented (cells are marked with arrows). Bar = 100μ m. (E) Graphic presentation of results from the Matrigel invasion chamber assays. Cells were counted from nine replicates and normalized to control samples. Invasion index represents the number of invading cells in each group relative to the number of cells invading in the control group. Bars represent means \pm SEM. Values significantly different from controls are indicated (*, *P* < 0.05).

invasion-vascular remodeling genes were not consistently affected by FOSL1 knockdown (*Ctsd*, *Faslg*, *Igf2*, and *Il17f*), while expression of *Adm* and *Serpine1* was significantly upregulated (Fig. 6).

FOSL1 regulation of *Mmp9***.** FOSL1 has been implicated as a direct regulator of *Mmp9* gene expression in cancer cells (16, 67), which prompted an assessment of FOSL1 interactions

with the *Mmp9* promoter in trophoblast cells. FOSL1 binds to AP-1 elements (28). The rat *Mmp9* promoter possesses two conserved AP-1 elements (site A: -107 to -101; site B: -533 to -527) (Fig. 7A). In differentiated Rcho-1 TS cells, FOSL1 was specifically bound to both conserved AP-1 elements, as determined by ChIP analysis (Fig. 7B). These two AP-1 elements are essential for the activation of the *Mmp9* promoter in



FIG. 3. Role of AKT isoforms in the regulation of the invasive trophoblast phenotype. Differentiated Rcho-1 TS cells expressing control, *Akt1, Akt2*, or *Akt3* shRNAs were evaluated. (A) qRT-PCR analysis for *Akt1, Akt2*, and *Akt3*. Experiments were performed in quadruplicate. (B) Western blot analysis of phosphorylated AKT (P-AKT), total AKT (pan-AKT), AKT1, and AKT2. ACTB was included as a loading control. (C) qRT-PCR analyses of invasion-vascular remodeling genes. All samples were normalized to control samples. Experiments were performed in quadruplicate. (D) Invasion was assessed using Matrigel chamber assays. Cells were counted from eight replicates and normalized to control samples. Bars represent means \pm SEM. Values significantly different from controls are indicated (*, *P* < 0.05).

differentiating Rcho-1 TS cells (Fig. 7C and D). Thus, the data are consistent with FOSL1 directly activating the *Mmp9* gene promoter during trophoblast cell differentiation.

FOSL1 modulates trophoblast cell invasion. The effects of FOSL1 on the regulation of a subset of invasion-vascular remodeling genes prompted an analysis of the role of FOSL1 in the regulation of *in vitro* trophoblast cell invasion. Similar to the modulation of the PI3K/AKT signaling pathway, the loss of FOSL1 decreased the *in vitro* invasive trophoblast ability as assessed using Matrigel chamber invasion assays (Fig. 8A and B). The results indicate that FOSL1 mediates some actions of the PI3K/AKT signaling pathway *in vitro*, including invasive cell behavior.

We next evaluated a role for FOSL1 in regulating *in vivo* trophoblast cell invasion. Histological analysis demonstrated that the trophoblast invasion index (depth of invasion/height of decidua) was significantly less at placentation sites with FOSL1 knockdown (Fig. 8C to F), implicating FOSL1 as a component of the *in vivo* regulatory pathway controlling trophoblast cell invasion.

Collectively, our findings indicate that a regulatory pathway involving PI3K/AKT and FOSL1 controls the invasive trophoblast phenotype.

DISCUSSION

Intrauterine trophoblast invasion and remodeling of the uterine vasculature are important features of hemochorial placentation. Blood vessels are restructured, permitting increased blood flow and nutrient delivery to the placenta and ultimately the fetus. In this study, we have identified a cellular signaling pathway involving PI3K/AKT and FOSL1 that regulates invasive trophoblast and vascular remodeling phenotypes. Small-molecule inhibitors were effective in interfering with the PI3K/AKT signaling pathway. This strategy of nullifying the actions of PI3K or AKT in trophoblast cells led to the inhibition of proinvasion and vascular remodeling genes. Blockade of the PI3K/AKT pathway also decreased nuclear accumulation of the FOSL1 protein. Disruptions of PI3K, AKT, or FOSL1 interfered with trophoblast cell invasion.

PI3K/AKT signaling activates the expression of a set of 10 genes encoding proteins implicated in cell invasion and/or vascular remodeling. The invasion-vascular remodeling set of genes can be placed into two groups, genes encoding proteins that potentially affect trophoblast movement and genes encoding proteins that are postulated to alter the vasculature. In the first group, the genes encode proteins that can induce cell movement (IGF2 [7, 40]), guide cells (SEMA6D [75]), and



FIG. 4. Expression of FOSL1 in Rcho-1 TS cells and rat placentation sites. (A) qRT-PCR analysis of *Fosl1* expression during Rcho-1 TS cell differentiation. Experiments were performed in triplicate. (B) Western blot analysis of FOSL1 protein in nuclear lysates of stem (Stem) or differentiating Rcho-1 TS cells treated with vehicle (Dif+V; 0.01% DMSO) or LY294002 (Dif+LY; 10 μ M). TBP was included as a loading control. (C) Western blot analysis of protein levels for FOSL1 in nuclear lysates from gestation days 11.5 and 18.5 placental tissues. Placental tissues isolated from gestation day 18.5 were dissected into junctional zone (JZ) and labyrinth zone (LZ) compartments. TBP was included as a loading control. (D) Immunohistochemical localization of FOSL1 in gestation days 7.5, 11.5, and 18.5 rat placentation sites. FOSL1 is localized to the nucleus of trophoblast giant cells (arrowheads). PRL3D1 (PL-I, day 7.5) and PRL3B1 (PL-II, days 11.5 and 18.5) are expressed in trophoblast giant cells (arrowheads) and were used as positive controls. Top bars = 1 mm; middle and bottom bars = 100 μ m. (E) Immunofluorescence colocalization of FOSL1 and DAPI; (right) cytokeratin and FOSL1. Please note that FOSL1 colocalizes with cytokeratin in invasive endovascular trophoblast cells. Bar = 100 μ m.

remodel the extracellular matrix (CTSD [59], MMP9 [53, 62], and SERPINE1 [33]). The second group consists of genes encoding proteins that directly affect endothelial cells and blood vessels (ADM [44, 81], CGM4 [82], FASLG [8, 37], and

IL17F [32]) or indirectly modulate blood vessel remodeling by altering immune cell cytokine production (CGM4 [34, 79, 80], IL17F [32], and PRL4A1 [4, 6, 60]). Although each of the 10 proteins encoded by these genes has the potential to influence



FIG. 5. Analysis of shRNA-mediated FOSL1 knockdown. (A and B) *In vitro* analyses. qRT-PCR measurements of *Fosl1* mRNA (A) and Western blot analysis of FOSL1 protein (B) in differentiated Rcho-1 TS cells expressing control or *Fosl1* shRNAs. The qRT-PCR measurements were performed in quadruplicate. (C to E) *In vivo* analyses. (C) Western blot analysis of FOSL1 protein in dissected placental samples exposed to control or *Fosl1* shRNAs. ACTB was included as a loading control. (D) qRT-PCR analysis of *Fosl1* mRNA in dissected placental samples exposed to control or *Fosl1* shRNAs. Dashed lines mark the means and the solid lines indicate the SEMs. Sample sizes: control, n = 14; *Fosl1* shRNA, n = 12. *Fosl1* shRNA treatment significantly decreased *Fosl1* mRNA expression (*, P < 0.05). The open circle represents a *Fosl1* shRNA sample that escaped knockdown and was not included in the statistical analysis. (E) FOSL1 immunohistochemistry of rat placentation sites exposed to control or *Fosl1*-shRNAs.



FIG. 6. Role of FOSL1 in the regulation of invasion-vascular remodeling gene expression. qRT-PCR analysis of invasion-vascular remodeling genes in differentiated Rcho-1 TS cells expressing control or *Fosl1* shRNAs. All samples were normalized to control samples. Experiments were performed in quadruplicate. Bars represent means \pm SEM. Values significantly different from controls are indicated (*, *P* < 0.05).

trophoblast invasion and/or vascular remodeling, the relative roles of each regulator and their specific orchestration during gestation are unknown.

Three AKT isoforms are expressed in trophoblast cells and regulate trophoblast cell biology in a unique and complicated overlapping manner. Knockdown of individual AKT isoforms affected subsets of PI3K/AKT-sensitive invasion-vascular remodeling genes and led to a unique expression profile. Expression of some invasion-vascular remodeling genes (*Adm, Ctsd,* and *Il17f*), nuclear accumulation of FOSL1 protein, and trophoblast cell invasion were not affected by interfering with the expression of a single AKT isoform. The retention of these functions in single-AKT-knockdown cells was associated with the presence of active cellular AKT, suggesting a potentially complex interplay of each AKT isoform.

Even though knockdown of individual AKT isoforms was not sufficient to decrease trophoblast cell invasion and FOSL1 nuclear protein accumulation, we gained some insight into the role of individual AKT isoforms in regulating expression of the invasion-vascular remodeling genes. Two of the 10 invasionvascular remodeling-related genes (Cgm4 and Sema6d) are sensitive to disruptions in any one of the three AKT isoforms, indicating that a high level of AKT activity is needed for regulating their expression. Other invasion-vascular remodeling genes were sensitive to disruption of a single isoform. Faslg and Serpine1 were sensitive only to interference of AKT1, implying that there must be some AKT1 substrate specificity associated with their regulation. Prl4a1 was sensitive to either AKT1 or AKT2 knockdown, and Mmp9 and Igf2 were sensitive to either AKT2 or AKT3 knockdown. These findings indicate a mixture of substrate specificity and promiscuity. Adm, Ctsd, and Il17f were exquisitely sensitive to small-molecule inhibition of PI3K or AKT but were not affected by knockdown of any single AKT isoform. Following the above logic, we would propose that the

activation of this subset of genes requires lower levels of AKT activity and/or exhibits complete AKT substrate promiscuity. The same conclusion could be made for the regulation of FOSL1 protein nuclear accumulation and trophoblast invasion. The balance of each individual AKT isoform may also be critical for the regulation of the trophoblast cell phenotype, as has been shown in breast cancer cells (39). Disruption of multiple AKT isoforms in the trophoblast cells may provide additional insights; however, based on initial experiments, this approach is challenging due to effects on cell survival. Finally, the subcellular location of AKT isoform in trophoblast cells (70). Intracellular AKT isoform location impacts access to substrates and the activation of specific regulatory pathways.

PI3K/AKT signaling regulates FOSL1 in trophoblast cells. Similar findings have been reported for other cell types (16, 18, 67). FOSL1 has also been shown to be a downstream mediator of the PI3K/AKT signaling pathway in regulating the expression of proinvasion genes, including Mmp9, and cellular invasion (16, 18, 67). The precise mechanism of PI3K/AKT regulation of FOSL1 has not been clarified, but here we show that loss of PI3K/AKT signaling results in decreased nuclear accumulation of FOSL1 protein. FOSL1 may be a substrate for AKT, and its phosphorylation may influence its nuclear accumulation and/or activity. Alternatively, the role of the PI3K/ AKT signaling pathway in regulating FOSL1 may be indirect. Cross talk between PI3K/AKT and other cellular signaling pathways in trophoblast has been described previously (65, 66, 74) and may contribute to the stabilization of FOSL1 protein in the nucleus (10). FOSL1 undergoes ubiquitin-independent degradation by the proteasome, which can be prevented by ERK1/2 phosphorylation within the C-terminal stabilizing region of the FOSL1 protein (10). Whether AKT can act simi-



FIG. 7. FOSL1 occupancy on AP-1 elements within the *Mmp9* promoter. (A) Schematic representation of putative AP-1 elements (black and white boxes) within *Mmp9* promoters of the rat (*rMmp9*), mouse (*mMmp9*), and human (*hMmp9*). Sites A and B (white boxes) were conserved in all three species. Putative AP-1 elements demarcated by black boxes were not conserved. TSS, translation start site. (B) ChIP determination of the occupancy of FOSL1 on sites A and B of the rat *Mmp9* promoter in Rcho-1 TS cells expressing control or *Fosl1* shRNAs. IgG was included as a negative control. Values significantly different from control shRNA are indicated (*, P < 0.05). (C and D) Activities of an empty luciferase reporter construct (control), a wild-type rat *Mmp9* promoter (-1024 to +47)-luciferase reporter construct (wt), or a mutant rat *Mmp9* promoter (-1024 to +47)-luciferase reporter was active and dependent upon AP-1 sites A and B (*, P < 0.05). All experiments were performed in triplicate. Bars represent the mean \pm SEM.

larly or whether it activates another kinase (such as extracellular signal-regulated kinase 1/2 [ERK1/2]) to stabilize FOSL1 remains to be determined.

FOSL1 has been implicated as a key regulator of placentation. A null mutation in the Fosl1 gene disrupts morphogenesis of the mouse placenta and results in prenatal lethality (72). Defective vascularization of the labyrinth zone of the mouse chorioallantoic placenta was viewed as the insult responsible for the prenatal demise of the fetus. In our report, invasive trophoblast cell populations were shown to express FOSL1, including those possessing an endovascular location, and FOSL1 was demonstrated to regulate the expression of a subset of trophoblast invasion-vascular remodeling genes (Cgm4, Mmp9, Prl4a1, and Sema6d). FOSL1-responsive genes include genes sensitive to knockdown of any AKT isoform (Cgm4 and Sema6d), a gene regulated by AKT1 and AKT2 (Prl4a1), and a gene regulated by AKT2 and AKT3 (Mmp9). Both in vitro and in vivo experiments implicated FOSL1 as a regulator of the invasive trophoblast phenotype. These FOSL1 actions are consistent with its prominent role in regulating cancer cell invasion (51, 58, 77, 86). FOSL1 occupies critical AP-1 elements within the Mmp9 promoter essential for gene regulation in trophoblast cells. *In vivo* trophoblast-specific knockdown of FOSL1 led to significantly decreased intrauterine endovascular trophoblast invasion. Disruptions in labyrinthine vascularization may have also been evident in the FOSL1 hypomorphs but were not examined in our report. Trophoblast invasion is less prominent in the mouse than in the rat (2, 3, 21). In the *Fosl1* null mouse, trophoblast invasion was probably of secondary significance to interruptions in nutrient and waste exchange (72).

Regulation of the invasive trophoblast lineage is complex, involving a network of signaling pathways and transcriptional regulators (48). Mitogen-activated protein kinases, PI3K/AKT, focal adhesion kinase, Rho proteins, and Wnt signaling pathways have all been implicated in regulating various aspects of trophoblast cell motility and invasion (1, 17, 38, 56, 65, 66, 73, 74). These signaling molecules potentially impinge on a set of transcription factors to direct the development and function of the invasive trophoblast lineage. Thus far, inhibitor of differentiation 2, signal transducer and activator of transcription 3, Ikaros, and peroxisome proliferator-activated receptor-gamma have been shown to possess putative modulatory roles on the invasive trophoblast phenotype (31, 41, 64, 83). In this report,



FIG. 8. FOSL1 regulates trophoblast invasion as assessed *in vitro* and *in vivo*. (A and B) The invasive abilities of Rcho-1 TS cells expressing control or *Fosl1* shRNAs were measured using Matrigel chamber assays. (A) Representative filters showing trophoblast invasion through Matrigel in control and FOSL1 knockdown cultures (cells are marked with arrows). Bar = 100 μ m. (B) Graphic presentation of results from the Matrigel invasion chamber assays. Cells were counted from nine replicates and normalized to control samples. (C to F) Evaluation of the impact of FOSL1 knockdown on trophoblast invasion *in vivo*. Control or *Fosl1* shRNAs were delivered to the trophectoderm of gestation day 4.5 blastocysts by use of a lentiviral delivery system and transferred to pseudopregnant rats, and placentation sites were evaluated on gestation day 11.5. (C to E) Immunohistochemistry of rat placentation sites was performed using anti-cytokeratin to identify trophoblast. Representative sections of gestation day 11.5 placentation of the invasion index (see Materials and Methods). Sample sizes: control, n = 12; *Fosl1* shRNA, n = 9. Bars represent means \pm SEM. Values significantly different from controls are indicated (*, P < 0.05).

we have generated experimental evidence indicating that the PI3K/AKT/FOSL1 signaling pathway is involved in regulating invasive trophoblast and vascular remodeling phenotypes. FOSL1 is shown to be integral to trophoblast-directed uterine spiral remodeling and establishment of the maternal-fetal interface.

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