

Conditional deletion of *Tsc1* in the female reproductive tract impedes normal oviductal and uterine function by enhancing mTORC1 signaling in mice

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ABSTRACT: Heightened mammalian target of rapamycin complex I (mTORC1) activity by genetic deletion of its direct inhibitor, *Tsc1*, is associated with aberrant development and dysfunction of the female reproductive tract in mice. Here, we compared the phenotypes of mice with conditional deletion of *Tsc1* in the female reproductive tract by either *progesterone receptor (PR)-Cre* (*Tsc1*^{PR(d/d)}), which inactivates *Tsc1* in all major cell types in the uterus (epithelium, stroma and myometrium), or *anti-Mullerian hormone type 2 receptor (Amhr2)-Cre* (*Tsc1*^{Amhr2(d/d)}), which inactivates stromal and myometrial *Tsc1*. *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)} females are infertile resulting from oviductal hyperplasia, retention of embryos in the oviduct and implantation failure. In contrast to the appropriate embryonic development after fertilization seen in *Tsc1*^{Amhr2(d/d)} females, embryo development was disrupted in *Tsc1*^{PR(d/d)} females. In addition, uteri in *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)} females showed epithelial hyperplasia but not endometrial cancer. In conclusion, *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)} have overlapping yet distinct phenotypes in the context of compartment-specific deletion of *Tsc1*.

Key words: *Tsc1* / uterus / ovary / oviduct / mouse

Introduction

There is evidence that heightened mammalian target of rapamycin complex I (mTORC1) activity induces dysfunction of the female reproductive tract (Fan *et al.*, 2008; Lu *et al.*, 2008; Reddy *et al.*, 2008; Adhikari *et al.*, 2009, 2010; Contreras *et al.*, 2010; Memarzadeh *et al.*, 2010; Hirota *et al.*, 2011; Tanaka *et al.*, 2012; Tanwar *et al.*, 2012). mTOR is a serine/threonine kinase that plays roles in cell proliferation and survival via multiple signaling pathways, and primarily contributes to two complexes which drive separate signaling pathways: mTORC1 with Raptor, mLST8 and Pras40, and mTORC2 with mSIN1, Rictor, mLST8 and Protor1 (Guertin and Sabatini, 2007; Zoncu *et al.*, 2011; Johnson *et al.*, 2013). The activation of mTORC1 by pAkt is suppressed by tuberous sclerosis complex I (*Tsc1*) and *Tsc2* by inhibiting ras homolog enriched in brain (Rheb), which is a direct activator of mTORC1 (Kwiatkowski and Manning, 2005; Zoncu *et al.*, 2011). mTORC1 activation leads to

phosphorylation of downstream targets, such as p70 S6 kinase leading to increased protein synthesis (Johnson *et al.*, 2013).

Tsc1 or *Tsc2* mutations in humans cause tuberous sclerosis, a rare, autosomal-dominant multisystem tumor syndrome embodied in the development of benign tumors in a variety of organs, including the brain, heart and kidneys (Cheadle *et al.*, 2000; Crino *et al.*, 2006; Curatolo *et al.*, 2008). While mice with homozygous deletion of *Tsc1* show embryonic lethality, mice missing one allele of *Tsc1* (*Tsc1*^{+/-}) develop renal cell carcinomas and benign tumors in several organs by 10 months of age. However, only ~7% of *Tsc1*^{+/-} mice show leiomyoma by 18 months in the uterus with no evidence of endometrial cancer (Kobayashi *et al.*, 2001). Considering these observations, mice with conditional deletion of *Tsc1* were utilized to examine its function in reproductive organs.

It was previously reported that oocyte-specific *Tsc1* deletion in mice prematurely activates the entire pool of primordial follicles enhancing follicular depletion in early adult life due to heightened mTORC1 signaling (Adhikari *et al.*, 2010). Furthermore, when *Tsc1* was deleted in

ovarian granulosa cells (GCs) using *anti-Mullerian hormone type 2 receptor (Amhr2)-Cre* ($Tsc1^{loxP/loxP}/Amhr2^{cre/+} = Tsc1^{Amhr2(d/d)}$), significantly fewer primordial follicles were found in deleted mice, suggesting premature ovarian insufficiency (Tanaka et al., 2012). After normal and superovulation, these mice also show compromised oocyte quality. *Amhr2-Cre* deletes *Tsc1* in oviductal stroma and myometrium, inducing oviductal epithelial hyperplasia and blocking oviductal transport of fertilized embryos into the uterus (Tanaka et al., 2012). Collectively, these results indicate that *Tsc1* has critical roles in preserving the ovarian reserve of oocytes in addition to maintaining the normal architecture of the oviduct for appropriate transport of preimplantation embryos to the uterus.

mTORC1 activity is also associated with endometrial cancer as a downstream target of *Pten* (phosphatase and tensin homolog) (Guertin and Sabatini, 2007; Lu et al., 2008; Zoncu et al., 2011). *Pten* is a tumor suppressor, inactivation of which increases phosphoinositide 3-kinase activity with enhanced Akt activation (Luo et al., 2003). Alteration of *Pten* is frequently observed in endometrial cancer, particularly in the epithelial compartment (Di Cristofano and Ellenson, 2007; Saal et al., 2007). Increased levels of activated Akt (phosphorylated Akt; pAkt) stimulate mTORC1 activity. In humans, however, mTORC1 activation is more frequently observed in advanced and high-grade endometrial cancer (Darb-Esfahani et al., 2009). This suggests that heightened mTORC1 signaling mainly affects endometrial cancer progression but not initiation.

Conditional deletion of genes in different uterine or oviductal tissue compartments may cause differential phenotypes. For example, conditional deletion of *Pten* in all major uterine compartments (epithelium, stroma and myometrium) using *PR-Cre* results in the development of endometrial cancer (Daikoku et al., 2008), whereas the deletion of *Pten* in the uterine stroma and myometrium by *Amhr2-Cre* fails to generate endometrial cancer even at the age of 5 months, but rather transforms myometrial cells to adipocytes (Daikoku et al., 2011). Therefore, we generated conditional deletion of *Tsc1* in the female reproductive tract with *PR-Cre* ($Tsc1^{loxP/loxP}/PR^{cre+/-} = Tsc1^{PR(d/d)}$) which deletes genes in the corpus luteum and oviductal epithelium, in addition to all major uterine tissue compartments. We found that $Tsc1^{PR(d/d)}$ females are totally infertile due to oviductal hyperplasia and retention of embryos similar to the phenotype found in $Tsc1^{Amhr2(d/d)}$ females. However, unlike $Tsc1^{Amhr2(d/d)}$ females, embryo development after fertilization was disrupted in $Tsc1^{PR(d/d)}$ females, but this was not due to poor oocyte quality. We also found that $Tsc1^{PR(d/d)}$ uteri showed epithelial hyperplasia but not cancer even after 300 days of age, but rather exhibited transformation of the uterine stroma and myometrium to benign tumors with heightened mTORC1 signaling in the epithelium, stroma and myometrium. In addition, many of them exhibited low locomotion, hunched-back posturing and vaginal bleeding by this age. These results provide evidence that heightened mTORC1 signaling alone by conditional deletion of *Tsc1* is insufficient to initiate endometrial cancer, but allows formation of benign tumors in the uterus, and a similar phenotype was noted when *Tsc1* was deleted in the stroma and myometrium by *Amhr2-Cre*.

Materials and Methods

Mice. $Tsc1^{loxP/loxP}$ mice (stock number 005680) were obtained from the Jackson Laboratory. *PR-Cre* mice were obtained from Francesco DeMayo

and John Lydon (Baylor College of Medicine), and *Amhr2-Cre* mice were obtained from Richard Behringer (MD Anderson Cancer Center). $Tsc1^{loxP/loxP}/PR^{cre/+}$ ($Tsc1^{PR(d/d)}$) mice and $Tsc1^{loxP/loxP}/Amhr2^{cre/+}$ ($Tsc1^{Amhr2(d/d)}$) were created by breeding $Tsc1^{loxP/loxP}$ mice with *PR-Cre* mice and *Amhr2-Cre* mice, respectively. For experiments, littermate floxed and gene-deleted mice were utilized within the same set of experiments to minimize the influence of genetic background variability and to ensure the validity of our results. All mice used in this investigation were housed in barrier facilities in the Cincinnati Children's Hospital Medical Center's Animal Care Facility according to National Institutes of Health and institutional guidelines for the use of laboratory animals. All protocols for the present study were reviewed and approved by the Cincinnati Children's Research Foundation's Institutional Animal Care and Use Committee. Mice were provided with autoclaved rodent Lab Diet 5010 (Purina) and UV light-sterilized RO/DI constant circulation water *ad libitum*.

LacZ staining

LacZ staining was performed as previously described (Daikoku et al., 2011). In brief, tissues were embedded in OCT after fixation in 0.2% paraformaldehyde and infusion in 30% sucrose at 4°C. Frozen sections were stained with 5-bromo-4-chloro-3-indolyl-β-D-galactopyranoside overnight at 37°C. The sections were counterstained with eosin.

Immunohistochemistry

Immunohistochemistry was performed as previously described (Daikoku et al., 2006). Tissues were fixed in PROTOCOL Safefix II (Thermo Fisher) and embedded in paraffin. Uterine sections (6 μm) were subjected to immunostaining using antibodies to phosphorylated Rps6 (pS6, Cell signaling), cytokeratin 8 (CK8; Developmental Studies Hybridoma Bank), α-smooth muscle actin (αSMA; Abcam) or Ki67 (Thermo Fisher). After deparaffinization and hydration, the sections were subjected to antigen retrieval by autoclaving in 10 mM sodium citrate solution (pH = 6) for 10 min. A diaminobenzidine kit (Invitrogen) was used to visualize antigens. The sections were counterstained with hematoxylin or eosin.

Western blot analysis

Tissue samples were prepared as previously described (Daikoku et al., 2006). After measuring protein concentrations, supernatants were boiled with SDS-PAGE sample buffer for 5 min. Samples were run on 10% SDS-PAGE gels and transferred onto PVDF membranes. Membranes were blocked with 10% milk or 5% bovine serum albumin in TBST and probed with antibodies to pS6 (Cell Signaling), Rps6 (S6, Cell Signaling), or actin (Santa Cruz) overnight at 4°C. After washing, blots were incubated in peroxidase-conjugated donkey anti-goat or donkey anti-rabbit IgG (Jackson Immuno Research Laboratories, Inc.). All signals were detected using chemiluminescent reagents (GE Healthcare). S6 and actin served as loading controls.

RNA extraction and RT-PCR

$Tsc1^{PR(f/f)}$ ($Tsc1^{loxP/loxP}/PR^{+/-}$) or $Tsc1^{PR(d/d)}$ uterine and oviductal RNA samples were prepared and analyzed as previously described (Daikoku et al., 2006). In brief, total RNA was extracted with Trizol (Invitrogen, Carlsbad, CA, USA) according to the manufacturer's protocol. After DNase treatment (Ambion, Austin, TX), 1 μg of total RNA was reverse transcribed with Superscript II (Invitrogen). A PCR was performed using primers 5'-GGAGTGAGCTTTGGAAGTGG-3' and 5'-CTTCTGAGA GACCTGGCTGAG-3' for *Tsc1*; 5'-CATTTGGGCCAGCTAAACAT-3' and 5'-CCCGGCAAAACAGGTAGTTA-3' for *Cre*; 5'-GTGGGCCGCC CTAGGCACCAG-3' and 5'-CTCTTTGATGTCACGCACGATTTC-3' for *Act β*.

In situ hybridization

Paraformaldehyde-fixed frozen sections were hybridized with ^{35}S -labeled cRNA probes as previously described (Daikoku *et al.*, 2006).

Analysis of ovulation, fertilization, embryo development, implantation and term pregnancy

Six to eight weeks old littermate $Tsc1^{PR(f/f)}$, $Tsc1^{PR(d/d)}$, $Tsc1^{Amhr2(f/f)}$ ($Tsc1^{loxP/loxP}/Amhr2^{+/+}$) or $Tsc1^{Amhr2(d/d)}$ females were mated with wild-type fertile males to induce pregnant mice. The day of vaginal plug formation was considered Day 1 of pregnancy. To examine fertilization, mice were sacrificed on Day 2 of pregnancy and oviducts were flushed with saline to recover embryos. To examine implantation, pregnant dams were assessed on Day 5 morning. Implantation sites were visualized by intravenous injection of a Chicago blue dye solution, and the number of implantation sites — demarcated by distinct blue bands—was recorded. For litter size analysis, pregnant females were monitored from Day 17 through Day 21 by observing mice daily for parturition timing and litter size.

In vitro fertilization

In vitro fertilization (IVF) was performed as previously described (Sun *et al.*, 2009). Wild-type or mutant females were superovulated by intraperitoneal injections of 5 IU of eCG (Sigma), followed by injections of 5 IU of hCG (Sigma) 48 h later. Cumulus–oocyte complexes from each female were collected from the oviduct ampulla 13–14 h after hCG injection and placed in 50- μl droplets of human tubal fluid (HTF) medium

(Chemicon). Sperms were collected from the cauda of the epididymis from mature WT male mice and placed into 0.2 ml of HTF medium for counting. After 2 h incubation in a humidified 5% CO_2 incubator at 37°C to allow capacitation, sperms ($\sim 1.2\text{--}1.5 \times 10^6$ sperm/ml) were then co-incubated with eggs to allow fertilization. After 6 h, eggs were washed and placed in 20- μl droplets of KSOM medium (Chemicon) for incubation. The cleavage rate (two-cell stage) after 24 h was used as an index of fertilization. Formation of blastocysts at 120 h indicated developmental potential of fertilized embryos.

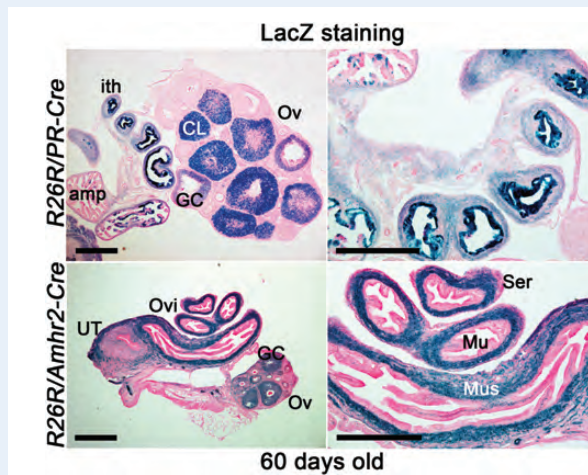


Figure 1 Localization of recombination in *PR-Cre* and *Amhr2-Cre* oviducts and ovaries. The conditional gene recombination by Cre recombinase induced by the *PR* promoter (upper panel) or *Amhr2* promoter (lower panel) was visualized by LacZ staining. *R26R/PR-Cre* females show recombination primarily in corpus luteum (CL), GCs and the isthmus of the oviduct (ith), while *R26R/Amhr2-Cre* females show recombination in the oviductal muscularis layer (Mus) and ovarian follicular GC. Tissues were harvested at the age of 60 days. Left panels are of lower magnification: left panel bars, 400 μm . Right panels are of higher magnification: bars, 100 μm . Ovi, oviduct; Ov, ovary; amp, oviductal ampulla; Mu, mucosa; Ser, serosa; UT, uterus.

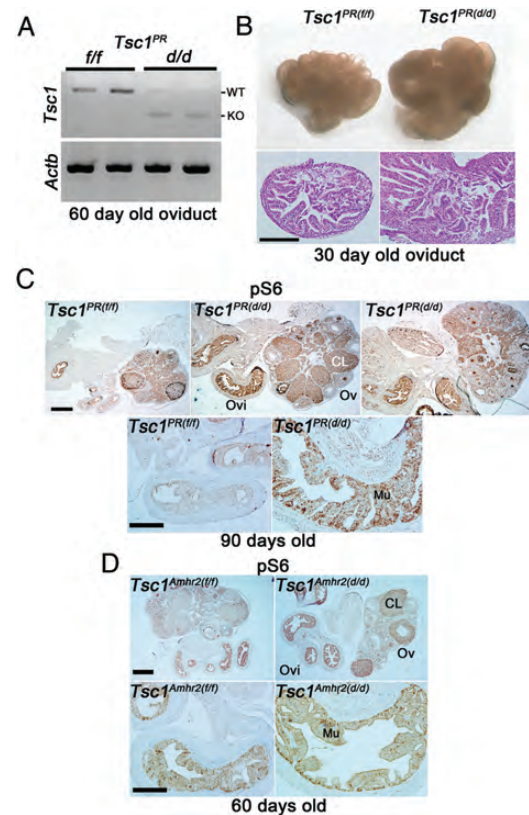


Figure 2 $Tsc1^{PR(d/d)}$ oviducts exhibit higher epithelial hyperplasia and increased mTORC1 signaling. (A) RT-PCR showing inactivation of *Tsc1* in $Tsc1^{PR(d/d)}$ oviducts compared with $Tsc1^{PR(f/f)}$ oviducts from mice at 60 days of age. *Actb* (encoding β -actin) served as a loading control. (B) Morphology and histology of oviduct and ovary of $Tsc1^{PR(f/f)}$ and $Tsc1^{PR(d/d)}$ from mice 6–8 weeks of age showing an increased oviductal size and epithelial hyperplasia in $Tsc1^{PR(d/d)}$ oviducts. Seven $Tsc1^{PR(f/f)}$ and nine $Tsc1^{PR(d/d)}$ females were assessed for oviductal and ovarian morphology, while one $Tsc1^{PR(f/f)}$ and three $Tsc1^{PR(d/d)}$ oviducts were assessed for histology. Bar, 100 μm . (C and D) Immunostaining for pS6 in oviducts and ovaries of $Tsc1^{PR(f/f)}$ ($n = 3$), $Tsc1^{PR(d/d)}$ ($n = 3$), $Tsc1^{Amhr2(f/f)}$ ($n = 4$), or $Tsc1^{Amhr2(d/d)}$ ($n = 4$) females indicated increased mTORC1 signaling primarily in $Tsc1^{PR(d/d)}$ oviductal mucosa (Mu). Ovaries and oviducts from $Tsc1^{Amhr2(f/f)}$ or $Tsc1^{Amhr2(d/d)}$ females were collected at 60 days ($n = 2$) or 90 days of age ($n = 2$). The intensity of staining was similar between these two groups. Top panels show lower magnification: bars, 400 μm ; bottom panels show higher magnification: bars, 100 μm . WT; wild-type, KO; knockout, Ovi, oviduct; Ov, ovary; CL, corpus luteum.

Measurement of serum estradiol and progesterone levels

Mouse blood samples were collected on Day 4 of pregnancy, and serum levels of estradiol-17 β (E₂) and progesterone (P₄) were measured using EIA kits (Cayman).

Statistical analysis

Student's *t*-test was performed to determine statistical significance between groups. *P* < 0.05 was considered significant.

Results

Differential recombination of *loxP* sites in the female reproductive organs by *PR-Cre* and *Amhr2-Cre*

Our major objective was to examine whether heightened mTORC1 by deletion of *Tsc1* by *PR-Cre* (*Tsc1*^{PR(d/d)}) shows differential phenotype compared with *Tsc1*-deleted mice by *Amhr2-Cre* (*Tsc1*^{Amhr2(d/d)}). As previously shown (Daikoku et al., 2011), both *PR-Cre* and *Amhr2-Cre* recombined *loxP* sites in female reproductive organs, however the recombination by each *Cre* is different depending on the cell type (Fig. 1). Using a *Rosa26-LacZ* reporter mouse line, we have shown that *PR-Cre* recombines *loxP* sites in all major uterine tissue compartments (epithelium, stroma and myometrium) of the uterus, while *Amhr2-Cre* did so only in the stroma and myometrium. In the ovary,

PR-Cre recombined *loxP* sites primarily in the corpora lutea (CL), while *Amhr2-Cre* showed recombination in follicular GCs and CL of the ovary. *PR-Cre* mainly recombined the *loxP* sites in the epithelium of the oviductal isthmus with some recombination also occurring in the stroma and muscle layer; signals were undetectable in the ampullary epithelium. On the other hand, *Amhr2-Cre* primarily recombined *loxP* sites in the stroma and muscle layer, although modest expression of *LacZ* was noted in epithelia of both the ampulla and isthmus. Taken together, we speculated that a set of distinct and overlapping phenotypes will be observed in *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)} females.

Pregnancy fails in *Tsc1*^{PR(d/d)} females due to oviductal retention of embryos

To evaluate fertility, we first examined pregnancy outcome in *Tsc1*^{PR(d/d)} females. Deletion of *Tsc1* in *Tsc1*^{PR(d/d)} oviducts was first confirmed by RT-PCR (Fig. 2A). These dams did not produce any pups (*n* = 6) compared with *Tsc1*^{PR(ff)} littermates (7.3 ± 0.8 pups/litter, *n* = 14). We then examined the status of implantation on Day 5 of pregnancy in 6- to 8-week-old *Tsc1*^{PR(ff)} and *Tsc1*^{PR(d/d)} littermate females. *Tsc1*^{PR(d/d)} uteri did not show any implantation sites as examined by the blue dye method and no unfertilized eggs or embryos were recovered from the uterus. Despite the recovery of comparable numbers of fertilized eggs between control and experimental groups on Day 2, morulae and blastocysts along with several degenerating eggs were recovered from the oviducts on Day 5 of pregnancy (Tables I and II). These results indicate that pregnancy failure was due to retention

Table I Normal fertilization in *Tsc1*-deleted mice. Day 2.

Genotype	No. of mice examined	Mice with fertilized eggs (%)	Total No. of recovered eggs	No. of degenerated eggs ^a	Total No. of fertilized eggs (% ^b)
<i>Tsc1</i> ^{PR(ff)}	7	7 (100%)	60	13	31 (65 ± 16)
<i>Tsc1</i> ^{PR(d/d)}	9	6 (67%)	176	86	51 (49 ± 14)
<i>Tsc1</i> ^{Amhr2(ff)}	8	7 (87.5%)	73	0	60 (95 ± 3)
<i>Tsc1</i> ^{Amhr2(d/d)}	3	3 (100%)	59	27	28 (82 ± 12)

^aDegenerated eggs from the last cycle.

^b% is shown by mean ± SEM and is not statistically different between the control and experimental groups (*P* > 0.05).

Table II Retention of eggs in the oviduct in *Tsc1*-deleted mice. Day 5.

Genotype	No. of mice examined	No. of mice with IS	No. of IS (mean ± SEM)	% of mice with oviductal eggs		Total No. of eggs from the oviduct	
				Eggs	Embryo	Embryo	Degenerated eggs
<i>Tsc1</i> ^{PR(ff)}	6	6 (100%)	8.0 ± 1.5	17% ^a	17%	1	6
<i>Tsc1</i> ^{PR(d/d)}	7	0 (0%)	0	100%	85.7%	26 ^b	71
<i>Tsc1</i> ^{Amhr2(ff)}	3	3 (100%)	9.0 ± 0.6	0%	0%	0	0
<i>Tsc1</i> ^{Amhr2(d/d)}	4	0 (0%)	0	100%	100%	9 ^c	28

IS, implantation sites.

^aOne mouse had one IS and seven eggs (six degenerated and one blastocyst) retained in the oviduct.

^bThese include 7 blastocysts, 16 morula, one 8-cell and two 2-cell.

^cThese include eight blastocysts and one morula.

Table III Differential development of IVF-derived embryos from *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)}.

Genotype	No. of donor mice	No. of eggs used for IVF	IVF rate No. of two-cell embryo (%)	Development No. of blastocysts/total No. of two-cell embryos (%)
<i>Tsc1</i> ^{PR(f/f)}	7	194	137 (70.6)	47/137 (34.3)*
<i>Tsc1</i> ^{PR(d/d)}	5	159	138 (86.8)	69/138 (50.0)* **
<i>Tsc1</i> ^{Amhr2(f/f)}	7	108	73 (67.6)	39/73 (53.4)**
<i>Tsc1</i> ^{Amhr2(d/d)}	7	45	35 (77.8)	2/35 (5.7)** **

*not statistically different ($P > 0.05$).** $P < 0.02$.*** $P < 0.02$.

of embryos in *Tsc1*^{PR(d/d)} oviducts similar to the phenotype seen in *Tsc1*^{Amhr2(d/d)} females (Tanaka et al., 2012).

We next compared the histology of oviducts between *Tsc1*^{PR(f/f)} and *Tsc1*^{PR(d/d)} females. *Tsc1*^{PR(d/d)} oviducts were morphologically larger than those of *Tsc1*^{PR(f/f)} females (Fig. 2B). Consistent with this observation, we found epithelial hyperplasia in *Tsc1*^{PR(d/d)} oviducts as early as 30 days of age, similar to the phenotype seen in *Tsc1*^{Amhr2(d/d)} oviducts (Tanaka et al., 2012). Most wild-type oviducts showed normal histology with no epithelial hyperplasia and timely passage of embryos through the oviduct into the uterus. These results suggest that this hyperplasia is a potential cause of oviductal retention of embryos in both *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)} females.

Heightened mTORC1 signaling after *Tsc1* deletion was examined by immunohistochemistry for the phosphorylated form of ribosomal protein Rps6 (pS6), a downstream target of mTORC1 (Fig. 2C). Indeed, intensity of pS6 signals was higher primarily in the mucosa in *Tsc1*^{PR(d/d)} oviducts; the signal was only slightly higher in the muscular layer. In comparison, deletion of *Tsc1* by *Amhr2-Cre* resulted in a modest increase of pS6 in the oviductal mucosa (Fig. 2D).

The above results led us to examine the ovulation and fertilization status in *Tsc1*^{PR(d/d)} and *Tsc1*^{Amhr2(d/d)} females. *Cre* is primarily expressed in oocytes and surrounding GCs in ovaries of *Amhr2-Cre* females, whereas it is primarily present in CL in *PR-Cre* ovaries (Fig. 1). The deletion of *Tsc1* in different compartments of the ovary did not affect ovulation in either model. The average number of two-cell embryos was comparable in *Tsc1*^{Amhr2(d/d)} and *Tsc1*^{PR(d/d)} females, suggesting that the process of ovulation is not adversely affected in both of these deleted females. In addition, the fertilization rates in these two models were comparable with those of their corresponding wild-type counterparts (Tables I and II). These results suggest that sperms can still travel into oviducts despite oviductal epithelial hyperplasia, and ovulated eggs are capable of fertilization in *Tsc1*^{Amhr2(d/d)} and *Tsc1*^{PR(d/d)} oviducts.

On Day 5 of pregnancy, implantation sites can be visualized by distinct blue bands after an intravenous blue dye injection, and the implantation sites are evenly distributed in wild-type uteri (Paria et al., 1993). Interestingly, we could not detect any implantation sites in uteri of *Tsc1*^{Amhr2(d/d)} and *Tsc1*^{PR(d/d)} females. Instead, embryos and eggs were recovered from oviducts, indicating persistent oviductal retention of embryos and eggs (Tables I and II). These results suggest that the hyperplasia resulting from deletion of *Tsc1* in epithelial cells,

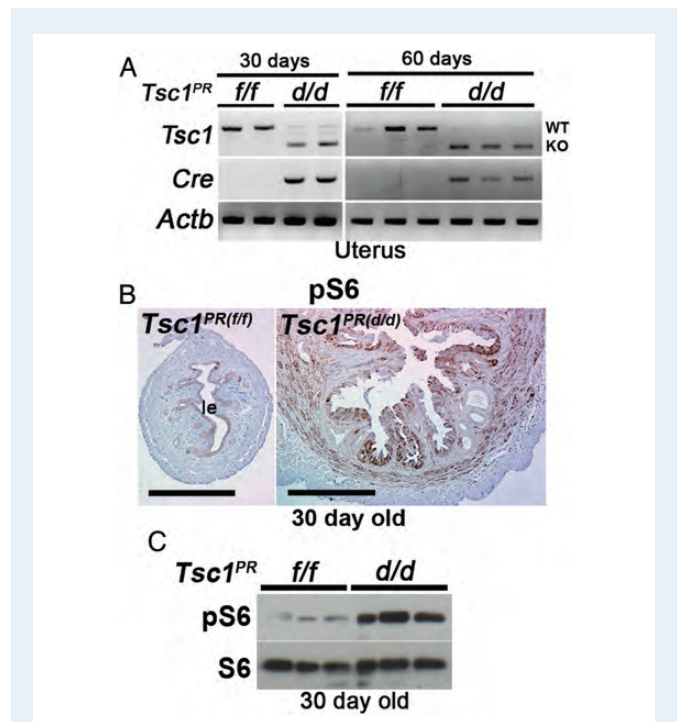


Figure 3 Heightened mTORC1 signaling in *Tsc1*-deleted uteri. (A) RT-PCR showing inactivation of *Tsc1* in *Tsc1*^{PR(d/d)} uteri compared with *Tsc1*^{PR(f/f)} uteri from mice at 30 or 60 days of age. *Actb* served as a loading control. (B) Immunohistochemistry of pS6 shows increased mTORC1 signaling in *Tsc1*^{PR(d/d)} uteri ($n = 4$ females) compared with control *Tsc1*^{PR(f/f)} uteri ($n = 4$ females) from mice at 30 days of age. Bars, 400 μ m. le, luminal epithelium. (C) Western blotting of pS6 indicates increased mTORC1 signaling in *Tsc1*^{PR(d/d)} uteri from mice at 30 days of age. S6 served as a loading control.

along with epithelial shedding into the oviductal lumen, causes defective oviductal embryo transport. Many of the recovered eggs were degenerate, presumably resulting from the previous ovulation; this may provide one possible explanation for recovery of degenerated eggs and embryos from deleted oviducts on Day 2 of pregnancy.

On Day 5 of pregnancy, examination of embryos recovered from oviducts showed compromised embryonic development in both *Tsc1*^{Amhr2(d/d)} and *Tsc1*^{PR(d/d)} females. Embryos ranging from two-cell

to blastocyst stages were recovered from deleted oviducts with a large number of embryos undergoing degeneration (Tables I and II). The results suggest that compromised embryo development is due to sub-optimal hyperplastic oviductal environment, inherent defects in oocyte competency or both. As shown in Table III, 50% of eggs from $Tsc1^{PR(d/d)}$ females fertilized *in vitro* developed into blastocysts, comparable with the rate of fertilization in $Tsc1^{PR(f/f)}$ females. These results suggest that the quality and fertilization competency of ovulated eggs from $Tsc1^{PR(d/d)}$ females were apparently normal and that the $Tsc1^{PR(d/d)}$ oviductal environment is not conducive to embryo development. In contrast, most IVF-derived embryos from $Tsc1^{Amhr2(d/d)}$ females did not develop into blastocysts, indicating inherent defects in oocyte quality in these mice.

Deletion of *Tsc1* increases mTORC1 signaling and alters uterine receptivity

Our next objective was to examine whether uterine deletion of *Tsc1* alters mTORC1 signaling and whether this alteration corresponds to changes in the expression of genes known to be critical for uterine receptivity since these mice show implantation failure. We first confirmed *Cre* expression and level of *Tsc1* deletion in non-pregnant $Tsc1^{PR(d/d)}$ uteri from 30-day old mice by RT-PCR analysis. As shown in Fig. 3A, uterine *Tsc1* mRNA levels are drastically reduced in these mice with low to undetectable levels by 60 days of age. We next examined whether a loss of *Tsc1* led to increased pS6 levels by western blotting and immunohistochemistry (Fig. 3B and C). pS6 was indeed upregulated in $Tsc1^{PR(d/d)}$ uteri with an increased

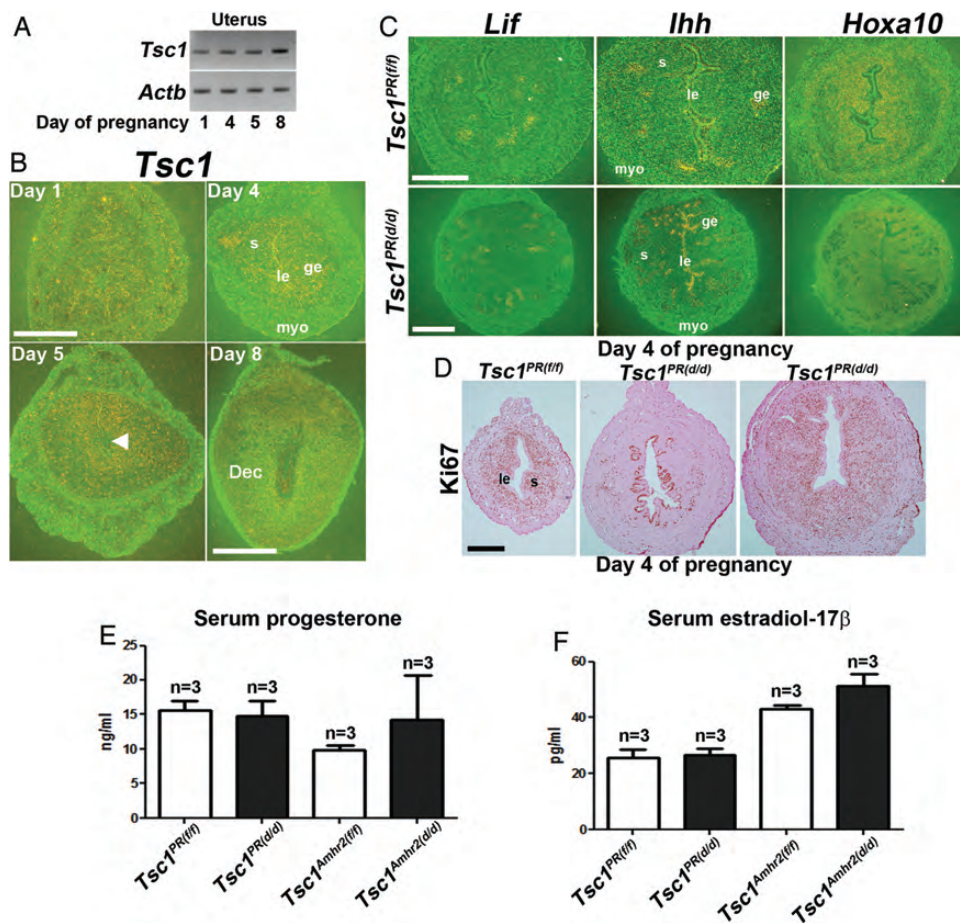


Figure 4 Uterine receptivity is compromised in $Tsc1^{PR(d/d)}$ uteri. (A and B) RT-PCR and *in situ* hybridization of *Tsc1* in wild-type peri-implantation uteri. *Tsc1* expression is primarily seen in the epithelium and myometrium (myo) on Day 1, glandular epithelium (ge) on Day 4 and in the stroma (s) on Days 5 and 8 of pregnancy ($n = 1$ per day of pregnancy). Magnification of Days 1, 4, and 5 of pregnancy: bar, 500 μm ; Magnification of Day 8 of pregnancy: bar, 1 mm. Arrowhead denotes the site of blastocyst. *Actb* served as a loading control. le, luminal epithelium; Dec, decidua. (C) *In situ* hybridization of *Lif*, *Ihh* and *Hoxa10* expression $Tsc1^{PR(f/f)}$ ($n = 3$ females) and $Tsc1^{PR(d/d)}$ ($n = 2$ females) uteri on Day 4 of pregnancy show comparable expression between the two groups. Upper bar, 250 μm ; lower bar, 500 μm . (D) Normal Ki67 staining is seen in $Tsc1^{PR(f/f)}$ females Day 4 ($n = 3$, left panel), while uteri of two of three $Tsc1^{PR(d/d)}$ females showed compromised epithelial differentiation and stromal proliferation (middle panel); uteri from one $Tsc1^{PR(d/d)}$ mouse showed a relatively normal expression pattern (right panel). Le, luminal epithelium; s, stroma. Bar, 400 μm . (E and F) Serum levels of E_2 and P_4 on Day 4 of pregnancy (mean \pm SEM). Three independent samples were examined in each group.

intensity of pS6 immunostaining observed in the epithelium and circular muscle with much less signals in the stroma. Unexpectedly, pS6 levels were not uniform in $Tsc1^{PR(d/d)}$ epithelia; a positive signal was primarily restricted to the luminal epithelium with low to undetectable levels in the glandular epithelium.

We next examined $Tsc1$ expression in the pregnant uterus during the peri-implantation period (Fig. 4A and B). $Tsc1$ was localized in the epithelium and myometrium on Day 1, glandular epithelium on Day 4 and in the stroma on Days 5 and 8 of pregnancy. However, the deletion of uterine $Tsc1$ in $Tsc1^{PR(d/d)}$ females did not affect estrogen-responsive leukemia inhibitor factor (*Lif*) expression or progesterone-responsive *Hoxa10* and Indian hedgehog (*Ihh*) expressions on Day 4 of pregnancy (Fig. 4C). We also examined uterine cell proliferation and differentiation status by Ki67 immunostaining on Day 4 when epithelial cells normally become differentiated and stromal cells undergo extensive proliferation in the preparation for implantation and decidualization in wild-type mice. Epithelial cells on Day 4 of pregnancy were still Ki67-positive in some $Tsc1^{PR(d/d)}$ uteri with sparse Ki67-positive cells in the stroma, indicating impaired epithelial differentiation and stromal cell proliferation in some females but not in others (Fig. 4D). Notably, serum progesterone and estradiol-17 β levels were not significantly altered between control

and deleted females on Day 4 of pregnancy (Fig. 4E and F). These results indicate that $Tsc1$ deficiency compromised epithelial–mesenchymal interactions required for uterine receptivity for implantation. In fact, a previous report showed failure of implantation after embryo transfer experiments using $Tsc1^{Amhr2(d/d)}$ mice (Tanaka et al., 2012).

Uterine deletion of $Tsc1$ induces epithelial hyperplasia but not cancer

Our next objective was to examine whether uterine deletion of $Tsc1$ in mice results in endometrial cancer. To address this question, we measured uterine wet weight as an index of uterine growth and compared uterine histology between $Tsc1^{PR(f/f)}$ and $Tsc1^{PR(d/d)}$ females. At 30 days of age, $Tsc1^{PR(d/d)}$ uteri became rigid and weighed considerably more (103.4 ± 26.0 mg, $n = 8$) compared with $Tsc1^{PR(f/f)}$ uteri (24.8 ± 5.3 mg, $n = 8$). At this age, $Tsc1^{PR(d/d)}$ uteri showed epithelial hyperplasia (Fig. 5A and Table IV). Although complex hyperplasia was noted in a few cases (2 of 8 females), most uteri showed simple hyperplasia relative to normal epithelial cells. Furthermore, the circular muscle layer appeared thicker with a disrupted architecture. At 90 days of age, the weight of $Tsc1^{PR(d/d)}$ uteri continued to increase

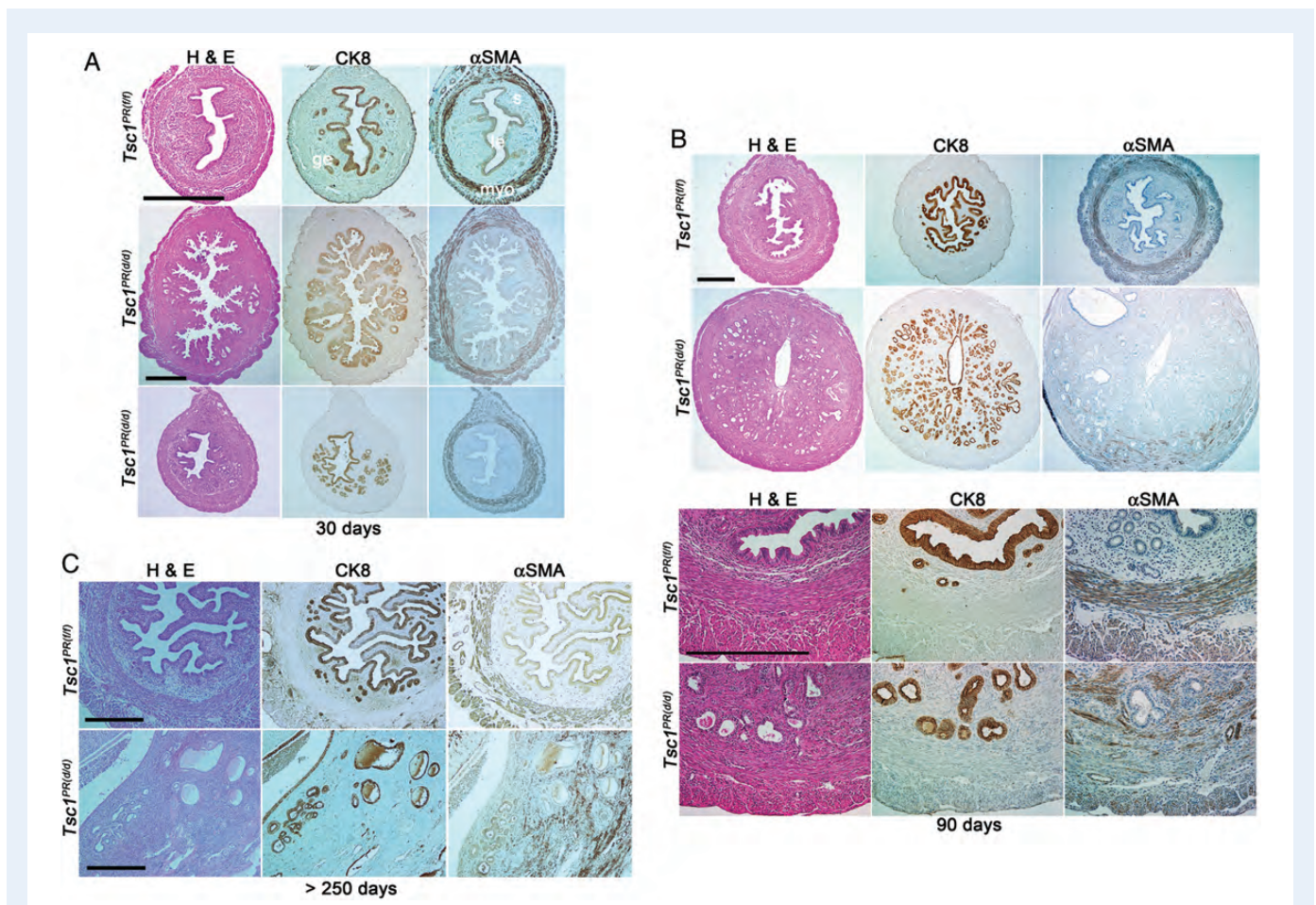


Figure 5 $Tsc1^{PR(d/d)}$ uteri show epithelial hyperplasia but not cancer. (A–C) Histology (H&E) and immunostaining of $Tsc1^{PR(f/f)}$ and $Tsc1^{PR(d/d)}$ uteri from mice at 30 days (A), 90 days (B) and more than 270 days of age (C). Bars, 400 μ m. CK8 and α SMA are markers for epithelia and myometrium, respectively. Le, luminal epithelium; ge, glandular epithelium; s, stroma; myo, myometrium. The number of females examined per genotype is indicated in Table IV.

Table IV Uterine histology of *Tsc1*-deleted mice.

Age (days)	Genotype	No. of mice examined	Uterine histology	Mice with indicated histology (%)
30	<i>Tsc1</i> ^{PR(f/f)}	8	Normal	8 (100%)
	<i>Tsc1</i> ^{PR(d/d)}	8	Simple hyperplasia ^a	5 (62.5%)
			Complex hyperplasia with luminal hyperplasia	2 (25%)
			Complex hyperplasia	1 (12.5%)
60	<i>Tsc1</i> ^{PR(f/f)}	7	Normal	7 (100%)
	<i>Tsc1</i> ^{PR(d/d)}	9	Moderate to marked simple hyperplasia	9 (100%)
90	<i>Tsc1</i> ^{PR(f/f)}	7	Normal	6 (85.8%)
			Mild simple hyperplasia	1 (14.2%)
Survival ^b	<i>Tsc1</i> ^{PR(d/d)}	8	Simple hyperplasia	8 (100%)
	<i>Tsc1</i> ^{PR(f/f)}	8	Normal	6 (75%)
			Simple hyperplasia	2 (25%)
	<i>Tsc1</i> ^{PR(d/d)}	6	Simple hyperplasia, stromal polyp with stromal overgrowth	4 (66.6%)
			Simple hyperplasia with focal atypia, stromal polyp with stromal overgrowth	1 (16.7%)
			Marked simple hyperplasia Leiomyosarcoma	1 (16.7%) ^c

^aRelative to normal epithelium.

^b*Tsc1*^{PR(f/f)}, 323–383 days old; *Tsc1*^{PR(d/d)}, 271–308 days old.

^cThis uterus possibly has a malignant spindle cell tumor.

(290.3 ± 28.4 mg, *n* = 8) compared with those of *Tsc1*^{PR(f/f)} uteri (78.3 ± 6.1 mg, *n* = 7). At this time, the circular muscle appeared to be more disrupted and the boundary between the stromal and circular muscle compartments was disturbed in *Tsc1*^{PR(d/d)} uteri (Fig. 5B). *Tsc1*^{PR(d/d)} glandular epithelia showed simple hyperplasia but no signs of cancer (Fig. 5B and Table IV). Furthermore, the luminal epithelium seemed to be intact. We also observed hyperplasia in *Tsc1*^{Amhr2(d/d)} uterine glandular epithelium (data not shown).

Long-term survival studies predicted shorter lifespan of *Tsc1* deleted mice. While control mice (*Tsc1*^{f/f}) survived for longer than 320 days of age, general malaise and vaginal bleeding were apparent in *Tsc1*^{PR(d/d)} females as early as 270 days and thus, experiments were terminated (Table IV). *Tsc1*^{PR(d/d)} uteri were grossly enlarged (*Tsc1*^{f/f}: 0.13 ± 0.02 g; *Tsc1*^{PR(d/d)}: 12.39 ± 0.49 g (including ovaries due to extensive adhesions), *n* = 6), showing excessive stromal overgrowth and polyp formation. However, the epithelium still exhibited simple hyperplasia and did not transform into cancer although the degree of hyperplasia increased with age in *Tsc1*^{PR(d/d)} uteri (Fig. 5C). Notably, only one *Tsc1*^{PR(d/d)} uterus exhibited leiomyosarcoma, while the others showed only benign tumors (Table IV). These results indicate that *Tsc1* deletion is not sufficient to initiate endometrial cancer.

Discussion

There is considerable evidence that mTORC1 signaling plays distinct roles in the female reproductive tract ranging from fertility to cancer. We have previously shown that conditional deletion of uterine *Pten* initiates endometrial cancer in mice with 100% penetrance (Daikoku et al., 2008). In this study, we have shown that *Tsc1* deletion in oviducts by *PR-Cre* causes retention of embryos in

the oviduct and interferes with their development after fertilization. The occurrence of oviductal embryo retention in *Tsc1* deleted oviduct by *Amhr2-Cre* has previously been reported and is confirmed here (Tanaka et al., 2012). Intriguingly, we find that the deletion of oviductal *Tsc1* compromises embryo development when trapped in the oviduct, suggesting that the etiology of early pregnancy failure in *Tsc1*-deleted females is 2-fold: compromised embryo passage through the oviduct and abnormal development within the oviduct. Although the exact nature of the causative factors that influence the quality of the oviductal environment and retention of embryos is not clear, hyperplasia of the epithelium and shedding of these epithelial cells into the lumen could be a potential cause. Further studies using these mouse models are warranted to understand the mechanisms underlying oviductal embryo retention. Phenotypical differences in embryo development *in vitro* between mice with deletion of *Tsc1* by *PR-Cre* and *Amhr2-Cre* are not clear at this time, but may result from *in vitro* conditions that perpetuated some inherent defects in oocytes from *Tsc1*^{Amhr2(d/d)} mice.

Compromised *Tsc* activity and increased mTORC1 signaling are associated with endometrial cancer in humans (Mak and Yeung, 2004; Lu et al., 2008; McCampbell et al., 2010). Although the importance of the mTORC1 pathway in endometrial cancer was previously shown in mice using conditional deletion of uterine epithelial *Lkb1* which indirectly inhibits the pathway mediated by *Tsc1* and *Tsc2* complexes, it is still unclear whether heightened mTORC1 in the uterine epithelium itself is sufficient to initiate endometrial cancer since *Lkb1* also regulates alternate pathways in addition to the mTORC1 pathway (Katajisto et al., 2007; Contreras et al., 2010). In the present study, we demonstrate that heightened mTORC1 signaling, resulting from deletion of *Tsc1*, is not sufficient to initiate cancer but can give rise to epithelial hyperplasia, suggesting that heightened

mTORC1 activation alone is not sufficient to initiate endometrial cancer but promotes its progression. In fact, mTORC1 activation is more frequently observed in advanced-stage and high-grade endometrial cancer in humans (Darb-Esfahani *et al.*, 2009).

Based on our present findings that mTORC1 activation did not uniformly occur in all uterine epithelia and that activation was not sustained past 90 days of age in *Tsc1*^{PR(d/d)} females, we believe that the deletion of *Tsc1* cannot initiate uterine cancer due to compensatory regulatory mechanisms limiting mTORC1 signaling. Another possibility is that an additional insult must be superimposed upon mTORC1 activation to initiate endometrial cancer. Accordingly, it has been reported that ~25% of patients with endometrial cancer previously had benign endometrial biopsies (Torres *et al.*, 2012). It would be interesting to examine whether deletion of a second tumor suppressor superimposed on *Tsc1* deletion can more readily initiate endometrial cancer. Alternatively, since the uterine epithelium is considered the site of origin for endometrial cancer, it is not yet known whether heightened mTORC1 signaling after uterine epithelial deletion of *Tsc1* contributes to the initiation and/or progression of this cancer.

Uterine deletion of *Tsc1* also increased mTORC1 signaling in the myometrium and stroma and transformed these tissue compartments into large yet benign tumors. Another interesting observation is that mesenchymal *Tsc1* deletion with *Amhr2-Cre* caused uterine epithelial hyperplasia, although at present, we do not understand the epithelial–mesenchymal interactions impacting this epithelial transformation. These results suggest that the effects of *Tsc1* deletion on mTORC1 activation vary depending on the cell type. Although the mechanisms governing these interactions remain unknown, further studies which add to our understanding of these disease processes using models with cell-type specific deletions of critical tumor suppressors may help to establish better strategies to treat endometrial cancer and other gynecologic diseases in which mTORC1 signaling is implicated.

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Authors' roles

T.D. and S.K.D. designed research; T.D., M.Y., H.X., X.S., J.C. and S.K.D. performed research; L.H.E. analyzed histology; T.D., M.Y., H.X., X.S., J.C., L.H.E. and S.K.D. analyzed data; and T.D., X.S., J.C., and S.K.D. wrote the paper.

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Conflict of interest

None declared.

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