

Signaling through retinoic acid receptors is essential for mammalian uterine receptivity and decidualization

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Retinoic Acid (RA) signaling has long been speculated to regulate embryo implantation, because many enzymes and proteins responsible for maintaining RA homeostasis and transducing RA signals are tightly regulated in the endometrium during this critical period. However, due to lack of genetic data, it was unclear whether RA signaling is truly required for implantation, and which specific RA signaling cascades are at play. Herein we utilize a genetic murine model that expresses a dominant negative form of RA receptor specifically in female reproductive organs to show that functional RA signaling is fundamental to female fertility, particularly implantation and decidualization. Reduction in RA signaling activity severely affects the ability of the uterus to achieve receptive status and decidualize, partially through dampening follistatin expression and downstream activin B/BMP2 signaling. To confirm translational relevance of these findings to humans, human endometrial stromal cells (hESCs) were treated with pan-RAR antagonist to show that *in vitro* decidualization is impaired. RNAi perturbation of individual *RAR* transcripts in hESCs revealed that *RAR α* in particular is essential for proper decidualization. These data provide direct functional evidence that uterine RAR-mediated RA signaling is crucial for mammalian embryo implantation, and its disruption leads to failure of uterine receptivity and decidualization resulting in severely compromised fertility.

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1 **Signaling through Retinoic Acid Receptors is Essential for**
2 **Mammalian Uterine Receptivity and Decidualization**

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19 **Abstract**

20 Retinoic Acid (RA) signaling has long been speculated to regulate embryo implantation, because
21 many enzymes and proteins responsible for maintaining RA homeostasis and transducing RA
22 signals are tightly regulated in the endometrium during this critical period. However, due to lack
23 of genetic data, it was unclear whether RA signaling is truly required for implantation, and which
24 specific RA signaling cascades are at play. Herein we utilize a genetic murine model that
25 expresses a dominant negative form of RA receptor specifically in female reproductive organs to
26 show that functional RA signaling is fundamental to female fertility, particularly implantation
27 and decidualization. Reduction in RA signaling activity severely affects the ability of the uterus
28 to achieve receptive status and decidualize, partially through dampening follistatin expression
29 and downstream activin B/BMP2 signaling. To confirm translational relevance of these findings
30 to humans, human endometrial stromal cells (hESCs) were treated with a pan-RAR antagonist to
31 show that in vitro decidualization is impaired. RNAi perturbation of individual *RAR* transcripts
32 in hESCs revealed that *RAR α* in particular is essential for proper decidualization. These data
33 provide direct functional evidence that uterine RAR-mediated RA signaling is crucial for
34 mammalian embryo implantation, and its disruption leads to failure of uterine receptivity and
35 decidualization resulting in severely compromised fertility.

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37

38 **Significance Statement:** Female infertility affects as many as 72 million individuals worldwide,
39 with 10% of cases remaining unsolved after clinical investigation. Retinoic acid is the
40 biologically functional metabolite of dietary vitamin A. The current study shows that in the event
41 that the mammalian uterus cannot respond properly to retinoic acid, it cannot properly receive an
42 otherwise healthy embryo for implantation, and pregnancy is unlikely to be achieved. A
43 functional uterine response to retinoic acid is therefore critical for early pregnancy success.

44 **Introduction**

45 During early pregnancy, the mammalian endometrium responds to changing ovarian hormones
46 and signaling cues from embryos. The endometrium undergoes extensive growth and
47 differentiation to become appropriately receptive to the incoming blastocysts for implantation (1,
48 2). Commonly referred to as the “window of implantation”, these complex endometrial events
49 must happen in rapid succession in a short time frame in order to achieve a receptive phase. In
50 the mouse, the implantation window starts the morning of 3.5 days post coitum (dpc, 0.5 dpc =
51 1200h of the day of vaginal plug), when fertilized eggs complete their development through the
52 blastocyst stage and descend into the uterus. By this time, the receiving luminal epithelium (LE)
53 ceases proliferation and initiates differentiation under the influence of rising progesterone (P4)
54 and a small estradiol (E2) surge, while the underlying stromal cells undergo extensive
55 proliferation and start to differentiate into morphologically and functionally distinct decidual
56 cells. Toward the evening of 4.5 dpc, the implantation window closes and the uterus enters the
57 refractory phase, during which embryos cannot implant. The rapid uterine changes that define
58 the implantation window are tightly regulated by a network of signaling pathways orchestrated
59 by hormones, growth factors, cytokines, and transcription factors. The importance of uterine
60 receptivity genes is evidenced by the severe implantation and/or decidualization defects that
61 occur in response to loss-of-function mutations (3-10).

62

63 Retinoic acid (RA), a physiologically active metabolite of its inactive precursor retinol (vitamin
64 A), is essential for many biological processes including cell survival, differentiation and
65 apoptosis (11). Vitamin A deficiency, as well as genetically disrupting RA function, leads to
66 defects in the development of many organs and tissues, including the nervous system, kidney,

67 skeleton, heart, lung and urogenital tracts (12). RA exerts its biological functions mainly through
68 binding to the nuclear RA receptors (RARs) facilitated by the Cellular Retinoic Acid-Binding
69 Proteins (CRABPs), or less frequently, to the non-canonical Peroxisome Proliferator-Activated
70 Receptor β/γ (PPAR β/γ) facilitated by Fatty Acid-Binding Protein 5 (FABP5). The ratio of
71 intracellular lipid-binding proteins CRABP2:FABP5 tips the balance toward one signaling
72 pathway or the other, which frequently have opposing effects (13). Both RARs and PPARs form
73 heterodimers with Retinoid X Receptors (RXRs), and they regulate target gene expression by
74 directly binding to RA response elements (RAREs) and peroxisome proliferator response
75 elements (PPREs), respectively. In addition to these two signaling cascades, RA can also bind to
76 cytoplasmic RARs and trigger rapid kinase phosphorylation, which in turn regulates downstream
77 signaling events (14).

78
79 Previous studies have implicated RA signaling in regulating female fertility. In both the rodent
80 and human endometrium, expressions of RA synthesizing (ALDHs/RALDHs) and metabolizing
81 enzymes (CYP26), as well as RA-binding proteins that regulate its bioavailability
82 (CRBPs/CRABPs/FABP5), are temporally and spatially controlled during early pregnancy (15-
83 17), suggesting their involvement in uterine receptivity and embryo implantation. In addition,
84 high expression of RA signaling receptors, including RARs, PPAR β/γ and RXRs, has been
85 reported at implantation sites of human and rodent endometrium (17-19). In cultured human
86 endometrial stromal cells (hESCs), gene silencing of CRABP2 and FABP5 by siRNA inhibits and
87 promotes decidualization, respectively (20), suggesting that RA-RAR signaling favors
88 decidualization. In seemingly contradictory in vitro data, treatment of RA at pharmacological
89 levels in hESC culture appears to impair decidualization (17, 21). Given the complex genomic and

90 non-genomic downstream events elicited by RA in various tissues and the absence of any uterine
91 data from genetic animal models, the definitive role of RA signaling in implantation requires
92 clarification.

93

94 In the current study, we utilized a previously characterized mouse strain that carries the *RaraT403*
95 truncated form of human *RARα* knocked in to the *Rosa26* locus to dissect the role of RA-RAR
96 signaling in embryo implantation (22). Cre-mediated recombination removes the floxed-STOP
97 sequence upstream of the *RaraT403*, resulting in expression of the dominant negative form of
98 RARα (hereinafter referred to as *RaraDN*), and subsequent inhibition of endogenous RAR-
99 mediated transcriptional regulation of target genes. We showed that functional RAR signaling is
100 required for mammalian uterine receptivity and decidualization both in the mouse model and in
101 cultured hESCs.

102

103 **Results**

104 ***RaraDN*^{Pgr} Females Are Severely Subfertile**

105 To generate mice with disrupted RA signaling in the female reproductive tracts, *RaraDN*^{f/+} mice
106 were mated to *Pgr-Cre* mice, resulting in offspring females carrying both alleles (*RaraDN*^{f/+};
107 *Pgr-Cre*, hereafter referred to as *RaraDN*^{Pgr}) and littermate controls (CTRL, no *Pgr-Cre*). *Pgr-*
108 *Cre*-mediated gene recombination in the uterus is first detected in the luminal and glandular
109 epithelia starting at two weeks of age, and gradually expands to the stroma and myometrium
110 (23). To assess disruption of RAR signaling, we examined the expression of several *Hox* genes,
111 known direct downstream targets of RAR, in the uteri of ovariectomized CTRL and *RaraDN*^{Pgr}

112 mice (**Figure S1**). Reduction in expression of the majority of the *Hox* genes in the *RaraDN^{Pgr}*
113 uteri demonstrates that RAR signaling is successfully disrupted in these animals. Female fertility
114 was evaluated by breeding test of wild-type males with either *RaraDN^{Pgr}* or CTRL females and
115 tracking the number of litters and pups produced by each female for 200 consecutive days. Six
116 out of seven *RaraDN^{Pgr}* females tested were completely sterile, producing zero litters over the
117 span of seven months (**Figure 1A**). The remaining *RaraDN^{Pgr}* female produced only two litters,
118 each consisting of only one pup after a long hiatus post mating setup (**Figure 1A and B**).
119 *RaraDN^{Pgr}* females as a whole are therefore severely subfertile, producing significantly fewer
120 litters and pups than controls (**Table 1**, $p=9.8\times 10^{-9}$ and 7.2×10^{-9} , respective, $n=5$ for CTRL and
121 $n=7$ for *RaraDN^{Pgr}*). No apparent developmental and behavioral abnormalities were observed in
122 the two pups born to the *RaraDN^{Pgr}* sterility escapee. Vaginal plugs were consistently observed
123 in the *RaraDN^{Pgr}* females, ruling out the possibility of behavioral issues preventing mating. As
124 these females never presented palpable pregnancies, early-stage pregnancy defects were
125 suspected.

126

127 ***RaraDN^{Pgr}* Females Exhibit Implantation Defects**

128 Mouse embryo implantation occurs between 3.5 and 4.5 dpc, when blastocysts attach to the
129 luminal epithelium which in turn triggers the underlying stromal cells to undergo
130 decidualization. Successful implantation is accompanied by increased local vascular
131 permeability, which can be visualized by tail-vein injection of Chicago blue dye. Distinct blue
132 dots indicating implantation sites were easily detectable along the uterine horns of CTRL mice at
133 4.5 dpc (**Figure 1D, arrows**, 9.7 ± 2.1 , $n=3$), but were completely absent in the *RaraDN^{Pgr}* uteri
134 (**Figure 1C, E**, 0.0 ± 0.0 , $n=3$, $p=0.0013$).

135
136 Many factors, alone or in combination, can contribute to failed pregnancy at an early stage,
137 including abnormal ovulation, irregular ovarian hormone levels, and poor uterine receptivity.
138 Since *Pgr*-Cre is also active in the adult ovary, including corpora lutea and hCG-stimulated
139 granulosa cells (23), it is essential to investigate whether disturbance of RA signaling affects
140 ovarian functions. Evaluation of the *Rara*DN^{Pgr} ovaries at 3.5 dpc revealed normal histology with
141 presence of multiple corpus lutea, which are the remnants of successfully matured vesicular
142 follicles after ovulation (**Figure S2A, B**). Morphologically normal blastocysts were recovered
143 from *Rara*DN^{Pgr} females by flushing the oviducts and uterine horns at 3.5 dpc (**Figure S2A,**
144 **inset**), and no significant difference was observed in the number of retrieved blastocysts at this
145 stage (**Figure S2C**, CTRL 8.5±3.5, n=3; vs. *Rara*DN^{Pgr} 6.5±2.1, n=3, $p=0.27$) indicating normal
146 fertilization rates. In addition, evaluation of serum ovarian hormone levels at this stage by
147 ELISA revealed no significant differences (**Figure S2D**). Together these data indicate normal
148 ovarian function in *Rara*DN^{Pgr} females.

149

150 ***Rara*DN^{Pgr} Females Exhibit Uterine Receptivity Defects**

151 Successful embryo implantation depends on the achievement of uterine receptivity through a
152 series of molecular, hormonal and morphological changes. In the 3.5 dpc mouse uterus, the
153 luminal epithelium typically ceases proliferation under the influence of decreased estrogen (E2)
154 and surging progesterone (P4) levels to prime for rapid remodeling and embryo embedding (24).
155 Meanwhile, luminal epithelial cells turn off genes for apical-basal polarity like Cadherin1
156 (CDH1, a.k.a. E-cadherin) to allow attachment of trophoblast cells to their apical pole (25). To
157 assess the status of uterine receptivity, we first examined gross uterine morphology at this stage

158 and found no overt abnormalities except for aberrant luminal closure (**Figure 1F, G**). Uteri of
159 control 3.5 dpc females exhibited typical signs of receptivity, i.e. halted epithelial proliferation
160 evidenced by limited phospho-Histone H3 staining (pHH3, **Figure 1G**) and reduced CDH1
161 expression exclusively in the luminal epithelium (**Figure 1K, arrowheads**). By contrast, the
162 *RaraDN^{Pgr}* uteri sustained high LE proliferating activity (**Figure 1H, arrowheads**) and high
163 CDH1 expression (**Figure 1J, arrowheads**) in the luminal epithelium, consistent with a pre- or
164 non-receptive uterus.

165
166 Quantitative RT-PCR was performed to further interrogate expression of genes involved in
167 uterine receptivity. Amphiregulin (*Areg*), a member of the epidermal growth factor family, is
168 upregulated exclusively in the uterine epithelium at 3.5 dpc surrounding the embedding embryos
169 in a P4-dependent manner (26). This up-regulation was absent in the *RaraDN^{Pgr}* mutant (**Figure**
170 **1L**). Expression of early growth response gene 1 (*Egr1*), a zinc finger transcription factor that is
171 crucial for cell proliferation and angiogenesis, was previously reported to be induced in the
172 subluminal stroma surrounding the blastocysts (27), but it is barely detectable in the mutant
173 uterus at 3.5 dpc. Previous studies in ovarian hormone-responsive cells including uterine
174 epithelial cells have shown that E2 signaling can promote luminal epithelium cell proliferation
175 by transactivating expression of the cell cycle gene cyclin D1 (*Ccnd1*), as well as facilitating its
176 nuclear translocation (28, 29). *RaraDN^{Pgr}* uteri exhibit a marked increase in *Ccnd1* mRNA,
177 which may contribute to the mutant's aberrant epithelial proliferation. Deregulation of
178 transcription factors essential for uterine receptivity and embryo implantation was also evident in
179 the *RaraDN^{Pgr}* uteri at 3.5 dpc, including those expressed in the epithelial compartment, such as
180 Forkhead Box O1 (*Foxo1*) (30), and those exclusively expressed in the stroma, such as

181 Homeobox A10 (*Hoxa10*) (31). RNAscope in situ hybridization of *Foxo1* revealed that its
182 elevation in the mutant is confined to the uterine epithelium (**Figure 1N, O**). In addition, analysis
183 of a subset of gold standard receptivity biomarkers used in customized endometrial receptivity
184 arrays (ERA) for clinical endometrial evaluation in humans (32, 33) revealed markedly reduced
185 expression of many receptivity biomarkers in the *RaraDN^{Pgr}* uteri during the peri-implantation
186 period (**Figure S3**). Even though many of these genes have been reported to be regulated by
187 ovarian hormones, the changes we observed in the *RaraDN^{Pgr}* mutant are unlikely to be elicited
188 solely by altered hormone signaling, because serum ovarian hormone level (**Figure S2**) as well as
189 uterine expression of ovarian hormone receptors and some of their well-established targets
190 remained unchanged (**Figure 1M**). Immunofluorescence of ESR1 and PR further confirmed that
191 the ovarian hormone receptors were expressed at normal locations and levels in *RaraDN^{Pgr}*
192 females comparable to their wildtype counterparts (**Figure 1P-S**).

193

194 **Decidualization is Compromised in *RaraDN^{Pgr}* Females**

195 Despite the absence of luminal closure, which is thought to help immobilize the embryos for
196 implantation, blastocyst attachment appears to successfully occur in the *RaraDN^{Pgr}* uteri at 4.5
197 dpc (**Figure 2B, D**), raising the possibility that failures in subsequent pregnancy events also
198 contribute to the fertility defects. As we and others have previously reported (34, 35), during
199 embryo attachment, strong CDH1 expression is only present in the apical poles of the uterine
200 epithelium and barely detectable on the basal side (**Figure 2C, arrows**). Interestingly, this
201 polarized localization of CDH1 is absent in the *RaraDN^{Pgr}* uteri; strong CDH1 staining was
202 observed on both sides (**Figure 2D, arrowheads**). Following embryo attachment, fibroblastic
203 uterine mesenchymal cells undergo decidualization. Decidualization is the rapid proliferation and

204 differentiation of these cells into morphologically distinct decidual cells, which provide a
205 plethora of growth factors and cytokines to support embryo development and serve an
206 immunoregulatory role during early pregnancy. To investigate whether decidualization is
207 affected in the *Rara*DN^{Pgr} females, we first examined the expression of known decidualization
208 markers during natural pregnancy in these mutants. Transcription factor heart-and neural crest
209 derivatives-expressed transcript 2 (HAND2) plays a critical role in uterine receptivity and
210 decidualization in the mouse, and its expression is induced in endometrial stromal cells starting
211 at 3.5 dpc and increases over time (7, 36). It modulates stromal-epithelial communications
212 through negative regulation of FGF signaling, and genetic ablation of *Hand2* in the mouse leads
213 to female infertility largely due to decidualization failure (7, 36). Immunofluorescence revealed
214 that HAND2 protein exhibits nuclear localization in the CTRL subepithelial stromal cells at 4.5
215 dpc (**Figure 2F, arrows**), but its level is dramatically reduced and its nuclear localization
216 undetectable in the mutant uterus (**Figure 2E**). This reduction in *Hand2* levels is confirmed at
217 the transcript level by qRT-PCR using RNA extracted from whole uterine tissues at 4.5 dpc
218 (**Figure 2G**). Expression of an array of genes involved in decidualization were evaluated by
219 qRT-PCR, and mutant uteri exhibit significant decreases in the majority of them, including *Add2*,
220 *Ereg*, *Gata2*, *Hbegf*, *Hsd11b1*, *Igfbp1*, *Lcn2*, *Prl* and *Wnt4*. Interestingly, a significant increase in
221 the transcript level of *Lpar3* was observed in the mutant 4.5 dpc uterus. *Lpar3* encodes
222 lysophosphatidic acid receptor 3, a G protein-coupled receptor for lysophosphatidic acid that
223 fine-tunes the local balance of P4 and E2 signaling during implantation (37, 38). Together these
224 results demonstrate a decidualization failure in *Rara*DN^{Pgr} females in the setting of natural
225 pregnancy.
226

227 To rule out the potential involvement of defective embryo attachment and/or defective hormone
228 regulation as a cause for decidualization failure, we performed an artificial decidualization assay.
229 In mice, decidualization of uterine stromal cells can be achieved by intra-luminal oil injection
230 into the uterine horns of ovariectomized and hormone-primed females followed by additional
231 hormone treatments post induction. As shown in **Figure 3A and B**, disruption of RA signaling in
232 *Rara*DN^{Pgr} uteri renders them non-responsive to decidual stimuli. Uterine weight gain due to
233 stimuli was completely abolished (**Figure 3C**), and differentiation markers like *Igfbp1*, *Prl* and
234 alkaline phosphatase (AP) failed to be induced in the mutant uteri (**Figure 3 D-F**). Genes
235 encoding RA receptors as well as some known downstream RAR signaling targets showed
236 differential expression in stimulated mutant uteri relative to controls (**Figure 3G**). Significant
237 reduction in transcript levels were observed in the mutants for RA receptors *Rara*, *Rarb*, *Rxra*
238 and *Rxrg*, as well as RA targets *Cdx1*, *Gbx2*, *Mmp9* and *Prrx2*. On the other hand, stimulated
239 mutant uteri exhibit drastic increases in mRNA levels for transcription factors *Msx1* and *Sox17*
240 relative to control decidua. Expression of *Msx1* was previously reported to sharply decline
241 following embryo attachment to prepare the uterus for implantation by modulating WNT and
242 FGF signaling between the epithelial and stromal compartments (39). In addition, persistent
243 *Msx1* expression was shown to be associated with uterine receptivity defects observed in *Lif*^{-/-}
244 mice (40). *Sox17* also plays critical roles during implantation through modulating the uterine
245 transcriptome (41). Most of the genes assayed including some of the RA receptors (**Figure 4A**)
246 and their downstream targets (**Figure 4B**) display the same trend of expression changes during
247 the peri-implantation period of natural pregnancy in *Rara*DN^{Pgr} uteri. These findings provide
248 further support that disrupted expression of RAR downstream targets likely contributes to the
249 decidualization defect observed in the *Rara*DN^{Pgr} mutants. We stress however, that at present we

250 cannot exclude the possibility that the decidualization defect in these mutants is secondary to the
251 observed uterine receptivity defect. Tissue-specific ablation of RA signaling in the implanting
252 uterus is required to address this point.

253

254 **Disrupted RAR-Signaling Leads to Reduced Follistatin and Aberrant Activin Signaling**

255 During gene expression analysis, we observed a striking decrease in the expression of follistatin
256 (*Fst*) in the *Rara*DN^{Pgr} at 3.5 dpc (**Figure 5A**). This is of particular interest as previous studies
257 revealed that *Fst* is a direct transcriptional target of RA signaling, containing RA responsive
258 elements (RAREs) in its promoter region (42). Additionally, *Pgr*-Cre-mediated genetic deletion
259 of *Fst* leads to female fertility defects very similar to our *Rara*DN^{Pgr} mutants (43).

260 Accompanying the sharp reduction of *Fst*, expression of inhibin β b (*Inhbb*), components of
261 activin B and downstream target of FST signaling, was significantly up-regulated (**Figure 5A**).

262 In the uterine-specific *Fst* knockout model, absence of FST and elevated activin B activity
263 caused reduction in BMP signaling, especially BMP2, through the Activin-SMAD signaling
264 pathway (43). In line with this notion, we observed a similar reduction in *Bmp2* expression at 4.5
265 dpc (**Figure 5B**), as well as reduction in phospho-SMAD1/5/8 (**Figure 5C**) in *Rara*DN^{Pgr}
266 mutants. RNAscope in situ hybridization was performed, which further confirmed the reduction
267 of *Fst* and *Bmp2* in *Rara*DN^{Pgr} uterus. *Fst* transcript was detected throughout the CTRL uterus at
268 3.5 dpc (**Figure 5D**) but was barely detectable in the mutant (**Figure 5E**). *Bmp2* transcript was
269 detected exclusively in the subepithelial stromal cells in CTRL uterus at 4.5 dpc (**Figure 5F**),
270 and its expression was markedly reduced in the mutant (**Figure 5G**). If the fertility defects
271 observed in the *Rara*DN^{Pgr} mice are indeed caused primarily by the loss of *Fst* expression, one
272 would expect the phenotype to be alleviated when FST is supplemented back to the mutant

273 uterus. To test this hypothesis, we isolated uterine stromal cells from 2.5 dpc mutants for in vitro
274 culture and added recombinant mouse FST to the medium at various concentrations. Forty-eight
275 hours after culture, expression of several decidualization markers is elevated by addition of FST;
276 for *Bmp2* and *Igfbp1* these changes are dose-dependent (**Figure 5H**). Similar restoration of
277 decidual marker expression were observed in a uterine organ culture system where either BSA-
278 or FST-soaked agarose beads were inserted into the lumens of 2.5 dpc *RaraDN^{Pgr}* uterine
279 segments and allowed to culture in vitro for two days (**Figure 5I**). To test whether FST is
280 sufficient to rescue the mutant implantation defects in vivo, FST was administered systemically
281 into 2.5 dpc *RaraDN^{Pgr}* females via tail vein injection. At 6.5 dpc, bulging regions along the
282 uterine horns resembling implantation sites were observed in these mutants (**Figure 5J**),
283 although they appeared smaller than normal implantation sites at this developmental stage.
284 Sections through the bulging regions revealed elevated *Bmp2* and *Hand2* transcripts (**Figure 5K**
285 **and 5L, respectively**), as well as extensive AP activity (**Figure 5M**), indicating restored
286 decidualization in the mutant by FST administration. However, histological analyses did not
287 reveal any uterine closure or embryo presence, suggesting that other aspects of implantation,
288 most likely uterine receptivity, cannot be rescued by FST alone. This partial rescue was observed
289 in two out of three *RaraDN^{Pgr}* females tested, with two and three bulging sites in each animal,
290 respectively. These data together demonstrate that RARs regulate uterine decidualization mainly
291 through FST.

292

293 **RAR Signaling Is Essential for Decidualization in hESCs**

294 In our previous study, we engineered a fluorescent reporter hESC line and performed genome
295 wide siRNA screening to identify genes required for normal decidualization (44). A total of 136

296 genes involved in the RA pathway were among the hits, including 29 that are upstream and 107
297 that are downstream of RAR signaling (**Figure S4**). To investigate the role of RAR signaling in
298 human endometrium, we performed individual siRNA knockdown against the three human *RAR*
299 genes (*RARA*, *RARB*, and *RARG*) in hESCs. In particular, knocking-down *RARA* significantly
300 inhibits in vitro decidualization of the hESCs, evidenced by decreased expression of
301 decidualization markers *IGFBP1* and *PRL* (**Figure 6A**). Successful knockdown of individual
302 *RAR* genes were confirmed by qRT-PCR (**Figure 6B**). Interestingly, siRNA against *RARA* not
303 only reduced *RARA* expression by more than 80%, but also simultaneously resulted in significant
304 increases in *RARB* and *RARG* expression, likely due to a compensatory signaling feedback loop
305 (**Figure 6B**). By contrast, knocking down *RARB* or *RARG* did not affect hESC decidualization,
306 nor did it elicit significant changes in the expression of other *RAR* genes (**Figure 6A, B**).

307
308 To further dissect the involvement of RAR genes in implantation, we evaluated the expression of
309 RA receptor genes by qRT-PCR in both mouse uterus during the peri-implantation period and
310 decidualized hESCs. As shown in **Figure S5**, in both model systems, *Rara/RARA* and
311 *Rarg/RARG* are the most abundant isotypes among the *Rar/RAR* genes, whereas *Rxra/RXRA* and
312 *Rxrb/RXRB* are the predominantly expressed *Rxr/RXR* genes. Hormonal regulation of the
313 receptor genes was also examined in cultured hESCs (**Figure S6**). Twenty-four hours of
314 exposure to MPA elevates *RARA* transcript level, and this effect is augmented by co-treatment of
315 E2+MPA, even though E2 alone does not elicit any changes. *RXRB*, on the other hand, is
316 induced and suppressed by E2 and MPA respectively, and co-treatment appears to counteract
317 each other and cancel out the effect. Expression levels of the other *RAR* and *RXR* genes are not
318 affected by hormone treatment within this time frame.

319

320 To further demonstrate dependency of human decidualization on RA signaling, hESCs were
321 treated with a pan-RAR antagonist, AGN194310 (45), at increasing concentrations. As shown in
322 **Figure 6C**, decidualization markers *IGFBP1* and *PRL* both exhibit a dose-dependent decrease in
323 expression upon drug treatment. Expression of endogenous *RAR* genes, including *RARA*, *RARG*,
324 *RXRA* and *RXRB*, also exhibits dose-dependent reductions in response to AGN194310 (**Figure**
325 **6D**). Taken together, these results strongly support the notion that RAR signaling, particularly
326 through *RARA*, is required for in vitro decidualization of hESCs.

327

328 **Discussion**

329 In the current study, we generated and characterized a mouse model with conditional disruption
330 of RA-RAR signaling specifically in female reproductive organs. The dominant negative
331 *RaraDN* allele used in this study has been previously shown to block endogenous RAR-
332 dependent signaling through competitively binding to RAREs (22). The vast majority of females
333 carrying only one copy of the *RaraDN* allele in *Pgr*-cre-expressing cells are sterile, whereas one
334 is severely subfertile, due to defective uterine receptivity and decidualization. Given that the
335 dominant negative receptor blocks RAR signaling in a dose-dependent manner, and that having
336 two alleles of *RaraDN* completely abolishes endogenous RA signaling (22), we expect the
337 detrimental effects on female fertility would be more severe in *Pgr*-cre; *RaraDN*^{*fllox/fllox*} females.
338 Our findings also indicate that RA signaling through PPAR β/γ and/or non-genomic pathways
339 cannot compensate for the loss of RAR signaling during implantation.

340

341 Even though *Pgr-cre* also mediates *Rara*DN expression in the ovary, there is no indication that
342 the mutant ovaries are affected. Not only do ovarian hormone levels remain unchanged in these
343 mutants, ovulation and fertilization also occur normally. Consistent with our data, genetic
344 ablation of all three *Rar* genes as well as that of all three RA synthesis enzymes (*Aldh1a1-3*) in
345 the developing mouse ovary, does not affect ovary differentiation or ovarian function (46). RAR
346 signaling endogenous to the embryos also does not appear to be required for uterine receptivity
347 or decidualization, as transgenic embryos carrying the *Rara*DN allele driven by an SV40 early
348 promoter implant and develop to term when transferred into wild-type recipient dams (47).
349
350 In the absence of ligand, RAR/RXR heterodimers can actively repress target genes by occupying
351 RAREs and complexing with corepressor proteins, such as nuclear-receptor corepressor (NCoR)
352 and silencing mediator of retinoic acid (SMRT), to prevent transcription (48-50). Presence of RA
353 induces conformational changes in the ligand-binding domain of RARs, resulting in
354 simultaneous attenuation of affinity for co-repressors and increased affinity for co-activators,
355 including histone acetyltransferases (HATs) and DRIP/TRAP/ARC coactivators and other
356 mediator-containing complexes, to decompress chromatin and transactivate target genes (51-53).
357 Rapid repression of target genes upon RA signaling activation has also been reported extensively
358 (54-56), however the molecular mechanism is less studied. It is believed that liganded
359 heterodimers recruit polycomb repressive complex 2 (PRC2), HDAC and co-regulator(s) to
360 actively inhibit target gene transcription, but the identity of the co-regulator(s) remains unknown.
361 Even though the three RAR genes share extensive homology and in many cases function
362 redundantly, unliganded heterodimers RXR/RAR β and RXR/RAR γ interact with SMRT co-
363 repressors differently from unliganded heterodimers of RXR/RAR α by mediating a substantial

364 level of transactivation rather than repression (57). The *Rara*DN mutant receptor used in this
365 study lacks the carboxyl terminal sequence of the human *RARA* gene, but is also highly efficient
366 at inhibiting the other two receptors (47). Dose-dependent blocking of transcription activation by
367 this receptor has been demonstrated in various RARE-reporters both in vitro and in vivo (22, 47,
368 58), however, little is known about its impact on relieving repression or active inhibition of
369 target genes. In the current study, we identified genes that are activated or repressed during the
370 peri-implantation window in *Rara*DN^{Pgr} uterus, suggesting both instructive and permissive roles
371 of RAR signaling. Whether these genes are direct transcriptional targets of RAR signaling, or
372 their expression reflects a manifestation of changes in a cohort of “master RAR targets”,
373 demands further investigation.

374

375 In this study, we reported that the receptivity and decidualization defects in the *Rara*DN^{Pgr} uterus
376 were partially caused by loss of *Fst* expression. FST, also known as activin-binding protein, is a
377 glycoprotein that regulates of TGF- β superfamily signaling, primarily through binding to activin
378 (59). Activin B, homodimer of Inhibin β B, binds to and activates ACVR2A/B and ALK4/7 and
379 in turn phosphorylates SMAD2/3. *Fst* is upregulated during peri-implantation in the mouse
380 uterus, which is believed to sequester activin B in order to allow BMP signaling activation (43).
381 Genetic ablation of *Fst* leads to severe female subfertility in mice with receptivity and
382 decidualization defects similar to *Rara*DN^{Pgr} mice (43), and aberrant expression of FST and
383 activins are associated with poor pregnancy outcome in IVF patients (60). In *Rara*DN^{Pgr} mice,
384 greatly reduced *Fst* expression was accompanied by increased *Inhbb* expression at 3.5 dpc and
385 loss of *Bmp2* induction at 4.5 dpc. Interestingly the loss of *Bmp2* and deregulation of other
386 decidualization markers were partially rescued by supplementation of FST protein in isolated

387 mutant endometrial stromal cells, in organ culture, as well as in vivo, suggesting that loss of FST
388 can largely account for the severity of decidualization defects in *Rara*DN^{Pgr} mice. However, *Fst*
389 down-regulation is unlikely to be the sole reason for the mutant impaired fertility for three
390 reasons. First, gene expression changes in the *Rara*DN^{Pgr} mutants including a wide array of
391 known RA targets were evident that have no known link to the FST and activin signaling
392 pathway. Second, the fertility defects, especially in terms of decidualization, are much more
393 severe in the *Rara*DN^{Pgr} mutants than in the *Fst*-cKO mice. Finally, not all decidualization genes
394 tested were rescued by FST supplementation, e.g. *Prl*. Thus, while FST is an important
395 downstream component of decidual RAR signaling, and while it is likely a direct RAR
396 transcriptional target (42), our findings suggest that a wider network of signaling pathways is at
397 play. Interestingly, regulation of BMPs by RAR signaling has been reported in other cellular
398 contexts. In the mouse testicular embryonal carcinoma cell line, RA induces *Bmp2*, while
399 simultaneously repressing *Bmp4*, specifically through RAR α and γ (61). In primary bone marrow
400 stromal cultures, addition of retinaldehyde stimulates *Bmp2* expression, and this induction is
401 dampened by co-treatment of RAR antagonist AGN193109 (62). Whether the regulation of
402 BMPs by RAR signaling is also mediated by FST in these specific cell types is not clear.

403

404 In addition to the mouse data, we also demonstrated the requirement of RAR signaling,
405 specifically through RAR α , in hESC decidualization. Knocking-down *RARA* in hESCs resulted
406 in significant down-regulation of decidualization markers, as well as elevation of *RARB/G*,
407 possibly by a compensatory mechanism in response to loss of RAR α signaling. Knockdown of
408 *RARB/G*, on the other hand, had no detectable effects on human in vitro decidualization.
409 Treatment of a pan-RAR antagonist also caused a dose-dependent reduction of decidualization

410 marker expression, as well the expression of major *RAR/RXR* genes. Infertility due to vitamin A
411 deficiency has been reported in humans, and fertility was restored after carefully titrated
412 supplementation of Vitamin A back to normal levels (63). A review of the vitamin A content of
413 the top 25 best-selling prenatal vitamins at the USA's top-grossing online store (amazon.com)
414 revealed that the percent daily value for pregnant and nursing individuals ranges from 0% to
415 185% from a variety of precursors, with the topmost best-seller having no vitamin A (**Table S1**).
416 Extreme excess maternal vitamin A is a documented teratogen, although this is shown to be
417 largely from feedback inhibition of native retinoic acid production in developing embryos (64).
418 Low maternal vitamin A intake can likewise cause birth defects such as diaphragmatic hernia
419 (65). The current study adds to the existing body of data to emphasize that not only is it essential
420 for maternal/fetal health to have biologically appropriate levels of maternal RA, but it is also
421 crucial for uterine receptivity and decidualization to have proper RA receptor signaling, as
422 shown herein in mice in vivo and in hESCs in vitro.

423

424 **Materials and Methods**

425 **Mice**

426 Generation of mice carrying *Rara*DN preceded by a floxed transcriptional/translational STOP
427 sequence was described previously (22), and cryopreserved sperms from mutant mice were
428 provided by Dr. Benjamin D Humphreys in the Division of Nephrology, Washington University
429 School of Medicine. Live mice carrying the *Rara*DN mutation were rederived via in vitro
430 fertilization (IVF) at the Mouse Genetics Core at Washington University. Pgr-Cre line was
431 provided by Dr. Francesco DeMayo at the National Institute of Environmental Health Sciences
432 (23) and were mated to *Rara*DN^{f/+} mice to generate offspring carrying both alleles (hereinafter

433 referred to as *Rara*DN^{Pgr}) and littermate controls (*Rara*DN^{f/+}, CTRL). Artificial decidualization
434 and tail vein injection were performed following standard procedures as described previously
435 (35, 66, 67). All mice used in this study were maintained in a barrier facility at Washington
436 University School of Medicine, Missouri, following the institution's regulations with an
437 approved protocol.

438

439 **Uterine Stromal Cell Isolation and Organ Culture**

440 Uteri of 2.5 dpc mice were collected, rinsed in cold Hank's Balanced Salt Solution (HBSS,
441 Gibco), cut into 2-3mm pieces, and digested in 1% trypsin (Sigma) in HBSS for 1 hour at RT
442 with gentle shaking. After incubation, luminal epithelium of each uterine segment was gently
443 squeezed out using fine forceps along the longitudinal axis of the uterus. The remaining uterine
444 tissues were transferred to a fresh tube, further digested in 0.25% trypsin and 1 mg/ml
445 collagenase (Sigma) in HBSS for 30 min at 37 °C with gentle shaking and dissociated by
446 pipetting several times following incubation. Cell suspension and tissue remnants were filtered
447 through a 70 µm nylon filter, and stromal cells were resuspended in hESC culture media (phenol
448 red-free DMEM/F12, 7.5% charcoal-stripped FBS, 1x non-essential amino acids, 1x antibiotic-
449 antimycotic). Stromal cells were seeded in 12-well plates, and follistatin (Sino Biological US
450 Inc) were added to the culture media at indicated concentrations. For organ culture, 2.5 dpc uteri
451 were cut into 2-3mm segments and placed on the membrane of multiwall inserts. The inserts
452 were then placed into 12-well culture plates containing 0.5 ml culture medium, and ten agarose
453 beads soaked in 100 ng/µl follistatin (Sino Biological US Inc., PA) or BSA were transferred into
454 the lumen of each segment (68). Both primary stromal cells and uterine organ cultures were
455 harvested 48 hours after for RNA isolation.

456

457 **Human Endometrial Stromal Cell (hESC) Culture**

458 Immortalized hESCs were characterized previously (69) and purchased from ATCC (the
459 American Type Culture Collection, #CRL-4003). hESCs were maintained in hESC culture media
460 plus 1x Insulin/Transferrin/Selenium (ITS, Gibco) in a humidified 37°C incubator supplied with
461 5% CO₂. To induce decidualization in vitro, culture medium was replaced with induction
462 medium (hESC medium plus 36 nM 17β-estradiol, 1 μM medroxyprogesterone/MPA, 0.1 mM
463 cAMP) and hESCs were allowed to decidualize for 96 hours before RNA extraction. Gene-
464 knockdown experiments were performed using DharmaFECT4 transfection reagent (GE
465 Healthcare Dharmacon, Inc.) and Silencer™ Select Validated siRNAs (ThermoFisher, Cat#
466 4427038, *RARA*, siRNA ID# s11801; *RARB*, siRNA ID# s534565; *RARG*, siRNA ID# s11807)
467 following manufacturer's instructions. The morning following transfection, 10x induction
468 cocktail topper was added to the culture to a final concentration of 1x and cells were cultured for
469 an additional 72 hours before harvest. For RAR-antagonist treatment, hESCs were treated with
470 induction medium with or without AGN194310 (Sigma) at indicated concentrations for 72 hours,
471 before harvest for RNA extraction.

472

473 **RNA Isolation and Real-time RT-PCR**

474 RNA was extracted in RNA STAT-60 reagent following manufacturer's instructions (Tel Test
475 Inc). Reverse transcription was performed using the High Capacity cDNA Reverse Transcription
476 kit (Applied Biosystems Inc., ABI), and qPCR performed on ViiA 7 Real-Time PCR System
477 (ABI) using PowerUp™ SYBR™ Green Master Mix (ABI). All results were repeated in three

478 biological replicates unless specified and relative gene expression changes were determined by
479 delta-delta Ct method (normalized to housekeeping gene, *Rpl7*). Primers are listed in **Table S2**.

480

481 **Histology, Immunofluorescence (IF) and Alkaline Phosphatase (AP) Activity Assay**

482 Tissue fixed in Bouin's fixative were processed for serial dehydration and embedding at the
483 Developmental Biology Histology Core at Washington University School of Medicine. Eight
484 micron paraffin sections were used for hematoxylin-eosin (H&E) staining and
485 immunofluorescence (70). All antibodies were used at 1:1000 dilution in blocking solution (1%
486 BSA, 3% normal goat serum in PBS): CDH1 (BD biosciences, San Jose, CA), phospho-Histone
487 H3 (Millipore), Alexa594 goat anti-rabbit and Alexa488 goat anti-mouse (Life Technologies
488 Corp., Carlsbad, CA). Paraformaldehyde (PFA)-fixed paraffin-embedded sections were used for
489 AP activity assay. The sections were dewaxed, rehydrated and washed in PBS. Rehydrated
490 sections were subsequently incubated in freshly-prepared AP staining solution containing 0.33
491 mg/ml NBT (nitro blue tetrazolium, Roche), 0.165 mg/ml BCIP (5-bromo-4-chloro-3-indolyl-
492 phosphate, Roche) in AP buffer (100mM Tris-Cl, pH 9.0, 150mM NaCl, 1mM MgCl₂) for color
493 development. Antibody catalog numbers are listed in **Table S3**.

494

495 **Ovarian Hormone Analyses**

496 Whole blood was collected by cardiac puncture and allowed to coagulate in 1.5ml Eppendorf
497 tubes at room temperature for 20 minutes. The blood samples were centrifuged at 3000 rpm for
498 10 min at 4°C, and supernatant (serum) aliquoted and stored at -80°C until use. Cayman
499 Chemical ELISA kits for detection of E2 (No. 501890) and P4 (No. 582601) were used to
500 determine serum hormone levels, following manufacturer's instructions. Plates were read on a

501 Bio-Rad 3550 microplate reader at the wavelength of 405nm, data processed in Microsoft Excel
502 and visualized in Graphpad Prism. Four biological replicates were tested and presented for each
503 genotype.

504

505 **In Situ Hybridization**

506 In Situ hybridization was performed on PFA-fixed paraffin-embedded 8 μ m tissue sections using
507 RNAscope® 2.5 HD Assay-RED kit (Advanced Cell Diagnostics, ACD, Newark, CA). Gene-
508 specific double-“Z” oligo probes compatible with the kit were ordered from ACD (probe-Mm-
509 *Bmp2*, Cat# 406661; probe-Mm-*Fst*, Cat# 454331) and detailed in situ procedure has been
510 described previously (35).

511

512 **Statistical Analysis**

513 All experimental groups contained three biological replicates, if not specified otherwise. Two-
514 tailed Student’s *t*-test assuming unequal variance was performed to compare means of the
515 experimental groups. For dose response experiments, one-way ANOVA with post-hoc Tukey
516 HSD test was performed. Data are presented as mean \pm SD, with raw individual experimental
517 data displayed as dot plots overlay, and *p*-value less than 0.05 was considered statistically
518 significant.

519

520 **Study Approval**

521 The animal studies included herein were reviewed and approved by the Institutional Animal Care
522 and Use Committee of Washington University in St. Louis, Missouri, USA. All studies were

523 performed to the current standards of the American Association for Laboratory Animal Science
524 so as to minimize pain, suffering, and total animals necessary for conclusive findings.

525 **Author contributions**

526 YY, MH, and LM designed the study. YY, MH, and SBC conducted experiments. YY, MH,
527 SBC, RK, and LM analyzed data and wrote the manuscript.

528

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536

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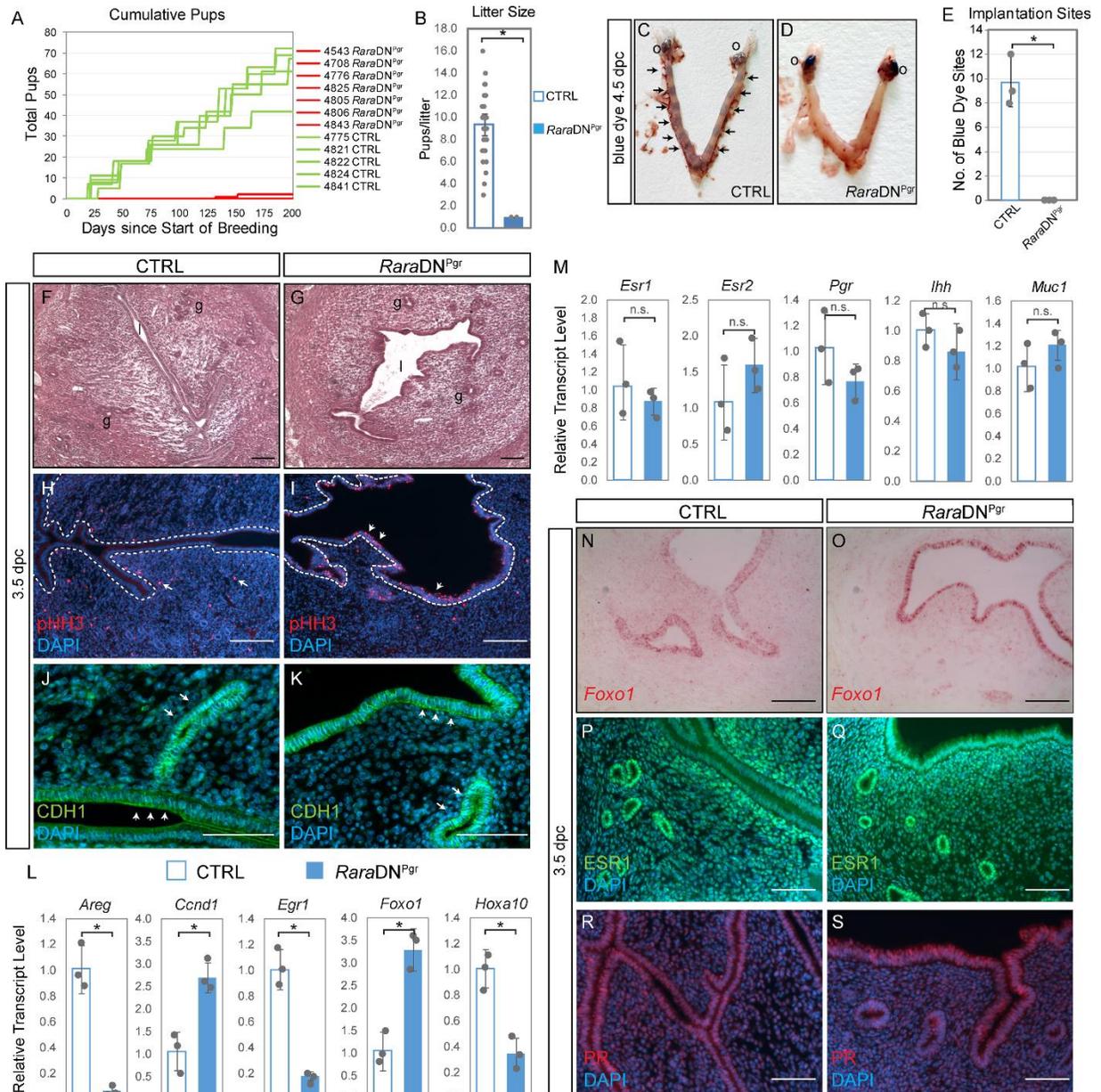
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735
- 736

737 **Table 1. 200-day Breeding Record**

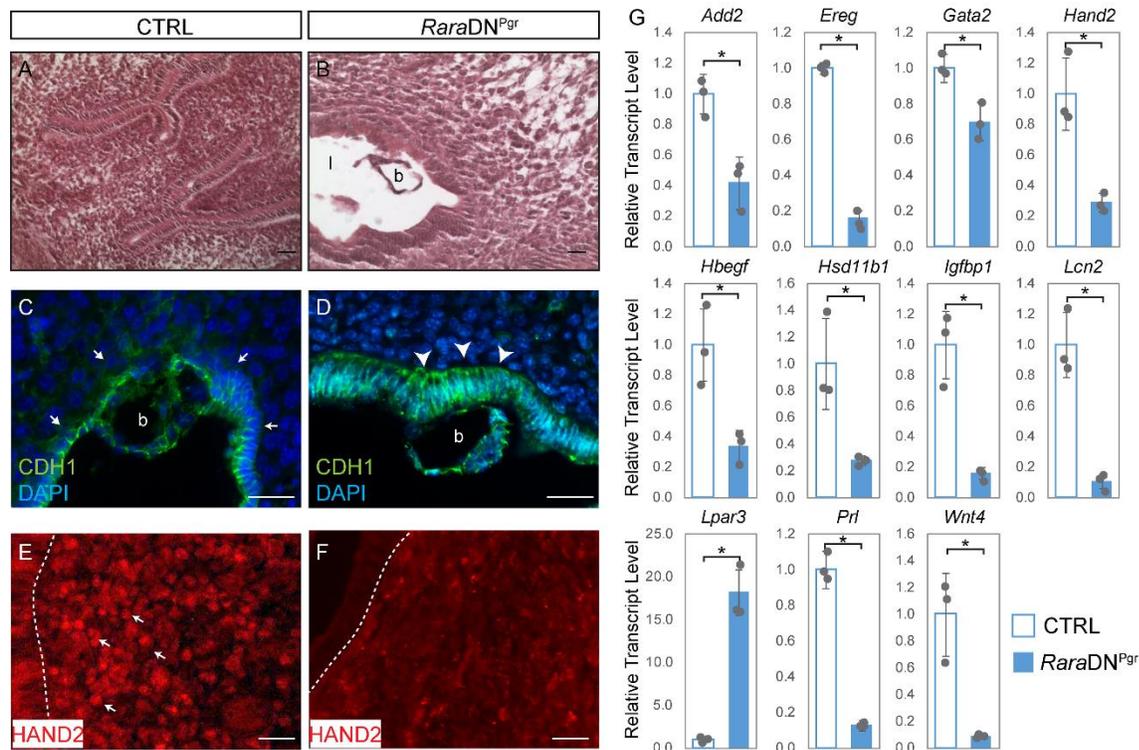
Genotype	Number of ♀ tested	Cumulative pups - total	Cumulative pups/female	Cumulative litters - total	Cumulative litters/female
CTRL	5	311	62.2	34	6.8
<i>RaraDN^{Pgr}</i>	7	2	0.286	2	0.286

738

739 **Figure Legends**

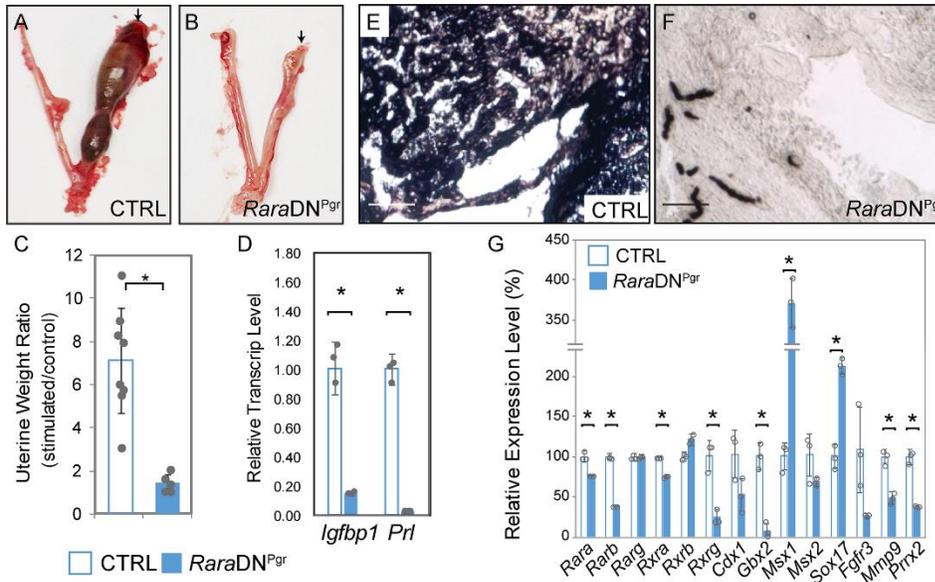


742 Cumulative number of pups produced by individual CTRL (green) and *RaraDN^{Pgr}* females (red)
743 over the period of 200 days. (B) Average number of pups per litter produced by the females
744 presented in A. (C, D) Representative images of visualization of implantation sites (arrows) by
745 Blue Dye injection at 4.5 dpc. O, ovaries. (E) Quantification of implantation sites from CTRL
746 (9.7 ± 2.1 , $n=3$) and *RaraDN^{Pgr}* females (0.0 ± 0.0 , $n=3$, $p=0.0013$). (F, G) H&E staining of
747 *RaraDN^{Pgr}* and CTRL uteri at 3.5 dpc. g, glands; l, lumen. (H, I) Immunofluorescence detecting
748 pHH3-positive proliferating cells at 3.5 dpc (arrowheads, uterine epithelial cells; arrows,
749 proliferating stromal cells; dashed lines outline the luminal epithelia). (J, K) CDH1
750 immunofluorescence on 3.5 dpc uterine sections. Note in the CTRL, reduced CDH1 level is
751 obvious in the luminal epithelium (J, arrowheads) when compared to glandular epithelium (J,
752 arrows); whereas this difference is negligible in *RaraDN^{Pgr}* uterus (K). (L, M) Gene expression
753 at 3.5 dpc determined by qRT-PCR, normalized to levels of housekeeping gene *Rpl7* and the
754 average transcript level of CTRL samples was set to one. (N, O) RNAscope in situ hybridization
755 of *Foxo1* showing elevated transcript level in the mutant epithelium. (P-S) IF staining of ESR1
756 (P,Q) and PR (R,S) revealed no apparent differences in expression. *, $p < 0.05$; n.s. not
757 significant. Scale bars: 50 μm .
758



759
 760 **Figure 2.** Failure of decidualization in 4.5 dpc *RaraDN^{Pgr}* uterus. (A, B) H&E staining of
 761 *RaraDN^{Pgr}* and CTRL uteri at 4.5dpc. l, lumen; b, blastocyst. (C, D) CDH1 immunofluorescence
 762 on 4.5dpc uterine sections. Note the presence of normal blastocyst in the mutant lumen (D), and
 763 persistent high CDH1 level in the underlying luminal epithelium especially at the basal side
 764 (arrowheads, compared to CTRL arrows in C). (E, F) Immunofluorescent detection of HAND2
 765 protein in the nuclei of decidual cells in the CTRL (arrows, E), which is absent in the mutant (F).
 766 (G) Relative transcript levels of genes involved in decidualization by qRT-PCR. Scale bars:
 767 50 μ m.

768



769

770 **Figure 3.** *RaraDN^{Pgr}* mutant uterus does not respond to artificial decidualization stimuli in vivo.

771 (A, B) Representative images of artificially decidualized uteri five days after stimulation. Arrows

772 indicate the uterine horn that received intrauterine oil infusion; contralateral horns serve as

773 controls. (C) Uterine weight ratio (wet weight stimulated/wet weight unstimulated) was

774 calculated for each animal, and graphed as mean \pm SD (CTRL, 7.1 ± 2.4 , $n=8$; *RaraDN^{Pgr}*, $1.4 \pm$

775 2.4 , $n=5$; $p=0.00036$). (D) Gene expression data of decidualization markers, *Igfbp1* and *Prl*, in

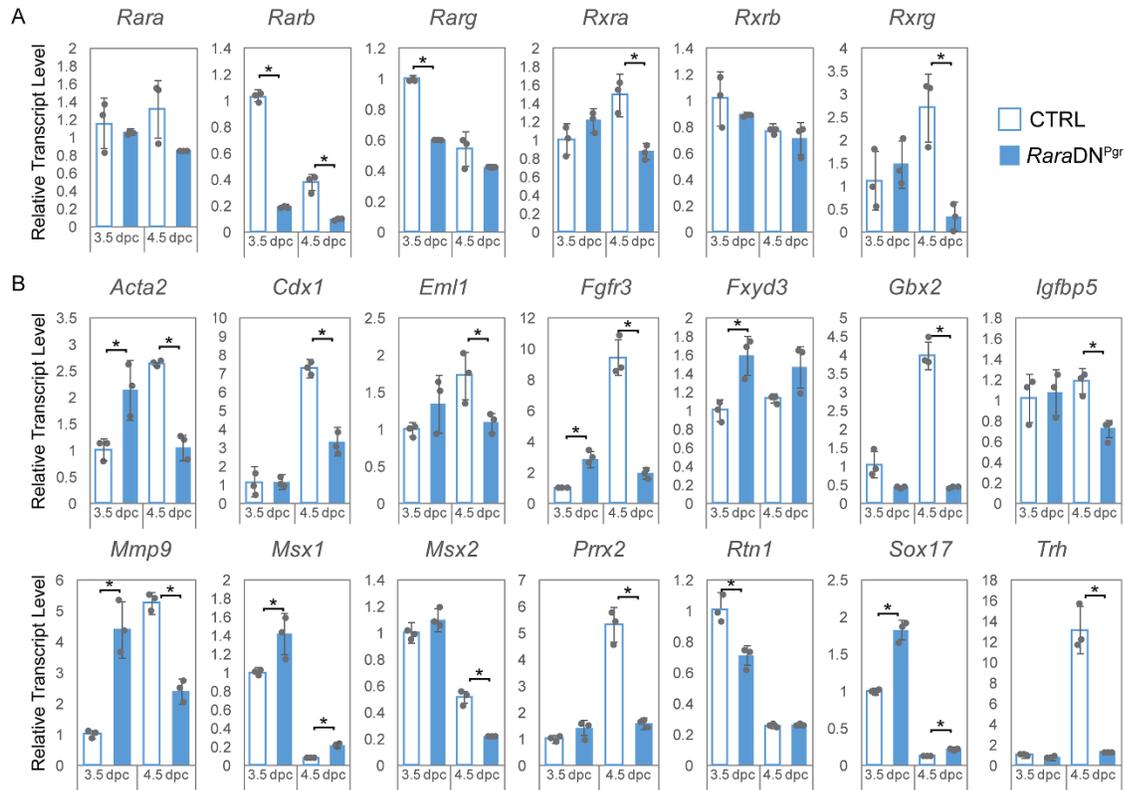
776 the stimulated uterine horn by qRT-PCR. (E, F) Alkaline phosphatase activity in the stimulated

777 uterine horns visualized by dark color development from AP substrate BCIP/NBT. (G) Gene

778 expression analyses of RA receptors and RAR targets comparing RNA extracted from stimulated

779 CTRL and *RaraDN^{Pgr}* uteri by qRT-PCR. Asterisks indicate $p < 0.05$. Scale bars: 50 μ m.

780



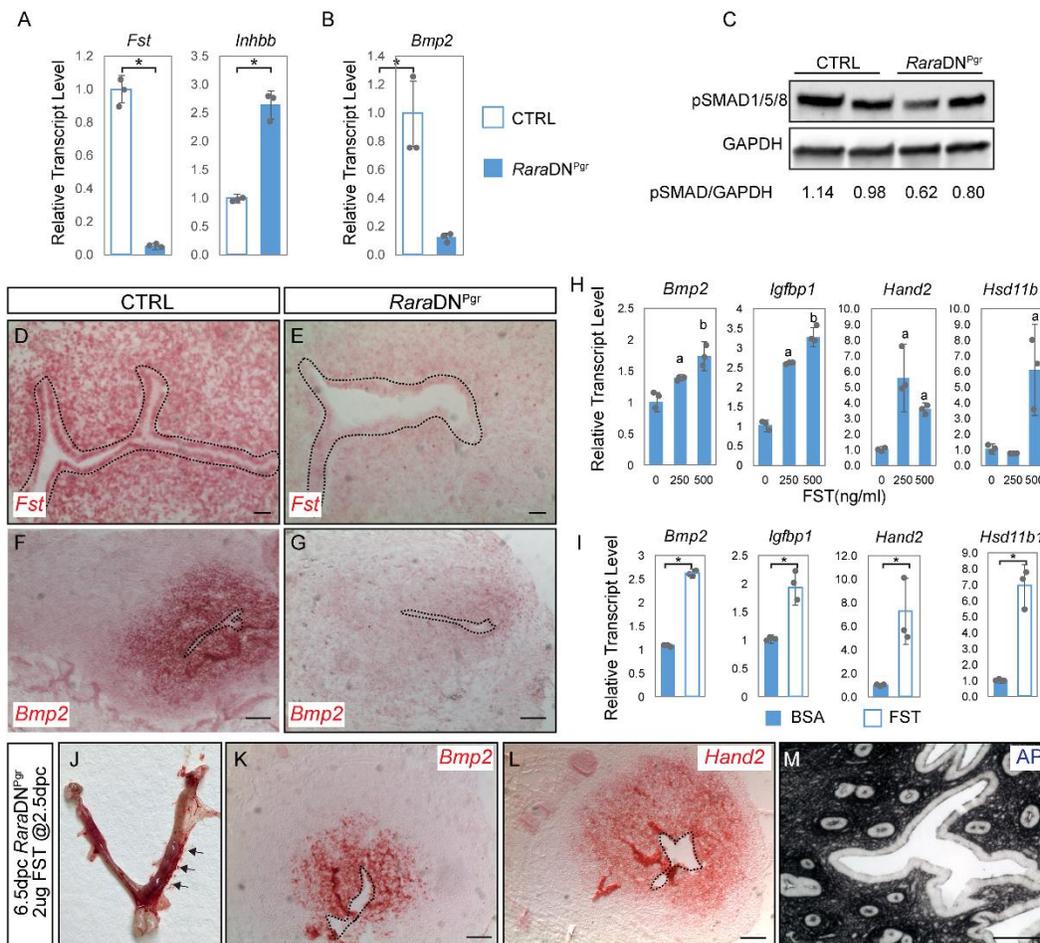
781

782 **Figure 4.** Changes in expression of RA receptor genes and known RAR targets in the *RaraDN^{Pgr}*

783 uterus during the peri-implantation period. Gene expression by qRT-PCR performed on whole

784 uterine RNA extract at 3.5 and 4.5 dpc detecting RA receptors (A) and targets (B). Asterisks

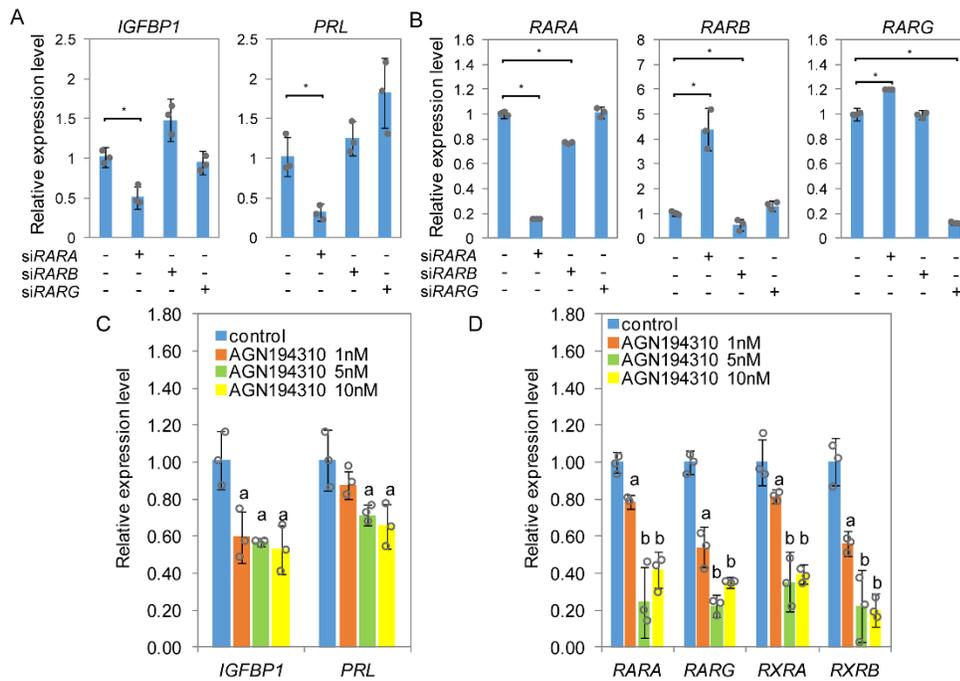
785 indicate $p < 0.05$ comparing to CTRL uteri of same timepoint.



786

787 **Figure 5.** Reduced follistatin expression and downstream changes in activin/BMP signaling are
 788 partially responsible for the fertility defects in *RaraDN^{Pgr}* uterus. (A, B) Relative expression
 789 levels of *Fst* and *Inhbb* at 3.5 dpc (A), and *Bmp2* at 4.5 dpc (B). (C) Western blot for phospho-
 790 Smad1/5/8 of whole uteri extract from CTRL and *RaraDN^{Pgr}* females at 3.5 dpc. WB band
 791 density was quantified in ImageJ and the relative density calculated as the ratio of
 792 pSMAD/GAPDH for each sample was listed. (D-G) RNAscope in situ hybridization of *Fst* at 3.5
 793 dpc (D, E) and *Bmp2* at 4.5 dpc (F, G). Positive results manifest as red staining; dotted lines
 794 outline the luminal epithelium. (H) Gene expression of decidualization markers in isolated
 795 endometrial stromal cells from 2.5 dpc *RaraDN^{Pgr}* uteri treated with recombinant mouse FST at
 796 indicated concentrations for 48 hours (n=2). (I) Gene expression of decidualization markers in

797 uterine segments dissected from 2.5 dpc *Rara*DN^{Pgr} and incubated with luminal agarose beads
 798 soaked with BSA or FST for two days (n=2). (J) Appearance of *Rara*DN^{Pgr} mutant uteri four
 799 days after receiving 2 µg FST via tail vein injection (arrows point to bulging regions resembling
 800 implantation sites). (K-L) RNAscope in situ hybridization of *Bmp2* (K) and *Hand2* (L) showed
 801 elevated expression in the bulging region. (M) BCIP-NBP staining of bulging regions shows
 802 extensive alkaline phosphatase activity. Asterisks indicate $p < 0.05$ by *t*-test; a indicates $p < 0.05$
 803 by one-way ANOVA between drug group and control group; and b indicates $p < 0.05$ by one-way
 804 ANOVA between different doses. Scale bars: 50µm.



805
 806 **Figure 6.** RAR signaling is essential for decidualization of hESCs. (A, B) Gene expression by
 807 qRT-PCR of human decidualization markers (A) and RAR genes (B) when individual RAR
 808 genes are silenced by siRNA. (C, D) Expression of decidualization markers (C) and selective
 809 RAR and RXR genes (D) when hESCs are treated with pan-RAR antagonist AGN194310 at
 810 indicated concentrations. Asterisks indicate $p < 0.05$ by *t*-test; a indicates $p < 0.05$ by one-way

811 ANOVA between drug group and control group; and b indicates $p < 0.05$ by one-way ANOVA

812 between different doses.

813