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Expression and Function of Macrophage-Inducible C-type Lectin (Mincle) in Inflammation Driven Parturition in Fetal Membranes and Myometrium

Short title: Mincle and labor

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1 **ABSTRACT**

2

3 The pivotal role of inflammatory processes in human parturition is well known, but not completely
4 understood. We have performed a study to examine the role of Macrophage-inducible C-type lectin
5 (Mincle) in inflammation-associated parturition. Using human samples, we show that spontaneous
6 labor is associated with upregulated Mincle expression in the myometrium and fetal membranes.
7 Mincle expression was also increased in fetal membranes and myometrium in the presence of pro-
8 labor mediators, the pro-inflammatory cytokines IL1B and TNF, and toll-like receptor (TLR)
9 ligands fsl-1, poly(I:C), LPS and flagellin. These clinical studies are supported by mouse studies,
10 where an inflammatory challenge in a mouse model of preterm birth increased Mincle expression in
11 the uterus. Importantly, elimination of Mincle decreased the effectiveness of pro-inflammatory
12 cytokines and TLR ligands to induce the expression of pro-labor mediators, namely pro-
13 inflammatory cytokines and chemokines, contraction-associated proteins and prostaglandins, and
14 extracellular matrix remodelling enzymes, matrix metalloproteinases. The data presented in this
15 study suggests that Mincle is required when inflammatory activation precipitates parturition.

16

17 **Keywords:** Mincle; human labor; myometrium; fetal membranes; inflammation

18 **INTRODUCTION**

19

20 Pattern recognition receptors (PRRs) are located on cell surfaces and recognise pathogen-associated
21 molecular patterns (PAMPs) and damage-associated molecular patterns (DAMPs) such as proteins,
22 nucleic acids, carbohydrates and lipids. PAMPs and DAMPs bind to PRRs to activate signal
23 transduction pathways resulting in activation of multiple signalling pathways including NF-κB.
24 Activation of these pathways leads to transcription of pro-inflammatory cytokines and chemokines,
25 which in turn attract leukocytes such as neutrophils to the site of pathogen invasion or tissue
26 damage. There are four main classes of PRRs; toll-like receptors (TLRs), NOD-like receptors
27 (NLRs), RIG-I-like receptors (RLRs) and C-type lectin receptors (CLRs). TLRs and NLRs have
28 been shown to play a role in regulating the inflammatory processes associated with human labor (1-
29 8). There are, however, no studies on the expression and function of CLRs in parturition.

30

31 CLRs are a large family of proteins that are divided into 17 groups based on functional and
32 structural characteristics. They contain an extracellular recognition domain that recognises a wide
33 range of microbial and fungal carbohydrates and lipids (9-11). CLRs also contain a signal
34 transmembrane domain and a cytoplasm domain. Macrophage-inducible C-type lectin (Mincle;
35 gene name CLEC4E) is a CLR that was first identified in 1999 as a downstream target of the

36 transcription factor NF-IL6 in macrophages (12). Mincle expression is strongly induced in response
37 to several inflammatory stimuli and stresses such as LPS, TNF, IL6 and saturated fatty acids (12-
38 15) and in various diseases characterised by increased inflammation including obesity, rheumatoid
39 arthritis, allergic contact dermatitis, ischemic stroke, traumatic brain injury, and hepatitis (13, 16-
40 20).

41
42 Mincle binding to its ligands initiates a cascade of events that leads to the recruitment of spleen
43 tyrosine kinase (Syk) which activates a cascade of signalling events that leads to the induction of
44 inflammatory genes via NF- κ B, mitogen activated protein kinase (MAPK), activator protein 1 (AP-
45 1) or nuclear factor of activated T-cells (NFAT) (21-27). Numerous studies have implicated Mincle
46 in regulating inflammation. Mincle deletion, blockade or pharmacologic inhibition attenuates the
47 inflammatory response in vivo (13-15, 18, 20, 28) and in vitro (14, 17, 18, 25).

48
49 Therefore, we hypothesised that Mincle expression would be upregulated during labor and with
50 infection in fetal membranes and myometrium. Furthermore, we hypothesised that pro-
51 inflammatory cytokines and TLR ligands would induce Mincle expression leading to upregulated
52 expression and secretion of mediators involved in the processes of labor. To examine this, we (i)
53 characterised the expression of Mincle in human myometrium and fetal membranes from laboring
54 and non-laboring women at term and preterm; (ii) investigated the effect of pro-inflammatory
55 cytokines, and bacterial and viral products on Mincle expression in fetal membranes and
56 myometrium; and (iii) determined the effect of Mincle siRNA knockdown on inflammation induced
57 pro-labor mediators in human primary myometrial and amnion cells.

58 **METHODS**

59

60 **Tissue collection**

61 Placenta (with attached fetal membranes) and myometrium was collected at the time of delivery
62 from women delivering a singleton baby. Written informed consent was obtained from all women.
63 Myometrium was obtained from the upper margin of the lower uterine segment incision during
64 Caesarean section. Tissue samples were obtained from women who delivered healthy, singleton
65 infants. Criteria for recruited patients were as we have previously described (29) and excluded
66 women with any underlying medical conditions (e.g. diabetes, asthma, preeclampsia).

67

68 Tissues were processed within 15 mins of delivery as previously described (30, 31). For the Mincle
69 expression studies, placenta and myometrium was obtained from laboring and non laboring women
70 at term or preterm as detailed below. Tissue samples were immediately snap frozen in liquid

71 nitrogen and stored at -80°C until processed by RT-qPCR as detailed below. For the tissue explant
72 and cell culture experiments, placenta and myometrium was obtained from non laboring women at
73 the time of term Caesarean section and used immediately as detailed below.

74

75 For the Mincle expression studies, myometrium was obtained from women at term Caesarean
76 section (i) in the absence of labor or (ii) during active spontaneous labor (n=8 patients per group) as
77 we have previously described (29). Labor was defined as the presence of regular uterine
78 contractions (every 3–4 min) resulting in cervical effacement and dilation. Fetal membranes
79 (amnion and choriodecidua) were obtained from women at preterm or term (i) undergoing elective
80 Caesarean section in the absence of labor or (ii) after spontaneous labor and vaginal delivery (n=9
81 patients per group) as we have previously described (29). Fetal membranes from the non-laboring
82 group were obtained from the area overlying the cervix (i.e. supracervical site, SCS) and from the
83 site of membrane rupture in the after labor group as previously described (32). For the term
84 samples, indications for Caesarean section in the absence of labor were breech presentation and/or
85 previous Caesarean section. Indications for Caesarean section in the term laboring samples were for
86 fetal malpresentation, fetal distress and delayed or failure to progress. Indications for preterm
87 delivery (in the absence of labor) were for placenta praevia, vasa praevia, placental abruption or
88 antepartum haemorrhage. None of the patients received any medications to augment or induce
89 labor, had pre-labor rupture of membranes, or any clinical signs of infection. The relevant clinical
90 characteristics of the patients used for these studies are detailed in Supplementary Tables 1 and 2.

91

92 To characterise infection associated changes in Mincle expression, amnion was collected from
93 women undergoing Caesarean section in the absence of labor (i) with histologically confirmed acute
94 chorioamnionitis or (ii) without histologically confirmed acute chorioamnionitis (n=8 patients per
95 group) as previously described (33). As previously described (33), choriodecidual tissue could not
96 be collected from the samples with histologically confirmed acute chorioamnionitis as it was
97 degraded. The relevant clinical characteristics of the patients used for these studies are detailed in
98 Supplementary Table 3.

99

100 **Tissue Explants**

101 To elucidate the effect of pro-inflammatory insults on Mincle expression, tissue explants were
102 performed as previously described (34). Briefly, fresh fetal membranes and myometrium (n=6
103 patients) were dissected, equilibrated in a humidified atmosphere of 8% O_2 (fetal membranes) or
104 21% O_2 (myometrium) for 1 h, and then incubated in DMEM containing 100 U/ml penicillin G and
105 100 $\mu\text{g}/\text{ml}$ streptomycin (50 mg tissue/ml) with or without 10 ng/ml IL1B (PeproTech; Rocky Hill,

106 NJ, USA), 10 ng/ml TNF (PeproTech; Rocky Hill, NJ, USA), 250 ng/ml fsl-1 (InVivoGen; San
107 Diego, California, USA), 20 µg/ml poly(I:C) (Sigma-Aldrich; St. Louis, MO, USA), 10 µg/ml LPS
108 (derived from *E. coli* strain 026:B6; Sigma-Aldrich; St. Louis, MO) or 1 µg/ml flagellin (purified
109 flagellin from *Salmonella typhimurium*; InVivoGen; San Diego, California, USA). After 20 h,
110 tissues were collected and stored at -80°C until assayed for Mincle mRNA expression by RT-qPCR
111 as detailed below.

112

113 **Primary myometrial and amnion epithelial and mesenchymal cell culture**

114 In order to elucidate if Mincle regulates pro-inflammatory mediators, we performed siRNA
115 knockdown experiments in primary myometrial and amnion cells (n=6-7 patients per experiment).
116 Myometrial cells were isolated and cultured as previously described (30). Isolation of amnion
117 epithelial cells and mesenchymal cells was performed as described (35) with minor modifications.
118 Briefly, approximately 3 g of amnion fragments were incubated in 10 ml of DMEM/F-12 with 1%
119 penicillin-streptomycin and 0.25% trypsin for 35 min at 37°C with gentle agitation. After straining
120 through a 100 µm cell strainer, the eluate was neutralised with 1% FBS and remaining fragments
121 were further digested in another 10 ml of DMEM/F-12 with 0.25% trypsin for 35 min at 37°C.
122 Eluate was combined with previous digest and epithelial cells were pelleted by centrifugation at 500
123 x g for 10 min, resuspended in complete media (DMEM/F-12 containing 10% FBS and 1%
124 penicillin-streptomycin) and plated in tissue culture flasks. Flasks were incubated at 37°C, 5% CO₂
125 and 8% O₂ with media changes 4 h after initial plating, then 24-48 h thereafter until confluent. The
126 de-epithelialized amnion fragments were washed extensively in PBS and then incubated in 10 ml
127 DMEM/F-12 containing 1% penicillin-streptomycin and 0.125% collagenase A at 37°C for 1 h with
128 gentle agitation, until fragments had dissolved. After straining, eluate was centrifuged at 500 x g for
129 10 min. Pelleted mesenchymal cells were plated in complete media (DMEM/F-12 containing 10%
130 FBS and 1% penicillin-streptomycin) in tissue culture flasks. Upon confluency (3-5 days), epithelial
131 and mesenchymal cells were trypsinized and plated in 48-well plates in complete media for siRNA
132 transfection experiments.

133

134 Mincle siRNA (siCLEC4E) and negative control siRNA (siCONT) (Ambion; Thermo Fisher
135 Scientific; Scoresby, Vic, Australia) were transfected in amnion and myometrial cells using
136 RNAiMax (Life Technologies; Mulgrave, Victoria, Australia) according to manufacturer's
137 guidelines. Cells were transfected with 50 nM siCLEC4E or 50 nM siCONT for 48 h, followed by
138 treatment with or without 1 ng/ml IL1B, 10 ng/ml TNF, 250 ng/ml fsl-1 or 5 µg/ml poly(I:C) for an
139 additional 20 h. After final incubation, cells and media were collected and stored at -80°C until

140 analysed as detailed below. MTT assay was used to assess cell viability as we have previously
141 described (36).

142

143 **Cell contraction assay**

144 To determine the effect of siCLEC4E on myometrial cell contractility, gel contraction assays were
145 performed as previously described (37). The area of the gel was determined using Image Lab
146 software (Bio-Rad Laboratories, Hercules, CA, USA). Experiments were performed in myometrium
147 obtained from 6 patients.

148

149 **Mice studies**

150 To determine the effect of preterm labor on Mincle mRNA expression, a mouse model was utilised,
151 injecting LPS on gestational day (gd) 15.5 mice and collecting myometrium at delivery of first pup.
152 Animal studies (conducted with approval from the Austin Health's Animal Ethics Committee) were
153 performed as we have previously described (38). Briefly, gd 15.5 mice were injected i.p. with LPS
154 (serotype O26:B6; 15 µg in 50 µl of PBS; Sigma) or sterile PBS (vehicle control). In the LPS group,
155 mice delivered the first pup between 18-22 h post-LPS treatment. None of the vehicle-injected mice
156 went into labor. The mice were killed on the birth of one pup, and time-matched controls were
157 killed directly afterward. Myometrial tissue was washed in PBS, flash frozen and stored at -80°C
158 until further analysis by RT-qPCR as detailed below.

159

160 **RNA extraction and RT-qPCR**

161 RNA extractions, cDNA synthesis and RT-qPCR was performed as previously described (39) using
162 100 nM of pre-designed and validated QuantiTect primers (primer sequences not available)
163 (Qiagen; Chadstone Centre, Vic, Australia). For normalisation of data, the average Ct of two
164 housekeeping gene, YWHAZ and succinate dehydrogenase (SDHA), was used and the fold
165 differences determined using the comparative Ct method.

166

167 **Enzyme immunoassays**

168 Commercial assays were used to measure the levels of IL6, CCL2, CXCL1, CXCL8, sICAM1
169 (R&D Systems; Minneapolis, MN, USA) and PGF_{2α} (Cayman Chemical Company; Ann Arbor, MI,
170 USA) in the incubation media. The interassay and intraassay coefficients of variation for all assays
171 were less than 10%.

172

173 **Gelatin zymography**

174 Incubation media was also collected and assessment of MMP9 was performed by gelatin
175 zymography as previously described (40). Gels were scanned using a ChemiDoc XRS system (Bio-
176 Rad Laboratories; Gladesville, NSW, Australia), inverted, and densitometry performed using
177 Quantity One image analysis software (Bio-Rad Laboratories; Gladesville, NSW, Australia).

178

179 **Statistical analysis**

180 All statistical analyses were undertaken using GraphPad Prism (GraphPad Software, La Jolla, CA,
181 USA). Normality of the data was assessed using the Shapiro-Wilk test. For Figure 1, an unpaired
182 Student's t-test was used to assess statistical significance between normally distributed data,
183 otherwise the Mann-Whitney U test was used. For Figure 2, the Wilcoxon test was used to assess
184 statistical significance. For all other comparisons, data were analysed by a repeated measures one-
185 way ANOVA (with LSD post-hoc testing to discriminate among the means); non-normally
186 distributed data were logarithmically transformed before analysis. Statistical significance was
187 ascribed to a P value ≤ 0.05 . Data is expressed as mean \pm SEM.

188 **RESULTS**

189

190 **Mincle expression is upregulated in laboring fetal membranes and myometrium**

191 The first aim was to determine whether the expression of Mincle mRNA changed with respect to
192 labor. In human fetal membranes, Mincle mRNA expression was significantly increased with term
193 labor (Figure 1A) and preterm labor (Figure 1B), compared to non-laboring tissues. There was also
194 a significant increase in Mincle mRNA expression in amnion collected from preterm women with
195 histological chorioamnionitis (Figure 1C), compared to preterm controls without histological
196 chorioamnionitis. In myometrium, there was a significant increase in Mincle mRNA with human
197 term labor (Figure 1D) compared to non-laboring tissue. To determine the effect of preterm labor on
198 Mincle mRNA expression, a mouse model was utilised, injecting LPS on gestational day 15.5 mice
199 and collecting myometrium at delivery of first pup. Figure 1E demonstrates that Mincle mRNA
200 expression is significantly increased in mouse myometrium with preterm (LPS) labor.

201

202 **Pro-inflammatory mediators increase Mincle expression in fetal membranes and myometrium**

203 To further elucidate the effect of labor on Mincle expression, tissue explants were incubated in the
204 presence of known labor mediators, namely pro-inflammatory cytokines IL1B and TNF, and
205 bacterial and viral products associated with TLR signalling; fsl-1 (TLR2/6), poly(I:C) (TLR3), LPS
206 (TLR4) and flagellin (TLR5). All six treatments significantly increased Mincle mRNA expression
207 in fetal membranes (Figure 2A) and myometrium (Figure 2B).

208

209 **Knockdown of Mincle suppresses pro-inflammatory cytokines and chemokines in human**
210 **primary myometrial and amnion mesenchymal cells *in vitro***

211 Subsequent experiments investigated the effect of Mincle suppression, using siRNA, in the presence
212 of inflammatory, pro-labor stimuli. To determine if Mincle regulates cytokine-induced expression
213 of pro-labor mediators, primary cells isolated from human myometrium and amnion (epithelial and
214 mesenchymal cells), were transfected with Mincle siRNA (siCLEC4E). The efficacy of transfection
215 was assessed by RT-qPCR and there was >75% decrease in Mincle mRNA expression in
216 siCLEC4E transfected cells. There was no effect of siCLEC4E on cell viability as determined by
217 MTT cell viability assay (data not shown).

218
219 Firstly, in amnion mesenchymal cells, treatment with IL1B significantly increased the expression of
220 pro-inflammatory cytokines IL1A and IL6 mRNA expression and the secretion of IL6 (Figure 3A-
221 C) and chemokines CCL2 and CXCL8 mRNA expression and secretion (Figure 3D-G), compared
222 to siCONT transfected cells. In siCLEC4E transfected cells, there was a significant decrease in
223 IL1B induced IL1A, IL6, CCL2 and CXCL8 mRNA expression (Figure 3A-B,D,F) and release of
224 IL6 and CXCL8 (Figure C,G), but not CCL2 (Figure 3E). Similarly in amnion mesenchymal cells
225 treated with TNF, siCLEC4E transfected cells displayed a significant decrease in TNF-induced
226 IL1A, IL1B, IL6, CCL2 and CXCL8 mRNA expression (Figure 3H-J,L,N) and release of IL6 and
227 CXCL8 (Figure 3K,O), but not CCL2 release (Figure 3M).

228
229 Figures 4-6 demonstrates the effect of siCLEC4E transfection in myometrial cells in the presence of
230 pro-labor mediators; pro-inflammatory cytokine IL1B (Figure 4), TLR2/6 ligand fsl-1 (Figure 5)
231 and TLR3 ligand poly(I:C) (Figure 6). Treatment with IL1B significantly increased TNF, IL1A,
232 IL6, CCL2, CXCL1 and CXCL8 mRNA expression (Figure 4A-C,E,G,I) and release of IL6, CCL2,
233 CXCL1 and CXCL8 (Figure 4D,F,H,J). While there was no effect of siCLEC4E transfection on
234 IL1B-induced TNF mRNA expression (Figure 4A), there was a significant decrease in IL1A mRNA
235 expression (Figure 4B) and IL6, CCL2, CXCL1 and CXCL8 mRNA expression and release (Figure
236 4C-J). Treatment of myometrial cells with fsl-1 (Figure 5) and poly(I:C) (Figure 6) displayed very
237 similar results; the expression of TNF, IL1A, IL1B, IL6, CCL2, CXCL1 and CXCL8 mRNA and
238 release of IL6, CCL2, CXCL1 and CXCL8 was significantly decreased in siCLEC4E transfected
239 cells, compared to siCONT cells treated with fsl-1 or poly(I:C).

240
241 **Knockdown of Mincle suppresses myometrial cell contractility *in vitro***

242 Pro-inflammatory cytokines can participate in myometrial activation by inducing contraction
243 associate proteins such as PTGS2 (41) and promote the production of the uterotonic prostaglandin

244 PGF_{2α} (42) which acts via the prostaglandin F receptor (PTGFR) (43). Thus, it was interest to
245 explore the effect of siCLEC4E on expression of contraction associated proteins (Cx43, OXTR,
246 PTGS and PTGFR), PGF_{2α} and myometrial cell contractility *in situ*. Treatment of siCONT cells
247 with IL1B and TNF significantly decreased Cx43 mRNA expression (Figure 7A,D), and
248 significantly increased PTGFR mRNA expression and secretion of PGF_{2α}, (Figure 7B-C,E-F),
249 compared to basal cells. While there was no effect of siCLEC4E on IL1B-induced Cx43 mRNA
250 expression (Figure 7A), there was a significant decrease in TNF-induced Cx43 mRNA expression
251 (Figure 7D). In siCLEC4E transfected cells, there was a significant decrease in IL1B- and TNF-
252 induced PTGFR mRNA expression and secretion of PGF_{2α} (Figure 7B-C,E-F). There was no
253 change with PTGS2 or OXTR mRNA expression in siCLEC4E transfected cells (data not shown).
254 To accompany these experiments, we determined whether Mincle can regulate myometrial
255 contractility *in situ* by utilising an *in vitro* gel contraction assay; primary myometrial cells deficient
256 in Mincle were embedded into 3D collagen gel matrices where the cells could contract in response
257 to TNF. As shown in Figure 7G, TNF induced cell contractility, evidenced by increased gel
258 shrinkage, compared to siCONT cells without TNF. In siCLEC4E transfected cells, TNF-induced
259 gel shrinkage was significantly suppressed.

260

261 **Knockdown of Mincle suppresses expression of ECM remodelling enzymes**

262 The pro-inflammatory cytokines IL1B and TNF play important roles in the initiation of labor,
263 particularly the rupture of membranes. The two constituent cell populations of the amnion,
264 epithelial and mesenchymal cells, were both investigated, as they independently respond to different
265 stimuli to produce different matrix metalloproteinases (MMPs); for example, amnion epithelial
266 cells, but not mesenchymal cells, participate in IL1B-induced MMP9 release (35), and in
267 mesenchymal cells, but not epithelial cells, thrombin can induce both MMP1 and MMP9 expression
268 (44).

269

270 In amnion epithelial cells, we found that IL1B significantly increased MMP9 mRNA expression
271 (Figure 8A) and release of pro MMP9 (Figure 8B). In siCLEC4E transfected cells, there was a
272 significant decrease in IL1B-induced MMP9 mRNA expression (Figure 8A) and pro MMP9
273 secretion (Figure 8B). Amnion mesenchymal cells were transfected with or without siCLEC4E and
274 treated with TNF. Treatment with TNF significantly increased MMP1, MMP8 and MMP9 mRNA
275 expression (Figure 8C-E). In siCLEC4E transfected cells, there was a significant attenuation in
276 TNF-induced MMP1 and MMP8 mRNA expression (Figure 8C,D), but there was no change with
277 MMP9 mRNA expression (Figure 8E).

278 **DISCUSSION**

279

280 This study adds to the growing literature of the role of PRRs in human labor. This study, however,
281 is the first to describe a role for the CLR in human myometrium and fetal membranes in the context
282 of parturition. Here, we show that the expression of the receptor Mincle (CLEC4E) is upregulated
283 in myometrium and/or fetal membranes after spontaneous labor at preterm and term and in response
284 to intrauterine infection. Furthermore, Mincle expression is highly induced by pro-inflammatory
285 and pro-labor insults such as pro-inflammatory cytokines (IL1B, TNF), bacterial products (LPS, fsl-
286 1, flagellin) and viral products (poly(I:C)). Knockdown of Mincle in primary amnion and
287 myometrial cells causes the suppression of mediators involved in regulating the processes of labor
288 and delivery including pro-inflammatory cytokines (TNF, IL1A, IL1B), chemokines (CCL2,
289 CXCL1, CXCL8) and ECM degrading enzymes (MMP1, MMP8, MMP9). In addition, our study
290 reveals a role for Mincle in regulating myometrial activation as assessed by suppression of
291 contraction associated proteins (Cx43, PTGFR) and secretion the uterotonic PGF_{2α} in cells deficient
292 in Mincle. Finally, using an *in situ* model of myometrial contractility, the absence of Mincle leads
293 to a significant augmentation of the magnitude of inflammation-induced collagen gel contractility.
294 Taken together, our results demonstrate a previously unconsidered important role of Mincle in
295 regulating inflammation induced pro-labor mediators.

296

297 Human labor at term in myometrium, and at preterm and term in fetal membranes was associated
298 with upregulated Mincle expression when compared to non-laboring tissues. Likewise, Mincle
299 expression was also upregulated in myometrium during labor in a mouse model of inflammation
300 induced preterm birth and in amnion from women with preterm chorioamnionitis compared to
301 amnion from women without preterm chorioamnionitis. Inflammation is a hallmark of labor, with
302 an influx of leukocytes and increased expression of cytokines, chemokines and other inflammatory
303 markers observed in uterus, cervix, placenta and fetal membranes during labor (45, 46).

304 Chorioamnionitis is an acute inflammation that occurs within the fetal membranes in response to the
305 presence of pathogenic organisms (47). To test the possibility that increased Mincle expression is a
306 result of these inflammatory responses, we evaluated the effect of the pro-inflammatory cytokines
307 on Mincle expression. We found that IL1B and TNF, two acute phase pro-inflammatory cytokines
308 that are secreted by leukocytes, significantly increased Mincle expression in fetal membranes and
309 myometrium. Collectively, these studies demonstrate that Mincle expression is regulated by
310 inflammatory insults in myometrium and fetal membranes, and may thus may play a role in human
311 labor.

312

313 IL1B and TNF are two acute phase cytokines that have vital roles during parturition. For example,
314 in various animal species, injection of IL1B or TNF can induce preterm labor (48-51) and increase
315 levels of pro-inflammatory cytokines, chemokines, prostaglandins, MMP9 and leukocytes in
316 amniotic fluid (49). These animal studies are supported by *in vitro* studies whereby IL1B or TNF
317 can further increase cytokine production, upregulate the expression of chemokines, activate ECM
318 matrix remodelling enzymes, induce expression of contraction associated proteins and stimulate the
319 ongoing release of prostaglandins (52-54). As Mincle was induced by IL1B and TNF, we
320 investigated if Mincle is involved in the genesis of pro-inflammatory and pro-labor mediators
321 induced by IL1B or TNF. Indeed, using siRNA knockdown experiments, we found that Mincle
322 regulates IL1B or TNF induced pro-inflammatory cytokines (TNF, IL1A, IL1B), chemokines
323 (CCL2, CXCL1, CXCL8) and ECM degrading enzymes (MMP1, MMP8, MMP9) in human
324 primary myometrial and amnion cells. Further, Mincle regulates IL1B or TNF induced expression
325 of contraction associated proteins (Cx43, PTGFR), secretion the uterotonic PGF_{2α}, and myometrial
326 contractility *in situ*. Taken together, these results suggest that Mincle is involved in the genesis of
327 IL1B or TNF induced mediators involved myometrial contractions and rupture of fetal membranes.

328

329 In addition to pro-inflammatory cytokines, TLRs can also induce Mincle expression in numerous
330 species (12, 14, 15, 55). Likewise, in this study, the TLR2/6 ligand fsl-1, TLR3 ligand poly(I:C),
331 TLR4 ligand LPS and the TLR5 ligand flagellin significantly increased Mincle expression in fetal
332 membranes and myometrium. This has important implications given that infection is the most
333 common cause of early preterm birth (56) and that pathogens and bacterial and viral products
334 engage TLRs to induce preterm labor in mice (6-8, 57).

335

336 Induction of Mincle by TLRs activates a cascade of events leading to the transcriptional regulation
337 of pro-inflammatory genes (12, 14, 15, 55). Our previous studies have shown that ligation of
338 TLR2/6 ligand fsl-1 and the TLR3 ligand poly(I:C) can induce pro-labor mediators in human
339 myometrial and amnion cells (1, 3). Thus, it was of interest to determine if Mincle is also involved
340 in TLR signalling. Using primary myometrial cells as a model system, and akin to the studies using
341 IL1B and TNF, we found that siRNA inhibition of Mincle decreased the effectiveness of the
342 TLR2/6 ligand fsl-1 and the TLR3 ligand poly(I:C) to induce the expression and secretion of pro-
343 inflammatory cytokines and chemokines. Collectively, the results presented in this paper suggest
344 that the expression and secretion of pro-labor mediators can be induced by TLRs through the
345 Mincle pathway.

346

347 The downstream pathways involved in Mincle induced regulation of pro-labor mediators is not
348 known. We found no effect of Mincle siRNA knockout on NF- κ B transcriptional activity in
349 myometrial cells, suggesting that Mincle regulates pro-inflammatory and pro-labor mediators in
350 gestational tissues by other downstream transcription factors. NFAT, MAPK and AP-1 have also
351 been shown to be involved in Mincle induced gene transcription (21-23). Notably, a few recent
352 studies indicate a potential role for NFAT in mediating stretch or oxytocin induced gene expression
353 in myometrium (58, 59). A role for MAPK and AP-1 proteins in regulating mediators involved in
354 myometrial contractility and degradation of fetal membranes has also been reported (60-63).
355 Further studies are therefore required to determine the functional role of NFAT, MAPK and AP-1 in
356 Mincle signalling.

357
358 A limitation of this study is that we could not assess Mincle protein expression due to non-
359 specificity of commercial antibodies (data not shown). Studies looking at Mincle expression and
360 localisation in myometrium throughout gestation would further clarify other potential functional
361 contributions Mincle makes in regulating myometrial contractility in pregnancy. Likewise, further
362 studies are required to determine the regional localisation of Mincle in fetal membranes, i.e. site
363 over the cervix prone to rupture versus distal areas, throughout gestational ages would shed light on
364 the role of Mincle in fetal membrane rupture. In this study, all experiments were performed in the
365 presence of an inflammatory insult. It would also be worth determining if Mincle plays a role in
366 parturition in the absence of inflammation.

367
368 In conclusion, the expression of Mincle is upregulated after labor and infection in fetal membranes
369 and myometrium. Furthermore, pro-inflammatory cytokines, and bacterial and viral products
370 increased Mincle expression. Knockdown of Mincle reduced the expression and secretion of
371 mediators involved in active labor and delivery. Importantly, in the absence of Mincle, the
372 effectiveness of pro-inflammatory cytokines and TLR ligands to induce the expression and or
373 secretion of pro-labor mediators was reduced. These data suggest that Mincle is part of the process
374 by which inflammation induces parturition.

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379

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386

387 **DISCLOSURE SUMMARY**

388 The authors have nothing to declare.

389

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FIGURE LEGENDS

Figure 1. Expression of Mincle in myometrium and fetal membranes

(A) Fetal membranes were obtained from women at term Caesarean section in the absence of labor (term no labor, n=9 patients) or from women after term spontaneous labor onset and delivery (term after labor, n=9 patients). (B) Fetal membranes were obtained from women at preterm Caesarean section in the absence of labor (preterm no labor, n=9 patients) or from women after preterm spontaneous labor onset and delivery (preterm after labor, n=9 patients). (C) Amnion was obtained from women at preterm Caesarean section without histological chorioamnionitis (preterm no CAM, n=8 patients) or from women at preterm Caesarean section with histological chorioamnionitis (preterm CAM, n=8 patients). (D) Human myometrium was obtained from women at term Caesarean section in the absence of labor (term no labor, n=8 patients) or from women at term Caesarean section during labor (term in labor, n=8 patients). (E) Myometrium was obtained from mice during LPS-induced labor at gestational day 15.5 and time-matched vehicle-injected mice (n=4 mice/group). For all data, Mincle mRNA expression was analysed by RT-qPCR. Individual data points represent different patients and the horizontal line represents the mean \pm SEM of each group. * $P \leq 0.05$, Mann-Whitney U test.

Figure 2. Effect of pro-inflammatory mediators on Mincle expression in fetal membranes and myometrium

Human (A) fetal membranes and (B) myometrium were incubated in the absence or presence of 10 ng/ml IL1B, 10 ng/ml TNF, 250 ng/ml fsl-1, 20 μ g/ml poly(I:C), 10 μ g/ml LPS or 1 μ g/ml flagellin for 20 h (n=6 patients/treatment). Mincle mRNA expression was analysed by RT-qPCR. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$, Wilcoxon test.

Figure 3. Effect of Mincle siRNA knockdown on pro-inflammatory cytokines and chemokines in human primary amnion mesenchymal cells

Human primary amnion mesenchymal cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with 1 ng/ml IL1B or 10 ng/ml TNF (n=6 patients). (A,B,D,F,H-J,L,N) IL1A, IL1B, IL6, CCL2 and CXCL8 mRNA expression was analysed by RT-qPCR. (C,E,G,K,M,O) The concentration of IL6, CCL2 and CXCL8 in the incubation medium was assayed by ELISA. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA).

Figure 4. Effect of Mincle siRNA knockdown on IL1B-induced pro-inflammatory cytokines and chemokines in human primary myometrial cells

Human primary myometrial cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with 1 ng/ml IL1B (n=7 patients). (A-C,E,G,I) TNF, IL1A, IL6, CCL2, CXCL1 and CXCL8 mRNA expression was analysed by RT-qPCR. (D,F,H,J) The concentration of IL6, CCL2, CXCL1 and CXCL8 in the incubation medium was assayed by ELISA. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA).

Figure 5. Effect of Mincle siRNA knockdown on fsl-1-induced pro-inflammatory cytokines and chemokines in human primary myometrial cells

Human primary myometrial cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with 250 ng/ml fsl-1 (n=6 patients). (A-D,F,H,J) TNF, IL1A, IL1B, IL6, CCL2, CXCL1 and CXCL8 mRNA expression was analysed by RT-qPCR. (E,G,I,K) The concentration of IL6, CCL2, CXCL1 and CXCL8 in the incubation medium was assayed by ELISA. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA).

Figure 6. Effect of Mincle siRNA knockdown on poly(I:C)-induced pro-inflammatory cytokines and chemokines in human primary myometrial cells

Human primary myometrial cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with 5 μ g/ml poly(I:C) (n=6 patients). (A-D,F,H,J) TNF, IL1A, IL1B, IL6, CCL2, CXCL1 and CXCL8 mRNA expression was analysed by RT-qPCR. (E,G,I,K) The concentration of IL6, CCL2, CXCL1 and CXCL8 in the incubation medium was assayed by ELISA. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA).

Figure 7. Effect of Mincle siRNA knockdown on myometrial contractility

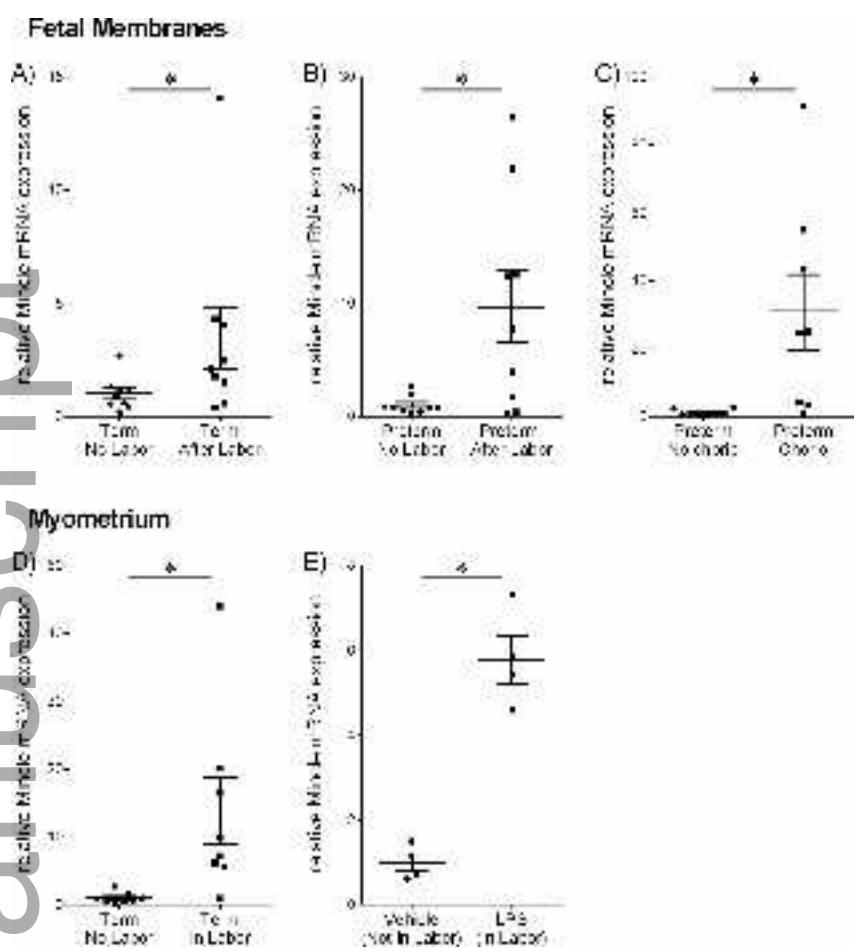
(A-F) Human primary myometrial cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with (A-C) 1 ng/ml IL1B or (D-F) 10 ng/ml TNF (n=8 patients). (A,B,D,E) PTGS2 and PTGFR mRNA expression was analysed by RT-qPCR. (C,F) The concentration of $\text{PGF}_{2\alpha}$ in the incubation medium was assayed by ELISA. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA). (G) Cell contraction assays were performed using collagen gels made from human primary myometrial cells transfected with 50 nM siCONT or 50 nM siCLEC4E for 48 h (n=6 patients). The collagen gels were then treated with or without 10 ng/ml TNF for 36 h, and the area of gel was determined.

Representative gel contraction image from 1 patient is also shown. * $P \leq 0.05$ (repeated measures one-way ANOVA).

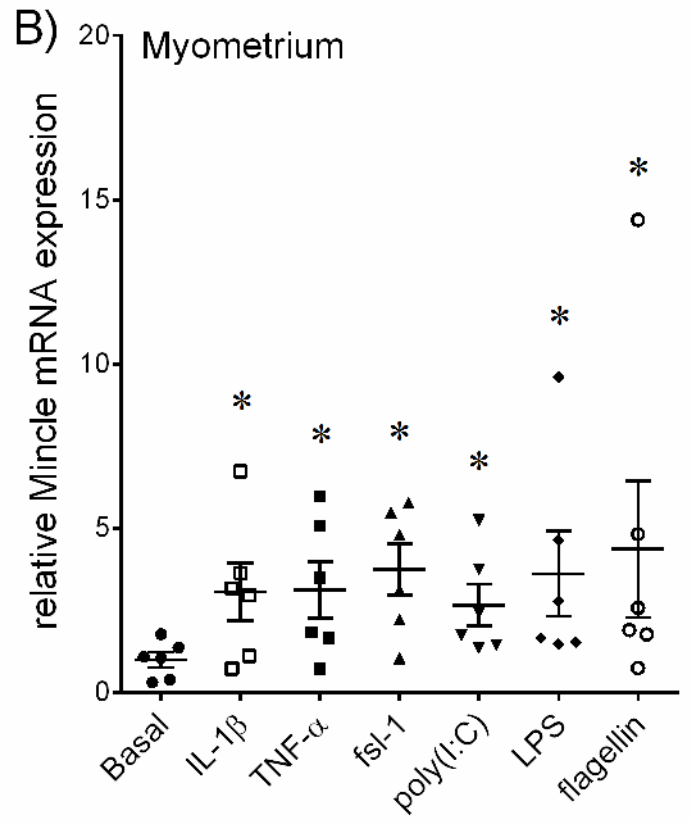
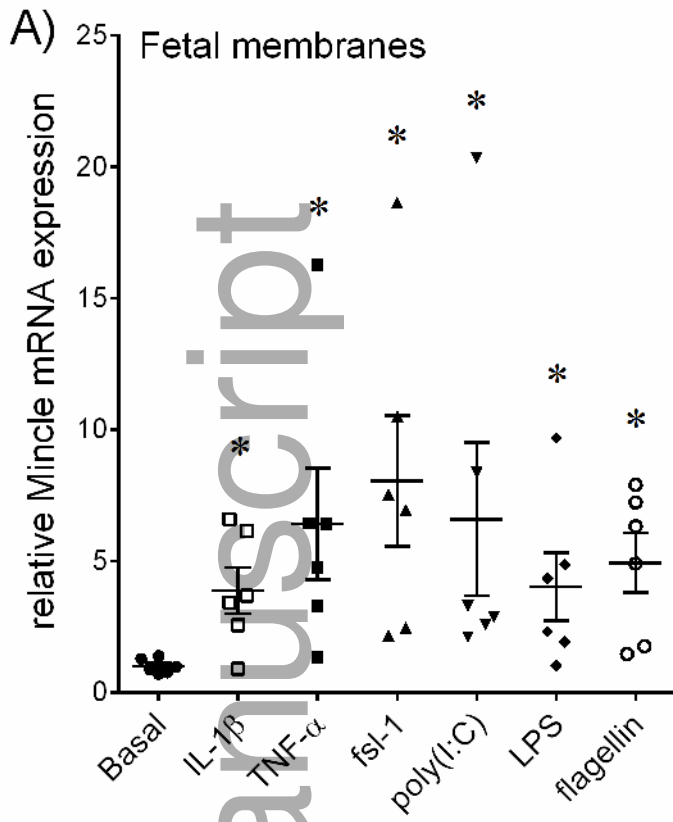
Figure 8. Effect of Mincle siRNA knockdown on the expression of ECM degrading enzymes in primary amnion cells

(A,B) Human primary amnion epithelial cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with 1 ng/ml IL1B (n=6 patients). (A) MMP9 mRNA expression was analysed by RT-qPCR. (B) The concentration of pro MMP9 in the incubation medium was assayed by zymography. Representative zymography image from one patient is also shown. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA). (C-E) Human primary amnion mesenchymal cells were transfected with 50 nM siCONT or 50 nM siCLEC4E and then treated with 10 ng/ml TNF (n=6 patients). (A) MMP1, (B) MMP8, and (C) MMP9 mRNA expression was analysed by RT-qPCR. Individual data points represent 6 independent experiments and displayed as mean \pm SEM. * $P \leq 0.05$ (repeated measures one-way ANOVA).

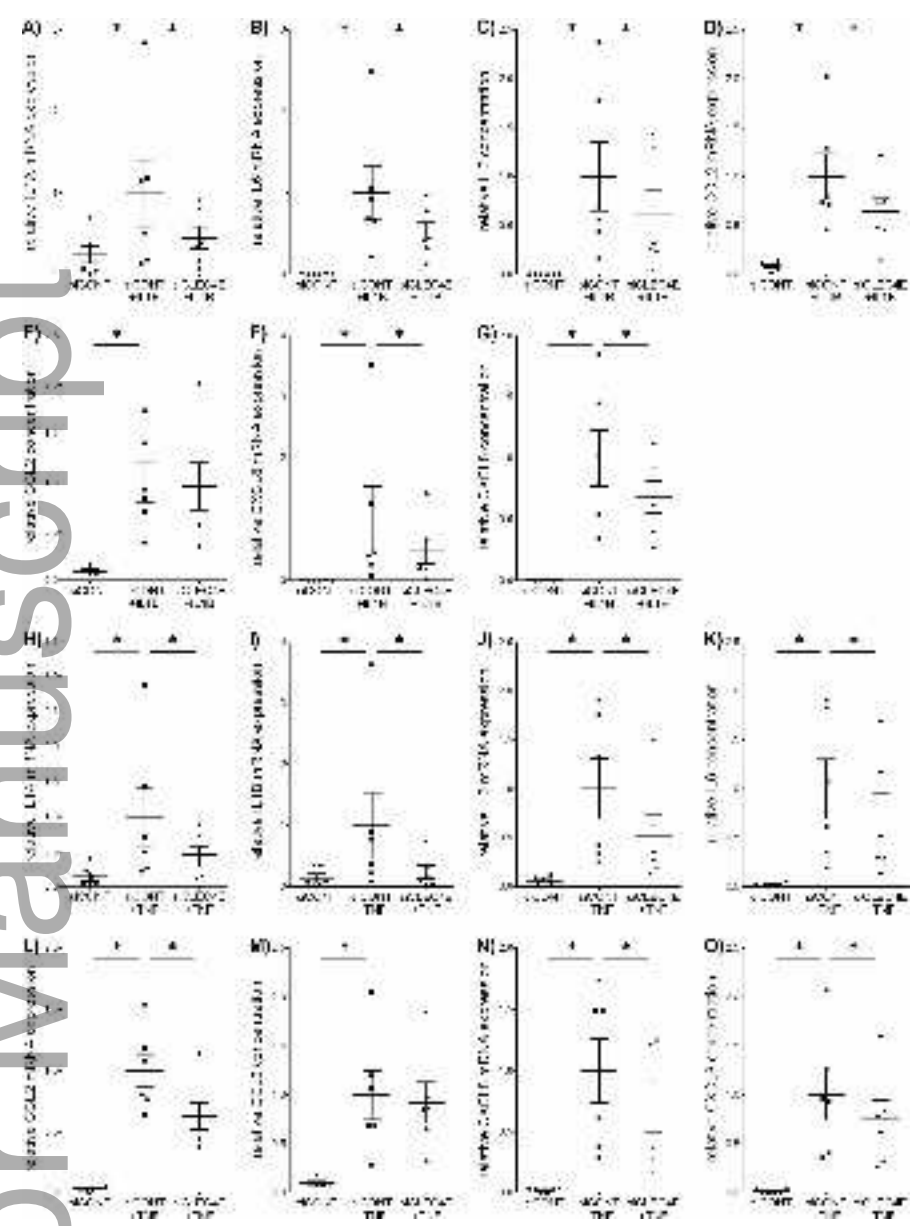
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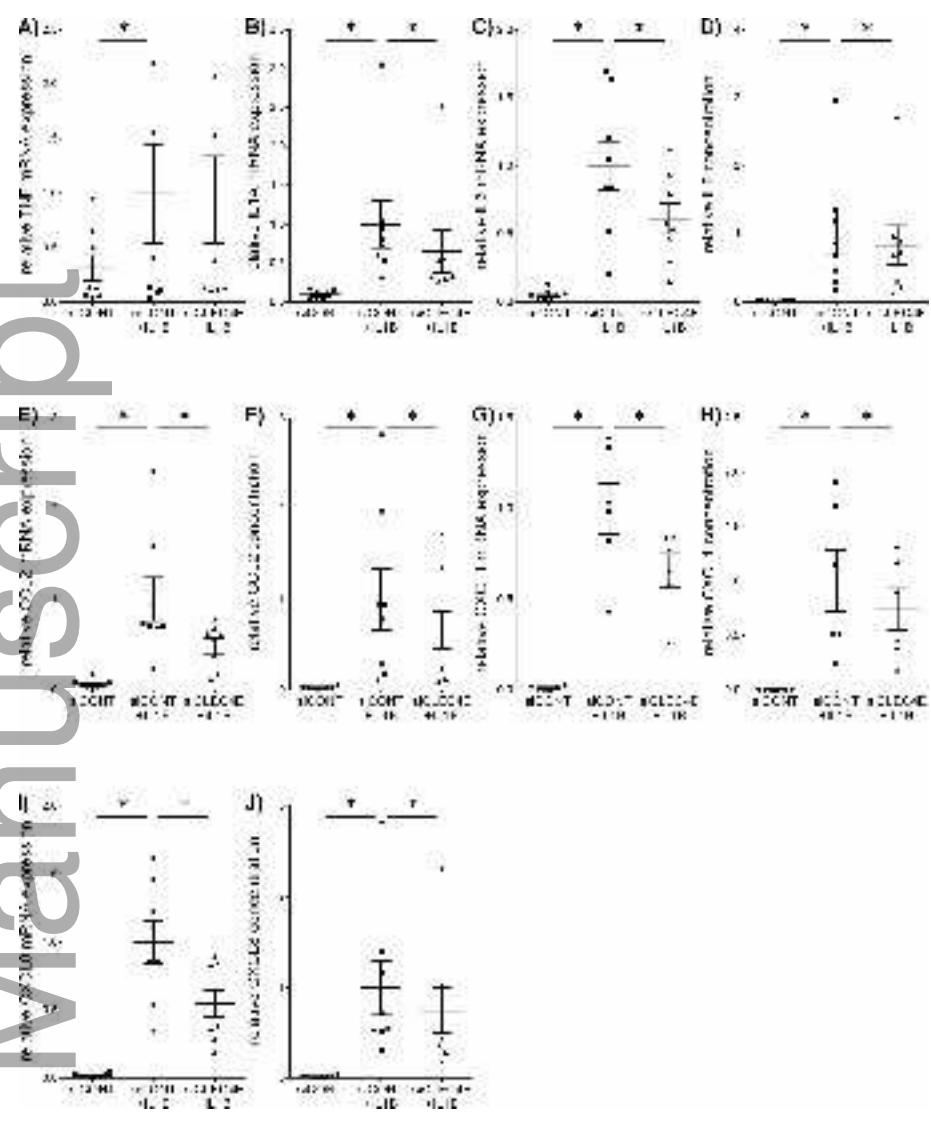
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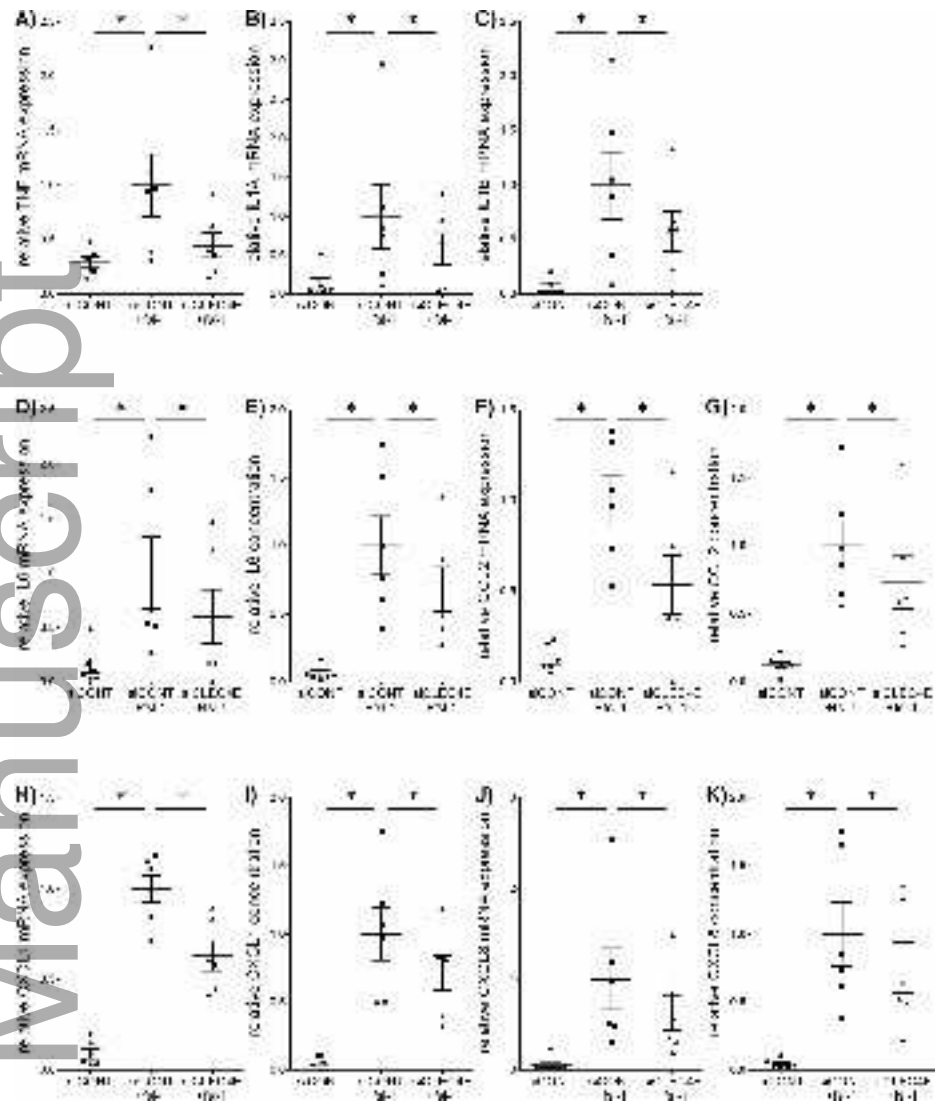
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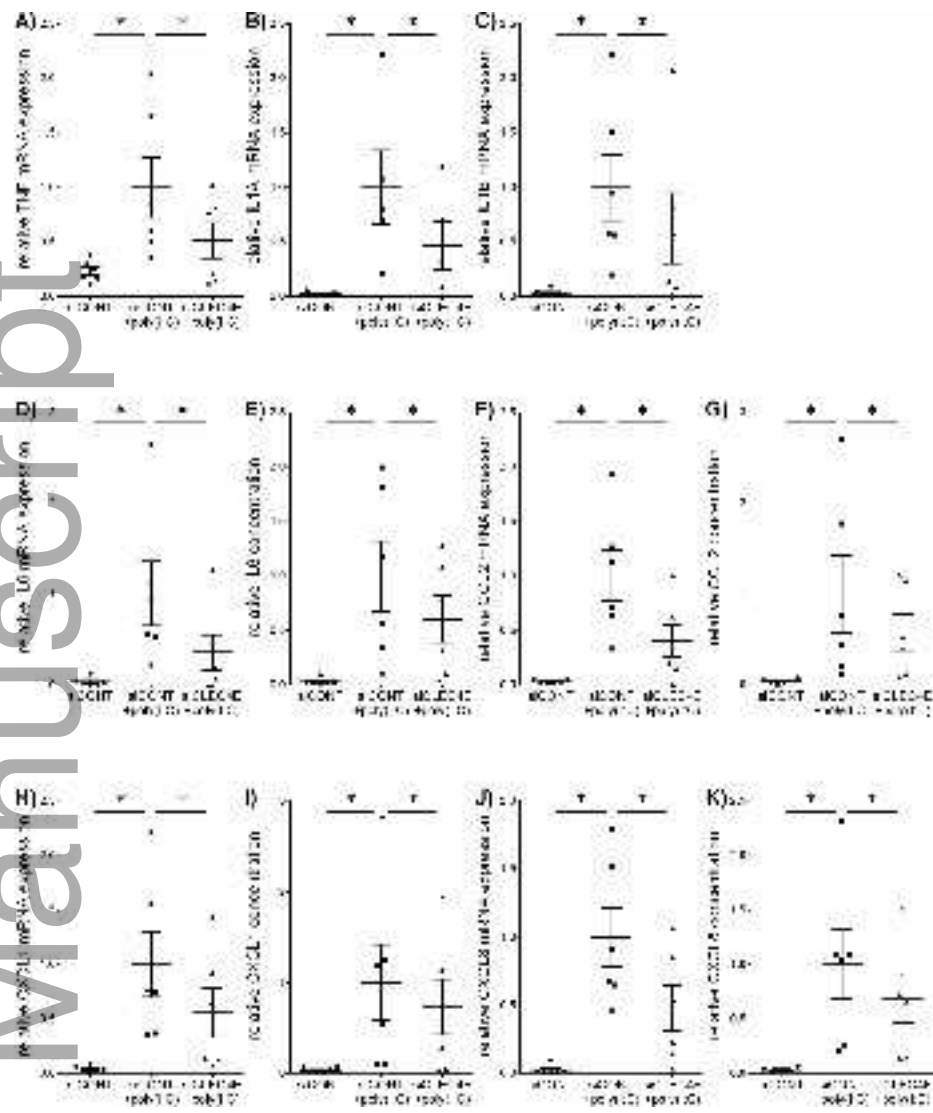
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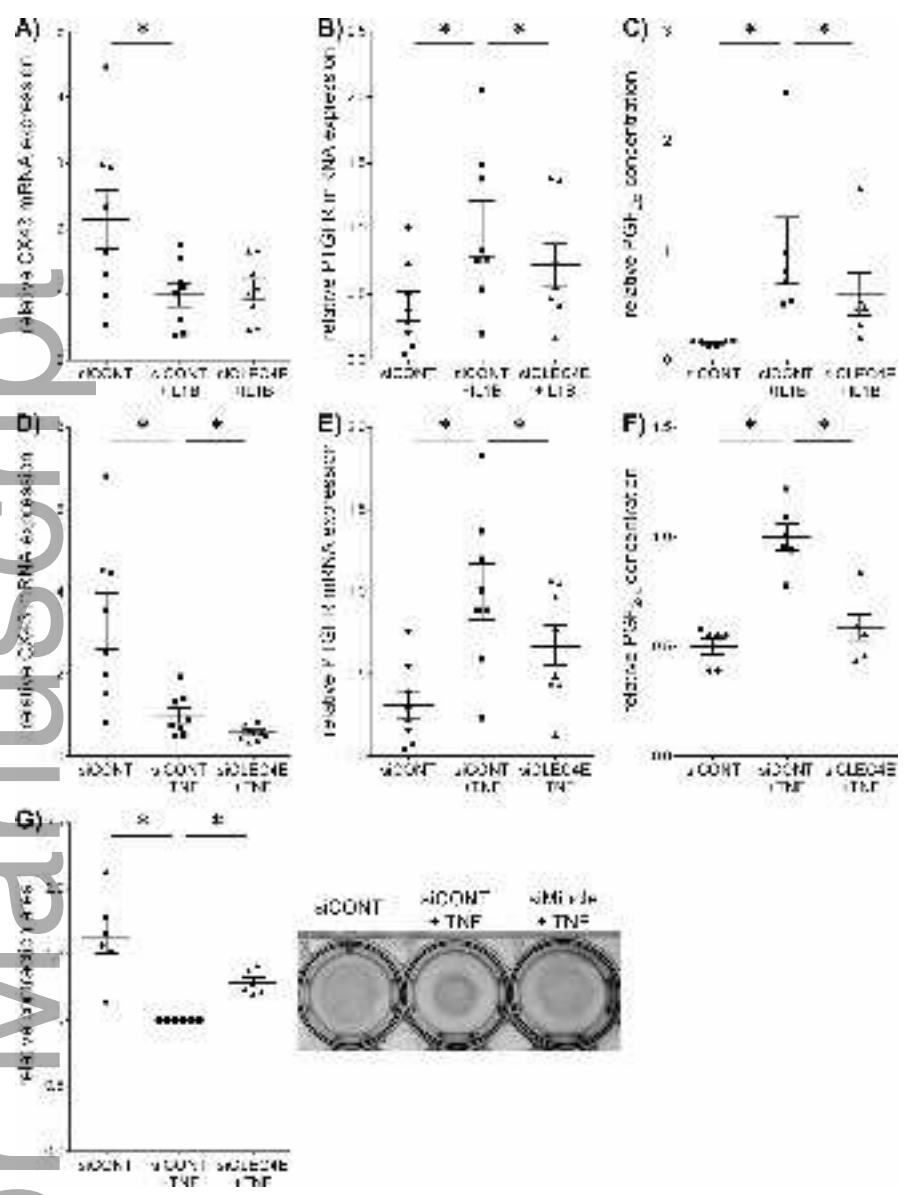
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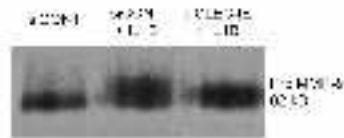
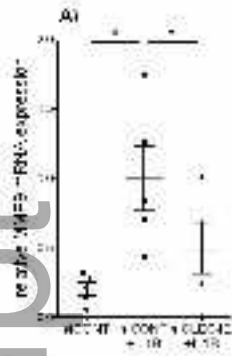


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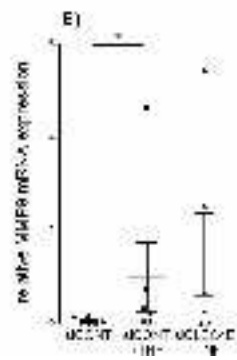


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Amnion epithelial cells



Amnion mesenchymal cells



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