

## Repurposing simvastatin as a therapy for preterm labor: evidence from preclinical models

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**ABSTRACT:** Preterm birth (PTB), the leading cause of neonatal morbidity and mortality, urgently requires novel therapeutic agents. Spontaneous PTB, resulting from preterm labor, is commonly caused by intrauterine infection/inflammation. Statins are well-established, cholesterol-lowering drugs that can reduce inflammation and inhibit vascular smooth muscle contraction. We show that simvastatin reduced the incidence of PTB in a validated intrauterine LPS-induced PTB mouse model, decreased uterine proinflammatory mRNA concentrations (IL-6, Cxcl1, and Ccl2), and reduced serum IL-6 concentration. In human myometrial cells, simvastatin reduced proinflammatory mediator mRNA and protein expression (IL-6 and IL-8) and increased anti-inflammatory cytokine mRNA expression (IL-10 and IL-13). Critically, simvastatin inhibited myometrial cell contraction, basally and during inflammation, and reduced phosphorylated myosin light chain concentration. Supplementation with mevalonate and geranylgeranyl pyrophosphate, but not farnesyl pyrophosphate, abolished these anticontractile effects, indicating that the Rho/Rho-associated protein kinase pathway is critically involved. Thus, simvastatin reduces PTB incidence in mice, inhibits myometrial contractions, and exhibits key anti-inflammatory effects, providing a rationale for investigation into the repurposing of statins to treat preterm labor in women.—Boyle, A. K., Rinaldi, S. F., Rossi, A. G., Saunders, P. T. K., Norman, J. E. Repurposing simvastatin as a therapy for preterm labor: evidence from preclinical models. *FASEB J.* 33, 2743–2758 (2019). www.fasebj.org

**KEY WORDS:** pregnancy · preterm birth · statins · inflammation · contraction

Spontaneous preterm labor (PTL) reportedly accounts for two thirds of cases of preterm birth (PTB). PTB is defined as delivery before 37 completed weeks of gestation and is responsible for 11.1% of annual births worldwide (1, 2). It is the leading cause of mortality in children aged <5 yr and has a substantial economic burden due to costs of neonatal intensive care and lifelong morbidity (3, 4). The etiology of PTL is poorly understood, but intrauterine infection/inflammation is believed to be the most common cause (5). Fetal exposure to this adverse intrauterine

environment can result in fetal organ injury, particularly to the brain, increasing the risk of cognitive and neurologic impairment (6, 7).

Human labor is an inflammatory process associated with the influx of immune cells into the gestational tissues and the increased production of proinflammatory mediators, which stimulates uterine contractility, cervical ripening, and fetal membrane rupture (8, 9). Contraction of the myometrium is facilitated by the cross-bridging of the myofilaments actin and myosin, instigated by the increase of intracellular calcium (Ca<sup>2+</sup>) concentration and completed by the phosphorylation of myosin light chain (MLC) (10). Although drugs are often administered to reduce uterine contractility, there is little evidence to suggest that these tocolytic agents have either a substantial impact on the timing of delivery or improve the health of premature neonates (11, 12). New treatments are urgently needed that both suppress myometrial contractions and the inflammation associated with PTL, to both delay delivery and prevent fetal injury.

Statins are 3-hydroxy-3-methylglutaryl-coenzyme A reductase inhibitors. These widely used drugs are potent inhibitors of cholesterol biosynthesis, and they are commonly used for the prevention of cardiovascular disease. In addition, they are reported to have immunomodulation, anti-inflammatory, and antioxidative stress properties (13,

**ABBREVIATIONS:** Cx, connexin; F-PP, farnesyl pyrophosphate; GG-PP, geranylgeranyl pyrophosphate; I-PP, isopentenyl pyrophosphate; MLC, myosin light chain; MTT, 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide; pMLC, phosphorylated myosin light chain; PTB, preterm birth; PTL, preterm labor; qRT-PCR, quantitative RT-PCR; ROCK, Rho-associated protein kinase

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(25, 100 ng/ml), mevalonate (200  $\mu$ M), GG-PP (10  $\mu$ M), or F-PP (10  $\mu$ M) for 24–48 h. MTT solution was applied for 4 h at 37°C. The medium and MTT solution were removed, and 100  $\mu$ l acidified isopropanol was added for 20 min; light absorbance was then measured at 540 nm (Multiskan EX, LabSystems).

### Reverse transcription and qRT-PCR for human cell lysate and mouse tissue

Total RNA was extracted from cell lysate and mouse uterus samples by using an RNeasy mini kit (Qiagen, Germantown, MD, USA) according to the manufacturer's instructions, and quantified by using a NanoDrop 2000c (Thermo Fisher Scientific, Hemel Hempstead, United Kingdom). Total RNA (300 ng/ $\mu$ l) was reverse transcribed by using the High Capacity cDNA Reverse Transcription Kit (Thermo Fisher Scientific). Pre-designed TaqMan gene expression assays (Thermo Fisher Scientific) were used to investigate genes of interest and are listed in Table 1. All qRT-PCR analyses were performed on a Thermo Fisher Scientific 7900HT instrument. Target mRNA expression was normalized for RNA loading using  $\beta$ -actin (Thermo Fisher Scientific), and the mRNA concentration in each sample was calculated relative to the vehicle control by using the  $2^{-\Delta\Delta C_t}$  method of analysis.

### Mouse and human ELISAs

Human myometrial cell supernatants were analyzed for IL-6 and IL-8 secretion by using DuoSet ELISAs (R&D Systems, Abingdon, United Kingdom). IL-6 was measured in mouse maternal serum and amniotic fluid (Quantikine; R&D Systems). Both assays were performed according to the manufacturer's instructions.

### Collagen gel contraction assay

A collagen gel contraction assay was used to assess the effect of simvastatin on the contraction of myometrial cells and was performed as previously described (26–28). Briefly, myometrial cells (10<sup>5</sup> cells/well) were embedded in type I rat tail collagen (Thermo Fisher Scientific). The collagen was allowed to polymerize overnight at 37°C. Treatments were prepared in 5% (v/v) fetal bovine serum DMEM, and wells were photographed at 0, 24, and 48 h by using a Leica MZ6 light microscope/camera (Leica Microsystems, Wetzlar, Germany). ImageJ software (National Institutes of Health, Bethesda, MD, USA) was used to analyze gel area. The gel area measurements at 24 and 48 h were calculated as

a percentage of the mean gel area at the 0-h time point for each independent experiment.

### In-cell Western assay

In-cell Western analyses were performed to quantify phosphorylated myosin light chain (pMLC), as previously described (26, 27, 29). Briefly, myometrial cells were seeded at  $2 \times 10^4$  cells/well in black, 96-well, clear base plates (PerkinElmer, Waltham, MA, USA). The treatments were prepared in 5% (v/v) fetal bovine serum DMEM and applied in triplicate for 48 h at 37°C. Cells were fixed with 4% formaldehyde (MilliporeSigma) for 15 min at room temperature, and plates were then washed and permeabilized 3 times with 0.1% Triton X-100 in PBS for 5 min. The plates were incubated with primary antibodies, polyclonal rabbit anti-pMLC 2 [Ser19] (3671; Cell Signaling Technology, Danvers, MA, USA), and anti- $\alpha$ -Tubulin mAb (T9026; Sigma-Aldrich), and then secondary antibodies, 800CW and 680RD (926-68072 and 926-32213; Li-Cor, Lincoln, NE, USA). The Odyssey CLx Imaging System (Li-Cor) was used to read the plates and measure the signal in each well. The intensity of pMLC fluorescence was quantified relative to the  $\alpha$ -Tubulin signal within the same well.

### Liquid chromatography tandem mass spectrometry quantification of serum progesterone

Progesterone concentration was measured in dam serum by liquid chromatography tandem mass spectrometry, using a QTrap 5500 (AB Sciex, Warrington, United Kingdom), with an Acquity Ultra Performance Liquid Chromatography (UPLC; Waters Corp., Manchester, United Kingdom). Mass spectral conditions are presented in Table 2. Analytes were extracted from serum (50  $\mu$ l) supported liquid extraction (400  $\mu$ l SLE+; Biotage, Uppsala, Sweden) with D9-progesterone (1.5 ng; C D N Isotopes, Pointe-Claire, QC, Canada) included as an internal standard. Separation took place at 40°C on an Ace Excel C18 (100  $\times$  2.1 mm, 1.7  $\mu$ m) column (Advanced Chromatography Technologies, Aberdeen, United Kingdom) using a gradient solvent system (50:50 of water with 0.1% formic acid and acetonitrile with 0.1% formic acid) with a gradient run of 6.5 min (Table 3).

### Protein extraction and quantification

Tissue was homogenized in lysis buffer (RIPA buffer; MilliporeSigma) containing 1 Complete Protease Inhibitor Cocktail Tablet (Roche, Basel, Switzerland) using a Tissue Lyser II (Qiagen) at 25 Hz. The samples were incubated on ice for 5 min and then centrifuged at 10,000g for 10 min at 4°C. The supernatant was then divided into aliquots and stored at  $-80^\circ\text{C}$ . Protein was quantified by using the DC (Bio-Rad, Hercules, CA, USA) protein assay according to the manufacturer's instructions.

### Western blot analyses

For connexin 43 (CX43) analysis, samples were loaded (20  $\mu$ g protein/sample) on 4–12% (12-well) Bis-Tris precast NuPAGE gels and run in 3-(N-morpholino)propanesulfonic acid running buffer (Thermo Fisher Scientific) at 180 V for 70 min. The protein was transferred with the use of a wet-transfer system (100 V for 90 min) to Immobilon-FL PVDF membranes (MilliporeSigma). The membranes were blocked with 5% nonfat dry milk in 0.5% Tween (MilliporeSigma) Tris-buffered saline (Thermo Fisher Scientific) and incubated with primary antibodies, polyclonal rabbit anti-CX43 (ab11370; Abcam, Inc., Cambridge, United Kingdom), and monoclonal mouse anti- $\alpha$ -Tubulin overnight at

TABLE 1. Pre-designed TaqMan gene expression assay IDs

| Gene                           | Species | Code          |
|--------------------------------|---------|---------------|
| <i>IL-6</i>                    | Human   | Hs00985639_m1 |
| <i>IL-8</i>                    | Human   | Hs00174103_m1 |
| <i>IL-10</i>                   | Human   | Hs00961622_m1 |
| <i>IL-13</i>                   | Human   | Hs00174379_m1 |
| <i>Ccl2</i>                    | Mouse   | Mm00441242_m1 |
| <i>Cox-2</i>                   | Mouse   | Mm00478374_m1 |
| <i>Cxcl1</i>                   | Mouse   | Mm04207460_m1 |
| <i>Cxcl2</i>                   | Mouse   | Mm00436450_m1 |
| <i>Cx43</i>                    | Mouse   | Mm01179639_s1 |
| <i>IL-1<math>\beta</math></i>  | Mouse   | Mm00434228_m1 |
| <i>IL-6</i>                    | Mouse   | Mm00446190_m1 |
| <i>IL-10</i>                   | Mouse   | Mm01288386_m1 |
| <i>TNF-<math>\alpha</math></i> | Mouse   | Mm00443258_m1 |

TABLE 2. Mass spectral conditions for analysis of progesterone and the internal standard, D9-progesterone, utilizing positive and negative electrospray

| Hormone         | MW (Da) | Precursor ion ( <i>m/z</i> ) | Production quan; qual | Declustering potential (V) | Collision energy (V) quan; qual | Cell exit potential (V) quan; qual |
|-----------------|---------|------------------------------|-----------------------|----------------------------|---------------------------------|------------------------------------|
| Progesterone    | 314.462 | 315.0                        | 97.0; 109.0           | 146                        | 29; 37                          | 10; 18                             |
| D9-Progesterone | 323.52  | 324.1                        | 100.0                 | 151                        | 31                              | 38                                 |

Quan, quantifier ion; qual, qualifier ion.

4°C. Membranes were incubated with secondary antibodies, 680RD and 800CW (Li-Cor), and scanned by using the Li-Cor Odyssey Fc Imaging System. The bands for CX43 and  $\alpha$ -Tubulin were quantified by using the Odyssey analysis software.

Chemiluminescence was performed as previously described, with minor alterations. Briefly, the protein (45  $\mu$ g protein/sample) was transferred to Immobilon-P PVDF membranes (MilliporeSigma) and blocked with 5% bovine serum albumin (MilliporeSigma) in 0.5% Tween Tris-buffered saline. After incubation with primary antibodies, monoclonal rabbit anti-IL-6 (12912; Cell Signaling Technology) and monoclonal mouse anti- $\alpha$ -Tubulin, and secondary antibodies, polyclonal swine anti-rabbit HRP (P0399; Agilent, Santa Clara, CA, USA) and polyclonal goat anti-mouse HRP (P0447; Agilent), the membranes were incubated with Amersham ECL Western blotting detection reagent (GE Healthcare Life Sciences, Marlborough, MA, USA) for up to 5 min, then scanned and analyzed as previously discussed.

## Statistics

Data are presented as means  $\pm$  SEM and were analyzed by using GraphPad Prism v.7 (GraphPad Software, La Jolla, CA, USA). For mouse studies, “*n*” represents the number of individual dams treated. For cell studies, “*n*” denotes the number of individual experiments performed, with the technical replicate number per experiment indicated in the figure legends. Time to delivery data were analyzed by using the Kruskal-Wallis test, with Dunn’s *post hoc* test. The percent data for live-born pups and collagen gel contraction were analyzed by performing an arcsine transformation on the proportions, to normalize the binomial distribution, followed by a 1-way ANOVA with either a Dunnett or Holm-Sidak *post hoc* test. ELISA and in-cell Western data were checked for normal distribution and square root transformed if necessary. These data, as well as qRT-PCR data, were then analyzed by using a 1-way ANOVA, followed by Dunnett’s *post hoc* test. Values of  $P < 0.05$  were considered statistically significant.

## Study approval

All animal experiments were reviewed with the Named Training and Competency Officer and Named Veterinary Surgeon at the University of Edinburgh before commencing these studies.

TABLE 3. Chromatographic conditions (flow rate 0.5 ml/min)

| Time (min) | Mobile phase A: water (0.1% formic acid, v/v) | Mobile phase B: acetonitrile (0.1% formic acid, v/v) |
|------------|---|--|
| 0          | 50  | 50   |
| 1          | 50  | 50   |
| 4          | 0   | 100  |
| 5          | 0   | 100  |
| 5.1        | 50  | 50   |
| 6.5        | 50  | 50   |

## RESULTS

### Simvastatin treatment reduces the incidence of PTB in an LPS-induced mouse model

To initiate PTB, mice received intrauterine LPS *via* ultrasound guidance. As expected, and as we have previously reported (24), LPS administration induced significantly earlier delivery than when mice were treated with PBS (mean time to delivery,  $29.7 \pm 3.6$  vs.  $63.5 \pm 2.3$  h;  $P < 0.0001$ ) (Table 4 and Supplemental Fig. 1A). However, when mice were treated with 20 or 40  $\mu$ g simvastatin in addition to LPS, the rate of PTB was reduced, and the mean time to delivery was significantly increased to  $46.9 \pm 6.5$  and  $45.3 \pm 4.7$  h, respectively ( $P = 0.0383$  and  $0.0469$  vs. LPS).

Consistent with adverse effects of inflammation-induced PTB, a substantial reduction in the percentage of live-born pups was observed after LPS administration: a mean of 18.4% of pups per litter survived delivery compared with 87.7% in the PBS group ( $P < 0.0001$ ) (Table 5 and Supplemental Fig. 1B). In mice exposed to LPS, simvastatin treatment resulted in more live-born pups, but this difference did not reach statistical significance. As expected, mean serum progesterone concentrations paralleled time to delivery (Supplemental Fig. 2A).

### Simvastatin treatment reduces systemic inflammation as well as uterine inflammation and contraction-associated gene expression

LPS stimulated a robust elevation of IL-6 in maternal serum ( $P < 0.0001$  vs. PBS alone) (Fig. 1A), which was significantly attenuated by simvastatin 40  $\mu$ g ( $P = 0.0213$  vs. LPS). In contrast, LPS administration did not alter IL-6 levels in the amniotic fluid, and simvastatin treatment had no additional effect (Fig. 1B).

We then investigated inflammatory and contraction-associated mRNA expression in the mouse uterus (Fig. 1C–H). Unsurprisingly, LPS treatment up-regulated the expression of *Il-6* ( $P = 0.0005$ ), *Il-10* ( $P = 0.0366$ ), *Cxcl1* ( $P = 0.0005$ ), and *Ccl2* ( $P = 0.0002$ ) in the mouse uterus. Treatment with simvastatin 40  $\mu$ g significantly down-regulated these genes compared with treatment with LPS alone: *Il-6*,  $P = 0.0095$ ; *Il-10*,  $P = 0.0328$ ; *Cxcl1*,  $P = 0.0464$ ; and *Ccl2*,  $P = 0.0342$ . In addition, the expression of gap junction gene *Cx43* was down-regulated with simvastatin 40  $\mu$ g treatment ( $P = 0.0143$  vs. LPS). *Cox-2* mRNA expression was unaltered by LPS and simvastatin treatment. Additional inflammatory genes (*Il-1 $\beta$* , *Tnf*, and *Cxcl2*) were

TABLE 4. Time to delivery

| Group                     | Time to delivery (h) vs. LPS |
|---------------------------|------------------------------|
| LPS                       | 29.74 ± 3.6                  |
| Simvastatin (20 µg) + LPS | 46.86 ± 6.5*                 |
| Simvastatin (40 µg) + LPS | 45.3 ± 4.7*                  |
| PBS                       | 63.54 ± 2.3****              |
| Simvastatin (20 µg) + PBS | 59.05 ± 2.2**                |
| Simvastatin (40 µg) + PBS | 52.84 ± 2.7                  |

\* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\*\* $P < 0.0001$ .

up-regulated by LPS but unaffected by simvastatin treatment (Supplemental Fig. 2). Protein expression of IL-6 and CX43 in the uterus was unchanged by LPS as well as by simvastatin treatment (Supplemental Fig. 3).

### Simvastatin significantly reduces proinflammatory mediator mRNA and protein expression in human myometrial cells

We then investigated the effect of simvastatin on inflammation *in vitro* in a human myometrial cell line treated with simvastatin (0.1, 10, 50 µM) either simultaneously with LPS (cotreatment) (Supplemental Fig. 4), 6 h before LPS (pretreatment) (Supplemental Fig. 5), or 6 h after LPS (post-treatment) (Fig. 2), for a total of 24 h. Cell metabolic activity, as a measure of cell viability, was unaffected by each of these treatments (Supplemental Fig. 6).

IL-6 mRNA expression was >6-fold greater in human myometrial cells after LPS stimulation ( $P < 0.0001$  vs. vehicle). This increase was significantly attenuated by 50 µM simvastatin in the cotreatment, pretreatment, and posttreatment groups ( $P < 0.0001$ ,  $P = 0.0007$ , and  $P = 0.0003$ , respectively) (Fig. 2A and Supplemental Figs. 4A and 5A). We found a robust upregulation (>50-fold) of IL-8 mRNA expression with LPS stimulation in each treatment group ( $P < 0.0001$  vs. vehicle) (Fig. 2B and Supplemental Figs. 4B and 5B). This up-regulation was significantly reduced with 50 µM simvastatin treatment in both the cotreatment ( $P = 0.0312$ ) and post-treatment ( $P = 0.041$ ) groups compared with LPS alone. We also observed a reduction in IL-6 secretion with 50 µM simvastatin and LPS cotreatment ( $P = 0.0015$  vs. LPS alone) (Supplemental Fig. 4C).

These effects on inflammatory mRNA expression were also observed at the protein level. We found a reduction in IL-6 secretion with 50 µM simvastatin and LPS cotreatment ( $P = 0.0015$  vs. LPS alone; Supplemental figure 4C). However, this reduction was not observed in the pre- and post-treatment groups (Supplemental Fig. 5C and Fig. 2C). There was a reduction in IL-8 secretion with both cotreatment ( $P = 0.0207$ ) and posttreatment but not pretreatment with 50 µM simvastatin ( $P = 0.0427$ ; Fig. 2D and Supplemental Figs. 4D and 5D).

### Simvastatin significantly increases anti-inflammatory cytokine mRNA expression in human myometrial cells

Interestingly, we observed that 10 µM simvastatin alone was able to up-regulate the expression of the anti-inflammatory

genes IL-10 and -13 ( $P = 0.0003$  and  $0.0008$  vs. LPS, respectively) (Supplemental Fig. 4E, F). Cotreatment with LPS and both 10 µM simvastatin and 50 µM simvastatin up-regulated IL-10 and -13 mRNA expression compared with LPS alone ( $P = 0.0064$  and  $0.0021$ ). These effects were similar regardless of treatment regimen, although some of the differences were not statistically significant (Fig. 2E, F and Supplemental Fig. 5E, F).

### Simvastatin treatment significantly inhibits both basal and LPS-induced contraction of human myometrial cells

Human myometrial cells were embedded in rat tail collagen to assess the effect of simvastatin treatment on the capacity of these cells to contract either alone or within an LPS-stimulated inflammatory environment. Vehicle-treated myometrial cells established a basal contraction, causing the gel to reduce in size. This scenario was evident within 24 h, as the mean gel size reduced to  $59.8 \pm 1.9\%$  of the baseline gel size (Fig. 3A–C). Simvastatin inhibited the basal contraction of the myometrial cells, resulting in substantially larger gels compared with the vehicle gels. This anti-contractile effect was observed with both 10 µM simvastatin ( $69.7 \pm 3.1\%$ ,  $P = 0.0059$  vs. vehicle) and 50 µM simvastatin treatment ( $67.5 \pm 2.0\%$ ,  $P = 0.027$  vs. vehicle). Stimulation with LPS induced further myometrial cell contraction, resulting in a 10.3% smaller gel area than the vehicle at 24 h ( $54.2 \pm 1.8\%$ ,  $P = 0.0293$  vs. vehicle). However, when simvastatin was coadministered with LPS, contraction was inhibited. Gel areas were significantly greater with 10 µM simvastatin ( $70.2 \pm 1.8\%$ ,  $P = 0.0059$ ) and 50 µM simvastatin treatment ( $70.3 \pm 1.7\%$ ,  $P = 0.0059$ ) compared with the vehicle. After 48 h, the vehicle mean gel size reduced further, and simvastatin treatment continued to inhibit contraction, both alone and in the presence of LPS (Fig. 3D).

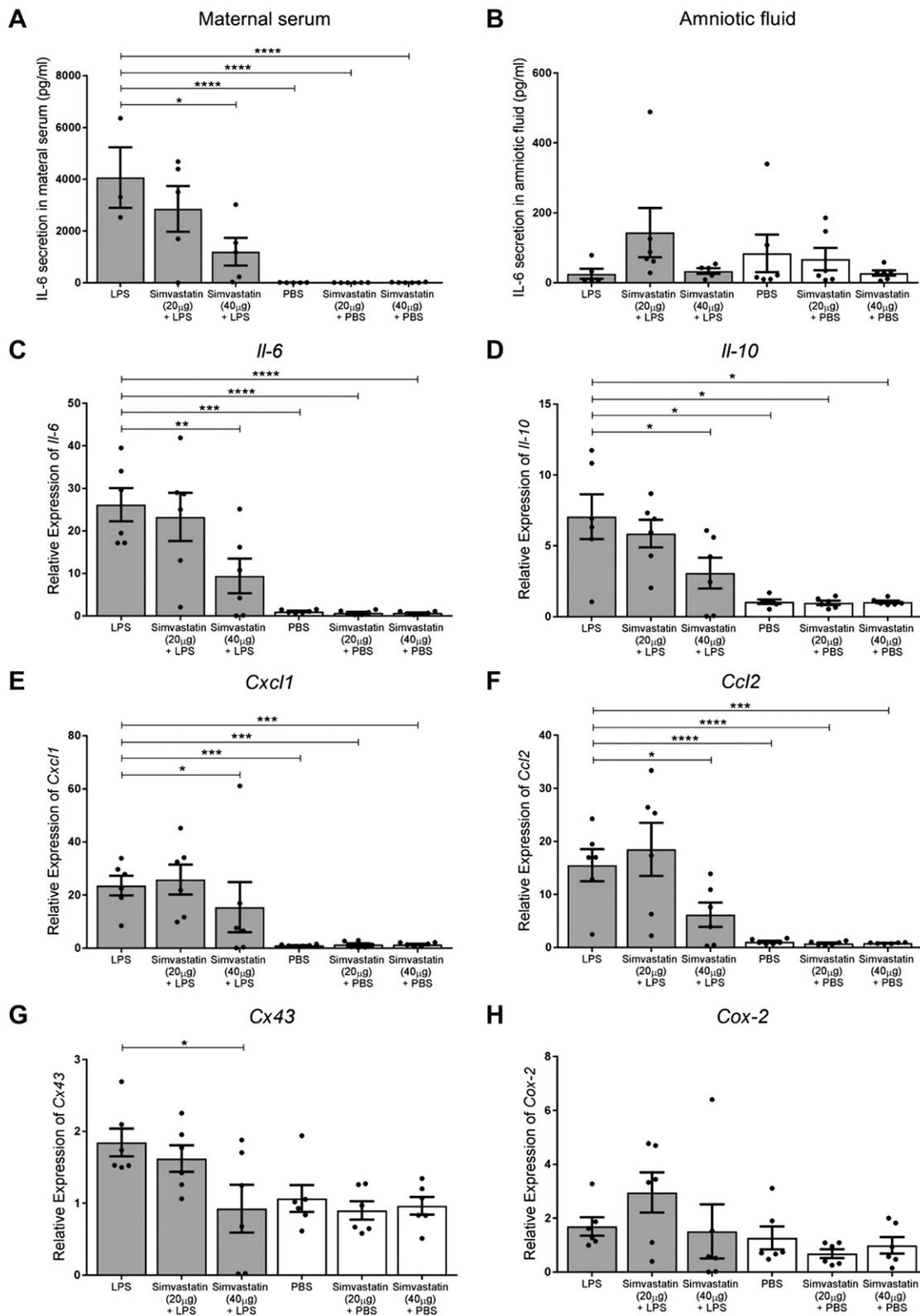
### Mevalonate and GG-PP but not F-PP supplementation abolish the anticontraction action of simvastatin

To investigate a mechanism by which simvastatin may be exerting its anticontraction effect, we supplemented the collagen gels with metabolites from the mevalonate pathway, which is the metabolic pathway that statins inhibit. When mevalonate was added with either 10 or 50 µM simvastatin, the gels contracted, abolishing the inhibitory effect of simvastatin on contractions. This effect was observed at both 24 and 48 h, with no significant

TABLE 5. Percentage of live-born pups

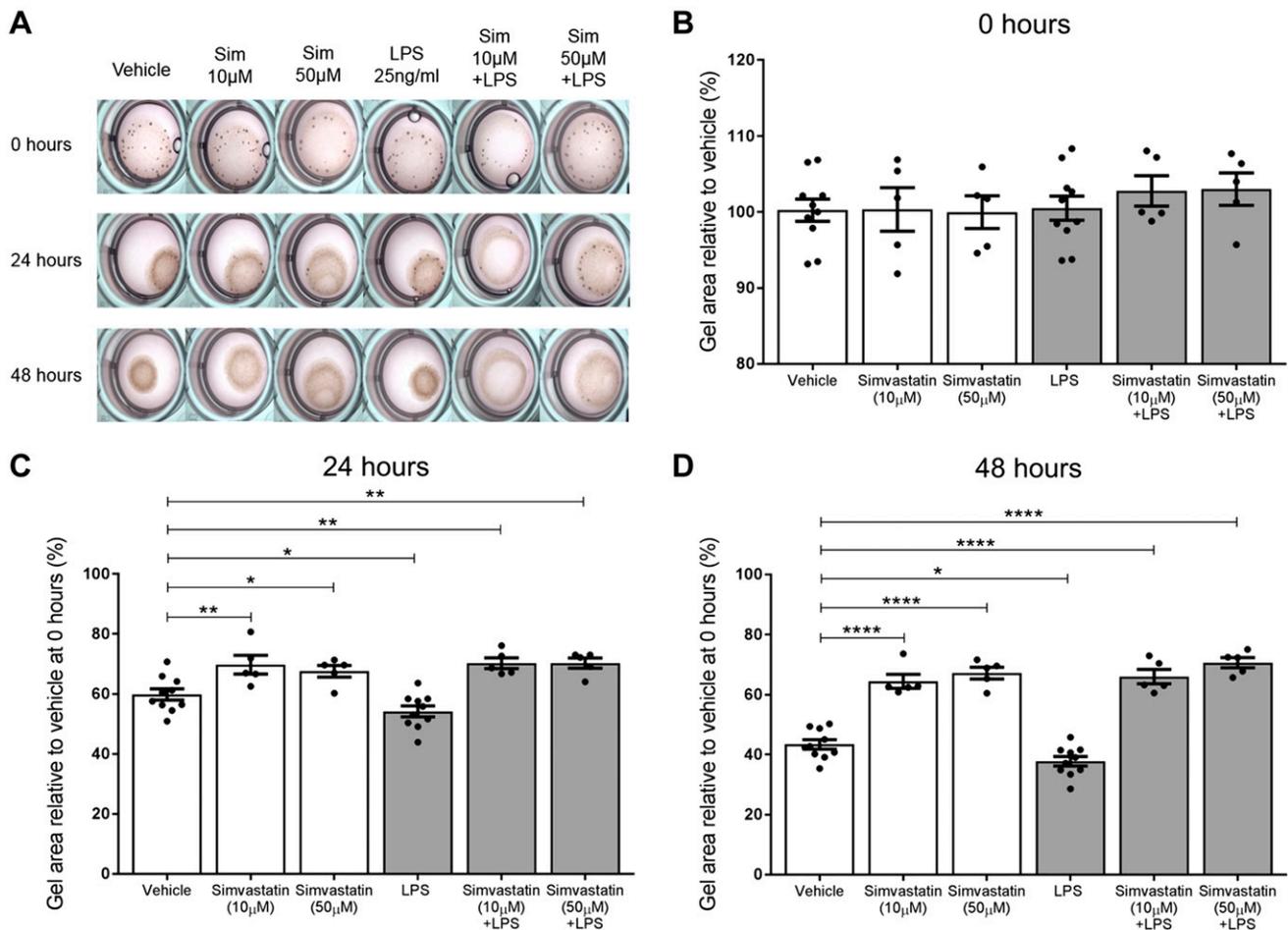
| Group                     | Live born pups (%) vs. LPS |
|---------------------------|----------------------------|
| LPS                       | 18.43 ± 7.6                |
| Simvastatin (20 µg) + LPS | 40.39 ± 10.2               |
| Simvastatin (40 µg) + LPS | 39.18 ± 9.1                |
| PBS                       | 87.7 ± 4.7****             |
| Simvastatin (20 µg) + PBS | 88.29 ± 7.6****            |
| Simvastatin (40 µg) + PBS | 76.22 ± 7.6****            |

\*\*\*\* $P < 0.0001$ .



**Figure 1.** Simvastatin reduces circulating IL-6 concentration and down-regulates uterine inflammatory and contraction-associated mRNA expression. *A*) IL-6 concentration in the maternal serum was elevated 6 h after LPS treatment ( $P < 0.0001$ ), which was attenuated by 40  $\mu$ g simvastatin ( $P = 0.0213$  vs. LPS);  $n = 3\text{--}6/\text{group}$ , 1-way ANOVA with Dunnett's *post hoc* test. *B*) IL-6 concentration in the amniotic fluid was unaffected by LPS or simvastatin;  $n = 5\text{--}6/\text{group}$ . *C*–*F*) Uterine inflammatory mRNA expression was significantly greater in LPS-treated mice: *Il-6*,  $P = 0.0005$  (*C*); *Il-10*,  $P = 0.0366$  (*D*); *Cxcl1*,  $P = 0.0005$  (*E*); and *Ccl2*,  $P = 0.0002$  (*F*). Simvastatin (40  $\mu$ g) significantly reduced the expression of these inflammatory mediators (*Il-6*,  $P = 0.0095$ ; *Il-10*,  $P = 0.0328$ ; *Cxcl1*,  $P = 0.0464$ ; and *Ccl2*,  $P = 0.0342$ ), as well as contraction-associated gene *Cx43* ( $P = 0.0143$ , *G*), compared with LPS alone. *H*) *Cox-2* expression was not altered by LPS or simvastatin;  $n = 6/\text{group}$ , 1-way ANOVA with Dunnett's *post hoc* test. All data are means  $\pm$  SEM. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$ .





**Figure 3.** Simvastatin treatment attenuates basal and LPS-induced contraction of myometrial cells embedded in collagen. *A, B*) Because these are smooth muscle cells, they established basal contraction, causing the gel size to reduce over time. *C, D*) Simvastatin inhibited basal contraction of the myometrial cells, resulting in substantially larger gels compared with the vehicle gels at both 24 h (10  $\mu$ M,  $P = 0.0059$ ; 50  $\mu$ M,  $P = 0.027$ ) and 48 h (10  $\mu$ M,  $P < 0.0001$ ; 50  $\mu$ M,  $P < 0.0001$ ). Stimulation with LPS induced further contraction, and the gel size was smaller than the vehicle at 24 h ( $P = 0.0293$ ) and 48 h ( $P = 0.0247$ ). When the gels were treated with simvastatin and LPS, contraction was inhibited, and the gel sizes were greater than the vehicle at 24 h (10  $\mu$ M,  $P = 0.0059$ ; 50  $\mu$ M,  $P = 0.0059$ ) and 48 h (10  $\mu$ M,  $P < 0.0001$ ; 50  $\mu$ M,  $P < 0.0001$ ). PBS and LPS groups,  $n = 10$ ; all other groups,  $n = 5$  (4–6 replicates). Mean % relative to vehicle mean at 0 h  $\pm$  SEM, 1-way ANOVA with Holm-Sidak *post hoc* test. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.0001$ .

expression of pMLC (Fig. 7A, B). Once again, when cells were treated with simvastatin and received either mevalonate or GG-PP supplementation, the effect of simvastatin was abolished, and the pMLC levels were akin to the vehicle-treated cells (Fig. 7A, C, D). When cells were treated with simvastatin and supplemented with F-PP, pMLC remained significantly reduced compared with the vehicle (10  $\mu$ M simvastatin,  $P = 0.0452$ ; 50  $\mu$ M simvastatin,  $P = 0.0008$ ). Thus, the addition of F-PP did not prevent simvastatin from inhibiting the phosphorylation of MLC (Fig. 7A, E), complementing the results of the collagen gel contraction assay.

## DISCUSSION

We present comprehensive evidence from preclinical models that statins (*e.g.*, simvastatin) should be considered as therapeutic agents for the treatment of PTL in women. Specifically, we have shown that treatment with simvastatin reduces the incidence of early delivery in a robust and

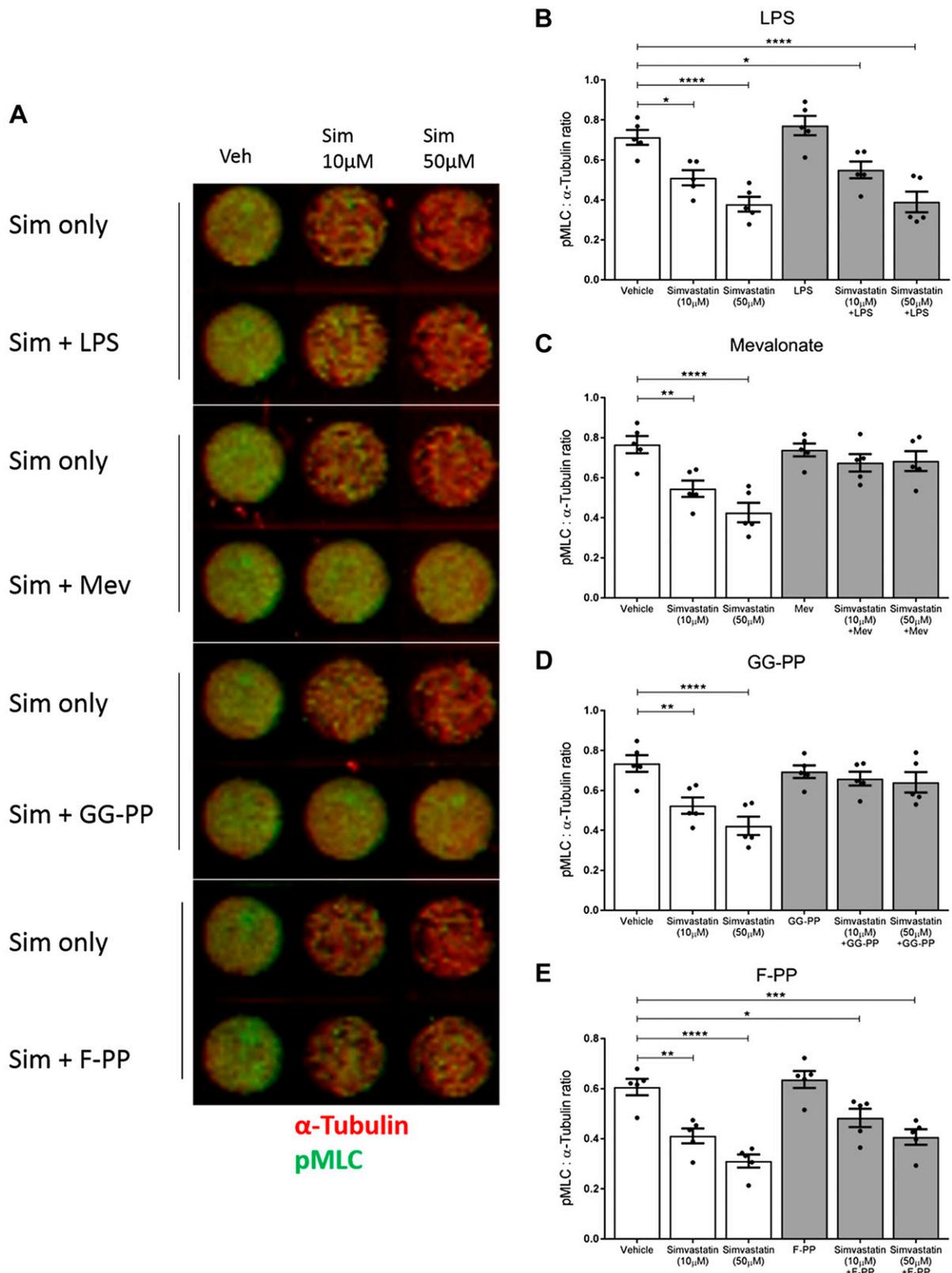
reproducible mouse model of PTB. We speculate that the ability of simvastatin to reduce systemic and uterine inflammation in this model may explain the lower incidence of PTB. These findings appear to agree with preliminary findings reported by Gonzalez *et al.* (16) whereby 20  $\mu$ g simvastatin was found to prevent PTB in a mouse model of PTB induced by intravaginal administration of LPS (serotype 055:B5). Unfortunately, neither we nor any other group has been able to replicate PTB through vaginal administration of LPS (24). Our study not only used a validated model of PTB in a larger cohort of mice, but we report on the survival of the pups immediately after delivery, identify an inhibitory effect of simvastatin on maternal inflammation, and defined the mechanism. These studies pave the way for definitive human studies of the effects of statins on PTL.

Interestingly, simvastatin did not have a dose-dependent effect on time to delivery or the percentage of live-born pups. However, the higher dose of 40  $\mu$ g simvastatin was more effective than the 20- $\mu$ g treatment at









**Figure 7.** Simvastatin reduces pMLC concentration, which is rescued by mevalonate and GG-PP but not F-PP supplementation. *A, B*) Simvastatin treatment reduced the phosphorylation of MLC both alone (10  $\mu$ M,  $P = 0.0112$ ; 50  $\mu$ M,  $P < 0.0001$ ) and in the presence of LPS (10  $\mu$ M,  $P = 0.0493$ ; 50  $\mu$ M,  $P < 0.0001$ ). LPS itself did not affect the phosphorylation of MLC. *A, C, D*) Mevalonate and GG-PP supplementation abolished the effect of simvastatin, and the pMLC concentrations were similar to the vehicle-treated cells. *A, E*) F-PP supplementation did not alter the action of simvastatin, and pMLC concentration remained significantly reduced compared with the vehicle (10  $\mu$ M,  $P = 0.0452$ ; 50  $\mu$ M,  $P = 0.0008$ ). All data:  $n = 5$ /group (in triplicate); means  $\pm$  SEM, 1-way ANOVA with Dunnett's *post hoc* test. \* $P < 0.05$ , \*\* $P < 0.01$ , \*\*\* $P < 0.001$ , \*\*\*\* $P < 0.0001$ .



In summary, the present study provides the rationale for further evaluation of the efficacy of simvastatin as a novel treatment for PTL. We have shown that simvastatin treatment reduced the incidence of PTB in an intrauterine, LPS-induced mouse model, as well as reduced inflammation in the uterus of these mice. Simvastatin also reduced LPS-induced inflammation in human myometrial cells and inhibited the contraction of these cells, both basally and within an inflammatory environment, *via* the inhibition of the Rho/ROCK pathway. Simvastatin treatment exhibited a number of useful properties, suggesting this drug would be an ideal candidate for the treatment of PTL by targeting underlying inflammation, inhibiting myometrial contractions, and subsequently preventing PTB. FJ

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## AUTHOR CONTRIBUTIONS

A. K. Boyle wrote the manuscript and was responsible for the investigation, methodology, and formal analysis of the study; S. F. Rinaldi contributed to the methodology, formal analysis, and supervision; A. G. Rossi and P. T. K. Saunders contributed to the conceptualization, formal analysis, supervision, and the review and editing of the manuscript; and J. E. Norman contributed to the conceptualization, formal analysis, supervision, funding acquisition, and the review and editing of the manuscript.

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