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The inwardly rectifying K⁺ channel KIR7.1 controls uterine excitability throughout pregnancy

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Abstract

Abnormal uterine activity in pregnancy causes a range of important clinical disorders, including preterm birth, dysfunctional labour and post-partum haemorrhage. Uterine contractile patterns are controlled by the generation of complex electrical signals at the myometrial smooth muscle plasma membrane. To identify novel targets to treat conditions associated with uterine dysfunction, we undertook a genome-wide screen of potassium channels that are enriched in myometrial smooth muscle. Computational modelling identified Kir7.1 as potentially important in regulating uterine excitability during pregnancy. We demonstrate Kir7.1 current hyper-polarizes uterine myocytes and promotes quiescence during gestation. Labour is associated with a decline, but not loss, of Kir7.1 expression. Knockdown of Kir7.1 by lentiviral expression of miRNA was sufficient to increase uterine contractile force and duration significantly. Conversely, overexpression of Kir7.1 inhibited uterine contractility. Finally, we demonstrate that the Kir7.1 inhibitor VUS90 as well as novel derivative compounds induces profound, long-lasting contractions in mouse and human myometrium; the activity of these inhibitors exceeds that of other uterotonic drugs. We conclude Kir7.1 regulates the transition from quiescence to contractions in the pregnant uterus and may be a target for therapies to control uterine contractility.

Keywords pregnancy; parturition; potassium channels; uterus; myometrium

Subject Categories Pharmacology & Drug Discovery; Urogenital System

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Introduction

At the end of pregnancy, the quiescent uterus must become highly contractile to mediate parturition. While the mechanisms initiating parturition in mammals are diverse (Smith, 2007), a final common pathway of uterine transition to a contractile phenotype appears to converge on the expression of a group of proteins, including the oxytocin receptor, prostaglandin endoperoxidase synthase 2, ion channels and connexin 43, that alter the uterine myometrial smooth muscle (MSM) cell from a state of low intrinsic excitability and refractoriness to stimulation, to a state that has high intrinsic excitability and is susceptible to stimulation (Garfield et al., 1977; Fuchs et al., 1982; Slater et al., 1995; Garfield & Maner, 2007). Control of stimulation is itself subject to complex gene-environment interactions, which act either independently of, or complimentary to, the underlying physiological changes of the uterus (Cha et al., 2013).

Irrespective of the underlying stimulus, the regulation of the duration and frequency of the myometrial contraction critically depends on control of calcium entry through voltage-gated L- and T-type calcium channels (Amedee et al., 1987; Blanks et al., 2007). Calcium is not only an important second messenger in the generation of force via calcium-calmodulin-dependent myosin light chain kinase, but also depolarizes the plasma membrane, allowing for activation of other voltage-dependent ion channels (Word et al., 1994; Brainard et al., 2007). This voltage-mediated control of uterine excitability is modulated in a gestation-dependent manner in all mammalian species. In particular, mid-gestation is characterized by a hyperpolarized membrane potential close to the reversal potential for potassium $E_k$ (Casteels & Kuriyama, 1965). As pregnancy progresses towards term, the myometrium becomes increasingly...
depolarized, to approximately −45 mV at parturition (Casteels & Kuriyama, 1965; Parkington et al, 1999). The mechanism underpinning this crucial, evolutionarily conserved process remains unknown.

A number of potassium channels have been demonstrated to play a role in shaping the myometrial action potential and modulating myometrial contractility (Brainard et al, 2007). The putative roles for the different channels are diverse and depend on the physiological environment. These roles range from voltage-dependent modulation of the action potential waveform (Knock et al, 1999), to modulating responses to intracellular calcium release through BKCa and SK3 (Khan et al, 1993; Pierce et al, 2008), intracellular ATP concentration through KATP (Khan et al, 1998) and uterine stretch through tandem pore channels (Tichenor et al, 2005).

Within the potassium channel super-family, inwardly rectifying potassium channels represents good candidates for the regulation of resting membrane potential. Kir7.1 is a member of the inwardly rectifying potassium channel sub-family and is only 38% identical to its closest relative, Kir4.2 (Doring et al, 1998; Krapivinsky et al, 1998; Pariseti et al, 1998). Kir7.1 is expressed in visceral tissues and some neurones within the CNS (Krapivinsky et al, 1998; Nakamura et al, 1999; Ookata et al, 2000; Shimura et al, 2001), although little is known about its functional role. Studies in retinal pigment epithelial cells indicate that the channel may play an integral role in setting resting membrane potential and modulating K+ recycling (Shimura et al, 2001). A rare genetic mutation in Kir7.1 causes autosomal-dominant snowflake vitreoretinal degeneration characterized by congenital degeneration of ocular tissues including the vitreous (Hejtmancik et al, 2008). Kir7.1 has also been linked to developmental pathways. For example, the jaguar/obelix mutation in zebra fish renders the Kir7.1 homologue non-functional. As a result, melanophores fail to respond appropriately to external cues, leading to melanosome aggregation and the phenotype of a broader striping pattern (Iwashita et al, 2006). Kir7.1 may also be involved in palate formation in mice. Kcnj13 was identified as one of 8 genes whose mis-expression correlates with formation of cleft palate in mice. Kir7.1 has also been linked to palate formation in mice.

To determine the functional significance of this K+ channel in the uterus, we first investigated its electrophysiological properties in freshly dissociated mouse MSMS. Under voltage-clamp conditions, an inwardly rectifying potassium current (Fig 2A,B) was inhibited by VU590, a known Kir7.1 inhibitor (Lewis et al, 2009). Consistent with the finding that Kir7.1 expression was higher at GD15 than GD18, the VU590 sensitive current density at −150 mV was significantly greater ($P < 0.05$) on GD15 when compared to GD18 (Fig 2C).

To understand the function of Kir7.1 in the generation of the myometrial action potential, we modelled the potential impact of changes in Kir7.1 channel density on myometrial electrogenesis using a Hodgkin-Huxley type current summing model. In free-running simulations of membrane potential, increasing Kir7.1 channel density (within the range measured in our experimental data) hyperpolarized resting membrane potential and decreased calcium entry during the action potential (Fig 3). Furthermore, simulations predicted that overall membrane conductance during the excited phase of the action potential is so finely balanced that small changes in Kir7.1 current density exerted large effects on membrane potential. Given its expression profile and biophysical properties, we hypothesized that Kir7.1 is a key regulator of myometrial membrane potential during gestation.

To assess the role of Kir7.1 in regulating uterine contractility experimentally, we used lentiviral vectors expressing miRNA targeting Kcnj13 or the human Kir7.1 channel to inhibit and over-express the channel in murine MSM, both in vitro and in vivo. Knockdown of Kir7.1, in vitro, significantly increased the contractile activity integral (2.5-fold) (area under the time-force curve), duration (2.2-fold) and maximum force (1.3-fold) of phasic contractions when compared to scrambled miRNA control (Fig 4 and Supplementary Fig S2). Conversely, over-expression of Kir7.1 significantly decreased all three parameters. To determine whether these alterations in contractility were due to electrogenic effects, we used sharp microelectrodes to measure membrane potentials in the treated myometrial strips. Knockdown of Kir7.1 depolarized resting membrane potential with extended excited periods (Fig 5A,B), whereas over-expression of Kir7.1 hyperpolarized resting membrane potential and suppressed excitability (Fig 5C). Mean resting membrane potentials differed significantly between the two treatment groups (Fig 5D).

Results

The aim of our study was to identify a potassium channel with the appropriate biophysical attributes to regulate the myometrial resting membrane potential during gestation. To identify candidates, we undertook a genome-wide qRT-PCR screen of all known K+ channels and associated subunits in cDNA pools generated from laser-capture micro-dissected MSM and whole myometrial tissue. Of seven transcripts enriched in MSM (Supplementary Table S1), only Kcnj13 coded for a K+ channel (Kir7.1) with appropriate biophysical attributes. Kcnj13 transcript levels increased markedly in the pregnant mouse uterus during mid-gestation, peaking on gestational day (GD) 15, and followed by a sharp decline towards term (C57BL/6J mice deliver in the morning of GD19; Fig 1A). Kcnj13 transcript levels (Fig 1B) were also significantly lower ($P < 0.05$) in samples taken from pregnant women in labour at term than in samples from women not in labour. Immunoblot of myometrial lysates from both labour and non-labour samples demonstrated a single immunoreactive band at −42 kDa. As positive and negative controls, we also tested human eye and adipose cell lysate, respectively (Fig 1C and Supplementary Fig S1). Furthermore, Kir7.1 immuno-reactivity was expressed in MSMs and was absent in the vasculature in both human and mouse myometrial samples (Fig 1D), supporting the specificity of the laser-capture screen.
Kir7.1 is expressed in uterine myocytes and is regulated in pregnancy in mice and humans.

A mRNA expression of Kcnj13 in mice (n = 5; mean ± SD, per GD) normalized to GD13. *P < 0.05, Student’s t-test.

B mRNA expression of KCNJ13 (plotted as arbitrary units relative to 18s rRNA) in human myometrial samples from women at term not in labour (NIL) and at term in labour (LAB) (n = 8, mean ± SD, per group). *P < 0.05, Student’s t-test.

C Immunoblot of pooled lysates from eye, adipose tissue, NIL and LAB myometrium (n = 4) probed with antibody to human Kir7.1 (full blot available in Supplementary Fig S1).

D Immunohistochemistry for Kir7.1 in human NIL myometrium (left panel), GD13 murine myometrium (centre panel) and GD15 murine myometrium (right panel). Arrow indicates absence of staining in blood vessel. Inset panels show tissue treated with pre-absorbed primary antibody control counterstained with haematoxylin.

Kir7.1 current in uterine myocytes decreases from mid-pregnancy to term.

A Measurement of inwardly rectifying, VU590-sensitive current in freshly dissociated GD15 murine myometrial cells. Shown are voltage-clamp recordings in the presence and absence of 10 μM VU590.

B Current-voltage relation (n = 5; mean ± SD, per data point) of current density [VU590 subtracted from control (vehicle alone)].

C Current density (pA/pF) at −150 mV and 500 ms in freshly dissociated murine myometrial cells from GD15 and GD18 (n = 5 mean ± SD, per GD). *P < 0.05, Student’s t-test.
Ion channel function in vivo may differ to that observed in vitro, because of either effects of the sample preparation, or influence of in vivo factors not captured in vitro. To account for these potential experimental confounders and to assess the phenotype of targeting Kir7.1 during pregnancy, we injected mice uteri on GD9 with anti-Kir7.1 miRNA or scrambled miRNA control. To record uterine activity, we surgically implanted a pressure catheter with remote telemeter into the injected horn of both knockdown and control mice. Intra-uterine pressure was recorded continuously, and the animals were monitored by video. Mice in which Kir7.1 was knocked down had significantly increased intrauterine pressure when compared to control mice (Fig 6 and Supplementary Fig S3). Consistent with our findings that Kir7.1 expression and channel activity were higher at GD15 than GD18, the effect on intrauterine pressure was more pronounced in midgestation.
The increased contractility associated with Kir7.1 knockdown raised the possibility that pharmacological targeting of this ion channel could be of clinical value, for example in the management of severe post-partum haemorrhage. To assess this possibility, we first performed current clamp microelectrode recordings of spontaneous action potentials in isolated murine myometrial strips. We observed slow, inter-contraction, depolarization of resting membrane potential followed by transient complex action potentials (Fig 7). Administration of 10 μM VU590 rapidly depolarized resting membrane potential to threshold (Fig 7Ai), followed by a sustained plateau potential that was reversible on wash out (Fig 7Aii and Aiii). In myometrial strips from women, application of VU590 increased the activity integral, largely due to an increase in contraction duration (Fig 7B), with contractions lasting several hours observed.

To assess the potential therapeutic benefit of inhibiting Kir7.1, we compared the effect of VUS90 with oxytocin, the established front-line treatment for post-partum haemorrhage. The effect of VUS90 on contractile force in samples taken from GD15 and GD18 mice was dependent on dose and gestation (Fig 8A) and when used in combination with oxytocin, increased activity integral by 172 ± 14 fold and 90 ± 42 fold on GD15 and GD18, respectively, as compared to 4 ± 2 fold and 8 ± 3 fold for OXT treatment alone (Fig 8B). The effect of VUS90 was similar in human term myometrium, which was also dose-dependent, with the observed increase in activity integral, largely due to an increase in contraction duration (Fig 8C,D). Importantly, sufficient channels remain at term to generate a significant phenotypic effect, underscoring the potential post-partum benefit of pharmacological intervention.
To expand the pharmacological tools for Kir7.1 inhibition, we used population patch-clamp technology to screen a library of known ion channel inhibitors in a Chinese Hamster Ovary cell line expressing Kir7.1 (Supplementary Fig S4 and Supplementary Methods). We identified compounds with different structures than VU590 to assess whether the phenotype of Kir7.1 inhibition was similar (Fig 9A). We also manufactured a control compound 1,3-Bis(5-nitro-1H-benzo[d]imidazol-2-yl)propane (BNBI), which is known to inhibit the structurally related Kir1.1 (Renal outer medulla potassium channel) but does not inhibit Kir7.1 (Bhave et al., 2011). Administration of 100 µM BNBI had no detectable effect on uterine contractility in the GD15 mouse (Fig 9B), suggesting that loss of efficacy for Kir7.1 inhibition is associated with loss of pro-contractile activity. Finally, administration of 10 µM MRT2000769, a potent Kir7.1 inhibitor that is structurally unrelated to VU590, induced long-lasting contractions similar to those observed with VU590 (Fig 9C). The effectiveness of MRT2000769 further supported the correlation between effective Kir7.1 inhibition and the phenotype of long-lasting contractions (Supplementary Fig S5).

Discussion

Inwardly rectifying potassium channels are known to regulate diverse but important physiological processes such as insulin secretion in the pancreas, regulation of the cardiac action potential, parasympathetic stimulation and potassium reuptake in the kidney (Hibino et al., 2010). In most cases, the channels act to hyperpolarize resting membrane potential by remaining persistently open, allowing the efflux of potassium. In excitable tissues, this dampens electrical activity, while in epithelial cells, the potassium gradient created in combination with energy-dependent ion pumps is used to transport other ions. In secretory cells, inhibition of the current causes depolarization and calcium entry leading to a secretion event (Liu et al., 2001; Ashcroft, 2005; Hebert et al., 2005). The important physiological roles of Kir channels are underscored by the many diseases that are associated with Kir channel malfunction, such as Bartter’s syndrome, Anderson syndrome, short Q-T syndrome and neonatal diabetes (Derst et al., 1997; Anderfingen et al., 2002; Edghill et al., 2004; Priori et al., 2005; Ellard et al., 2007).

Here we present the novel finding that Kir7.1 is a crucial regulator of membrane potential in uterine myocytes during pregnancy in both mice and humans. In mid-gestation, high expression of Kir7.1 keeps the resting membrane potential close to the reversal potential for potassium, increasing the depolarizing drive required to initiate an action potential, calcium entry and subsequent contraction. At term, this damping of excitability is lost by reduction, but not complete loss, of Kir7.1. Our results also indicate that inhibition of Kir7.1 when combined with oxytocin administration is synergistic, a mechanism that may be related to the sensitivity of Kir7.1 to intracellular phosphatidylinositol 4,5-bisphosphate depletion (Pattnaik & Hughes, 2009). Such a mechanism, when acting in conjunction with a decreased gap junction density and decreased receptors to stimulatory ligands, could provide a robust means of maintaining uterine quiescence during gestation (Garfield et al., 1977; Fuchs et al., 1982; Smith, 2007).

Within the Kir superfamily, Kir7.1 displays several unique properties such as low sensitivity to Ba²⁺ and Cs⁺, low single channel conductance, no internal block by Mg²⁺ ions, and a relative insensitivity to external K⁺ concentration, tetraethylammonium (IC₅₀ > 10 mM) or 4-aminopyridine (IC₅₀ ~10 mM) (Krapivinsky et al., 1998). In in vitro expression systems, Kir7.1 exhibits rapid
activation kinetics and is essentially non-inactivating at potentials negative to 40 mV with a small single channel conductance in both recombinant and native cells (Doring et al., 1998; Krapivinsky et al., 1998; Shimura et al., 2001). Since the channel has a low conductance and little voltage or time dependence in physiological ranges, high expression of this channel gives rise to a stable and hyperpolarized resting membrane potential. Thus, the biophysics of Kir7.1 is ideally suited to regulating myometrial smooth muscle cells during mid-gestation quiescence.

Transgenic mouse models have demonstrated the critical nature of control of myometrial membrane potential during parturition. For example, mice overexpressing the small conductance potassium channel SK3 suffer acute uterine dystocia, and both mother and pups die during delivery (Bond et al., 2000). Similarly, mice harbouring a smooth muscle specific deletion of the important uterine gap junction protein connexin43 demonstrate a significant delay in delivery and increased mortality of pups (Doring et al., 2006). These data, in conjunction with our in vitro observations demonstrating a significant increase in intraterine pressure in the absence of endocrine changes, underline the importance of the development of uterine excitability during gestation to the overall delivery process. It is notable that the decrease in Kir7.1 expression precedes progesterone withdrawal in the mouse suggesting that modulation of excitability is, at least in part, independent of progesterone. These biophysical factors act in addition to, and in concert with, other endocrine/paracrine changes that alter uterine stimulants (Cha et al., 2013) and lead to uterine disorders such as preterm labour or dysfunctional labour. The overall role of membrane potential in the control of parturition in the context of other controlling factors is summarized in Fig 10.

In addition to a clear effect on resting membrane potential, our computer modelling predicted that increasing Kir7.1 activity
modulates calcium entry during the plateau phase of the complex action potential, resulting in a decrease in contractile force and inefficient contractions. Decreasing Kir7.1 has the effect of stressing the repolarizing drive such that increased contributions from other K^+ channels are required to ensure action potential repolarization (Supplementary Fig S6) (Fink et al., 2006; Greenwood et al., 2009; McCallum et al., 2011). These effects explain the longer action potentials observed during pharmacological inhibition and experimentally induced reduction in the expression of Kir7.1 protein. When Kir7.1 current density is reduced to very low levels, the myometrial smooth muscle effectively loses phasic behaviour and manifests a new tonic-like tone. The greatly increased contraction length induced by Kir7.1 inhibition may be useful for reducing blood loss in an atonic uterus. As a treatment for post-partum haemorrhage, Kir7.1 block would have the advantage of being more potent than current first line treatments. The mechanism of action also circumvents the agonist pathways (Fig 10) targeted by current treatments that are prone to desensitization during failed labour inductions.

Materials and Methods

Ethical approval

All procedures involving women were conducted within the guidelines of The Declaration of Helsinki and were subject to local ethical approval (REC-05/Q2802/107). Prior to surgery, informed written consent for sample collection was obtained.

All animal procedures complied with the guidelines for the care and use of animals set forth by the National Institutes of Health. The Animal Studies Committee at Washington University in St. Louis approved all protocols (protocol number 20110138 to Sarah K. England). Adult C57BL/6J (Jackson Laboratory) female mice were mated at 8 weeks of age until 6 months of age. Mice were mated for 2-h time periods, and the presence of a copulatory plug was marked 0 days postcoitum (dpc). Animals were housed in Washington University School of Medicine vivarium in the BJCIH building, which is an AAALAC (Association for Assessment and Accreditation of Laboratory Animal Care International)
approved animal care facility. For in vivo intrauterine pressure measurements, eight pregnant mice were injected with scrambled miRNA and eight pregnant mice were injected with Kir7.1 miRNA.

**Subject criteria and selection**

Subjects were recruited into two groups, spontaneous labour (LAB) and elective caesarean section not in labour (NIL) between 38 and 40 weeks gestation. The LAB group was undergoing caesarean section for reasons of undiagnosed breech. LAB was defined as regular contractions (< 3 min apart), membrane rupture and cervical dilatation (> 2 cm) with no augmentation.

**Sample collection**

At caesarean section, samples were collected before syntocin administration by knife biopsy from the lower uterine segment incision. Samples were washed briefly in saline and flash-frozen in liquid nitrogen for mRNA, immunoblot or immunohistochemistry analyses. Samples for cell isolation or contractility experiments were placed in ice-cold modified Krebs–Henseleit (m-KHS) solution and used the same day.

**Laser capture screen**

mRNA was extracted from 100 mg of frozen human myometrium using Trizol reagent (Invitrogen), and further column purified by RNeasy kit (Qiagen) according to the manufacturer's instructions. Total RNA was quantified by spectrophotometer and further tested for quality and purity by bioanalyser (Agilent Technologies) according to the manufacturer's instructions and stored at −80°C until qRT-PCR analysis.

Laser Capture Microdissection (LCM): All slides, LCM caps, Haematoxylin and Eosin (H&E) Staining Kit for LCM and solutions were obtained from Molecular Machines & Industries. Briefly, cryomold-mounted myometrial samples were transferred from −70°C and equilibrated in a pre-cooled cryostat (−30°C) for 10 min. Sections (8 μm) were cut and H&E stained according to

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**Figure 9. Stimulation of uterine contractions correlates with Kir7.1 inhibitory potency.**

A The structures of the three compounds tested in this study. BNBI, a potent Kir7.1 inhibitor does not inhibit Kir7.1. MRT2000769 is structurally unrelated to VU590 and was identified as inhibiting Kir7.1 in a high throughput electrophysiology screen. VU590 is the first described inhibitor of Kir7.1.

B, C Effects of BNBI (B) and MRT2000769 (C) on murine GD15 myometrial contractility.
the manufacturer’s instructions. Slides were dehydrated in graded alcohols and xylene and placed in a desiccator until laser capture. At least 50,000 μm² of human uterine myocytes was captured for each gene. mRNA was extracted from each isolation by using the Picopure RNA isolation kit (Arcturus) according to the manufacturer’s instructions. Reverse transcription was as previously described. PCR primer design and qRT-PCR are described in the Supplementary Methods.

**Immunohistochemistry**

Frozen sections (8 μm) were fixed in ice-cold acetone for 5 min prior to incubation in phosphate-buffered saline (PBS). Endogenous peroxidase activity was blocked by immersing the slides for 30 min in freshly prepared 0.3% hydrogen peroxide in PBS. Slides were pre-incubated in 1.5% non-immune goat serum in PBS for 30 min at room temperature and then in primary anti-Kir7.1 antibody (1:100; Alomone) overnight at 4°C in a humidified chamber. Primary
antibody pre-absorbed for 24 h with recombinant antigen served as negative control. Staining was visualized using the Vectastain Elite ABC rabbit IgG kit (Vector Laboratories) according to the manufacturer’s instructions.

**Immunoblot**

After tissue was suspended in RIPA lysis buffer containing protease inhibitor cocktail tablets, mechanically homogenized and cleared by centrifugation, the supernatant was aliquoted and frozen. Whole tissue and protein lysates from adult human eye, adult mouse adipose and adult mouse brain were purchased from Novus Biologicals (Cambridge, UK) at a stock concentration of 5 mg/ml. All protein concentrations were confirmed with the BioRad assay (Bio-Rad laboratories, Hemel Hempstead, UK). A total of 40 µg of protein per sample (pooled from 4 biopsies each for NIL and LAB samples) was subjected to SDS-PAGE according to standard protocols. The membrane was blocked in 5% milk protein solution (Marvel, Lincs, UK) for 1 h at room temperature, incubated with primary rabbit polyclonal anti Kir7.1 antibody (1:200; Alomone Labs, Jerusalem) overnight at 4°C in blocking buffer and then incubated with polyclonal goat anti-rabbit HRP secondary antibody (1:100; Dako, Ely, UK). ECL Plus (GE Healthcare LTD, Amersham place, UK) was used to detect signal. To confirm equal loading, the blot was treated with Restore western blot stripping buffer (Thermo Fisher Scientific, Hemel Hempstead, UK) for 15 min at room temperature, washed, blocked and re-probed with an antibody to human β-actin.

**Electrophysiology**

**Cell isolation**

Strips of myometrium from the longitudinal layer (2 × 2 × 20 mm) of time-mated C57BL/6j mice were isolated and washed in Ca²⁺ and Mg²⁺ free Hank’s balanced salt solution (HBSS) at 37°C for 10, 20 and 30 min, respectively, followed by 45 min incubations in digestion solution (Roche Blendzyme 3) at 37°C according to the manufacturer’s instructions. Digestion was terminated by several dilutions with fresh HBSS. Cells were dispersed by slow trituration through a wide-bore fire-polished glass pipette in HBSS. Single myometrial cells were impaled with glass microelectrodes filled with 1 M KCl of resistance 80 MΩ. Transmembrane potentials were recorded using the perforated patch configuration of the whole cell patch-clamp technique (Rae et al, 1991). The antibiotic amphotericin B (720 µg/ml) was used to perforate the cell membrane. Currents were elicited by stepping to a range of potentials between −150 mV and +80 mV from a holding potential of −60 mV. To isolate currents that were sensitive to inhibition by drug application, difference currents were obtained by electronic subtraction of traces. Currents were filtered at 10 kHz and sampled at 5 kHz. Voltage protocols were delivered via a Digidata 1440a computer interface using pCLAMP 9.0 software (Molecular Devices, Sunnyvale, CA, USA).

**Current clamp**

Strips (5 × 10 mm) of murine myometrium from the longitudinal layer were pinned out on a sylgard base and perfused with m-KHS containing (in mM: NaCl, 133; KCl, 4.7; Tes, 10; glucose, 11.1; MgSO₄, 1.2; KH₂PO₄, 1.2; CaC₂, 2.5; adjusted to pH 7.4 at 25°C with NaOH) at 37°C on an upright microscope (MVX10, Olympus). Tissue was incubated with 5 µM wortmannin (Sigma) to prevent spontaneous contractions from dislodging impalements. Smooth muscle cells were impaled with glass microelectrodes filled with 2 M KCl of resistance 80–120 MΩ. Transmembrane potentials were recorded with an amplifier (Axopatch 700b; Axon Instruments) and a Digidata 1440a computer interface running pCLAMP 9.0 software.

**In vitro knockdown and contractility**

Strips of GD15 or GD18 myometrium from the longitudinal layer (2 × 2 × 20 mm) were washed in sterile Ca²⁺ and Mg²⁺ free HBSS DMEM/F-12. Strips were placed (in triplicate) under 1.5× slack length tension in media containing 2% dextran-coated charcoal-treated foetal bovine serum with 0.5 mM 8-bromo-cAMP (Sigma), 10⁻⁶ M medroxyprogesterone acetate (Sigma) and pLenti6-cppt-CMV-mCherry-mouse 543A miRNA (Anti-Kir7.1), pLenti6-cppt-CMV-mCherry-neg miRNA (Scrambled), or pLenti6-cppt-CMV/TOR-humKir7.1-IRES-mCherry-opre (+Kir7.1). Construction of lentiviral vectors is detailed in the Supplementary Methods.

On day five, strips were placed under 2mN tension in a four channel flatbed organ bath (DMT) in m-KHS solution. Isometric force was recorded on ADI Instruments LABCHART software. Activity integral was measured as the area under the time-force curve (mN/s) over a 20-min period. Contraction duration was determined as the mean duration (rise and fall to baseline) of contractions within a 20-min period. Maximum force was determined as the peak force measurement within a 20-min period. All measurements were made over the same time period for all strips.

**In vivo intrauterine pressure measurements**

C57BL/6j time-mated mice were anesthetized on GD8 to GD10 by intraperitoneal injection of ketamine (100 mg/kg) plus xylazine (10 mg/kg, IP), and a PhysioTel PA-C10 transmitter (Data Sciences International) was implanted in one horn of the pregnant uterus between the uterine wall and foetal sacs. At this time, 3.7 × 10⁷ particles/ml of pLenti6-cppt-CMV-mCherry-Neg miRNA (scrambled) or pLenti6-cppt-CMV-mCherry-miRNA mouse Kir7.1 (Anti-Kir7.1) were injected into the uterine muscle of the implanted horn. Five days later, uterine pressures were continuously measured every 10 s at 500 Hz with Dataquest A.R.T. data acquisition system version 4.31 (DSI) for 4–8 days.
The paper explained

Problem
Abnormal uterine activity has profound health consequences for both mother and infant as well as health services and national wealth. Clinical conditions associated with abnormal uterine activity are preterm labour (PTL), dysfunctional labour and post-partum haemorrhage (PPH). Preterm birth is the biggest cause of neonatal mortality and morbidity. Dysfunctional labour leads to operative vaginal and abdominal delivery with its inherent maternal risks, and PPH is a major global cause of maternal morbidity and mortality accounting for around 25% of deaths in post-partum mothers in developing nations. The challenge is to understand both the heterogeneous aetiology of these conditions and to develop new methods to manipulate uterine function.

Results
Uterine contractions are fundamentally controlled by complex electrical signals that regulate calcium entry at the plasma membrane of the uterine myocytes. Our findings have identified Kir7.1 as a novel regulator of electrical activity of the uterus during gestation in both mice and humans. We found that the expression of Kir7.1 was regulated in pregnancy and that decreasing Kir7.1 activity in vitro and in vivo greatly increased uterine activity. Conversely, increasing Kir7.1 expression inhibited uterine contractions. We found that this effect was mediated by altering the uterine resting membrane potential and action potential. Finally, we identified novel pharmacological inhibitors of Kir7.1 that stimulated uterine contractions with greater efficiency than the currently used uterine stimulant oxytocin.

Impact
Our findings have identified Kir7.1 as a key regulator of uterine activity during gestation. The results of this study add to our understanding of the underlying mechanisms that control uterine activity during gestation. The development of new inhibitors to Kir7.1 may be a novel tool for manipulating uterine function for the treatment of dystocia and post-partum haemorrhage.

Data analysis and statistics for telemetry studies
Longitudinal changes in intrauterine pressure after infection with pLenti6-cppt-CMV-mCherry-Neg miRNA (scrambled) or pLenti6-cppt-CMV-mCherry-miRNA mouse Kir7.1 (Anti-Kir7.1) were analysed by using a linear-mixed effect (LME) model implemented by PROC MIXED in SAS 9.3. Before analysis, data were first temporally aligned according to gestational days. The dependent (response) variable was taken as the hourly average intrauterine pressure. Fixed effects were treatment group (scrambled vs miRNA), gestation day and their interaction. Random intercepts were also used to incorporate mouse-specific effects. In addition, a first-order autoregressive correlation structure was used to account for the repeated measurements over time, which implies that the temporal correlation among repeated measures decays as a power function of the time lag. Residuals from initial LME analysis displayed a skewed distribution, and a square-root transformation of the response variable was found to be effective to correct the non-normality in the residuals. A few mice had negative hourly average pressures in certain hours, and 58 such observations (around 3% of the entire data set and split equally between the scrambled and miRNA group) were excluded from the analysis.

Compound screening
Cell line generation, cell culture, and Kir7.1 automated electrophysiology assay and data analysis are all described in the Supplementary Methods. The MRCT ion channel focused compound file was selected in collaboration with the Dundee Hit Finding Unit and consisting of ~4,000 compounds. The set covered 119 bioactive templates from nine categories of ion channel targets and molecular weights of the screening compounds ranged between 150 and 450.

Computational simulations
Computational simulations are described in detail the Supplementary data.

Supplementary information for this article is available online: http://embomolmed.embopress.org

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Author contributions
HAB, DAR, SKE, JSD, MJT, CK and AMB designed the research; CM, CR, EB, SM, JA, AS, YC, JZ, DT, JJ, PW, TD, SK, DJT, PB, RC and AMB carried out the research; SQ, MV and ST, phenotyped the subjects and provided the samples; HAB, CM, NL, SKE and AMB analysed the data; and AS, HAB, MJT, SKE, JJB and AMB wrote the paper.

Conflict of interest
The authors declare that they have no conflict of interest.

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