# and after $17\beta$ -estradiol stimulation

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Received 25 April 2002; revised 14 June 2002; accepted 14 June 2002

First published online 2 July 2002

Edited by Ned Mantei

Abstract Myometrial maxi-K channels are modulated by  $\beta$  subunits. We aimed to determine whether  $\beta$  subunits are modulated to affect uterine excitability during gestation. RNase protection analyses revealed that mouse  $\beta 1$  subunit transcripts are regulated during gestation with peak expression at day 14 of pregnancy. Immunohistochemical analysis indicates an increase of this subunit during gestation. Upregulation of the  $\beta 1$  transcript occurs with 4-day exposure to 17\beta-estradiol but not progesterone, and acute estradiol exposure has no effect on  $\beta 1$  transcript expression. These findings verify that  $\beta 1$  subunit transcript is regulated in mouse myometrium during gestation and estrogens may contribute to this increase. © 2002 Federation of European Biochemical Societies. Published by Elsevier Science B.V. All rights reserved.

*Key words:* Mouse; Uterus; Maxi-K;  $\beta$  Subunit; Gestation; Estrogen; Progesterone

### 1. Introduction

Large-conductance calcium- and voltage-activated potassium channels (maxi-K channels) play a prominent role in uterine contractility [1] by providing a potent repolarizing current that is essential for uterine quiescence. RNase protection assays (RPAs) and immunoblotting studies have shown that myometrial maxi-K channel  $\alpha$  subunit expression is modulated in mouse during gestation with expression levels increasing significantly at late gestation and decreasing post-partum [2]. However, using electrophysiological methods Benkusky et al. found a lower current density in 19-day pregnant mice [2]. Other studies by Wang et al., in late gestation rats, detected either no change in maxi-K current density in some cells, a change in the proportional amount the maxi-K current contributed to total current, or total absence of maxi-K in other cells [3], suggesting that the maxi-K channel current contributes less to the repolarizing current in uterine smooth muscle cells. One explanation for these findings is altered regulation of the maxi-K channel accessory  $\beta$  subunit, which enhances the channel's sensitivity to voltage and  $Ca^{2+}$  [4,5]. While studies of the maxi-K channel  $\alpha$  subunit indicate that this transcript is regulated by hormones involved in stress and pregnancy [2,6], recent evidence suggests that activation of the maxi-K channel by 17β-estradiol occurs when the hormone

binds to the channel's accessory  $\beta 1$  subunit [7]. Since estrogen levels increase at late gestation in the mouse [8], an increase in the  $\beta 1$  subunit could contribute to the potent repolarizing buffering current needed from late gestation until parturition commences. We provide evidence that the maxi-K channel  $\beta 1$ subunit transcript and protein is modulated during gestation in the mouse myometrium. Upregulation of the maxi-K channel  $\beta 1$  subunit transcript expression occurs with 4-day exposure to  $17\beta$ -estradiol in both ovariectomized and non-ovariectomized mice and is unaffected by progesterone.

#### 2. Materials and methods

2.1. Mouse breeding and  $17\beta$ -estradiol and progesterone stimulation

Adult C57BL/6J mice were mated at 8–10 weeks of age as previously described [2]. The uteri were excised and flash-frozen in liquid nitrogen.

For estradiol experiments, female C57BL/6J mice 8–10 weeks of age were divided into two groups. The non-stimulated group received injections of vegetable oil (vehicle) and the estrogen-stimulated group received 8.5  $\mu$ g injections of 17 $\beta$ -estradiol per day (Sigma, St. Louis, MO, USA) for 4 days based on a previously published protocol in rats [1]. The same protocol was administered to ovariectomized mice from Harlan Laboratories (Indianapolis, IN, USA). For acute stimulation studies, ovariectomized mice were injected with similar 17 $\beta$ estradiol concentrations and uteri excised 4 h after injection [9,10].

For progesterone experiments, the non-stimulated group received injections of vegetable oil, twice daily for 4 days. The progesteronestimulated group received a single injection of 8.5 µg 17β-estradiol, to upregulate progesterone receptors, followed by injections of 0.2 mg of progesterone (Sigma, St. Louis, MO, USA) for 4 days, twice daily [11,12]. All injections were administered subcutaneously. Mice were euthanized by CO<sub>2</sub> exposure either 24 h after (4-day exposure) or 4 h (acute) after the last injection and the uteri flash-frozen.

#### 2.2. RT-PCR

Oligonucleotide primer pairs flanking the coding region of the maxi-K channel  $\beta$  subunits ( $\beta$ 1– $\beta$ 4) were created. Mouse brain and myometrial RNA were extracted and separated as described previously [2]. Human brain RNA was purchased (Clontech, Palo Alto, CA, USA). The RNA was used for RT-PCR of the  $\beta$  subunits (Stratagene Prostar kit, La Jolla, CA, USA). The detected products were purified and sequenced.

#### 2.3. Generation of antisense RNA probes and RPAs

A biotin-labeled antisense probe corresponding to nt 1–195 of the  $\beta$ 1 subunit was synthesized using Ambion's Maxiscript in vitro transcription kit. The mouse cyclophilin loading control construct was a 103 nt fragment (Ambion). The RPAs were performed as previously described [2] using 25 µg of total RNA electrophoresed through 7.5% polyacrylamide/8 M urea denaturing gels at 250 V for 4 h. The blots were detected using chemiluminescence. Optical densities of the protected fragments were measured using LabWorks 3.0 (Ultra-Violet Products, Inc.) and normalized to the most intense signal or the loading controls. Mean normalized values are plotted  $\pm$  S.E.M.

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Fig. 1. Determination of transcript expression of the maxi-K channel  $\beta$ 1 transcript during gestation. 25 µg of total RNA isolated from mouse uteri at five gestational stages was hybridized with 1 ng of a biotin-labeled antisense RNA probe of the  $\beta$ 1 subunit. A: A representative blot shows the  $\beta$ 1 transcript (arrow) peaks at midgestation (P14) and decreases at post-partum day 2 (PP2). A probe against cyclophilin (Cyc) was used as a loading control. B: A densitometric summary shows the transcript expression level of the  $\beta$ 1 subunit during gestation. Optical densities of the protected fragments were measured and normalized to the most intense band (*n*=4). The asterisk denotes statistical significance of differences from the non-pregnant (NP) controls.

#### 2.4. Immunoblotting and immunohistochemistry

For immunoblotting and immunofluorescent experiments, the mouse maxi-K channel  $\beta$ 1 subunit polyclonal antibody (Affinity Bioreagents) was used at a 1:500 dilution. Western blot analysis was performed as described previously [2]. For immunofluorescence experiments, uteri isolated from mice were fixed in 4% paraformaldehyde and embedded in OCT. Cryopreserved tissues were sectioned in

10 nm sections, blocked in buffer containing 5% heat-inactivated fetal bovine serum and 0.3% bovine serum albumin and incubated with the maxi-K channel ß1 subunit (Affinity Bioreagents) for 2 h at room temperature. Sections were subsequently incubated with biotin-conjugated donkey anti-rabbit IgG (1:1000) for 30 min at room temperature (Jackson ImmunoResearch) and streptavidin-conjugate Cy5 (1 mg/ml) for 20 min at room temperature (Jackson Immuno-Research). Myometrial smooth muscle tissue was identified by Cy2 conjugate of mouse monoclonal anti- $\alpha$ -smooth muscle actin (Sigma, 1:500). Images were visualized and recorded using a laser confocal microscope and quantified with image analysis software (Zeiss). Control experiments were performed using a commercial competing peptide (100 µg/ml; Affinity Bioreagents) and by immunocytochemistry of cells transfected with a control or maxi-K channel ß1 adenoviral construct, both containing green fluorescent protein (GFP) as a reporter gene. Immunofluorescence intensity of the maxi-K channel ß1 signal was measured in fields of similar size at different stages of gestation. Results are expressed as average pixel intensity.

#### 2.5. Statistics

Significant differences between groups in the gestational RPAs were determined by one-way analysis of variance followed by Tukey–Kramer multiple comparisons test. Statistical significance of differences between groups in the estrogen-stimulated RPAs were determined by Student's *t*-test. Differences were considered significant at P < 0.05.

#### 3. Results

# 3.1. The maxi-K channel $\beta$ 1 subunit transcript peaks at mid-gestation

PCR analyses detected the presence of maxi-K channel B1 and  $\beta 2$  subunit transcripts in the mouse uterus, however the  $\beta$ 3 and  $\beta$ 4 subunit transcripts were not detected (data not shown). RPAs were performed to assess whether differential transcript expressions of the  $\beta 1$  and  $\beta 2$  subunits are potential mechanisms for changes in uterine excitability during gestation. Total RNA from the five gestational stages was hybridized with an antisense biotin-labeled RNA probe of the  $\beta$ 1 subunit. A representative RPA blot demonstrates that the  $\beta$ 1 subunit transcript peaks at mid-gestation (P14) and decreases by day 19 (arrow, Fig. 1A). This trend in transcript regulation differs from the regulation of the maxi-K channel  $\alpha$  subunit, which peaks at term gestation (P19) [2]. These data are summarized in Fig. 1B where the densities of the protected fragments of the  $\beta$ 1 subunit were measured and normalized to the most intense signal (n=4). Quantification confirms that the



Fig. 2. Maxi-K channel  $\beta$ 1 subunit antibody specificity. 25 µg of membrane protein from cells transfected with maxi-K channel  $\beta$ 1 and shamtransfected cells were probed with a polyclonal antibody for this protein (A). A band corresponding to ~30 kDa was detected in the  $\beta$ 1-transfected cells ( $\beta$ 1; arrow).  $\beta$ 1 was detected in cells transfected with a  $\beta$ 1 expression construct (red fluorescence; C) but not in cells transfected with a control vector (B). Both vectors express GFP (green fluorescence).





















Fig. 3. Immunohistochemical analyses demonstrate  $\beta$ 1 subunit protein localization and regulation in mouse myometrium during pregnancy. Cryosections of mouse uteri isolated at various stages of pregnancy were incubated with rabbit polyclonal  $\beta$ 1 antibody and donkey anti-rabbit Cy5 IgG (red). Myometrial smooth muscle tissue was identified by  $\alpha$ -smooth muscle actin fluorescence (green). Maxi-K channel  $\beta$ 1 subunit expression appears to increase from the non-pregnant state to higher levels mid-gestation. Expressions in both the longitudinal smooth muscle (L) and circular smooth muscle layer (C) are similar. The bars represent 20  $\mu$ m. Addition of the competing peptide block detected no  $\beta$ 1 (red fluorescence).





Fig. 5. 17β-Estradiol and progesterone regulation of the maxi-K channel ß1 transcript expression in mouse myometrium. RPAs compared transcript levels of the maxi-K channel β1 subunits following 17β-estradiol and progesterone stimulation (arrows). Representative blots (upper panels) and densitometric analyses (lower panels) show that the  $\beta$ 1 transcript increases in both non-ovariectomized (A) and ovariectomized (B) mouse myometrium after stimulation for 4 days with 17 $\beta$ -estradiol (ES or ES<sub>ovx</sub>) as compared to vehicle control (NS or NS<sub>ovx</sub>). B1 transcript expression does not change significantly with acute 17β-estradiol stimulation (C; ES<sub>ovx-a</sub>) or 4-day progesterone stimulation (D; PS) as compared to their respective vehicle controls (NS<sub>oyx-a</sub>, and NS). For all experiments, total mouse uterine RNA was hybridized with 1 ng of a biotin-labeled antisense RNA probe to the ß1 subunit. In densitometric analyses cyclophilin (Cyc) was used as the loading control and bands were normalized to cyclophilin (n = 3-8). Asterisks show statistically significant differences from the vehicle controls.



Fig. 4. Quantification of maxi-K channel  $\beta$ 1 subunit expression in mouse myometrium during gestation. The bar graph represents the average pixel intensities from three separate immunohistochemical experiments. Maxi-K channel  $\beta$ 1 subunit expression is increased in P7, P14, and P18 mice as compared to non-pregnant and post-partum day 2 stages.

transcript level of the  $\beta 1$  subunit is significantly increased throughout gestation (P7–P19) and decreases by post-partum day 2.

Although the  $\beta 2$  subunit transcript was detected by RT-PCR, it was not detected by RPA indicating that the  $\beta 2$  subunit is not present at high levels in the mouse myometrium during gestation or amplification occurred from contaminating non-myometrial tissue.

# 3.2. The maxi-K channel $\beta$ 1 subunit protein is upregulated during mid-gestation

Immunohistochemical experiments were performed to detect both  $\beta$ 1 subunit protein expression levels and localization. The antibody used reacted specifically with the  $\beta 1$  subunit in Western blots (Fig. 2A) and in cells transfected with a  $\beta$ 1 subunit expressing construct (Fig. 2B,C). As seen in Fig. 3, longitudinal smooth muscle (L) and circular smooth muscle (C) layers both express similar levels of the  $\beta$ 1 subunit (red) at different gestational stages. Similar to the transcript measurements, the  $\beta$ 1 protein expression is increased in pregnancy days 7, 14, and 18 as compared to non-pregnant and postpartum day 2 stages.  $\alpha$ -Actin levels (green) change during gestation, however, the signal from  $\beta$ 1 protein was exceedingly high compared to  $\alpha$ -actin, and  $\alpha$ -actin could be detected at all stages. Inclusion of the competing peptide abolished B1 fluorescence in myometrial tissue (PP2 peptide). The immunohistochemical results suggest that maxi-K channel B1 subunit protein is present in the mouse myometrium and is upregulated during pregnancy as compared to non-pregnant mice uteri.

Quantification of maxi-K channel  $\beta$ 1 using fluorescence microscopy demonstrates that this protein is upregulated during pregnancy (Fig. 4). There is a slight decrease at P14 compared to P7 and P18, which was not predicted by transcript levels (n= 3).

## 3.3. The maxi-K channel β1 subunit transcript is upregulated by 17β-estradiol

We sought to determine whether estrogen could contribute to increases in transcript expression of the  $\beta 1$  subunit seen during gestation. RPAs were performed on RNA from myometrium of mice that were stimulated 4 days with 17 $\beta$ -estradiol (ES) or the vehicle control (NS). Representative data hormones as RPAs performed on uteri from ovariectomized mice stimulated with  $17\beta$ -estradiol (ES<sub>ovx</sub>) or vehicle control (NS<sub>ovx</sub>) demonstrated similar results (n = 8, Fig. 5B). These data suggest that estrogen is a possible initiator of the increase in the  $\beta$ 1 subunit transcript during gestation.

# 3.4. Maxi-K channel $\beta$ I subunit transcript is not regulated by acute exposure to 17 $\beta$ -estradiol or progesterone

Previous studies have shown that 17 $\beta$ -estradiol has an acute effect on the maxi-K channel  $\beta$ 1 subunit [7]. To determine whether the increase in the  $\beta$ 1 subunit transcript following stimulation by 17 $\beta$ -estradiol is an acute effect, RPAs were performed on uterine RNA from ovariectomized mice stimulated for 4 h with 17 $\beta$ -estradiol (ES<sub>ovx-a</sub>) or vehicle control (NS<sub>ovx-a</sub>). Representative data shown in Fig. 5C indicate that the  $\beta$ 1 subunit transcript level does not change after 4-h stimulation with 17 $\beta$ -estradiol in ovariectomized mice compared to non-stimulated ovariectomized mice. Densitometric analyses of the blots (Fig. 5C, lower panel) demonstrate that short-term stimulation by 17 $\beta$ -estradiol has no significant effect on the  $\beta$ 1 transcript in contrast to the increase seen after longer stimulation (*n* = 3).

To assess whether the increase in the  $\beta$ 1 subunit transcript seen at late gestation was a general effect of hormonal changes, RPAs were performed on uterine RNA from mice treated with progesterone (PS) or vehicle control (NS). Fig. 5D shows a representative blot indicating that  $\beta$ 1 subunit transcript does not increase following progesterone stimulation. Densitometric analysis of the blots (bottom panel) in which the densities of the  $\beta$ 1 subunit bands were measured and normalized to its loading control (n = 3) shows that stimulation by progesterone has no significant effect on the transcript level of the  $\beta$ 1 subunit. Song et al. [13] have reported similar results with another myometrial channel transcript, Kv4.3, which is regulated by 17 $\beta$ -estradiol but not by progesterone.

### 4. Discussion

These experiments provide the first evidence that regulation of the modulatory maxi-K channel B1 subunit occurs during gestation in the mouse myometrium. During gestation, myometrial cell permeability to Ca<sup>2+</sup> increases, and membrane potential increases to more depolarized potentials at term [14]. Therefore, a maxi-K channel with increased sensitivity to voltage and Ca<sup>2+</sup> would maintain K<sup>+</sup> efflux promoting uterine quiescence during gestation. Alternatively spliced forms of the maxi-K channel  $\alpha$  subunits, which are sensitive to  $Ca^{2+}$  and voltage, are regulated during gestation [2]. Another mechanism for altering uterine excitability is by  $\beta 1$  subunit regulation. We have determined that one potential mechanism for the increase in  $\beta 1$  subunit transcript during gestation is via genomic effects of estrogen. Although previous results have shown short-term effects of estrogen on the  $\beta$ 1 subunit [7], acute stimulation had no effect on  $\beta$ 1 subunit transcript, suggesting that non-genomic effects are also involved in channel regulation. This appears to be estrogen specific as no effect on  $\beta 1$  transcript was observed after progesterone stimulation. In previous studies, doses of  $0.5-17 \ \mu g$ of estrogen administered to mice led to serum levels of 17β-estradiol of 45.6–513 pg/ml [15]. These levels are comparable to circulating plasma levels of 17β-estradiol during pregnancy [16], suggesting a physiological role of estrogen on maxi-K channel  $\beta$ 1 subunit regulation.

Protein regulation of the  $\beta$ 1 subunit may be similar to transcript regulation, however immunohistochemical analyses suggest that  $\beta$ 1 protein is slightly reduced at P14 as compared to P7 and P18. This is likely a result of the difficulty in accurately measuring fluorescent signals in a tissue that drastically changes anatomically. Immunohistochemical analyses suggest that the ratio of the maxi-K channel  $\alpha$  and  $\beta$ 1 subunits would likely change during gestation whereby the  $\alpha$  subunit increases throughout gestation peaking at day 19 [2] while the B1 subunit peaks earlier during pregnancy. Recent studies have demonstrated that in brain tissue maxi-K channel  $\alpha$  subunits can form functional channels devoid of the  $\beta$  subunit [17], while in vascular smooth muscle it is usually associated with the  $\beta$ 1 subunit [18], indicating that altered  $\alpha$  and  $\beta$  subunit association can occur. This stoichiometric change may indicate more maxi-K channels which are not associated with  $\beta$  subunits later in pregnancy thereby producing channels that are less sensitive to depolarization and increased [Ca<sup>2+</sup>]<sub>i</sub> caused by myometrial stretch. This decrease in the channel's sensitivity to voltage and  $Ca^{2+}$  may be a mechanism to increase the uterine excitability prior to parturition. The  $\beta$ 1 subunit transcript is regulated in the mouse myometrium during gestation, in part due to estrogen. The ability of estrogen to regulate this channel after both acute and longer exposure suggests a multifaceted role of this hormone. Further examination of the association of these channel subunit proteins throughout pregnancy will determine whether association of maxi-K channel  $\alpha$ and  $\beta$  subunits is altered and stoichiometric differences occur to modulate uterine excitability.

Acknowledgements: The authors thank Mr. Daniel Fergus for critical review of the manuscript. This work was supported by National Institutes of Health Grants HD-37831 and HD-01371 (to S.K.E.) and American Heart Association Postdoctoral Fellowship 0120580Z (to V.P.K.). Reagents for these studies were provided by the Diabetes and Endocrinology Research Center (DK-25295) and the Gene Transfer and Vector Core (DK-54759) at the University of Iowa.

#### References

- Anwer, K., Oberti, C., Perez, G.J., Perez-Reyes, N., McDougall, J.K., Monga, M., Sanborn, B.M., Stefani, E. and Toro, L. (1993) Am. J. Physiol. 265, C976–C985.
- Benkusky, N., Fergus, D., Zucchero, T. and England, S. (2000) J. Biol. Chem. 275, 27712–27719.
- [3] Wang, S., Yoshino, M., Sui, J., Wakui, M., Kao, P. and Kao, C. (1998) J. Gen. Physiol. 112, 737–756.
- [4] McManus, O., Helms, L., Pallanck, L., Ganetzky, B., Swanson, R. and Leonard, R. (1995) Neuron 14, 645–650.
- [5] Uebele, V., Lagrutta, A., Wade, T., Figueroa, D.J., Liu, Y., McKenna, E., Austin, C.P., Bennett, P.B. and Swanson, R. (2000) J. Biol. Chem. 275, 23211–23218.
- [6] Xie, J. and McCobb, D. (1998) Science 280, 443-446.
- [7] Valverde, M., Rojas, P., Amigo, J., Cosmelli, D., Orio, P., Bahamonde, M.I., Mann, G.E., Vergara, C. and Latorre, R. (1999) Science 285, 1929–1931.
- [8] Barkley, M., Geschwind, I. and Bradford, G. (1979) Biol. Reprod. 20, 733–738.
- [9] Parsons, B., Rainbow, T., Snyder, L. and McEwen, B. (1984) Neuroendocrinology 39, 25–30.
- [10] Sircar, R. and Kim, D. (1999) J. Pharmacol. Exp. Ther. 289, 54– 65.
- [11] Kurita, T., Lee, K., Saunders, P.T., Cooke, P.S., Taylor, J.A., Lubahn, D.B., Zhao, C., Makela, S., Gustafsson, J.A., Dahiya, R. and Chuha, G.R. (2001) Biol. Reprod. 64, 272–283.

- [12] Paria, B., Das, N., Das, S., Zhao, X., Dileepan, K. and Dey, S. (1998) Endocrinol. J. 139, 3958–3966.
   [13] Song, M., Helguera, G., Eghbali, M., Zhu, N., Zarei, M., Olcese,
- R., Toro, L. and Stefani, E. (2001) J. Biol. Chem.
  [14] Wray, S. (1993) Am. J. Physiol. 264, C1–C18.
- [15] Pelleymounter, M.A., Baker, M.B. and McCaleb, M. (1999) Am. J. Physiol. 276, E955–E963.
- [16] Parkening, T.A., Lau, I.-F., Saksena, S.K. and Chang, M.-C. (1978) J. Gerontol. 33, 191–196.
   [17] Chang, C.-P., Dworetzky, S., Wang, J. and Goldstein, M. (1997)
- Mol. Brain Res. 45, 33-40.
- [18] Tanaka, Y., Meera, P., Song, M., Knaus, H.-G. and Toro, L. (1997) J. Physiol. 502, 545–557.