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Expression of voltage-dependent potassium channels in first trimester human placentae			
Hiten D. Mistry <sup>1,2*</sup> , Lesia O.	Kurlak <sup>3</sup> , Guy S. Whitley <sup>4</sup> , Judith E. Cartwright <sup>4</sup> , Fiona Broughton		
Pipkin <sup>3</sup> & Rachel M. Tribe <sup>1</sup>			
<sup>1</sup> Division of Women's Health, King	's College London, Women's Health Academic Centre, KHP, SE1 7EH, UK;		
<sup>2</sup> Department of Nephrology, Hype	ertension, Clinical Pharmacology and of Clinical Research, University of Bern,		
CH-3010 Berne, Switzerland;			
<sup>3</sup> Department of Obstetrics & Gynaecology, School of Medicine, University of Nottingham, NG5 1PB, UK; <sup>4</sup> Division			
of Biomedical Sciences, St George	's University of London, SW17 ORE, UK.		
*Correspondence address:	Dr. Hiten D. Mistry		
	Division of Women's Health		
	King's College London		
	Women's Health Academic Centre, KHP		
	St Thomas' Hospital		
	Westminster Bridge Road		
	London, SE1 7EH, UK		
	Tel: 020 71888151; Fax: 020 7620 1227		
	Email: <u>hiten.mistry@kcl.ac.uk</u>		
<b>Running Title:</b> Placental K <sub>v</sub>	7 channels in early pregnancy		
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	Expression of voltage-dependence Hiten D. Mistry <sup>1,2*</sup> , Lesia O. Pipkin <sup>3</sup> & Rachel M. Tribe <sup>1</sup> <sup>1</sup> Division of Women's Health, King <sup>2</sup> Department of Nephrology, Hype CH-3010 Berne, Switzerland; <sup>3</sup> Department of Obstetrics & Gyna of Biomedical Sciences, St George <b>*Correspondence address:</b> <b>*Correspondence address:</b>		

# 23 Abstract

- 24 Potassium channel α-subunits encoded by KCNQ1-5 genes form voltage-dependent channels
- 25 (Kv7), modulated by KCNE1-5 encoded accessory proteins. The aim was to determine KCNQ
- 26 and KCNE mRNA expression and assess protein expression/localisation of the KCNQ3 and
- 27 KCNE5 isoforms in first trimester placental tissue. Placentae were obtained from women
- undergoing elective surgical termination of pregnancy (TOP) at  $\leq 10$  weeks' (early TOP) and >10
- 29 weeks' (mid TOP) gestations. KCNQ1-5 expression was unchanged during the first trimester.
- 30 KCNE5 expression increased in mid TOP vs. early TOP samples (P=0.022). This novel study
- 31 reports mRNA and protein expression of Kv7 channels in first trimester placentae.

32

### 34 Introduction

35 Potassium  $(K^+)$  channel expression is essential for normal physiological functions of endothelial and smooth muscle cells in a variety of vascular beds [1]. Members of the Kv7 voltage-gated 36 37 potassium channel subfamily Kv7.1-7.5 are encoded by KCNQ1-5 genes; the KCNQ-encoded α-38 subunits can form channel complexes with KCNE-encoded β-subunits (KCNE1-5). Alterations 39 in  $K_V7$  channel expression and their activation properties affect cell function, cell proliferation 40 and differentiation [2]. 41 42 Knowledge of  $K_V$  channels in the feto-placental circulation is limited [3, 4]. The presence of 43 functional  $K_V7$  channels in human chorionic plate arteries [5], suggests a role in control of 44 vascular tone. Perfusion studies in placental allantochorial blood vessels incubated with K<sup>+</sup> 45 channel blockers, exhibit responses to altered oxygenation, support this further [6-9]. 46 47 We have previously reported raised Kv7 mRNA and protein expression in placental tissue from 48 pre-eclamptic women compared to normotensive controls near term [3]. The reason for this 49 difference is unknown; however these channels may be involved in early placentation, which is 50 disrupted in pre-eclampsia. The aims of this study were to establish placental KCNQ/KCNE 51 mRNA expression profiles in early pregnancy and to compare with previous observations from 52 normotensive and pre-eclamptic women. 53 54 55

3

### 57 Materials and Methods

58 After local Ethical committee approval (Wandsworth Local Research Ethics Committee) and 59 with appropriate informed consent, placental chorionic villous tissue, was obtained from women 60 undergoing elective surgical TOP at St. George's Hospital, London during the early 1st trimester 61 (early-TOP; <10 weeks', ([mean  $\pm$  SD] 8.8  $\pm$  0.9 weeks); n = 6) and late 1<sup>st</sup> trimester (mid-TOP, 62 gestational age >10 weeks' ( $12.9 \pm 0.9$  weeks); n = 7). Samples were divided and either placed in 63 RNAlater (Qiagen, UK), stored at -80°C or fixed in formalin and wax embedded. 64 We compared the early pregnancy observations with data previously obtained using identical 65 methodology, from placental tissue collected at delivery from 24 women with normotensive 66 pregnancy (40.1  $\pm$  1.2 weeks) and 22 women with pre-eclampsia (36.8  $\pm$  3.6 weeks) [3]. 67 Total RNA extraction, reverse transcription and real-time PCR were conducted as previously 68 described [3]. Immunohistochemical staining was performed using goat polyclonal antibodies 69 (KCNQ3 and KCNE5 (4 µg/ml for both), as previously described [3]. Goat IgG was used as a 70 negative control. All slides were assessed by the same observer (HDM) and quantified using the 71 Positive Pixel Algorithm of Aperio ImageScope software [3, 10]. 72 73 74 75 76 77 78

79	Results
19	Results

- 80 Expression of mRNA of KCNQ and KCNE genes was observed in first trimester tissues (Table
- 81 1). KNCQ4 and KCNQ5 were low or undetectable, whereas KCNQ3 and KCNQ1 showed the
- 82 greatest expression. KCNE1, KCNE2 and KCNE4 expression was low in all TOP samples;
- 83 KCNE5 was highly expressed isoform in mid-TOP and significantly greater than in early-TOP
- 84 (P=0.022; Table 1).

85

- 86 High protein expression for both KCNQ3 and KCNE5 was observed, with staining being
- 87 localised predominantly to the syncytiotrophoblast, cytotrophoblast and mesenchyme (Figure 1).
- 88 No significant differences were observed between early and mid-TOP.

89

Lower mRNA expression of KCNQ2, KCNQ4, KCNQ5 and KCNE1 was observed in the TOP
samples compared to the term placentae (Table 1). KCNQ3 and KCNE5 mRNA expression was
significantly decreased in both early and mid-TOP and normotensive controls compared to preeclamptic placentae (Table 1).

94

- 95 Positive immunostaining for both KCNQ3 and KCNE5 was significantly lower in the
- 96 normotensive and pre-eclamptic compared to both the early- and mid-TOP placentae (Table 1).

97

99 **Discussion** 

100 This study presents novel data concerning placental KCNQ/KCNE mRNA expression profiles

101 early in pregnancy. KCNQ3 and KNCE5 were the predominant isoforms, localised to

102 syncytiotrophoblast and mesenchyme. This is similar to the profile in third trimester

103 normotensive and pre-eclamptic placentae [3].

104

The only KCNQ isoforms to be significantly expressed in early pregnancy were KCNQ1 and
KCNQ3. Both are known to be important for steroid production and inhibition of cell
proliferation, essential factors in placentation [11-15]. KCNE expression was low in all early
pregnancy, with the exception of KCNE5 in late-TOP placenta. In other studies, KCNE5 mRNA
expression is markedly reduced in human 2<sup>nd</sup> trimester trophoblast cells cultured under hypoxic
conditions [16] and oxygen sensitive K<sup>+</sup> channels, including Kv channels, may be important for
the detection and response to a oxygenation stimulus [7, 17].

112

113 KCNQ3 and KCNE5 protein expression in early pregnancy was particularly high in the 114 cytotrophoblast and syncytiotrophoblast. The presence of these proteins in the mesenchyme, the 115 site of angiogenesis [18, 19] during this critical window of feto-placental vascular development, 116 suggests a possible role for these proteins in vessel remodelling. Comparison with our third 117 trimester normotensive and pre-eclamptic data indicate that KCNQ3 and KCNE5 proteins show 118 similar localisation throughout pregnancy. KCNQ3 and KCNE5 expression was lower in tissues 119 taken at delivery, but still raised in tissue from pre-eclampsia compared to normotensive 120 controls. This lower expression, at term, could be due to changes in placental structure as 121 pregnancy progresses, where reticulum cells and fibroblasts are the major cell types [20].

122 Contrasting data between mRNA and protein may be due to mRNA being less stable than protein
123 and since such high expression of protein was observed in TOP samples, subtle differences may
124 not be detected.

126	We had access to a limited number of samples at early gestations and unavoidably, the use of
127	such samples precludes knowing whether these pregnancies may have developed pre-eclampsia.
128	Nevertheless, taken together with our previous work on third trimester placentae, we have
129	provided novel data suggesting a potential role for $K_V7$ channels in early placentation. Future
130	work is needed to characterise the functional impact of KCNQ3 and KCNE5 co-expression both
131	in the development of the early pregnancy placenta and in pre-eclampsia.
132	
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141	
142	Conflict of Interest Statement: There are no conflicts of interest.
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206	<b>Figure</b>	Legends

207	Figure 1: A) KCNQ3 B) KCNE5 immunostaining in 1) early-TOP, 2) mid-TOP and 3) IgG
208	negative controls. In photomicrographs, positive cells appear brown; magnification x400. High
209	protein expression and was localised mainly to the syncytiotrophoblast (red arrows), but was also
210	evident in the mesenchyme (blue arrows). In graphs, data are presented as median [IQR]
211	positivity; scale bar = $100 \ \mu m$ .
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mRNA expression (normalised GAPDH, copy number) x100 <sup>#</sup>	Early TOP (n = 6)	Mid TOP (n = 7)	Normotensive Control (n = 24)	Pre-eclampsia (n = 22)
KCN01	0.4	0.3	3.54	5.6
Kenqi	[0.3, 0.5]	[0.3, 0.5]	[2.0, 10.4]	[2.0, 11.0]
KCNQ2	0.2 [0 1 0 4] <sup>e</sup>	0 <sup>b</sup>	0.7 [0 2, 1, 6]	1 [0 3 2]
KCNQ3	0.3 [0.2, 0.4] <sup>e</sup>	0.4 [0.2, 0.8] <sup>f</sup>	2.3 [0.8, 6.8] <sup>c</sup>	46,967 [26,457, 10,2570]
KCNQ4	0 [0, 0.03]ª	0	0.4 [0.2, 0.7]	0.2 [0.1, 0.6]
KCNQ5	0.01 [0, 0.02] <sup>a</sup>	0.04 [0, 0.08] <sup>b</sup>	8.3 [2.9, 20.3]°	2.2 [0.6, 7.2]
KCNE1	0.02 [0, 0.02]ª	0.04 [0, 0.08] <sup>b</sup>	1.5 [1, 4.3]	0.8 [0.5, 2.2]
KCNE2	0.07 [0.03, 0.08]	0.09 [0.06, 0.1]	0.04 [0, 0.08]	0.02 [0, 0.1]
KCNE3	0.9 [0.6, 1.0]	0.7 [0.6, 1]	1.5 [0.8, 3.2]	1.2 [0.7, 3]
KCNE4	0.2 [0.2, 0.5]	0.2 [0.2, 0.3]	0.2 [0.1, 0.5]	0.3 [0.2, 0.6]
KCNE5	0.6 [0.4, 1] <sup>a</sup>	2 [1, 3] <sup>g</sup>	16 [9, 29] <sup>c</sup>	194 [91, 338]
Positive immunostaining (arbitrary units) <sup>#</sup>	Early TOP (n = 7)	Mid TOP (n = 5)	Normotensive Control (n = 6)	Pre-eclampsia (n = 6)
KCNQ3	0.97 [0.96, 0.99]ª	0.99 [0.96, 0.99] <sup>b</sup>	0.18 [0.1, 0.2] <sup>c</sup>	0.31 [0.2, 0.35]
KCNE5	0.98 [0.94, 0.99] <sup>a</sup>	0.99 [0.98, 0.99] <sup>b</sup>	0.13 [0.06, 0.16] <sup>c</sup>	0.35 [0.28, 0.43]

**Table 1:** KCNQ and KCNE mRNA and protein expression isoform placental.

230

<sup>a</sup>: P<0.05 early TOP vs. normotensive controls and pre-eclampsia; <sup>b</sup>: P<0.05 mid TOP vs.

232 normotensive controls and pre-eclampsia; <sup>c</sup>: P<0.05 normotensive controls vs. pre-eclampsia; <sup>d</sup>:

233 P<0.05 early TOP vs. normotensive controls; <sup>e</sup>: P<0.05 early TOP vs. pre-eclampsia; <sup>f</sup>: P<0.05

234 mid TOP vs. normotensive controls; <sup>g</sup>: P<0.05 mid TOP vs. pre-eclampsia. Data presented as

235 median [IQR]. <sup>#</sup>Normotensive control and pre-eclampsia data previously published [3].