

The Role and Interaction of Imprinted Genes in Human Fetal Growth

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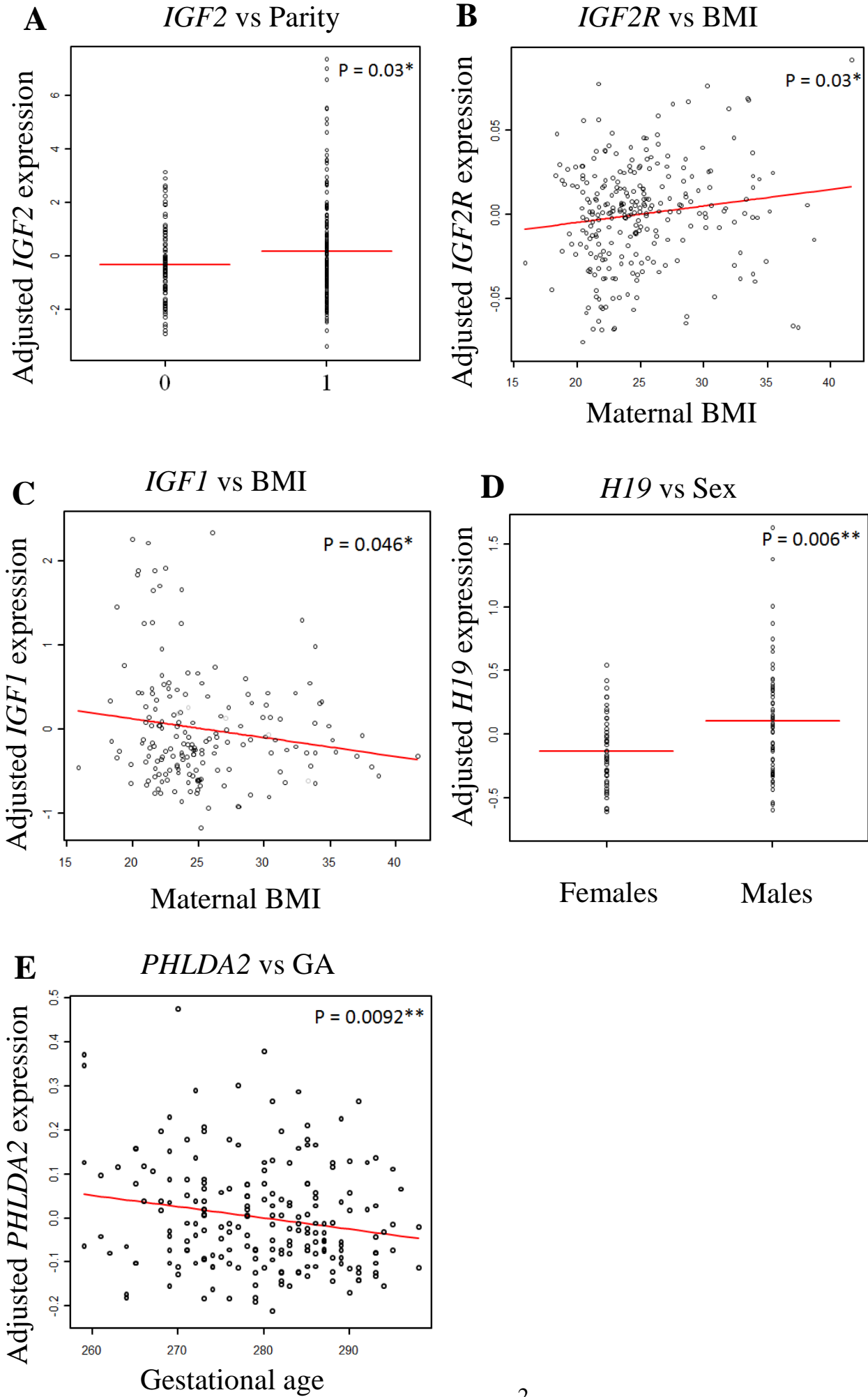
Journal: Philosophical Transactions B

DOI: 10.1098/rstb.2014.0074

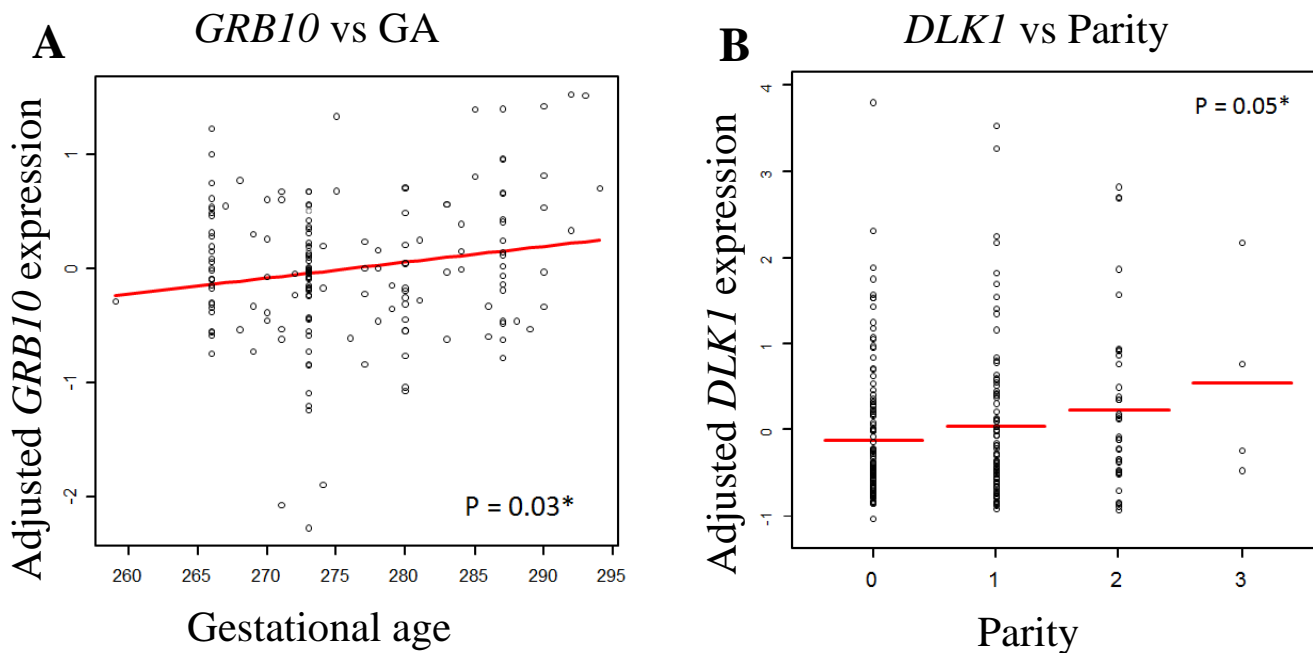
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Supplementary Figure 1. Association between CVS gene expression and factors influencing fetal growth. The associations between gene expression in CVS tissue and the following variables were tested in the regression model; baby's sex, parity, gestational age at birth, maternal BMI and smoking. (A) *IGF2* is expressed higher in the parity > 1 group ($P = 0.03$). Maternal BMI is positively and negatively correlated to (B) *IGF2R* ($P = 0.03$) and (C) *IGF1* ($P = 0.046$). (D) *H19* is expressed higher in males ($P = 0.006$). (E) *PHLDA2* expression is higher as gestational age at birth (in days) advances ($P = 0.0092$).



Supplementary Figure 2. Association between term placenta gene expression and factors influencing fetal growth. The influence of baby's sex, parity, gestational age at birth, maternal weight and smoking on imprinted genes expression in term placenta was tested using a regression model. Partial residual plots showing a positive correlation between gestational age (in days) at birth and (A) *GRB10* expression ($P = 0.03$). Parity is positively correlated to (B) *DLK1* expression (para 1 $P = 0.47$, para 2 $P = 0.05$ and para 3 $P = 0.12$).

Supplementary tables

Supplementary Table 1. Medical details of participants in the Moore and the KCL CVS cohorts.

Cohorts	Moore cohort	KCL CVS cohort
Sample number	302 trios	355
<u>Maternal information:</u>		
Pre-pregnancy weight (Kg)	67 ± 11.7	67 ± 13
Maternal BMI	NA	25 ± 4.4
Maternal age (years)	34 ± 5	35.7 ± 5.6
Smoking (% smokers)	27 (8.9 %)	33 (9 %)
Parity	0: 148 (49 %)	0: 93 (26 %)
	1: 99 (32 %)	>1: 177 (50 %)
	>2: 55 (18 %)	NA: 85 (24 %)
<u>Baby's information:</u>		
Ethnicity:		
White European	302 (100 %)	251 (71 %)
Black		19 (5%)
Asian		11 (3%)
Oriental		9 (3%)
Unknown		65 (18 %)
Gestational age at birth	39 ± 1 (weeks)	279.4 ± 9 (days)
Gestational age at CVS	NA	89.6 ± 4.2
Gender (% males)	51 %	54 %
Crown rump length (mm)	NA	66.9 ± 8
Birth weight (g)	3,432 ± 453	3,507 ± 636
Placental weight (g)	702 ± 137	NA
Head circumference (cm)	34.9 ± 1.5	NA

Mean ± standard deviations. NA = information not available.

Supplementary Table 2. Correlation between gene expression in CVS and variables used in the regression model.

Gene	Mat BMI	Sex	Parity	GA at term	Smoking
<i>IGF2</i>	0.45	0.38	0.03*	0.39	0.42
<i>IGF2R</i>	0.03*	0.62	0.56	0.33	0.48
<i>IGF1</i>	0.046*	0.13	0.07	0.89	0.99
<i>IGF1R</i>	0.067	0.97	0.87	0.5	0.45
<i>PHLDA2</i>	0.92	0.62	0.63	0.0092**	0.28
<i>DLK1</i>	0.74	0.28	0.71	0.11	0.88
<i>H19</i>	0.14	0.006*	0.95	0.65	0.74

GA = gestational age (days), BMI = body mass index.

Supplementary Table 3. Correlation between gene expression in term placenta and variables used in the regression model.

Gene	MW	Sex	Para 1	Para 2	Para 3	GA	Smoking
<i>IGF2</i>	0.31	0.42	0.12	0.098	0.85	0.54	0.29
<i>IGF2R</i>	0.88	0.56	0.2	0.71	0.43	0.53	0.41
<i>PHLDA2</i>	0.099	0.084	0.12	0.83	0.56	0.51	0.25
<i>MEST</i>	0.99	0.96	0.33	0.93	0.26	0.94	0.27
<i>H19</i>	0.64	0.65	0.065	0.78	0.64	0.84	0.81
<i>GRB10</i>	0.63	0.55	0.35	0.31	0.62	0.036*	0.46
<i>DLK1</i>	0.87	0.71	0.47	0.05*	0.13	0.16	0.099
<i>MEG3</i>	0.622	0.262	0.700	0.165	0.696	0.939	0.913
<i>PEG10</i>	0.2	0.7	0.92	0.22	0.99	0.53	0.47
<i>PEG3</i>	0.4	0.95	0.0019**	0.014*	0.061	0.18	0.16
<i>SLC22A18</i>	0.57	0.4	0.97	0.09	0.7	0.99	0.12
<i>CDKN1C</i>	0.41	0.085	0.14	0.85	0.14	0.94	0.96
<i>PLAGL1_all</i>	0.43	0.83	0.66	0.38	0.23	0.3	0.77
<i>PLAGL1_imp</i>	0.4	0.83	0.39	0.64	0.78	0.018*	0.64
<i>IGFBP3</i>	0.33	0.84	0.002	0.62	0.43	0.56	0.8

GA = gestational age (days), MW = maternal weight, Para = parity.

Supplementary Table 4. Primer sequences for RT-qPCR.

Gene	Sequence 5' to 3'	Size (bp)	Accession	Ref
<i>L19</i>	F: GCGGAAGGGTACAGCCAAT R: CAGGCTGTGATACATGTGGCG	130	NM_000981	(1)
<i>PEG10</i>	F: TGTCTTCGCAGAGGAGTCCTCG R: ACTTCTGTGGGGATGGAGGCC	105	NM_001040152	(2)
<i>PEG3</i>	F: TGCAGCTTGCTGAAGACGATG R: AAGATTCATCTTCACAAATCCC	211	NM_006210	(2)
<i>DLK1</i>	F: AGCACCTATGGGGCTGAATG R: TCCCAGCCGTCGGTGCAAATGC	82	NM_003836	(2)
<i>MEG3</i>	F: ACGGCGGAGAGCAGAGAG R: AGGAGAGACCCGGGTGAG	106	NR_002766	MI
<i>H19</i> (Term)	F: GGAGTTGTGGGAGACGGCCTTGAGT R: CCAGTCACCCGGCCCAGATGGAG	109	NR_002196	(3)
<i>H19</i> (CVS)	F: ACACTATGGCTGCCCTCTG R: AGAGGGTTTTGTGTCCGGAT	100	NR_002196	MI
<i>PHLDA2</i>	F: GAGCGCACGGGCAAGTA R: CAGCGGAAGTCGATCTCCTT	68	NM_003311	(1)
<i>GRB10</i>	F: CACGAGCCCTTGCTGTTTA R: AAGTCCCCTCCACCTTCTA	115	NM_001001555	(2)
<i>IGFBP3</i>	F: CGGTCTTCCTCCGACTCAC R: GTCAACGCTAGTGCCGTCAG	91	NM_000598.4	JMF
<i>SLC22A18</i>	F: GCAGGATGAGCGCTTAGGC R: GCACCCCGAAGGTGGTTTGC	168	NM_183233	(2)
<i>CDKN1C</i>	F: CAGCTGCACTCGGGGATTTTC R: GATCTCTTGCGCTTGCGGAAG	125	NM_000076	(2)
<i>PLAGL1_Imp</i>	F: AGCCGTGCTCACAGCTCAG R: AGGCAGCAGCCACATTAGAC	200	NM_001080954	II
<i>PLAGL1_All</i>	F: GTGAGGAGTGTGGGAAGAAG R: GAGGTGGTCCAGTAGCACCTC	140	NM_001080951	II

Annealing temperature was set to 60°C for the all qPCR primers. MI = primer designed by Miho Ishida. JMF = primer designed by Jennifer May Frost. II = primer designed by Isabel Iglesias-Platas. Imp = imprinted isoform. All = all isoforms.

Supplementary Table 5. Primer sequences for the imprinting analysis.

Gene	Sequence 5' to 3'	Tm (°C)	Polymorphism	Accession	Ref
<i>DLK1</i> geno/RT*	F: CACCTGCGTGGATGATGAG R: GATGGTGAAGCAGATGGCCTG	58	rs1802710	NM_003836	(2)
<i>MEG3</i> geno	F: TCACCTGCTAGCAAAGTGGGA R: GGCTGCCTCGTTTACATACA	58	rs45617834 rs941575	NR_002766	(2)
<i>MEG3</i> RT	F: AAGGACCACCTCCTCTCCAT R: AGGAGGTTGCTCTCATCGAC	56			
<i>PEG3</i> geno/RT*	F: GAACCCTGTCAATATCTTTTTGAG R: CATGAGAACCACCTTCAACAAAC	57	rs1055359	NM_001146184	(2)
<i>ACTB</i> geno/RT	F: GTCTTCCCCTCCATCGTG R: GGTCATCTTCTCGCGGTTG	59	NA	NM_001101	ACT
<i>PEG10</i>	F: ACAGAGATGTAAGAGGCAGGC R: ATTCACAGCATTGTAAGAAGTTCAC	60	rs13073 rs13226637	NM_015068	MI
<i>H19</i>	F: CTCACCCACCGCAATTCATT R: CCCTGGATGCTGTACTGTCT	59	rs2067051		MI
<i>SLC22A18</i>	F: CCCGATCAACTGGACTTTTG R (Geno): GGCACGATGGAGAAGTGCATG R (RT): GCACCCGAAGGTGGTTTGCA	58	rs1048046, rs1048047	NM_183233	(4)
<i>CDKN1C</i>	F: CTAGCCAGCAGGCATCGAG R: CTCATCGTGGATGTGCTG	58	PAPn repeat	NM_001122631	(4)
<i>PLAGL1</i>	F: GAGCACCATTTAAAGGAACG R: GGTTCTACCCAGACATGGACC	55	rs2076684	NM_006718_P1 NM_001080951_P2	II

MI = primer designed by Miho Ishida. JMF = primer designed by Jennifer May Frost. II = primer designed by Isabel Iglesias-Platas. ACT = primer designed by Anna Clare Thomas.

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