Supplementary material for He P, et al. 2014. Analysis of gene expression identifies candidate markers and pathways in pre-eclampsia. *Journal of Obstetrics and Gynaecology* DOI: 10.3109/01443615.2014.990430.

Supplementary Table I. Clinical information on the study population.

Sample (original number)	Gestational Age (weeks)	Birth Weight (g)	SD of birthweight	Sex	Outcome of the pregnancies
Group of EOPET					
PM12	31.72	1305	−1.5	male	IUGR
PM21	33.72	1650	−1.87	male	IUGR
PM39	32	1700	0.31	male	IUGR
PM43	31.72	1440	−0.88	female	AVM
PM51	34	1400	−2.92	female	IUGR, oligohydramnios, AVM
PM86	24.86	545	−1.64	male	IUGR
PM116	32.42	1480	−0.7	male	IUGR
PM129	37.28	1840	−2.17	male	HELLP, AVM
Mean value	32.21	1420			
Group of control					
PL113	25	978	0.66	male	
PL64	26	758	−0.5	female	
PL65	31.42	1849	0.99	male	
PL56	32.56	2100	0.69	female	PPROM
PL33	32.86	2230	1.26	male	
PL11	33.72	2495	1.68	male	Abruption
PL76	36	2630	0.48	male	Chorioamionitis, PPROM
PL96	34	2040	−0.23	male	PPROM
Mean value	31.45	1885			