








Supplementary information to:

Original article:

METABOLOMIC ANALYSIS OF THE HUMAN PLACENTA REVEALS PERTURBATIONS IN AMINO ACIDS, PURINE METABOLITES, AND SMALL ORGANIC ACIDS IN SPONTANEOUS PRETERM BIRTH

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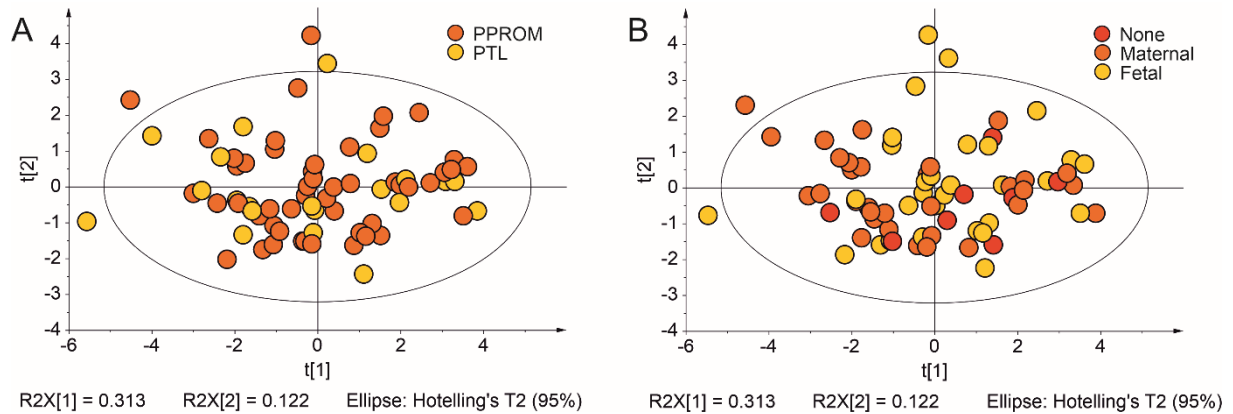
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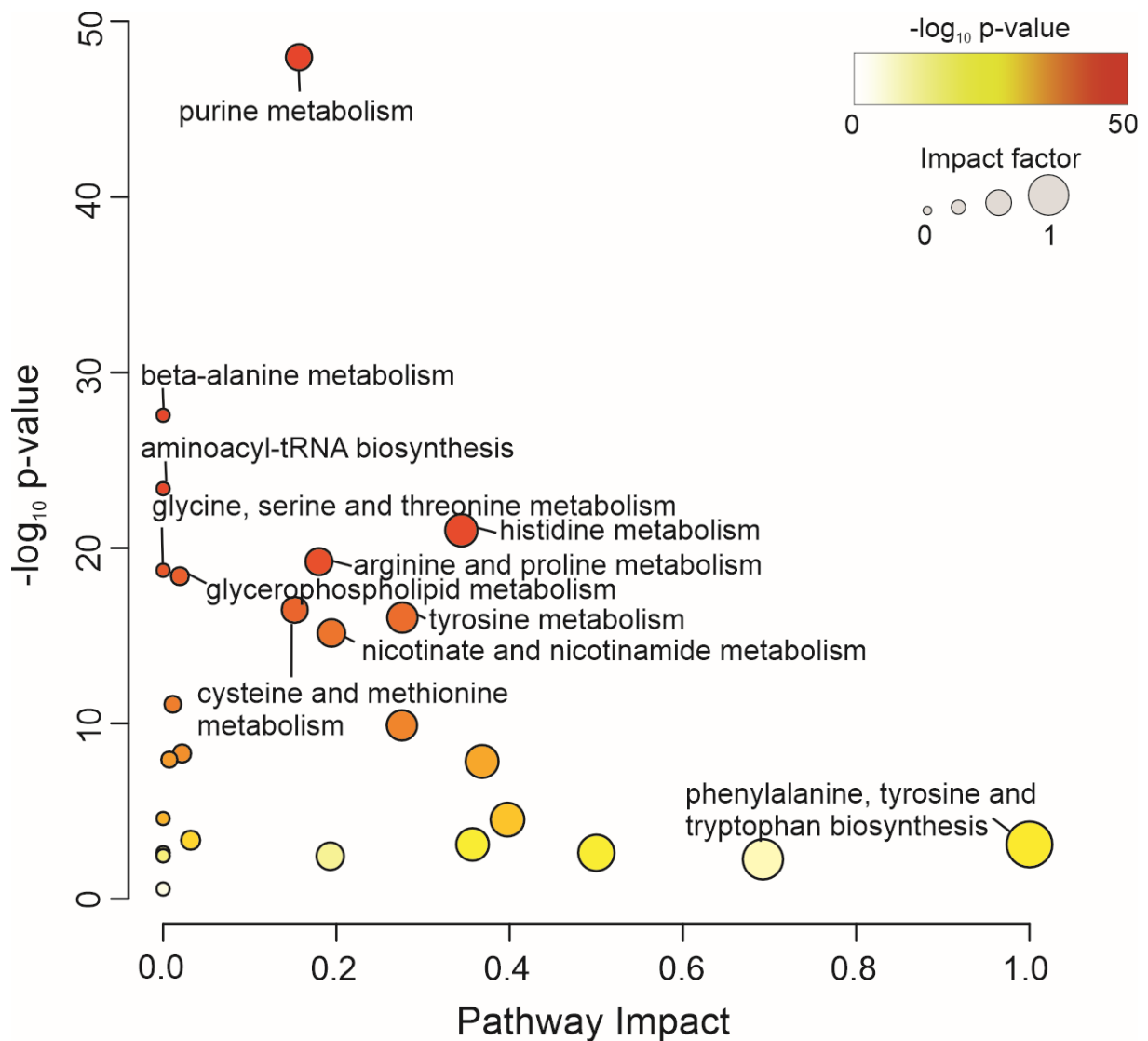
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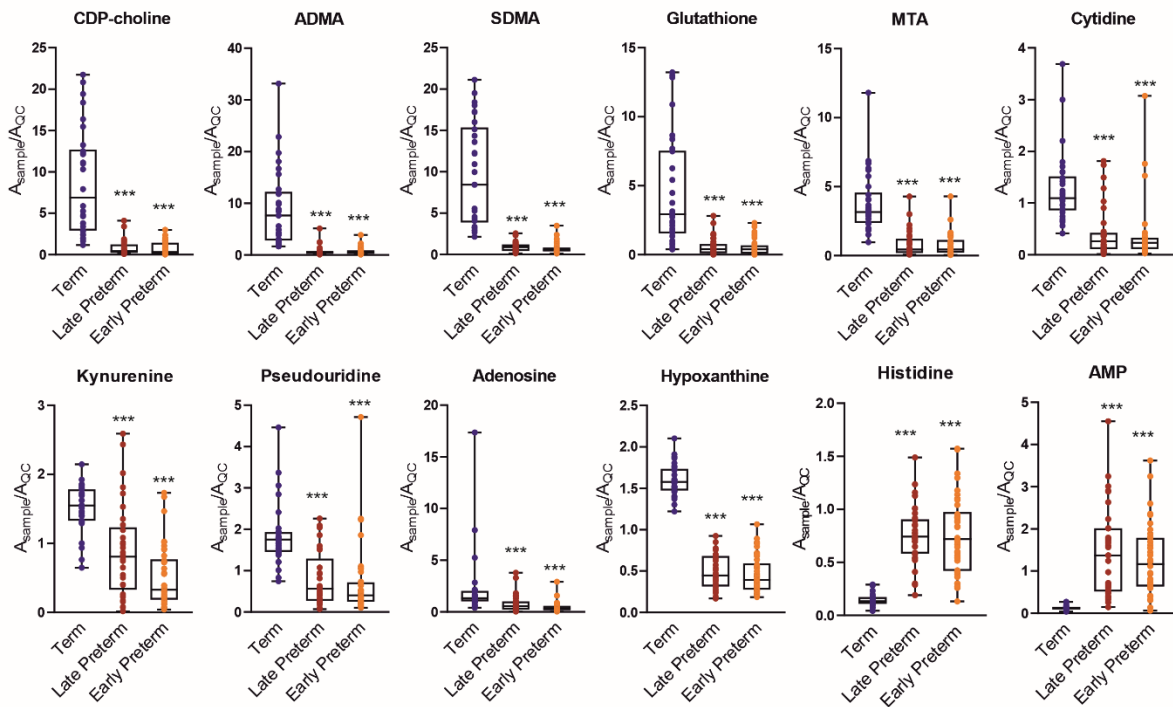
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Supplementary Figure 1: Principal component analysis of metabolomic profiles for preterm birth placentas. Correlations between placentas associated with PPROM and PTL preterm birth phenotypes (A) and indicated HCA grades (B).



Supplementary Figure 2: Pathway enrichment analysis of significantly altered metabolites in spontaneous preterm birth placentas. The node color gradient indicates the negative log p-value and the node radius indicates the pathway impact score. The 10 most significantly affected metabolic pathways are identified by name.



Supplementary Figure 3: Box plots of up- and down-regulated metabolites in early and late spontaneous preterm birth deliveries. Statistical analysis was performed using the nonparametric Mann-Whitney test, and the false discovery rate was performed using the Bonferroni and Benjamini-Hochberg corrections.

Supplementary Table 1: Clinical and demographic information on preterm pregnancies included in the study

	PTL (n = 22)	PPROM (n = 50)	-log ₁₀ p value
Gestational age at delivery, week	32.0 (27.2 - 34.1)	34.4 (31.9 - 36.0)	2.09
Newborn weight, kg	1.7 (1.0 - 2.2)	2.2 (1.7 - 2.6)	2.21
Maternal BMI prepregnancy, kg/m ²	21.0 (19.0 - 22.5)	23.1 (19.7 - 25.0)	1.12
Maternal BMI at admission, kg/m ²	23.1 (22.5 - 25.6)	26.5 (23.4 - 28.4)	1.48
Early preterm (<34 wk), n (%)	16 (73)	23 (46)	1.36
Late preterm (34-37 wk), n (%)	6 (27)	27 (54)	1.36
MIAC, n (%)	4 (18)	16 (32)	0.57
HCA, n (%)	22 (100)	43 (86)	1.03
Funisitis, n (%)	3 (14)	16 (32)	0.83
Male fetal sex, n (%)	14 (64)	31 (62)	0.00
Caesarian delivery, n (%)	8 (36)	15 (30)	0.22
Vaginal delivery, n (%)	14 (64)	35 (70)	0.22
Primipara, n (%)	11 (50)	26 (52)	0.00
Administration of corticosteroids, n (%)	16 (73)	30 (60)	0.37
Administration of antibiotics, n (%)	19 (86)	48 (96)	0.79
Administration of tocolytics, n (%)	11 (50)	5 (10)	3.40
Amniotic fluid IL-6 conc. at admission, ng/mL	4.7 (1.8 - 11.2)	1.0 (0.4 - 2.0)	2.74
Maternal serum CRP conc. at delivery, mg/L	7.5 (3.6 - 14.9)	4.8 (2.4 - 13.1)	0.49
Maternal serum WBC count at delivery, ×10 ⁹ L	13.8 (11.3 - 16.2)	13.4 (11.0 - 16.6)	0.01
Apgar score at 5 min <7, n (%)	5 (23)	1 (2)	2.05
Apgar score at 10 min <7, n (%)	3 (14)	2 (4)	0.79

Presented data are medians with interquartile range, numbers, or percentages. Statistical analysis was performed using the nonparametric Mann-Whitney test or Fisher's exact test where appropriate.

Supplementary Table 2: Composition of the internal standard mixture used for metabolomic analysis

Internal standard	Concentration (ng/μL)
GABA[D6]	1050.29
Aspartic acid[D3]	1113.94
Dopamine[D4]	0.32
Glutamic acid[D3]	1050.29
Kynurenic acid[D5]	3.18
Kynurenine[D4]	3.18
L-dopa[D3]	2.55
Leucine[D3]	159.13
Lysine[D4]	286.44
Serotonin[D4]	0.64
Epinephrine[D3]	95.48
3-methoxytyramine[D4]	0.32

Supplementary Table 3: UHPLC/MS method validation results for metabolomic analysis of human placenta samples

	Without matrix		With matrix		Linearity range [pmol/mL]	LOD [pmol/mL]	Accuracy [%]			Precision [%]			Extraction recovery [%]
	Slope	R ²	Slope	R ²			Intraday		Inter-day	Intraday		Inter-day	
							LL	HL	HL	LL	HL	HL	
GABA[D6]	0.1058	0.988	0.0064	0.982	6110-122211	109	100	101	95	9	4	9	78
Aspartic acid[D3]	53.250	0.997	0.002	0.980	48999–489993	8166	102	101	92	7	7	13	52
Dopamine[D4]	49.841	0.993	73.169	0.999	4–42435	1	100	95	98	9	3	7	51
Glutamic acid[D3]	7.871	0.998	0.070	0.996	1777–444231	223	101	100	95	5	3	8	78
Kynurenic acid[D5]	158.820	0.996	101.690	0.999	3–34351	1	102	92	97	9	4	9	48
Kynurenine[D4]	172.520	0.999	0.975	0.999	3–31430	1	106	105	91	15	11	16	63
L-dopa[D3]	24.067	0.989	33.966	0.996	3–33319	1	102	89	97	18	4	11	58
Leucine[D3]	0.469	0.993	N/A	N/A	99-9942	1657	101	102	93	10	8	13	71
Lysine[D4]	2.966	0.989	N/A	N/A	44-1776	1974	107	105	97	17	11	15	32
Serotonin[D4]	53.248	0.997	10.872	0.996	37-37012	2	102	97	94	8	6	9	31
Epinephrine[D3]	168.170	0.992	52.181	0.993	37-35821	1	99	95	97	7	5	6	74
3-methoxytyra- mine[D4]	87.138	0.995	8.166	0.999	39-38959	3	103	99	95	12	8	11	53

LL – low concentration level, HL – high concentration level

Supplementary Table 4: Metabolite fold change and p-value after FDR corrections for identified metabolites

Rank	Metabolite	Negative log (p-value); significance after Bonferroni correction $\alpha = -\log(0.05; 0.01; 0.001 / 43) = 2.9 (*)$; $3.6 (**)$; $4.6 (***)$			Negative log (p-value); significance after Benjamini and Hochberg correction $\alpha = -\log(\text{RANK} * 0.05; 0.01; 0.001 / 43) = 1.39 (*)$; $2.11 (**)$; $3.16 (***)$			Median fold change		
		Preterm vs. Term	Late Pre-term vs. Term	Early Preterm vs. Term	Preterm vs. Term	Late Pre-term vs. Term	Early Preterm vs. Term	Preterm vs. Term	Late Pre-term vs. Term	Early Preterm vs. Term
41	5-Hydroxytryptophan	0.21	0.07	0.58	0.21	0.07	0.58	0.11	-0.03	0.27
12	Acetylcarnitine ^a	13.06	9.41	10.27	13.06	9.41	10.27	-0.38	-0.64	-0.70
18	Acetylneuraminate ^a	9.66	8.30	6.87	9.66	8.30	6.87	-0.51	-1.05	-1.03
26	Adenine ^a	5.72	3.22	5.59	5.72	3.22	5.59	-0.21	-0.31	-0.35
19	Adenosine	9.25	4.74	11.12	9.25	4.74	11.12	-0.72	-1.27	-2.05
1	ADMA	15.00	13.89	15.00	15.00	13.89	15.00	-0.93	-3.88	-4.04
3	AMP ^a	15.00	15.00	13.85	15.00	15.00	13.85	9.03	3.49	3.26
33	Arginine	1.98	1.98	1.15	1.98	1.98	1.15	0.23	0.31	0.29
40	Caffeine ^a	0.33	0.14	0.90	0.33	0.14	0.90	0.07	0.78	-0.20
25	Carnitine	6.40	5.00	5.11	6.40	5.00	5.11	-0.31	-0.52	-0.56
4	CDP-choline ^a	15.00	13.28	14.52	15.00	13.28	14.52	-0.95	-4.04	-4.46
38	CMP ^a	0.62	0.29	0.90	0.62	0.29	0.90	0.20	0.24	0.32
2	Creatine	15.00	11.53	15.00	15.00	11.53	15.00	-0.66	-1.47	-1.66
14	Cytidine	11.19	7.64	10.83	11.19	7.64	10.83	-0.76	-2.04	-2.25
21	GABA	8.46	5.67	8.17	8.46	5.67	8.17	-0.16	-0.24	-0.26
39	Glutamic acid	0.44	0.69	0.16	0.44	0.69	0.16	0.10	0.19	0.05
29	Glutamine	3.40	2.89	2.70	3.40	2.89	2.70	-0.26	-0.42	-0.45
11	Glutathione ^a	14.00	9.93	11.37	14.00	9.93	11.37	-0.87	-2.91	-2.96
42	GMP ^a	0.11	0.08	0.39	0.11	0.08	0.39	-0.11	0.19	-0.34
34	Guanosine	1.88	1.91	1.17	1.88	1.91	1.17	0.26	0.45	0.32
5	Histidine	15.00	15.00	15.00	15.00	15.00	15.00	4.54	2.48	2.44
6	Hypoxanthine	15.00	15.00	15.00	15.00	15.00	15.00	-0.73	-1.82	-2.01
24	Inosine	7.46	6.03	5.88	7.46	6.03	5.88	1.27	1.20	1.16
20	Kynurenic acid	8.79	7.28	6.69	8.79	7.28	6.69	-0.61	-1.12	-1.47
16	Kynurenine	9.94	4.80	11.31	9.94	4.80	11.31	-0.64	-0.94	-2.24
7	L-DOPA	15.00	12.06	11.24	15.00	12.06	11.24	-0.63	-1.35	-1.63
32	Leucine	2.61	1.13	3.05	2.61	1.13	3.05	-0.27	-0.16	-0.56

Rank	Metabolite	Negative log (p-value); significance after Bonferroni correction $\alpha = -\log(0.05; 0.01; 0.001 / 43) = 2.9 (*)$; $3.6 (**)$; $4.6 (***)$			Negative log (p-value); significance after Benjamini and Hochberg correction $\alpha = -\log(\text{RANK} * 0.05; 0.01; 0.001 / 43) = 1.39 (*)$; $2.11 (**)$; $3.16 (***)$			Median fold change		
		Preterm vs. Term	Late Pre-term vs. Term	Early Preterm vs. Term	Preterm vs. Term	Late Pre-term vs. Term	Early Preterm vs. Term	Preterm vs. Term	Late Pre-term vs. Term	Early Preterm vs. Term
27	Metanephrine	5.29	3.28	5.02	5.29	3.28	5.02	-0.29	-0.47	-0.51
36	Methionine	0.87	1.07	0.50	0.87	1.07	0.50	-0.05	-0.09	-0.05
8	MTA ^a	15.00	11.53	13.46	15.00	11.53	13.46	-0.86	-2.85	-2.83
13	Nicotinamide	12.26	8.71	10.06	12.26	8.71	10.06	-0.47	-0.89	-0.94
23	Pantothenate ^a	7.87	4.53	7.63	7.87	4.53	7.63	-0.55	-0.94	-1.21
30	Paraxanthine	3.10	1.63	3.23	3.10	1.63	3.23	-0.55	-0.87	-1.51
43	Phenylalanine	0.00	0.48	0.31	0.00	0.48	0.31	-0.06	0.10	-0.34
17	Pseudouridine	9.91	6.03	9.26	9.91	6.03	9.26	-0.74	-1.63	-2.11
22	Quinolinic acid	8.44	5.89	8.00	8.44	5.89	8.00	1.26	0.98	1.33
15	SAH ^a	9.96	6.65	9.26	9.96	6.65	9.26	-0.39	-0.63	-0.91
9	SDMA	15.00	15.00	15.00	15.00	15.00	15.00	-0.92	-3.25	-3.92
31	Tryptophan	2.89	1.24	3.40	2.89	1.24	3.40	1.11	0.82	1.41
35	Tyramine	1.22	1.19	1.01	1.22	1.19	1.01	-0.10	-0.09	-0.16
28	Tyrosine	4.60	3.70	3.70	4.60	3.70	3.70	-0.25	-0.38	-0.51
10	Urate ^a	15.00	9.30	13.96	15.00	9.30	13.96	-0.45	-0.76	-0.92
37	Urocanate ^a	0.72	1.31	0.23	0.72	1.31	0.23	0.19	0.37	0.23

^a Annotated metabolites without identical standard

Supplementary Table 5: Spearman's correlation analysis

Metabolite	GA		CRP		IL6		Fetal weight		BMI at delivery		WBC at delivery	
	r	p-value	r	p-value	r	p-value	r	p-value	r	p-value	r	p-value
5-Hydroxytryptophan	-0.245	0.039	0.144	0.235	0.192	0.111	-0.292	0.013	-0.104	0.388	0.289	0.015
Acetylcarnitine ^a	0.087	0.469	0.071	0.559	-0.001	0.995	0.083	0.489	0.238	0.046	0.035	0.775
Acetylneuraminat ^a	-0.053	0.660	0.194	0.108	0.067	0.582	0.020	0.870	0.285	0.016	0.100	0.411
Adenine ^a	0.064	0.594	0.157	0.196	0.013	0.918	0.038	0.751	0.006	0.960	0.213	0.077
Adenosine	0.304	0.010	0.014	0.909	0.007	0.954	0.295	0.013	0.054	0.653	-0.036	0.767
ADMA	0.185	0.122	0.086	0.479	-0.091	0.453	0.145	0.229	0.042	0.725	-0.057	0.641
AMP ^a	0.067	0.576	0.137	0.259	-0.024	0.841	0.090	0.457	0.194	0.106	-0.052	0.670
Arginine	0.152	0.205	-0.043	0.722	-0.121	0.320	0.143	0.234	-0.065	0.592	-0.064	0.596
Caffeine ^a	0.234	0.050	-0.137	0.258	-0.282	0.018	0.210	0.079	0.055	0.652	-0.255	0.033
Carnitine	0.109	0.366	0.052	0.671	-0.067	0.582	0.082	0.498	0.182	0.129	-0.015	0.905
CDP-choline ^a	0.064	0.593	0.104	0.391	-0.002	0.987	0.090	0.456	0.197	0.100	0.029	0.812
CMP ^a	-0.037	0.761	0.200	0.096	0.053	0.662	-0.016	0.893	0.187	0.119	0.079	0.517
Creatine	0.149	0.214	0.035	0.776	0.023	0.853	0.189	0.114	0.055	0.650	-0.101	0.404
Cytidine	0.165	0.168	-0.032	0.792	0.130	0.284	0.138	0.251	-0.028	0.820	-0.033	0.786
GABA	0.163	0.173	0.136	0.262	-0.015	0.905	0.064	0.593	0.022	0.854	-0.124	0.309
Glutamic acid	0.083	0.491	0.053	0.664	-0.093	0.444	0.115	0.340	0.144	0.232	-0.032	0.793
Glutamine	-0.039	0.750	0.042	0.728	0.208	0.084	0.036	0.766	-0.089	0.461	-0.022	0.859
Glutathione ^a	-0.107	0.373	0.264	0.027	0.099	0.415	-0.115	0.339	0.019	0.877	-0.009	0.943
GMP ^a	0.201	0.093	0.079	0.516	-0.008	0.947	0.153	0.203	-0.024	0.845	-0.097	0.425
Guanosine	0.261	0.028	0.080	0.512	0.091	0.454	0.271	0.022	0.106	0.381	0.112	0.356
Histidine	0.029	0.809	0.091	0.454	-0.117	0.335	0.025	0.838	0.008	0.946	0.008	0.948
Hypoxanthine	0.201	0.093	-0.026	0.829	-0.040	0.743	0.182	0.129	0.027	0.825	-0.007	0.956
Inosine	0.117	0.333	0.091	0.454	0.042	0.730	0.130	0.278	0.078	0.519	0.081	0.506
Kynurenic acid	0.102	0.395	-0.123	0.309	0.182	0.132	0.120	0.320	0.088	0.466	0.097	0.425
Kynurenine	0.411	0.000	0.123	0.311	-0.072	0.554	0.380	0.001	0.067	0.579	-0.043	0.722

Metabolite	GA		CRP		IL6		Fetal weight		BMI at delivery		WBC at delivery	
	r	p-value	r	p-value	r	p-value	r	p-value	r	p-value	r	p-value
L-DOPA	0.076	0.526	-0.135	0.265	0.014	0.909	0.118	0.327	0.016	0.894	-0.038	0.753
Leucine	0.197	0.100	0.157	0.195	0.016	0.896	0.176	0.142	0.095	0.432	-0.071	0.559
Metanephrine	0.214	0.074	-0.050	0.681	-0.020	0.867	0.250	0.035	0.113	0.348	0.013	0.912
Methionine	-0.069	0.570	0.088	0.467	0.050	0.684	-0.078	0.519	-0.138	0.251	0.087	0.475
MTA ^a	0.007	0.954	0.128	0.293	-0.006	0.964	0.073	0.545	0.315	0.007	0.035	0.777
Nicotinamide	0.220	0.066	0.133	0.272	0.030	0.806	0.147	0.222	0.110	0.360	0.029	0.809
Pantothenate ^a	0.148	0.219	0.072	0.553	-0.084	0.491	0.167	0.164	0.147	0.221	-0.164	0.174
Paraxanthine	0.134	0.267	0.018	0.884	-0.099	0.414	0.060	0.617	-0.087	0.470	-0.079	0.515
Phenylalanine	0.267	0.025	-0.145	0.232	0.120	0.322	0.239	0.045	-0.049	0.684	0.015	0.900
Pseudouridine	0.227	0.057	-0.092	0.450	0.008	0.945	0.224	0.060	0.063	0.602	0.004	0.974
Quinolinic acid	-0.149	0.215	0.158	0.190	0.079	0.515	-0.221	0.064	0.197	0.099	0.124	0.306
SAH ^a	0.300	0.011	0.188	0.119	-0.046	0.708	0.266	0.025	0.089	0.462	-0.045	0.712
SDMA	0.270	0.023	-0.021	0.866	-0.099	0.417	0.310	0.009	0.217	0.069	-0.063	0.604
Tryptophan	-0.206	0.084	-0.069	0.570	0.201	0.095	-0.224	0.060	-0.001	0.992	0.074	0.542
Tyramine	0.015	0.900	-0.026	0.829	0.018	0.881	-0.017	0.890	-0.140	0.243	-0.063	0.605
Tyrosine	0.123	0.309	-0.155	0.202	0.006	0.960	0.095	0.429	-0.255	0.032	0.064	0.599
Urate ^a	0.269	0.023	0.205	0.089	-0.092	0.449	0.286	0.016	0.309	0.009	0.006	0.963
Urocanate ^a	0.255	0.032	-0.011	0.929	0.089	0.466	0.169	0.159	-0.264	0.026	0.200	0.096

^a Annotated metabolites without identical standard