

gene	SNP	alleles	location	mean +/- SD	Min-Max	number of fetuses	mean +/- SD	Min-Max	number of fetuses
					Kidney			Liver	
LOC166785	rs1068	A+,G	4q31.21	5.39 +/- 3.81	1.47-10.86	5	3.83 +/- 3.41	1.24-9.59	5
DCTD	rs978	A,G	4q35.1	2.25 +/- 1.70	1.30-5.28	5	4.27 +/- 2.70	1.17-6.85	5
RAI16	rs7490	C,T+	8p21.3	1.21 +/- 0.10	1.14-1.38	5	2.12 +/- 1.27	1.09-3.81	5
SETBP1	rs3324	G,T+	18q21.1	1.24 +/- 0.40	1.02-1.95	5	4.28 +/- 4.95	1.16-12.97	5
PSMD4	rs3539	A,C+	1q21.2	1.18 +/- 0.13	1.01-1.33	5	1.60 +/- 0.66	1.05-2.42	5
KIAA0252	rs1942	C+,T	15	1.49 +/- 0.30	1.07-1.83	5	2.59 +/- 2.75	1.27-7.50	5
?	rs3601	A,G+	?	1.31 +/- 0.13	1.21-1.50	4	1.53 +/- 0.55	1.03-2.31	6
?	rs1491	G,T+	?	1.96 +/- 1.30	1.12-4.23	5	2.30 +/- 2.39	1.03-6.57	5
KIAA0854	rs2833	C,T	8q24.13	1.89 +/- 1.21	1.02-3.72	5	2.21 +/- 2.14	1.04-6.02	5
ROS1	rs1759	A,T+	6q22	1.23 +/- 0.30	1.00-1.74	5	1.60 +/- 1.02	1.05-3.42	5
BN51T	rs2645	C,T	8q21	2.87 +/- 2.35	1.28-6.99	5	1.63 +/- 0.64	1.08-2.35	4
?	rs3075	C,T+	?	1.41 +/- 0.50	1.07-2.16	4	3.14 +/- 4.18	1.08-10.61	5
CYP11B2	rs3097	A+,G	8q21-q22	8.07 +/- 7.19	2.23-20.11	5	6.79 +/- 6.11	2.40-15.47	4
LOC120425	rs1805	A+,C	11q23.3	3.82 +/- 1.59	1.60-5.40	4	3.22 +/- 1.97	1.06-5.63	5
DDX31	rs3433	A,G	9q34.3	4.48 +/- 6.59	1.19-16.26	5	1.85 +/- 0.84	1.26-3.09	4
C3	rs17030	A,G+	19p13.3-p13.2	3.67 +/- 3.15	1.03-7.34	4	1.07 +/- 0.10	1.01-1.22	4
FBXO5	rs1712	C+,T	6q25-q26	1.35 +/- 0.28	1.11-1.74	4	1.60 +/- 0.76	1.11-2.72	4
KIAA0761	rs3517	C,T	1p13.1	2.36 +/- 1.57	1.04-4.34	4	3.68 +/- 1.83	1.03-4.93	4
OAS1	rs2660	A+,G	12q24.1	1.63 +/- 0.55	1.20-2.42	4	1.29 +/- 0.14	1.09-1.41	4
SERPINB5	rs1004	A,C+	18q21.3	1.89 +/- 0.65	1.29-2.48	4	2.36 +/- 0.89	1.56-3.63	4
DYT1	rs1182	A,C+	9q34	1.42 +/- 0.12	1.31-1.58	4	1.71 +/- 0.75	1.07-2.75	4
FBXO5	rs1713	C,T+	6q25-q26	1.86 +/- 0.74	1.10-2.70	4	1.74 +/- 0.66	1.08-2.56	4
TCF2	rs2689	A+,T	17cen-q21.3	1.18 +/- 0.13	1.04-1.36	4	2.39 +/- 1.71	1.08-4.75	4
LOC123568	rs16923	C+,T	15q13.2	4.66 +/- 4.10	1.57-10.71	4	4.84 +/- 5.47	1.73-13.04	4
ARHGAP8	rs3329	A,G	22q13.31	2.40 +/- 0.48	1.79-2.95	4	1.95 +/- 1.27	1.18-3.86	4
FDFT1	rs3258	C+,T	8p23.1-p22	1.40 +/- 0.54	1.02-2.20	4	1.28 +/- 0.28	1.03-1.67	4
?	rs3069	C,T+	?	1.34 +/- 0.13	1.19-1.48	4	2.04 +/- 0.83	1.26-3.02	4
?	rs1009	C+,T	?	1.14 +/- 0.08	1.03-1.21	4	1.59 +/- 0.69	1.00-2.59	4
LOC60626	rs3216	C+,G	11p15.5	1.26 +/- 0.16	1.05-1.44	4	1.92 +/- 1.29	1.07-3.81	4
VAMP8	rs16988	A,G	2p12-p11.2	1.54 +/- 0.89	1.03-2.87	4	1.52 +/- 0.51	1.12-2.21	4
MET	rs1621	A+,G	7q31	1.76 +/- 0.75	1.03-2.79	4	1.30 +/- 0.43	1.06-1.94	4
RANGAP1	rs1953	A,G	22q13	3.75 +/- 3.83	1.08-9.26	4	2.32 +/- 1.22	1.09-3.80	4
DKFZP727G0	rs1837	C,T	9q33.3	2.00 +/- 1.46	1.17-4.20	4	1.68 +/- 0.66	1.13-2.65	4
KIAA0254	rs1781	C,T+	11q25	1.83 +/- 0.92	1.11-3.05	4	1.21 +/- 0.24	1.04-1.55	4

PPP2R5C	rs3405	C,T+	3p21	2.54 +/- 2.12	1.12-5.66	4	2.22 +/- 0.15	2.06-2.35	3
?	rs3261	A+,G	?	1.74 +/- 0.74	1.11-2.56	3	2.35 +/- 1.19	1.16-3.89	4
?	rs3175	C,T+	?	2.37 +/- 1.27	1.53-3.83	3	2.84 +/- 2.21	1.06-5.99	4
SMUG1	rs971	C+,T	12q13.11-q13.3	1.17 +/- 0.12	1.06-1.34	4	2.13 +/- 1.88	1.04-4.30	3
?	rs3364	A,G+	?	1.66 +/- 0.59	1.15-2.43	4	1.87 +/- 0.72	1.44-2.70	3
DKFZP564F0	rs3023	C,T	2p23.1	4.84 +/- 7.38	1.06-15.92	4	2.21 +/- 0.73	1.58-3.02	3
LOC126299	rs3046	A,G	19q13.32	1.40 +/- 0.55	1.04-2.20	4	2.11 +/- 1.38	1.08-3.68	3
?	rs1280	A+,G	?	3.29 +/- 3.01	1.53-6.77	3	1.04 +/- 0.07	1.00-1.14	4
LOC143915	rs1078	A,G	11q23.2	2.02 +/- 1.38	1.11-3.61	3	2.88 +/- 1.66	1.07-5.06	4
?	rs17145	C+,T	?	3.48 +/- 2.99	1.45-7.90	4	1.49 +/- 0.30	1.24-1.82	3
VAV2	rs16763	A,C+	9q34.1	3.14 +/- 1.85	1.12-4.74	3	4.18 +/- 2.13	1.31-6.44	4
KRT13	rs1031	G,T	17q21-q23	3.92 +/- 3.25	1.05-7.45	3	4.00 +/- 2.55	1.12-5.98	3
?	rs1169	C+,T	?	2.03 +/- 1.20	1.10-3.75	4	1.05 +/- 0.02	1.04-1.07	2
C21orf25	rs3169	A+,G	21q22.3	1.12 +/- 0.04	1.07-1.16	3	3.11 +/- 3.18	1.03-6.78	3
?	rs17035	A,T	?	1.47 +/- 0.61	1.09-2.18	3	1.87 +/- 0.66	1.12-2.37	3
FLJ20701	rs1498	A,G+	2q37.1	5.05 +/- 6.93	1.01-13.04	3	1.75 +/- 0.69	1.18-2.52	3
LAMR1	rs2723	C,T+	3p21.3	1.29 +/- 0.41	1.04-1.76	3	1.82 +/- 1.18	1.10-3.19	3
DHCR7	rs7690	C,T+	11q13.2-q13.5	1.53 +/- 0.58	1.02-2.17	3	1.13 +/- 0.10	1.02-1.22	3
CHAF1A	rs2992	C,T	19p13.3	2.28 +/- 1.24	1.03-3.51	3	1.95 +/- 0.89	1.17-2.92	3
LOC162007	rs2993	G,T	16p13.2	7.20 +/- 5.99	1.59-13.51	3	3.51 +/- 2.99	1.35-6.92	3
KIAA0582	rs1228	A+,G	2p13.3	1.16 +/- 0.19	1.04-1.38	3	2.87 +/- 2.01	1.52-5.17	3
?	rs1233	C+,T	?	5.99 +/- 9.17	1.24-19.74	4	1.22 +/- 0.10	1.15-1.29	2
MATN2	rs2615	A,G+	8q22	1.08 +/- 0.06	1.02-1.15	3	2.04 +/- 0.81	1.12-2.67	3
SERPINA1	rs1303	G+,T	14q32.1	1.77 +/- 0.74	1.27-2.62	3	1.24 +/- 0.21	1.10-1.48	3
?	rs3133	A+,G	?	2.14 +/- 1.01	1.43-3.29	3	1.64 +/- 0.71	1.07-2.43	3
FLJ20051	rs2783	C,T+	5p15.2	2.34 +/- 1.66	1.39-4.26	3	1.35 +/- 0.23	1.17-1.60	3
?	rs1225	A,T+	?	1.66 +/- 0.68	1.23-2.68	4	3.17 +/- 2.71	1.25-5.08	2
PEPD	rs3556	C+,T	19q12-q13.2	1.14 +/- 0.14	1.05-1.30	3	1.51 +/- 0.66	1.11-2.27	3
?	rs17036	A,G	?	3.67 +/- 1.65	1.83-5.02	3	4.18 +/- 2.27	2.18-6.65	3
?	rs2976	A,G	?	3.40 +/- 2.74	1.78-6.55	3	3.35 +/- 1.86	2.24-5.50	3
LOC153527	rs6848	A,T+	5q31.3	1.66 +/- 1.02	1.01-2.83	3	1.45 +/- 0.27	1.15-1.68	3
PEPD	rs1038	C+,T	19q12-q13.2	1.11 +/- 0.08	1.05-1.20	3	1.68 +/- 0.61	1.03-2.23	3
SEC22A	rs1944	C+,T	3q21.1	1.48 +/- 0.29	1.22-1.80	3	2.82 +/- 1.89	1.06-4.81	3
FLJ11125	rs1456	A,G	8p21.3	3.00 +/- 1.95	1.07-4.96	3	2.15 +/- 1.29	1.20-3.62	3
LOC59346	rs3064	C,G+	8p21.2	1.25 +/- 0.16	1.07-1.37	3	2.80 +/- 2.91	1.11-6.16	3
?	rs1958	C+,T	?	2.40 +/- 1.10	1.36-3.56	3	3.28 +/- 1.81	1.40-5.02	3
LOC165581	rs2947	A,G	2q35	3.65 +/- 2.67	1.43-7.17	4	4.43 +/- 0.03	4.41-4.45	2

?	rs2649	A,G+	?	1.35 +/- 0.23	1.09-1.52	3	1.72 +/- 0.45	1.43-2.24	3
RAI14	rs1390	A,G+	5p13.3-p13.2	3.38 +/- 3.28	1.23-7.15	3	5.64 +/- 3.06	2.81-8.88	3
?	rs2891	C,T+	?	2.52 +/- 2.14	1.27-4.99	3	1.34 +/- 0.21	1.21-1.58	3
?	rs1318	C,T+	?	1.69 +/- 0.92	1.08-3.06	4	1.33 +/- 0.04	1.30-1.35	2
VEZATIN	rs3596	C+,T	12q22	1.98 +/- 1.21	1.27-3.38	3	1.21 +/- 0.19	1.05-1.43	3
TAP2	rs17034	A,G	6p21.3	3.40 +/- 2.97	1.40-7.79	4	1.96 +/- 0.61	1.53-2.40	2
CNTNAP2	rs3194	G+,T	7q35-q36	1.12 +/- 0.12	1.02-1.26	4	6.67 +/- 6.84	1.83-11.51	2
LOC153277	rs3533	C,T	5q23.2	2.26 +/- 1.34	1.46-3.81	3	2.08 +/- 1.17	1.09-3.37	3
ADORA3	rs3393	A,G+	1p21-p13	1.77 +/- 0.65	1.19-2.47	3	1.58 +/- 0.62	1.17-2.30	3
LOC93538	rs2221	A+,G	10q11.21	2.65 +/- 2.16	1.12-4.17	2	2.07 +/- 1.21	1.22-3.45	3
LOC169515	rs1917	A,G	9p22.2	2.94 +/- 1.95	1.11-4.99	3	7.14 +/- 2.57	5.32-8.95	2
TG	rs3161	C,T+	8q24.2-q24.3	2.08 +/- 0.47	1.75-2.42	2	1.36 +/- 0.49	1.03-1.93	3
LOC131022	rs2800	C+,T	3q24	4.11 +/- 4.39	1.01-7.22	2	3.79 +/- 2.35	1.20-5.78	3
?	rs1978	A,T+	?	3.21 +/- 3.33	1.27-7.06	3	3.11 +/- 2.95	1.03-5.19	2
FBLN1	rs1119	C+,G	22q13.31	3.08 +/- 2.11	1.52-5.49	3	2.00 +/- 1.37	1.03-2.98	2
?	rs2872	C,G+	?	2.16 +/- 1.20	1.32-3.01	2	2.04 +/- 1.57	1.04-3.85	3
?	rs1434	G+,T	?	1.90 +/- 0.30	1.69-2.12	2	2.31 +/- 0.36	1.90-2.60	3
SDC4	rs3529	A,G+	20q12	1.85 +/- 1.15	1.04-2.67	2	1.22 +/- 0.21	1.06-1.46	3
?	rs1444	A+,G	?	3.13 +/- 2.07	1.10-5.24	3	4.72 +/- 3.12	2.51-6.93	2
DISC1	rs3524	C,T	1q42.1	2.09 +/- 1.42	1.08-3.09	2	1.78 +/- 0.85	1.06-2.71	3
LOC51008	rs3312	C,T+	10pter-q25.3	1.72 +/- 0.44	1.26-2.14	3	1.33 +/- 0.37	1.07-1.59	2
PAPPA	rs1405	C,T	9q33.1	2.77 +/- 2.00	1.01-4.93	4	6.85	6.85-6.85	1
?	rs3466	C,T	?	1.07 +/- 0.03	1.05-1.09	2	2.39 +/- 0.99	1.43-3.42	3
PAXIP1L	rs3501	A+,G	7q36	1.48 +/- 0.62	1.04-1.91	2	2.35 +/- 1.30	1.20-3.77	3
VELI1	rs1537	C,G	12q21	1.64 +/- 0.14	1.54-1.73	2	5.05 +/- 2.87	3.01-8.33	3
LIPC	rs690	A,C	15q21-q23	7.00 +/- 8.01	1.33-12.67	2	2.80 +/- 1.49	1.11-3.89	3
LSS	rs2968	C,T	21q22.3	2.36 +/- 1.86	1.04-3.67	2	1.96 +/- 1.60	1.02-3.81	3
LOC148281	rs3838	A+,C	1p11.2	3.07 +/- 1.29	1.60-4.01	3	4.25 +/- 1.48	3.20-5.29	2
LOC123429	rs3080	A,T+	15q22.2	2.46 +/- 1.48	1.45-4.15	3	7.12 +/- 0.54	6.74-7.50	2
KIAA1867	rs1874	A+,G	11q24	1.90 +/- 0.92	1.00-2.84	3	1.54 +/- 0.61	1.11-1.97	2
FLJ14594	rs3144	C,T+	15q24.1	2.26 +/- 2.14	1.01-4.74	3	1.49 +/- 0.12	1.40-1.57	2
KRN1	rs2664	C,T	11q13.5	12.87 +/- 16.43	1.01-31.62	3	1.55 +/- 0.59	1.13-1.97	2
THPO	rs6141	A+,G	3q27	14.16 +/- 18.20	1.29-27.02	2	1.44 +/- 0.24	1.28-1.61	2
?	rs3366	C,T+	?	1.66 +/- 0.19	1.53-1.80	2	2.58 +/- 1.27	1.68-3.47	2
LOC81691	rs6654	A,G	16p12.2	5.37 +/- 4.64	2.09-8.65	2	4.47 +/- 4.80	1.08-7.87	2
SCA1	rs16885	C,T	6p23	1.93 +/- 1.23	1.07-2.80	2	3.35 +/- 2.78	1.39-5.32	2
FLJ10509	rs2347	A,C+	16p11.2	1.97 +/- 0.57	1.57-2.37	2	1.64 +/- 0.06	1.59-1.68	2

KIAA0409	rs2890	C,T+	11p15.4	1.88 +/- 1.01	1.17-2.60	2	1.19 +/- 0.06	1.14-1.23	2
?	rs16810	C+,T	?	2.78 +/- 0.56	2.38-3.18	2	4.27 +/- 2.36	2.60-5.94	2
GP	rs3719	C,T+	5q35.3	4.33 +/- 2.23	2.75-5.90	2	1.93 +/- 0.88	1.30-2.56	2
?	rs691	C,T+	?	6.44 +/- 4.24	1.88-10.26	3	3.73	3.73-3.73	1
LHFP	rs3108	C,T	13q12	4.31 +/- 2.44	2.58-6.03	2	3.55 +/- 2.83	1.54-5.55	2
PIP5K1B	rs2017	C+,T	9q13	1.93 +/- 1.12	1.14-2.72	2	1.16 +/- 0.03	1.14-1.19	2
PB1	rs1570	A,T	3p21	2.72 +/- 1.50	1.03-3.90	3	1.18	1.18-1.18	1
DKFZP434N1	rs2975	C,T+	3p25.1-p24.3	4.52 +/- 4.96	1.15-10.21	3	1.27	1.27-1.27	1
?	rs3293	A+,C	?	3.32 +/- 2.64	1.45-5.18	2	4.59 +/- 4.54	1.38-7.79	2
LOC158326	rs1724	C+,T	9p22.3	1.30 +/- 0.11	1.22-1.37	2	5.89 +/- 5.23	2.19-9.58	2
?	rs3751	A,C	?	7.71 +/- 9.45	2.01-18.61	3	1.42	1.42-1.42	1
ACOX1	rs3682	A+,G	17q24-17q25	4.89 +/- 5.30	1.15-8.64	2	1.71 +/- 0.95	1.03-2.38	2
SMURF1	rs3294	C,G	7q21.1-q31.1	1.76 +/- 0.98	1.07-2.45	2	2.33 +/- 1.41	1.33-3.32	2
COL21A1	rs1480	A+,G	6p12.3-p11.2	2.97 +/- 3.20	1.10-6.66	3	1.33	1.33-1.33	1
?	rs1509	A,C	?	3.23 +/- 0.62	2.79-3.68	2	2.93 +/- 2.67	1.04-4.82	2
?	rs17026	C+,T	?	6.35 +/- 7.41	1.11-11.59	2	6.43 +/- 7.32	1.25-11.60	2
HEF1	rs2034	A,C+	6p25-p24	8.29 +/- 10.20	1.07-15.50	2	2.25 +/- 1.23	1.37-3.12	2
?	rs17050	C,T+	?	4.38 +/- 4.48	1.54-9.54	3	1.48	1.48-1.48	1
FLJ23027	rs2943	A,G+	14q32.32	1.81 +/- 0.99	1.11-2.52	2	2.10 +/- 1.51	1.03-3.17	2
FXVD5	rs3458	C,T	19q12-q13.1	4.98 +/- 1.56	3.87-6.08	2	3.23 +/- 2.87	1.20-5.26	2
?	rs3222	G+,T	?	11.49 +/- 6.23	7.08-15.89	2	1.85 +/- 0.22	1.69-2.00	2
LOC164439	rs1533	A,G+	21q11.2	1.45 +/- 0.57	1.05-1.86	2	1.85 +/- 1.05	1.11-2.59	2
?	rs3039	A,G+	?	1.07 +/- 0.01	1.06-1.07	2	2.31 +/- 0.59	1.89-2.73	2
ALDH1B1	rs3043	C,G+	9p13	2.33 +/- 1.52	1.26-3.40	2	2.17 +/- 0.10	2.10-2.25	2
TNFRSF19	rs1630	A+,G	13q12.11-q12.3	1.07 +/- 0.09	1.01-1.13	2	2.67 +/- 2.23	1.09-4.24	2
SLC5A6	rs1395	A+,G	2p23	1.73 +/- 0.92	1.08-2.38	2	1.13 +/- 0.06	1.08-1.17	2
LOC120425	rs1806	A,G	11q23.3	2.52	2.52-2.52	1	1.84 +/- 0.82	1.02-2.66	3
TNKS	rs1393	C,G+	8p23.1	1.70 +/- 0.47	1.36-2.03	2	2.49 +/- 1.13	1.69-3.29	2
LOC90075	rs1811	A,G+	19q13.12	2.74 +/- 2.14	1.23-4.25	2	1.62 +/- 0.19	1.49-1.76	2
?	rs3480	A+,G	?	1.90 +/- 1.16	1.07-2.72	2	2.16 +/- 0.78	1.61-2.71	2
ADORA3	rs923	C,T+	1p21-p13	1.93 +/- 1.28	1.02-2.84	2	2.32 +/- 1.52	1.25-3.39	2
UGDH	rs1450	C+,T	4p15.1	5.54 +/- 6.36	1.04-10.03	2	1.54 +/- 0.27	1.35-1.74	2
LAMB3	rs2566	C+,T	1q32	1.27 +/- 0.08	1.21-1.33	2	3.59 +/- 3.41	1.18-6.01	2
C11orf25	rs3550	A,C	11p14.3-p14.2	5.29 +/- 4.89	1.09-10.66	3	1.46	1.46-1.46	1
LEPROTL1	rs2657	A+,G	8p21.2-p21.1	1.30 +/- 0.26	1.12-1.48	2	2.13 +/- 1.12	1.35-2.92	2
?	rs1900	C+,T	?	1.18 +/- 0.19	1.05-1.32	2	2.19 +/- 0.22	2.03-2.35	2
SCLY	rs17069	A+,C	2q37.3	2.80 +/- 2.17	1.48-5.30	3	1.43	1.43-1.43	1

?	rs3469	C,T+	?	1.33 +/- 0.03	1.31-1.34	2	2.42 +/- 1.40	1.43-3.41	2
?	rs3306	A+,G	?	1.69 +/- 0.76	1.15-2.23	2	1.68 +/- 0.34	1.44-1.92	2
LOC161986	rs1224	C+,T	16q12.1	1.97 +/- 1.09	1.19-2.74	2	1.98 +/- 1.10	1.21-2.76	2
ARHA	rs3448	A,G+	3p21.3	1.13 +/- 0.13	1.04-1.23	2	1.75 +/- 0.99	1.05-2.45	2
?	rs1913	A+,G	?	1.34 +/- 0.14	1.24-1.44	2	2.37 +/- 1.68	1.18-3.56	2
SRP19	rs3317	A+,G	5q21-q22	1.63 +/- 0.66	1.00-2.31	3	1.38	1.38-1.38	1
FLJ21324	rs1299	A,G+	21q22.3	1.49 +/- 0.60	1.07-1.91	2	2.04 +/- 1.32	1.10-2.97	2
FANCA	rs1230	A+,G	16q24.3	1.23 +/- 0.09	1.17-1.29	2	3.45 +/- 3.01	1.32-5.58	2
CRADD	rs1585	C,T	12q21.33-q23.1	1.04	1.04-1.04	1	1.98 +/- 0.79	1.12-2.67	3
KIAA0751	rs2003	A,G+	8q22.3-q23.1	1.66 +/- 0.22	1.50-1.82	2	1.69 +/- 0.72	1.17-2.20	2
USP19	rs3212	C+,T		1.93 +/- 0.34	1.65-2.30	3	1.27	1.27-1.27	1
DSCAML1	rs16867	C,T	11q23	6.06 +/- 3.37	2.94-9.62	3	5.45	5.45-5.45	1
?	rs16931	C+,G	?	2.76 +/- 2.20	1.20-4.31	2	1.60 +/- 0.23	1.43-1.76	2
?	rs1935	C,G+	?	1.39 +/- 0.32	1.16-1.62	2	5.78 +/- 5.75	1.71-9.84	2
SH3BGRL2	rs1503	G,T+	6q13-15	1.46 +/- 0.47	1.13-1.79	2	2.54 +/- 1.99	1.13-3.94	2
LOC133080	rs2726	C+,T	4q22.1	1.10 +/- 0.09	1.03-1.16	2	2.63 +/- 1.91	1.28-3.98	2
IFNAR2	rs3153	C+,T	21q22.11	1.80 +/- 0.34	1.56-2.04	2	2.83 +/- 0.97	2.14-3.52	2
FLJ22344	rs2636	A,T	5q14.3	3.10 +/- 1.55	2.00-4.20	2	2.88 +/- 2.48	1.13-4.64	2
PDP	rs911	C+,G	8q22.2	3.86 +/- 4.92	1.01-9.54	3	1.76	1.76-1.76	1
SELL	rs12938	C+,T	1q23-q25	1.92 +/- 0.82	1.33-2.50	2	4.77 +/- 2.00	3.36-6.19	2
PMPCB	rs2878	C,T+	7q22-q32	1.25 +/- 0.27	1.06-1.44	2	2.65 +/- 1.81	1.38-3.93	2
?	rs3611	C,T+	?	2.55	2.55-2.55	1	1.12 +/- 0.02	1.10-1.15	3
?	rs3817	A,C	?	1.47	1.47-1.47	1	4.76 +/- 2.02	3.33-6.19	2
FLJ10569	rs2934	A+,C	8p21.2	5.52 +/- 5.16	1.87-9.17	2	4.03	4.03-4.03	1
?	rs3792	C,T+	?	2.08 +/- 0.88	1.46-2.71	2	1.25	1.25-1.25	1
KIAA0753	rs2009	C+,T	17p13.2	2.12 +/- 0.50	1.77-2.48	2	1.85	1.85-1.85	1
MEOX2	rs17199	A+,G	7p22.1-p21.3	1.17	1.17-1.17	1	3.55 +/- 0.54	3.17-3.94	2
INADL	rs1363	C,T+	1p31.3	2.02	2.02-2.02	1	2.16 +/- 1.47	1.13-3.20	2
F3	rs3354	A+,G	1p22-p21	1.50 +/- 0.35	1.25-1.74	2	7.61	7.61-7.61	1
FACL3	rs1485	C+,T	2q34-q35	1.23	1.23-1.23	1	2.27 +/- 1.57	1.16-3.38	2
PSMF1	rs1217	C+,T	20p12.2-p13	1.37	1.37-1.37	1	1.67 +/- 0.61	1.24-2.10	2
SLC18A2	rs3523	A,T+	10q25	2.64 +/- 1.72	1.43-3.86	2	1.61	1.61-1.61	1
SPTBN1	rs3287	A+,G	2p21	1.41 +/- 0.19	1.27-1.54	2	2.23	2.23-2.23	1
BRCA1	rs16940	C,T+	17q21	2.06 +/- 1.44	1.04-3.09	2	1.21	1.21-1.21	1
MAPK4	rs3288	C+,T	18q12-q21	1.70 +/- 0.59	1.28-2.12	2	1.57	1.57-1.57	1
OPCML	rs3360	C+,T	11q25	1.14	1.14-1.14	1	2.20 +/- 0.26	2.02-2.39	2
LOC142827	rs16872	A,C+	10q23.33	2.17 +/- 0.22	2.02-2.32	2	1.18	1.18-1.18	1

MGC10327	rs3726	A+,G	15q26	2.15 +/- 0.20	2.01-2.29	2	1.51	1.51-1.51	1
LOC127133	rs2752	G,T	1q42.3	2.61	2.61-2.61	1	4.75 +/- 1.95	3.38-6.13	2
RASGRF2	rs1589	C,T+	5q13	4.53 +/- 4.62	1.26-7.80	2	1.58	1.58-1.58	1
?	rs3072	A+,G	?	6.36 +/- 6.67	1.65-11.08	2	2.6	2.60-2.60	1
LOC91496	rs1590	A+,C	9q22.33	1.51	1.51-1.51	1	4.20 +/- 0.29	3.99-4.41	2
CDH13	rs1461	C,T+	16q24.2-q24.3	6.97 +/- 1.77	5.72-8.22	2	2.94	2.94-2.94	1
?	rs1458	A,G+	?	1.95	1.95-1.95	1	2.86 +/- 2.47	1.11-4.60	2
ISG15	rs1921	A,G+	1p36.33	1.87 +/- 1.02	1.15-2.59	2	1.89	1.89-1.89	1
?	rs1863	A+,T	?	2.99 +/- 1.42	1.98-3.99	2	1.32	1.32-1.32	1
LOC128344	rs1668	C,T+	1p12	6.22 +/- 6.86	1.75-14.12	3		1.75-14.12	-
ITPR3	rs3227	C,G+	6p21	1.97 +/- 1.22	1.11-2.83	2	2.94	2.94-2.94	1
FLJ10103	rs1735	C+,T	15q25.3	3.49 +/- 1.94	2.12-4.86	2	1.29	1.29-1.29	1
?	rs1504	G+,T	?	1.92 +/- 1.08	1.09-3.15	3		1.09-3.15	-
KIAA1069	rs2352	A,G+	3q25.31	1.58 +/- 0.62	1.14-2.03	2	3.98	3.98-3.98	1
ONECUT1	rs2460	A,G+	15q21.1-q21.2		-	-	3.16 +/- 3.30	1.07-6.97	3
?	rs1348	A,G	?		1.07-6.97	-	2.40 +/- 0.71	1.78-3.18	3
AMPH	rs2810	A+,G	7p14-p13	1.43 +/- 0.07	1.38-1.48	2	4.32	4.32-4.32	1
EXT1	rs1188	A,G+	8q24.11-q24.13	1.24	1.24-1.24	1	6.56 +/- 7.63	1.16-11.96	2
?	rs1429	C,T+	?	1.34 +/- 0.47	1.00-1.68	2	3.39	3.39-3.39	1
?	rs1180	A,G+	?	1.96 +/- 1.06	1.21-2.71	2	1.63	1.63-1.63	1
TEM7	rs3254	C+,T	17q21.1	1.88 +/- 0.89	1.25-2.51	2	1.9	1.90-1.90	1
?	rs3439	A+,C	?	2.52 +/- 2.12	1.02-4.01	2	1.08	1.08-1.08	1
FLJ20105	rs2302	A+,G	xq12	2.10 +/- 0.89	1.47-2.73	2	2.01	2.01-2.01	1
FLJ11598	rs3371	C+,T	15q15.1	1.07	1.07-1.07	1	2.29 +/- 0.66	1.82-2.76	2
?	rs1272	C,G	?	4.04	4.04-4.04	1	7.03 +/- 4.22	4.04-10.01	2
?	rs1692	A,C+	?	2.51 +/- 2.12	1.01-4.01	2	1.12	1.12-1.12	1
DKFZP727C0	rs1815	C,T+	17q21.31	2.06	2.06-2.06	1	3.83 +/- 3.29	1.50-6.16	2
?	rs2804	A+,T	?	1.33 +/- 0.46	1.01-1.66	2	5.69	5.69-5.69	1
FLJ13639	rs2735	C,G+	13q14.2	3.17 +/- 2.03	1.74-4.61	2	4.57	4.57-4.57	1
LOC166005	rs3092	A,G	3p21.33	2.02 +/- 0.17	1.89-2.14	2	2.82	2.82-2.82	1
KIAA1172	rs1544	A,G	21q22.1	4.87	4.87-4.87	1	3.28 +/- 0.06	3.23-3.32	2
COL1A2	rs17073	C+,T	7q22.1	2.20 +/- 1.66	1.02-3.37	2	15.06	15.06-15.06	1
ZIM2	rs3143	A,G+	19q13.4	4.59 +/- 2.34	2.93-6.24	2	5.72	5.72-5.72	1
NUP133	rs2904	A,C+	1q42.2	1.11	1.11-1.11	1	3.65 +/- 3.31	1.31-5.99	2
FLJ10483	rs2905	A+,G	14q32.13	2.52 +/- 0.32	2.29-2.75	2	1.13	1.13-1.13	1
LOC167262	rs1500	C,G+	5q14.1	1.05	1.05-1.05	1	5.93 +/- 6.82	1.11-10.76	2
DKFZp761D2	rs16844	C,T+	1p31.2	6.94	6.94-6.94	1	1.66	1.66-1.66	1

?	rs1786	C,T+	?	1.12	1.12-1.12	1	2.97	2.97-2.97	1
?	rs1791	A,C+	?	1.51	1.51-1.51	1	4.69	4.69-4.69	1
CHERP	rs17029	A,G+	19p13.1	1.14	1.14-1.14	1	2.85	2.85-2.85	1
TAF2	rs2634	C+,G	8q24.12	1.01	1.01-1.01	1	2.14	2.14-2.14	1
?	rs2589	A,G	?	3.39	3.39-3.39	1	2.82	2.82-2.82	1
KIAA1237	rs2635	C+,G	3q21.2	1.62	1.62-1.62	1	2.51	2.51-2.51	1
IL19	rs1798	C,G+	1q32.2	3.08	3.08-3.08	1	1.14	1.14-1.14	1
CA8	rs3140	A,T+	8q11-q12	5	5.00-5.00	1	1.29	1.29-1.29	1
MGC11349	rs3231	C+,T	3q21.3	1.16	1.16-1.16	1	2.13	2.13-2.13	1
?	rs3671	C,T+	?	2.57 +/- 0.08	2.51-2.63	2			
LOC137446	rs1998	A,G+	8q13.3	12.57 +/- 10.50	5.15-20.00	2			
SLC25A21	rs17191	C+,T	14q11.2	1.91	1.91-1.91	1	2.54	2.54-2.54	1
H-L(3)MBT	rs3205	A+,G	20q13.12	2.74	2.74-2.74	1	2.68	2.68-2.68	1
LOC51131	rs2980	A+,G	13q14.12	1.29	1.29-1.29	1	6.96	6.96-6.96	1
CNR2	rs1105	A+,G	1p36.11	2.27	2.27-2.27	1	15.65	15.65-15.65	1
LOC90139	rs2911	A,T+	11p11.2	1.54	1.54-1.54	1	2.84	2.84-2.84	1
RPS6KA2	rs2984	A,G+	6q27	3.36 +/- 3.03	1.21-5.50	2		1.21-5.50	-
TAF11	rs2985	C,T+	6p21.31	1	1.00-1.00	1	3.22	3.22-3.22	1
PROL3	rs2988	A,G+	4q13.3	5.32	5.32-5.32	1	4.37	4.37-4.37	1
LOC166173	rs2917	C+,G	3q26.1	31.06	31.06-31.06	1	2.38	2.38-2.38	1
PTP4A1	rs1879	C,T+	6q12	4.11	4.11-4.11	1	1.53	1.53-1.53	1
FLJ21877	rs1322	C,T+	2p22.1	1.38	1.38-1.38	1	5.33	5.33-5.33	1
?	rs3622	A,G+	?	2.14 +/- 0.09	2.07-2.20	2			
LOC166670	rs17139	A,G+	4q35.1	2	2.00-2.00	1	5.97	5.97-5.97	1
EIF2B1	rs3704	G+,T	12q24.31	1.05	1.05-1.05	1	3.96	3.96-3.96	1
FLJ14566	rs2867	G+,T	3p21.32	1.52	1.52-1.52	1	15.63	15.63-15.63	1
KIAA0662	rs1963	A,G	9q22.2	4.99	4.99-4.99	1	6.28	6.28-6.28	1
GLP1R	rs1820	A,T+	6p21	3.87	3.87-3.87	1	2.14	2.14-2.14	1
?	rs1207	A+,G	?	2.63 +/- 2.28	1.01-4.24	2			
CSF2RB	rs16845	C,G	22q13.1	2.8	2.80-2.80	1	3.31	3.31-3.31	1
SNRPN	rs705	C+,T	15q12	5.4	5.40-5.40	1	5.21	5.21-5.21	1
?	rs1973	A,C+	?	5.66	5.66-5.66	1	1.15	1.15-1.15	1
HNF3B	rs1974	A,G+	20p11				1.72 +/- 0.63	1.27-2.17	2
LOC145622	rs17068	C+,G	15q11.2	2.42	2.42-2.42	1	1.86	1.86-1.86	1
FANCC	rs3277	A,C+	9q22.3	2.18	2.18-2.18	1	1.65	1.65-1.65	1
LOC132789	rs2709	A+,G	4p13	2.1	2.10-2.10	1	4.25	4.25-4.25	1
MAP3K8	rs3034	A,G+	10p11.2	2.57	2.57-2.57	1	1.4	1.40-1.40	1

LOC160822	rs1292	C,T+	13q31.1	2.91	2.91-2.91	1	2	2.00-2.00	1
KIFAP3	rs4519	A,T+	1q23.2	4.5	4.50-4.50	1	7.28	7.28-7.28	1
?	rs907	A,G	?	14.38	14.38-14.38	1	2.41	2.41-2.41	1
HERC1	rs2318	C,G+	15q22	3.32	3.32-3.32	1	9.54	9.54-9.54	1
RGS6	rs3291	A+,G	14q24.3	3.29	3.29-3.29	1	1.47	1.47-1.47	1
KIAA0205	rs3292	C+,T	1p36.13-q42.3	5.95	5.95-5.95	1	1.98	1.98-1.98	1
?	rs3041	A+,G	?	1.55	1.55-1.55	1	3.69	3.69-3.69	1
ITGB5	rs2675	A+,C	3q21.2	1.04	1.04-1.04	1	6.33	6.33-6.33	1
ITGB5	rs2676	C+,T	3q21.2	3.45	3.45-3.45	1	1.88	1.88-1.88	1
SNTB1	rs1301	C+,T	8q23-q24	3.2	3.20-3.20	1	1.63	1.63-1.63	1
BTAf1	rs1437	C,T+	10q22-q23	2.06	2.06-2.06	1	2.6	2.60-2.60	1
?	rs3376	C,T+	?	3.69	3.69-3.69	1	2.09	2.09-2.09	1
ELAC2	rs2523	C,T+	17p11.2	4.07	4.07-4.07	1	2.28	2.28-2.28	1
?	rs1167	C+,T	?	1.89	1.89-1.89	1	57.32	57.32-57.32	1
XRCC5	rs3834	C,T+	2q35	2.33	2.33-2.33	1	1.97	1.97-1.97	1
?	rs16870	C,T+	?	4.62 +/- 3.33	2.27-6.98	2			
PRKCB1	rs3380	A,T+	16p11.2	2.44	2.44-2.44	1	1.27	1.27-1.27	1
?	rs3077	C,T+	?	3.6	3.60-3.60	1	1.55	1.55-1.55	1
FLJ11598	rs16858	C+,T	15q15.1	1.35	1.35-1.35	1	8.38	8.38-8.38	1
?	rs3455	A+,G	?	3.62 +/- 3.23	1.34-5.91	2			
MYBPC1	rs2764	A,G+	12q23.2	1.2	1.20-1.20	1	11.29	11.29-11.29	1
?	rs1745	A,G+	?	3.12	3.12-3.12	1	1.24	1.24-1.24	1
DELGEF	rs1528	C+,T	11p14.3	13.04	13.04-13.04	1	5.08	5.08-5.08	1
CA3	rs17042	C+,T	8q13-q22	5.14	5.14-5.14	1	1.26	1.26-1.26	1
LOC169166	rs1609	C,T+	8q22.3	2.24 +/- 1.55	1.15-3.34	2			
IFI27	rs2799	C+,G	14q32	1.05	1.05-1.05	1	4.11	4.11-4.11	1
?	rs3481	A+,G	?	1.54	1.54-1.54	1	7.65	7.65-7.65	1
GYG2	rs1268	C,T+	xp22.3	2.01	2.01-2.01	1	5.84	5.84-5.84	1
NUP88	rs2407	C,T+	17p13	2.64	2.64-2.64	1	2.37	2.37-2.37	1
FLJ22761	rs2611	C+,G	10q22.1	1.01	1.01-1.01	1	2.56	2.56-2.56	1
LOC127253	rs16910	G,T+	1p31.1	1.35	1.35-1.35	1	4.41	4.41-4.41	1
LOC131022	rs2801	C+,T	3q24	2.18	2.18-2.18	1	1.2	1.20-1.20	1
LOC128809	rs1761	A,G+	20p11.21	1.66	1.66-1.66	1	2.82	2.82-2.82	1
LOC163096	rs1081	A,T	19q13.11	4.33 +/- 1.94	2.96-5.70	2			
?	rs3113	A+,G	?	4.11	4.11-4.11	1	1.58	1.58-1.58	1
?	rs3130	C+,T	?	1.19	1.19-1.19	1	5.12	5.12-5.12	1
?	rs3132	A+,G	?	1.42	1.42-1.42	1	4.6	4.60-4.60	1

GRF2	rs1772	A,G+	9q34.3	1.47	1.47-1.47	1	6.09	6.09-6.09	1
FSTL1	rs1700	A,G+	3q13.33	1.04	1.04-1.04	1	2.25	2.25-2.25	1
HTR2A	rs3125	C+,G	13q14-q21	1.06	1.06-1.06	1	16.46	16.46-16.46	1
KIAA0699	rs556	A,G+	9q22.2	2.27	2.27-2.27	1	1.22	1.22-1.22	1
KIAA1058	rs2899	A,T+	13q32.3	66.01	66.01-66.01	1			
LOC140404	rs3367	C+,T	21q22.3	5.42	5.42-5.42	1			
?	rs1495	C,T+	?	6.1	6.10-6.10	1			
?	rs1802	C,T+	?	3.06	3.06-3.06	1			
MGA	rs2616	A,G+	15q15	2.4	2.40-2.40	1			
KCNB2	rs3357	A,G+	8q13.2	4.02	4.02-4.02	1			
?	rs3285	A+,G	?				2.52	2.52-2.52	1
SMPD3	rs1750	A+,G	16q22.1	4.36	4.36-4.36	1			
?	rs1774	C,T+	?	4.58	4.58-4.58	1			
MYF6	rs3121	C+,T	12q21	2.59	2.59-2.59	1			
MYO3B	rs17083	C+,T	2q31.1-2q31.2	2.27	2.27-2.27	1			
PRKCM	rs16895	A,G+	14q11	2.49	2.49-2.49	1			
KIAA1867	rs2697	A,T+	11q24				5.22	5.22-5.22	1
NRAP	rs1140	A,G+	10q24-q26				7.77	7.77-7.77	1
FLJ23142	rs3396	G+,T	2q31.1				5.32	5.32-5.32	1
CEBPG	rs2772	A+,C	19q13.12	2.42	2.42-2.42	1			
?	rs3067	C+,T	?				13.58	13.58-13.58	1
?	rs16785	C,T+	?				5.56	5.56-5.56	1
?	rs2931	C+,T	?	2.13	2.13-2.13	1			
FLJ11264	rs3185	G,T+	8p21.2				6.58	6.58-6.58	1
?	rs3309	A+,T	?	3.42	3.42-3.42	1			
?	rs1857	G,T+	?	27.34	27.34-27.34	1			
LOC165141	rs1897	A,G+	2p22.1				3.29	3.29-3.29	1
LOC130582	rs2261	C+,T	2p22.3	2.53	2.53-2.53	1			
RGR	rs3526	A+,G	10q23				4.34	4.34-4.34	1
LOC159888	rs3131	G+,T	11q14.1	3.06	3.06-3.06	1			
LOC163094	rs2909	C+,T	19q13.11	5.39	5.39-5.39	1			
?	rs1089	C+,T	?	2.17	2.17-2.17	1			