

## ERRATUM

# Response to Belgard *et al.*

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*Molecular Psychiatry* (2014) **19**, 743–744; doi:10.1038/mp.2014.27; published online 18 March 2014

**Correction to:** *Molecular Psychiatry* (2014) **19**, 407–409; doi:10.1038/mp.2013.186; published online 14 January 2014

should have been listed as 10 242. In addition, 107 SNPs were omitted. The corrected table appears in its entirety below:

Following publication of this paper, the authors noticed errors in Table 1. The gene number for SNP rs9832697 was missing; it

**Table 1.** List of all 237 SNPs for ASD classifier in the CEU Cohort,<sup>2</sup> organised from highest to lowest median weightings

SNP	Weight lower	Weight median	Weight upper	Gene no.	Gene symbol
rs968122	1.5465	1.5555	1.5645	27 345	KCNMB4
rs876619	0.9476	1.2092	1.4708	2775	GNAO1
rs11020772	0.8553	0.8641	0.8729	2915	GRM5
rs9288685	0.5856	0.5998	0.614	3635	INPP5D
rs10193128	0.5836	0.5946	0.6056	3635	INPP5D
rs7842798	0.5298	0.5386	0.5474	114	ADCY8
rs3773540	0.5125	0.5208	0.5291	55 799	CACNA2D3
rs1818106	0.5002	0.5161	0.532	80 310	PDGFD
rs2384061	0.4195	0.4306	0.4417	109	ADCY3
rs12582971	0.3983	0.4295	0.4607	5288	PIK3C2G
rs10409541	0.4067	0.4189	0.4311	773	CACNA1A
rs2300497	0.3782	0.3889	0.3996	801	CALM1
rs7562445	0.3741	0.3843	0.3945	2066	ERBB4
rs7313997	0.3382	0.3567	0.3752	5801	PTPRR
rs2239118	0.3348	0.3552	0.3756	775	CACNA1C
rs4688054	0.1801	0.3476	0.515	2932	GSK3B
rs10823195	0.2597	0.3445	0.4294	1763	DNA2
rs9798267	0.2759	0.3388	0.4017	84 083	ZRANB3
rs1075354	0.4236	0.3177	0.6402	55 799	CACNA2D3
rs1942052	0.2641	0.3088	0.3535	130 013	ACMSD
rs4696443	0.2525	0.3047	0.3569	23 321	TRIM2
rs243196	0.2402	0.2976	0.3549	1112	FOXP3
rs16929470	0.1854	0.2712	0.3571	775	CACNA1C
rs7580690	0.1647	0.2248	0.285	83 439	TCF7L1
rs7145618	0.1515	0.2238	0.296	5528	PPP2R5C
rs3770132	0.1514	0.2093	0.2673	3676	ITGA4
rs3790095	0.1215	0.2017	0.2819	2775	GNAO1
rs1013459	0.1417	0.1969	0.2522	2774	GNAL
rs11001056	0.1519	0.1891	0.2263	5592	PRKG1
rs10952662	0.148	0.1868	0.2257	26 047	CNTNAP2
rs7756516	0.152	0.1853	0.2186	3120	HLA-DQB2
rs8054767	0.1322	0.1803	0.2284	5579	PRKCB
rs2239028	0.1121	0.1763	0.2405	775	CACNA1C
rs3935743	0.0969	0.1737	0.2505	5336	PLCG2
rs4324526	0.1239	0.16	0.196	53 343	NUDT9
rs11736177	0.1047	0.1525	0.2004	5593	PRKG2
rs3020827	0.1094	0.144	0.1786	3815	KIT
rs976266	0.0975	0.1373	0.1771	114	ADCY8
rs884080	0.0999	0.136	0.172	5590	PRKCZ
rs7903424	0.0927	0.1319	0.1711	6934	TCF7L2
rs4465567	0.0837	0.1224	0.1611	10 125	RASGRP1
rs2270838	0.059	0.1221	0.1852	11 060	WWP2
rs1029088	0.0662	0.1202	0.1741	6262	RYR2
rs1873423	0.0555	0.1193	0.1831	5579	PRKCB
rs16931011	0.0708	0.1177	0.1646	3709	ITPR2
rs2920022	0.0779	0.1164	0.1548	5908	RAP1B
rs10153211	-0.0367	0.1154	0.2675	2903	GRIN2A
rs9469718	0.0241	0.1139	0.2037	2914	GRM4
rs2036109	0.0782	0.1133	0.1484	148	ADRA1A
rs2684777	0.0727	0.1133	0.1539	3480	IGF1R
rs629720	0.065	0.1106	0.1562	80 310	PDGFD

**Table 1.** (Continued)

SNP	Weight lower	Weight median	Weight upper	Gene no.	Gene symbol
rs831550	0.0734	0.1096	0.1458	1716	DGUOK
rs2394538	0.0242	0.1093	0.1945	3098	HK1
rs948992	0.068	0.1068	0.1456	9965	FGF19
rs2174257	0.0668	0.105	0.1432	5592	PRKG1
rs11755661	0.0412	0.1041	0.167	5071	PARK2
rs1928168	0.0657	0.099	0.1322	401 237	LINC00340
rs7100765	0.0434	0.0935	0.1436	5593	PRKG2
rs1369450	0.0563	0.0924	0.1285	114	ADCY8
rs1040336	-0.0615	0.091	0.2435	2272	FHIT
rs10407144	0.0434	0.0872	0.131	773	CACNA1A
rs10794197	0.045	0.0869	0.1287	1488	CTBP2
rs3734464	0.0247	0.0868	0.149	5071	PARK2
rs7864216	-0.0072	0.0863	0.1798	9630	GNA14
rs4254056	0.0432	0.0846	0.126	338 751	ORS2L1
rs988920	0.0453	0.0842	0.1232	9229	DLGAP1
rs12393998	0.0536	0.0839	0.1142	8450	CUL4B
rs872794	0.0413	0.0813	0.1213	3778	KCNMA1
rs2503220	-0.0527	0.0806	0.214	5142	PDE4B
rs10468681	0.0356	0.08	0.1243	2774	GNAL
rs7258489	0.0428	0.079	0.1152	808	CALM3
rs153968	0.0379	0.0765	0.115	5144	PDE4D
rs944761	0.0361	0.076	0.1159	9568	GABBR2
rs2161630	0.0232	0.0754	0.1276	10 725	NFAT5
rs7097311	0.0294	0.0703	0.1111	5593	PRKG2
rs2088747	-0.0137	0.0693	0.1522	11 060	WWP2
rs9832697	-0.0766	0.0689	0.2144	10 242	KCNMB2
rs7731023	0.0343	0.0683	0.1023	6502	SKP2
rs7120612	0.0224	0.0659	0.1094	390 055	ORS2A6
rs2033655	0.0277	0.0647	0.1017	109	ADCY3
rs10768450	0.0261	0.0645	0.1029	119 682	ORS1L1
rs3121309	0.018	0.0631	0.1083	89 780	WNT3A
rs7067880	0.0035	0.0629	0.1222	5593	PRKG2
rs2296328	0.0202	0.0614	0.1027	3710	ITPR3
rs10208384	0.0154	0.0591	0.1027	55 821	ALLC
rs5909975	0.029	0.0577	0.0864	2892	GRIA3
rs10041275	-0.003	0.0573	0.1177	2890	GRIA1
rs2779531	0.0118	0.0559	0.1	9568	GABBR2
rs1869901	0.0159	0.0539	0.0918	5330	PLCB2
rs7809503	0.0119	0.0526	0.0932	51 422	PRKAG2
rs1881638	0.0125	0.0525	0.0924	51 422	PRKAG2
rs13359392	-0.0036	0.0511	0.1058	2890	GRIA1
rs2716191	-0.0002	0.0466	0.0934	5608	MAP2K6
rs11123677	0.0031	0.0462	0.0893	1261	CNGA3
rs2937127	-0.0358	0.0414	0.1187	11 060	WWP2
rs2305347	-0.0055	0.0392	0.0839	23 265	EXOC7
rs2471226	-0.0052	0.0389	0.083	107	ADCY1
rs1881628	-0.0107	0.0368	0.0843	51 422	PRKAG2
rs362249	-0.0042	0.0361	0.0764	51 807	TUBA8
rs4708928	-0.0031	0.036	0.0751	5071	PARK2
rs1266348	0.0028	0.0359	0.0689	5634	PRPS2
rs1186055	-0.0032	0.0358	0.0747	5027	P2RX7
rs1453541	-0.1057	0.0354	0.1766	219 983	OR4D6

Table 1. (Continued)

SNP	Weight lower	Weight median	Weight upper	Gene no.	Gene symbol
rs3746821	-0.0262	0.0335	0.0932	958	CD40
rs220740	-0.0085	0.0332	0.0749	10 846	PDE10A
rs2299679	-0.014	0.0331	0.0801	5332	PLCB4
rs887387	-0.0028	0.0317	0.0662	489	ATP2A3
rs7174459	-0.0092	0.0288	0.0669	4735	NEDD5
rs884399	-0.0073	0.0281	0.0634	5581	PRKCE
rs5021051	-0.0146	0.027	0.0686	2895	GRID2
rs2903813	-0.0208	0.0252	0.0711	3315	HSPB1
rs1062935	-0.0207	0.0245	0.0697	57 521	RPTOR
rs9347553	-0.0154	0.0228	0.0609	5071	PARK2
rs11072416	-0.0259	0.0222	0.0703	6263	RYR3
rs4553343	-0.0304	0.0204	0.0712	2977	GUCY1A2
rs7146234	-0.0132	0.0202	0.0535	5495	PPM1A
rs848282	-0.0191	0.0172	0.0536	55 120	FANCL
rs7962764	-0.0495	0.0126	0.0748	5801	PTPRR
rs12726519	-0.0377	0.0098	0.0572	5321	PLA2G4A
rs718949	-0.0303	0.0093	0.0489	1488	CTBP2
rs1954787	-0.0264	0.0089	0.0441	2900	GRIK4
rs2238079	-0.0283	0.0084	0.045	775	CACNA1C
rs1337420	-0.0398	0.008	0.0558	2898	GRIK2
rs917948	-0.0553	0.0075	0.0704	5536	PPP5C
rs3817222	-0.1848	0.0055	0.1957	4660	PPP1R12B
rs17531147	-0.0612	0.003	0.0672	55 970	GNL1
rs6420481	-0.0462	0.0014	0.0489	57 521	RPTOR
rs38557	-0.0428	-0.0018	0.0392	781	CACNA2D1
rs1430158	-0.0542	-0.0055	0.0432	5136	PDE1A
rs1480645	-0.0542	-0.0055	0.0432	27 115	PDE7B
rs9173	-0.0433	-0.0061	0.0311	661	POLR3D
rs3924047	-0.171	-0.0072	0.1566	22 941	SHANK2
rs756944	-0.0596	-0.0077	0.0442	5608	MAP2K6
rs2248068	-0.0467	-0.0099	0.0269	83 439	TCF7L1
rs12906456	-0.0506	-0.0116	0.0274	399 694	SHC4
rs6593958	-0.0485	-0.0121	0.0243	2005	ELK4
rs4647992	-0.0995	-0.0145	0.0706	4790	NFKB1
rs17193	-0.0551	-0.0149	0.0253	10 846	PDE10A
rs702543	-0.0508	-0.0161	0.0186	5145	PDE4D
rs2283492	-0.0595	-0.0191	0.0213	115	ADCY9
rs542413	-0.0566	-0.0201	0.0165	5581	PRKCE
rs731571	-0.0608	-0.0201	0.0207	4293	MAP3K9
rs2069812	-0.0559	-0.0201	0.0157	3567	IL5
rs697851	-0.0648	-0.0215	0.0219	3708	ITPKB
rs933150	-0.0947	-0.0244	0.0459	326	AIRE
rs4748444	-0.0611	-0.0246	0.012	783	CACNB2
rs7235163	-0.0738	-0.0375	-0.0011	9229	DLGAP1
rs4575213	-0.0731	-0.038	-0.0029	5592	PRKG1
rs11048476	-0.0801	-0.0384	0.0033	3709	ITPR2
rs4145903	-0.0762	-0.0395	-0.0028	783	CACNB2
rs10505029	-0.1011	-0.0404	0.0203	51 366	UBR5
rs1122838	-0.1213	-0.0408	0.0396	9630	GNA14
rs1993477	-0.0818	-0.0434	-0.0049	51 366	UBR5
rs2179871	-0.0912	-0.0454	0.0005	10 369	CACNG2
rs10740244	-0.0892	-0.0467	-0.0041	5592	PRKG1
rs2503220	-0.1151	-0.0472	0.0207	5142	PDE4B
rs1065657	-0.0838	-0.0488	-0.0139	51 465	UBE2J1
rs12714137	-0.1234	-0.0528	0.0179	83 439	TCF7L1
rs7176475	-0.1275	-0.0537	0.0201	123 746	PLA2G4E
rs1937671	-0.0953	-0.0545	-0.0138	5592	PRKG1
rs7079293	-0.0902	-0.0549	-0.0196	10 581	SORBS2
rs1003854	-0.1288	-0.0551	0.0187	326	AIRE
rs919741	-0.0962	-0.0565	-0.0169	815	CAMK2A
rs750438	-0.1075	-0.0574	-0.0074	11 184	MAP4K1
rs6139034	-0.0997	-0.0576	-0.0154	3704	ITPA
rs1554606	-0.1087	-0.0599	-0.0111	6018	IL6
rs7108524	-0.0938	-0.0603	-0.0267	81 286	OR51E3
rs1002424	-0.1023	-0.0626	-0.0229	5562	PRKAA1
rs2239316	-0.1033	-0.0631	-0.0228	1387	CREBBP
rs5030949	-0.157	-0.0653	0.0264	3098	HK1
rs17682073	-0.1006	-0.066	-0.0315	6262	RYR2
rs1872902	-0.1108	-0.0665	-0.0221	80 310	PDGFD
rs11602256	-0.1035	-0.0671	-0.0308	4041	LRPS

Table 1. (Continued)

SNP	Weight lower	Weight median	Weight upper	Gene no.	Gene symbol
rs10770675	-0.1088	-0.0675	-0.0262	5139	PDE3A
rs9347587	-0.1058	-0.0696	-0.0334	5071	PARK2
rs4643498	-0.1246	-0.0778	-0.0309	5136	PDE1A
rs2451078	-0.2263	-0.0802	0.0659	93 492	TPSTE2
rs3771832	-0.2236	-0.0803	0.063	6869	TACR1
rs6508808	-0.1506	-0.0809	-0.0112	6261	RYR1
rs3748386	-0.1475	-0.0831	-0.0187	11 060	WWP2
rs12207523	-0.1622	-0.0831	-0.0041	5169	ENPP3
rs2779556	-0.1702	-0.0867	-0.0031	9568	GABBR2
rs7918241	-0.1351	-0.0898	-0.0444	1488	CTBP2
rs2069837	-0.1527	-0.0924	-0.0321	6018	IL6
rs12972670	-0.1359	-0.0926	-0.0493	390 892	OR7A10
rs2652452	-0.1316	-0.0989	-0.0661	3760	KCNJ3
rs9469529	-0.1532	-0.1004	-0.0475	3710	ITPR3
rs7677751	-0.1496	-0.1023	-0.0551	5156	PDGFRA
rs7926083	-0.1395	-0.1028	-0.0661	341 276	OR10A2
rs12577638	-0.1598	-0.11	-0.0601	2900	GRIK4
rs11102321	-0.1838	-0.1104	-0.037	5906	RAP1A
rs1050395	-0.1576	-0.1154	-0.0732	490	ATP2B1
rs8106399	-0.1586	-0.1196	-0.0805	773	CACNA1A
rs7971175	-0.1566	-0.1201	-0.0836	5502	PPP1R1A
rs11602535	-0.166	-0.1236	-0.0812	219 981	OR5A2
rs11644436	-0.1733	-0.1253	-0.0774	5336	PLCG2
rs10762342	-0.1909	-0.1283	-0.0658	5592	PRKG1
rs11583646	-0.2023	-0.1311	-0.0599	6262	RYR2
rs6118611	-0.1819	-0.1321	-0.0822	5332	PLCB4
rs2587891	-0.1722	-0.1322	-0.0922	2775	GNAO1
rs4651343	-0.1739	-0.1333	-0.0926	5321	PLA2G4A
rs1659506	-0.1761	-0.1363	-0.0966	23 295	MGRN1
rs2271986	-0.1968	-0.1367	-0.0767	4842	NOS1
rs2302898	-0.1775	-0.1375	-0.0975	10 381	TUBB3
rs6971999	-0.2088	-0.1425	-0.0763	26 212	OR2F2
rs2272197	-0.1896	-0.1485	-0.1073	4216	MAP3K4
rs4947963	-0.1867	-0.1493	-0.1119	1956	EGFR
rs7536307	-0.1876	-0.1507	-0.1138	26 289	AK5
rs12462609	-0.2085	-0.151	-0.0936	773	CACNA1A
rs1517521	-0.2925	-0.152	-0.0114	23 180	RFTN1
rs8063461	-0.1865	-0.1534	-0.1203	7249	TSC2
rs888817	-0.1937	-0.1604	-0.1272	5924	RASGRF2
rs922445	-0.2435	-0.1659	-0.0883	2775	GNAO1
rs339408	-0.203	-0.167	-0.131	9322	TRIP10
rs7512378	-0.2068	-0.1691	-0.1314	55 811	ADCY10
rs7870040	-0.2408	-0.1892	-0.1376	774	CACNA1B
rs3904668	-0.2423	-0.2069	-0.1715	29 993	PACS1N1
rs12716928	-0.2784	-0.2073	-0.1362	5336	PLCG2
rs6679454	-0.262	-0.2305	-0.199	1601	DAB2
rs10783235	-0.2951	-0.2549	-0.2147	121 275	OR10AD1
rs3910363	-0.3063	-0.2611	-0.2159	57 495	K1AA1239
rs16853387	-0.3799	-0.3367	-0.2935	26 154	ABCA12
rs17629494	-0.5242	-0.507	-0.4898	5592	PRKG1
rs4648135	-0.5807	-0.526	-0.4713	4790	NFKB1
rs17643974	-0.5527	-0.5424	-0.5321	1488	CTBP2
rs1243679	-0.5771	-0.5674	-0.5577	341 799	OR6S1
rs2240228	-0.5942	-0.5816	-0.569	26 532	OR10H3
rs260808	-0.5938	-0.5836	-0.5734	80 310	PDGFD
rs4128941	-0.6166	-0.6082	-0.5998	8313	AXIN2
rs769052	-0.6321	-0.6235	-0.6149	7322	UBE2D2
rs984371	-0.7273	-0.7181	-0.7089	219 437	OR5L1
rs4308342	-1.0196	-0.8938	-0.768	1633	DCK
rs11145506	-0.94	-0.9172	-0.8944	9630	GNA14
rs905646	-0.97	-0.9624	-0.9548	2915	GRM5
rs6483362	-0.9894	-0.9661	-0.9428	2915	GRM5
rs12317962	-1.4869	-1.32	-1.1531	27 345	KCNMB4
rs8053370	-1.7162	-1.6956	-1.675	2775	GNAO1

Abbreviations: ASD, autism spectrum disorder; CEU, Central European; SNP, single-nucleotide polymorphism. Weight indicates the contribution of each SNP to ASD clinical status. The lower and upper weights represent the 95% confidence intervals (CIs) of the distribution of weights for each SNP.