

Supplementary Table 3: Differential expression Asian Vs Caucasian

Ensembl	HUGO	Log2FC	Log2FC.SE	p.value	FDR
ENSG00000243955	GSTA1.protein_coding	4.13	0.57	6.01E-13	1.74E-08
ENSG00000101842	VSIG1.protein_coding	2.93	0.44	3.42E-11	4.96E-07
ENSG00000104722	NEFM.protein_coding	3.16	0.48	5.64E-11	5.67E-07
ENSG00000105852	PON3.protein_coding	2.75	0.46	2.67E-09	1.55E-05
ENSG00000198914	POU3F3.protein_coding	-5.49	0.93	3.94E-09	1.73E-05
ENSG00000110680	CALCA.protein_coding	-4.73	0.84	1.52E-08	5.17E-05
ENSG00000197888	UGT2B17.protein_coding	-3.95	0.73	6.25E-08	1.45E-04
ENSG00000168269	FOXI1.protein_coding	-4.25	0.79	8.58E-08	1.84E-04
ENSG00000267594	CYP4F24P.unprocessed_pseudogene	3.14	0.59	9.38E-08	1.94E-04
ENSG00000160181	TFF2.protein_coding	-3.77	0.72	1.35E-07	2.45E-04
ENSG00000171102	OBP2B.protein_coding	-4.41	0.85	2.06E-07	3.62E-04
ENSG00000172551	MUCL1.protein_coding	-3.99	0.79	4.33E-07	6.33E-04
ENSG00000256618	MTRNR2L1.protein_coding	-3.25	0.65	6.03E-07	8.12E-04
ENSG00000163286	ALPPL2.protein_coding	-3.08	0.63	1.08E-06	1.26E-03
ENSG00000080293	SCTR.protein_coding	-2.64	0.54	1.21E-06	1.35E-03
ENSG00000166736	HTR3A.protein_coding	-3.00	0.62	1.21E-06	1.35E-03
ENSG00000225329	LHFPL3-AS2.lincRNA	-2.44	0.50	1.32E-06	1.43E-03
ENSG00000185247	MAGEA11.protein_coding	-4.60	0.96	1.62E-06	1.62E-03
ENSG00000273079	GRIN2B.protein_coding	-2.79	0.59	1.82E-06	1.76E-03
ENSG00000231290	APCDD1L-AS1.lincRNA	-2.59	0.55	2.14E-06	1.94E-03
ENSG00000137090	DMRT1.protein_coding	-2.51	0.53	2.46E-06	2.16E-03
ENSG00000198883	PNMA5.protein_coding	-3.87	0.82	2.64E-06	2.25E-03
ENSG00000129988	LBP.protein_coding	-3.50	0.75	3.26E-06	2.68E-03
ENSG00000263711	RP11-169F17.1.lincRNA	-5.22	1.12	3.29E-06	2.68E-03
ENSG00000117148	ACTL8.protein_coding	-3.05	0.66	3.54E-06	2.81E-03
ENSG00000234068	PAGE2.protein_coding	-4.95	1.08	4.28E-06	3.14E-03
ENSG00000131668	BARX1.protein_coding	-2.86	0.63	5.31E-06	3.77E-03
ENSG00000143452	HORMAD1.protein_coding	-3.09	0.68	6.17E-06	4.22E-03
ENSG00000183454	GRIN2A.protein_coding	-2.39	0.53	7.49E-06	4.91E-03
ENSG00000174899	PQLC2L.protein_coding	-3.16	0.71	8.14E-06	5.13E-03
ENSG00000124260	MAGEA10.protein_coding	-5.44	1.22	8.29E-06	5.16E-03
ENSG00000095713	CRTAC1.protein_coding	-2.83	0.64	8.58E-06	5.21E-03
ENSG00000227195	MIR663AHG.lincRNA	-4.96	1.12	8.68E-06	5.21E-03
ENSG00000101441	CST4.protein_coding	-3.53	0.80	8.91E-06	5.26E-03
ENSG00000146013	GFRA3.protein_coding	-2.54	0.57	9.17E-06	5.37E-03
ENSG00000101210	EEF1A2.protein_coding	-2.95	0.67	1.24E-05	6.84E-03
ENSG00000157005	SST.protein_coding	-3.46	0.79	1.26E-05	6.90E-03
ENSG00000163440	PDCL2.protein_coding	-3.70	0.85	1.36E-05	7.12E-03
ENSG00000204511	MCCD1.protein_coding	-3.77	0.87	1.40E-05	7.23E-03
ENSG00000161055	SCGB3A1.protein_coding	-2.71	0.63	1.44E-05	7.33E-03

ENSG00000075043	KCNQ2.protein_coding	-3.36	0.78	1.73E-05	8.35E-03
ENSG00000259803	SLC22A31.protein_coding	-2.33	0.55	2.11E-05	9.79E-03
ENSG00000181433	SAGE1.protein_coding	-3.86	0.91	2.30E-05	1.04E-02
ENSG00000163064	EN1.protein_coding	-3.75	0.90	3.10E-05	1.34E-02
ENSG00000101746	NOL4.protein_coding	-2.59	0.63	3.51E-05	1.50E-02
ENSG00000188162	OTOG.protein_coding	-2.25	0.55	3.73E-05	1.54E-02
ENSG00000138083	SIX3.protein_coding	-2.69	0.65	3.89E-05	1.60E-02
ENSG00000132874	SLC14A2.protein_coding	-2.35	0.57	4.16E-05	1.65E-02
ENSG00000204421	LY6G6C.protein_coding	-2.76	0.67	4.12E-05	1.65E-02
ENSG00000261409	RP6-24A23.7.sense_overlapping	2.25	0.55	4.57E-05	1.75E-02
ENSG00000078898	BPIFB2.protein_coding	-3.25	0.80	4.86E-05	1.80E-02
ENSG00000197587	DMBX1.protein_coding	-2.24	0.55	5.29E-05	1.93E-02
ENSG00000177468	OLIG3.protein_coding	-3.39	0.84	5.39E-05	1.94E-02
ENSG00000189127	ANKRD34B.protein_coding	-2.28	0.56	5.50E-05	1.94E-02
ENSG00000165588	OTX2.protein_coding	-3.83	0.95	5.48E-05	1.94E-02
ENSG00000175868	CALCB.protein_coding	-2.79	0.69	6.07E-05	2.09E-02
ENSG00000253642	RP11-317N12.1.lincRNA	-4.54	1.13	6.25E-05	2.13E-02
ENSG00000144481	TRPM8.protein_coding	-2.02	0.51	6.76E-05	2.26E-02
ENSG00000197616	MYH6.protein_coding	-2.72	0.68	6.78E-05	2.26E-02
ENSG00000229743	LINC01159.lincRNA	-4.22	1.06	6.94E-05	2.28E-02
ENSG00000243137	PSG4.protein_coding	-2.62	0.66	7.33E-05	2.39E-02
ENSG00000134398	ERN2.protein_coding	-2.15	0.55	8.00E-05	2.56E-02
ENSG00000168878	SFTPB.protein_coding	-2.65	0.67	8.45E-05	2.69E-02
ENSG00000163207	IVL.protein_coding	-3.14	0.80	9.17E-05	2.86E-02
ENSG00000164434	FABP7.protein_coding	-3.71	0.95	9.84E-05	3.03E-02
ENSG00000147381	MAGEA4.protein_coding	-4.52	1.17	1.11E-04	3.32E-02
ENSG00000142549	IGLON5.protein_coding	-2.17	0.56	1.22E-04	3.46E-02
ENSG00000164089	ETNPPL.protein_coding	-2.27	0.59	1.22E-04	3.46E-02
ENSG00000143556	S100A7.protein_coding	-2.95	0.77	1.20E-04	3.46E-02
ENSG00000225667	ENSG00000225667	-3.28	0.85	1.23E-04	3.46E-02
ENSG00000178522	AMBN.protein_coding	-4.35	1.13	1.23E-04	3.46E-02
ENSG00000085465	OVGP1.protein_coding	-2.59	0.68	1.24E-04	3.48E-02
ENSG00000261122	FLJ26245.lincRNA	-3.46	0.91	1.40E-04	3.73E-02
ENSG00000081248	CACNA1S.protein_coding	-3.03	0.80	1.66E-04	4.21E-02
ENSG00000215644	GCGR.protein_coding	-2.71	0.72	1.68E-04	4.21E-02
ENSG00000197172	MAGEA6.protein_coding	-4.37	1.17	1.83E-04	4.52E-02
ENSG00000007350	TKTL1.protein_coding	-2.42	0.65	1.89E-04	4.63E-02
ENSG00000177984	LCN15.protein_coding	-2.59	0.69	1.93E-04	4.66E-02