

	Illumina identifier	HGNC gene symbol	Refseq ID	Ex1 LR(0.08uM+TGFb/UT+DMSO+TGFb)	Ex1 -log10(pVAL(0.08uM+TGFb/UT+DMSO+TGFb))	Ex1 LR(2uM+TGFb/UT+DMSO+TGFb)	Ex1 -log10(pVAL(2uM+TGFb/UT+DMSO+TGFb))	Ex1 LR(0.08uM-TGFb/UT+DMSO)	Ex1 -log10(pVAL(0.08uM-TGFb/UT+DMSO))	Ex1 LR(2uM-TGFb/UT+DMSO)	Ex1 -log10(pVAL(2uM-TGFb/UT+DMSO))	Ex1 LR(2uM-TGFb/0.08uM-TGFb)	Ex1 -log10(pVAL(2uM-TGFb/0.08uM-TGFb))	Ex1 LR(2uM+TGFb/0.08uM+TGFb)	Ex1 -log10(pVAL(2uM+TGFb/0.08uM+TGFb))
ILMN_1766054	ABCA1	NM_005502.2	0.1	0.13	2.39	21.54	-0.01	0.01	1.65	18.49	1.66	16.77	2.29	18.78	
ILMN_1801767	ABHD3	NM_138340.3	-1.08	7.75	0.17	0.27	-1.56	15.36	-0.1	0.11	1.45	11.8	1.25	8.3	
ILMN_1731433	ABP1	NM_001091.2	-0.16	0.21	2.93	22.43	-1.49	6.1	0.78	1.59	2.27	10.97	3.09	21.66	
ILMN_1708672	ACAT2	NM_005891.2	-0.94	7.73	0.2	0.49	-0.41	0.68	0.81	2.54	1.22	4.88	1.14	8.98	
ILMN_1769694	ACCN2	NM_001095.2	1.09	11	1.61	17.8	1.39	13.66	1	7.36	-0.39	0.87	0.51	2.58	
ILMN_1749014	ACLY	NM_001096.2	-0.62	3.92	0.4	1.75	-0.02	0.02	1.04	7.44	1.07	6.74	1.01	7.44	
ILMN_1684585	ACSL1	NM_001995.2	-0.08	0.12	1.59	18.18	-1.25	11.8	0.32	0.78	1.56	14.84	1.66	17.31	
ILMN_1752269	ACSS1	NM_032501.2	0.58	3.49	1.54	15.23	-0.17	0.26	0.83	5.37	1.01	6.66	0.96	6.88	
ILMN_2336595	ACSS2	NM_001076552.1	-0.34	2.48	1.44	20.81	-0.57	4.74	0.73	7.3	1.3	17.39	1.78	23.36	
ILMN_1714197	ACSS2	NM_018677.2	-0.26	1.52	1.36	19.51	-0.51	4	0.77	8.36	1.28	17.56	1.62	21.29	
ILMN_1702858	ADHFE1	NM_144650.2	0.33	1.18	1.59	15.43	-1.21	10.15	-0.08	0.07	1.13	7.6	1.26	10.05	
ILMN_2363634	ADHFE1	NM_001077593.1	0.3	1	1.19	10.63	-0.8	5.14	0.35	0.94	1.15	8.48	0.89	5.75	
ILMN_1778796	ADSS	NM_001126.2	1.79	10.05	1.41	7.15	1.29	4.47	0.92	2.12	-0.38	0.26	-0.39	0.31	
ILMN_2333107	AES	NM_198970.1	1.21	9.86	1.12	9.03	0.83	5.25	1.16	9.28	0.33	0.59	-0.08	0.01	
ILMN_1674759	AGPAT7	NM_153613.2	0.34	1.1	1.23	10.01	0.86	6.17	1.3	12.43	0.45	1.27	0.89	5.02	
ILMN_1809139	AHCTF1	NM_015446.3	1.22	11.68	0.45	2.41	1.08	8.89	0.11	0.12	-0.97	6.14	-0.77	4.91	
ILMN_1669557	AIM1L	NM_001039775.1	0.5	1.65	1.24	8.06	0.39	0.89	1.03	5.73	0.63	1.79	0.74	2.64	
ILMN_1728662	ALDH3B1	NM_000694.2	1.87	16.01	3.07	26.6	0.47	1.33	1.94	17.88	1.47	10.01	1.2	7.85	
ILMN_2363880	ALDH3B2	NM_001031615.1	0.42	1.44	1.55	12.52	0.43	1.46	1.2	10.15	0.77	3.86	1.13	6.79	
ILMN_1763666	ALDH3B2	NM_001031615.1	2.01	17.54	3.74	31.24	1.7	14.8	2.76	29.53	1.06	5.34	1.73	13.29	
ILMN_1755974	ALDOC	NM_005165.2	-1.13	11.41	0.43	2.45	-1.01	11.05	0.43	2.05	1.44	17.13	1.56	15.4	
ILMN_1690252	ALKBH2	NM_001001655.1	1.12	11.83	0.59	4.58	1.9	26.92	0.92	8.06	-0.99	8.06	-0.53	2.9	
ILMN_1692332	ALOX12B	NM_001139.2	1.24	13.37	1.45	16.43	1.49	21.93	1.45	20.54	-0.04	0.03	0.22	0.32	
ILMN_1801851	ALPPL2	NM_031313.1	-0.07	0.1	1.13	10.49	0.14	0.17	1.59	15.71	1.45	12.11	1.2	9.94	
ILMN_1792409	AMOT	NM_133265.2	0.38	0.83	2.71	19.33	-1.56	7.59	-0.09	0.05	1.48	5.66	2.33	14.68	
ILMN_2215824	ANKRD20A1	NM_032250.1	-0.06	0.09	1.06	11.39	-0.5	2.16	0.56	2.55	1.07	7.6	1.12	10.74	
ILMN_2232520	ANKRD28	NM_015199.2	0.89	9.59	1.48	18.23	0.79	5.57	1.06	9.16	0.27	0.47	0.59	4.13	
ILMN_2217601	ANXA9	NM_003568.1	0.21	0.54	1.07	9.89	-0.41	0.96	0.73	2.86	1.14	5.93	0.86	6.02	
ILMN_1709820	AOAH	NM_001637.1	0.2	0.55	1.15	11.51	-0.05	0.05	1.01	7.96	1.06	7.69	0.95	7.52	
ILMN_2408851	ARHGAP30	NM_181720.2	0.28	0.94	1.3	12.59	0.29	0.57	1.3	10.15	1.01	5.6	1.02	7.69	
ILMN_1666076	ARHGEF4	NM_015320.2	1.99	19.99	2.05	20.99	1.65	12.61	1.66	12.24	0.01	0	0.06	0	
ILMN_1721626	ARID5B	NM_032199.1	1.81	17.71	3.47	32.17	-1.89	13.44	-0.03	0.02	1.85	11.21	1.66	14.55	
ILMN_2232463	ARL14	NM_025047.1	0.64	3.06	2.02	16.88	-0.5	1.66	0.96	5.57	1.46	10.61	1.38	9.05	
ILMN_1718046	ARNT2	NM_014862.3	0.87	7.36	2.01	21.24	0.97	8.44	1.92	24.52	0.94	6.64	1.14	9.49	
ILMN_1723412	ASCL2	NM_005170.2	1.48	13.46	1.12	9.42	1.73	20.61	1.11	9.59	-0.62	2.69	-0.36	0.82	
ILMN_1708778	ASS1	NM_000050.4	0.81	5.79	1.73	16.68	0.61	3.85	1.17	12.17	0.57	2.61	0.92	6.02	
ILMN_1748970	ATAD4	NM_024320.2	-0.24	0.82	1.46	15.9	-0.02	0.02	1.31	18.14	1.33	16.7	1.71	17.05	
ILMN_1654322	ATP1B3	NM_001679.2	-0.51	3.23	0.63	4.63	-0.61	4.04	0.68	4.64	1.29	13.18	1.14	9.97	
ILMN_2376771	ATR	NM_001184.2	0.23	0.43	1.51	11.37	-1.09	5.88	0.35	0.57	1.44	8.18	1.28	7.67	
ILMN_1671731	AVPI1	NM_021732.1	1.86	25.43	2.29	30.62	1.21	15.95	1.67	25.02	0.45	2.03	0.44	2.97	
ILMN_1733970	BAG1	NM_004323.4	0.69	4.2	1.14	9.31	0.36	0.67	1.06	5.52	0.7	1.99	0.45	1.31	
ILMN_1771652	BAIAP2L2	NM_025045.4	1.76	24.46	1.96	27.22	1.52	24.88	1.32	19.63	-0.21	0.42	0.2	0.47	
ILMN_1686626	BAT1	NM_080598.4	1.19	11.85	1.09	10.74	1.4	12.68	1.57	14.63	0.16	0.14	-0.1	0.02	
ILMN_17110514	BCL3	NM_005178.2	1.28	12.15	0.74	5.51	1.64	24.11	0.95	9.82	-0.68	4.57	-0.54	2.41	
ILMN_1737314	BCL6	NM_001706.2	0.65	4.38	2.25	23.06	-0.53	1.5	0.52	1.35	1.05	4.74	1.6	14.46	
ILMN_1814686	BCMO1	NM_017429.2	0.15	0.39	2.42	27.66	-1.16	10.71	0.87	6.07	2.03	22.8	2.27	24.17	
ILMN_1814808	BFAR	NM_016561.1	1.16	13.69	0.95	10.82	0.96	8.57	1.05	9.61	0.09	0.07	-0.2	0.35	
ILMN_1712608	BFSP1	NM_001195.2	1.13	10.97	0.87	7.69	0.69	3.93	1	7.52	0.31	0.54	-0.26	0.44	
ILMN_1699989	BNIPL	NM_138278.2	0.47	1.7	1.71	13.89	-1.1	6.31	0.33	0.53	1.43	8.52	1.24	7.74	
ILMN_1677684	BTBD16	NM_144587.2	1	7.23	1.22	9.78	0.75	6.04	1.42	17.59	0.67	4.01	0.22	0.17	

ILMN_2401878	DUSP10	NM_144729.1	-0.31	1.1	1.47	14.7	-1.39	11.73	0.82	4.25	2.21	21.39	1.78	16.44
ILMN_1808391	DUSP4	NM_001394.5	1.46	16.26	0.65	5.39	1.73	19.49	1.05	8.12	-0.68	3.03	-0.81	6.29
ILMN_1652163	DVL2	NM_004422.2	0.12	0.35	1.04	14.16	0.32	1.32	1.11	13.54	0.79	6.55	0.92	10.62
ILMN_1769201	ELF3	NM_004433.3	1.22	9.33	4.88	37.37	-1.4	7.71	1.32	6.59	2.72	20.17	3.65	28.57
ILMN_1658685	ELFN2	NM_052906.3	1.33	13.4	1.03	9.75	1.74	22.58	1.26	13.04	-0.49	1.79	-0.29	0.63
ILMN_1718297	EML4	NM_019063.2	1.49	13.08	1.07	8.43	1.21	10.99	0.82	5.13	-0.39	0.91	-0.42	1.12
ILMN_2091792	ENTPD6	NM_001247.1	1.08	8.83	0.66	4.14	1.08	8.43	0.55	2.18	-0.53	1.63	-0.41	1.17
ILMN_1716507	EPB41L1	NM_012156.2	0.12	0.23	1.13	11.6	-1	6.96	0.33	0.73	1.33	9.82	1.01	8.58
ILMN_1664265	EPHA1	NM_005232.3	1.19	14.53	1.04	12.43	1.04	13.11	0.9	9.75	-0.14	0.2	-0.15	0.15
ILMN_1802646	EPHB6	NM_004445.2	1.43	14.66	1.25	12.56	1.49	14.9	1.52	14.77	0.03	0.01	-0.18	0.15
ILMN_1651699	EPS8	NM_004447.4	-0.26	0.84	1.13	10.85	-2.25	21.53	-1.18	7.04	1.07	5.09	1.39	12.54
ILMN_1803367	EVI1	NM_005241.1	0.06	0.09	1.63	18.76	-0.62	2.09	0.69	2.42	1.31	7.29	1.56	16.22
ILMN_1671766	F12	NM_000505.3	1.21	17.29	0.61	7.05	1.25	21.37	0.44	3.22	-0.81	9.1	-0.6	5.8
ILMN_1758672	FAM107B	NM_031453.2	1.24	9.46	2.1	18.41	0.5	1.88	1.25	10.3	0.75	3.4	0.86	4.41
ILMN_1691760	FAM45A	NM_207009.2	0.83	8	1.86	21.77	0.07	0.09	1.03	9.95	0.96	7.69	1.03	9.5
ILMN_1740466	FAM46A	NM_017633.2	-1.08	13.34	0.22	0.94	-0.64	5.81	0.65	5.79	1.29	16.95	1.3	15.15
ILMN_1757440	FAM69B	XM_001130258.1	0.71	6.09	1.13	12.17	1.04	12.4	1.25	16.23	0.21	0.39	0.42	1.85
ILMN_2239774	FAM83A	NM_032899.4	1.37	12.59	0.54	2.99	1.07	5.23	0.42	0.75	-0.65	1.46	-0.83	5.09
ILMN_1670158	FAM83A	NM_207006.1	1.17	9.44	0.25	0.59	1.27	11.12	0.71	3.66	-0.55	1.79	-0.92	5.57
ILMN_1670807	FAM84B	NM_174911.3	2.03	26.25	3.13	36.7	1.23	14.09	1.71	22.94	0.48	1.93	1.1	12.55
ILMN_1784871	FASN	NM_004104.4	-0.85	4.14	0.45	1.3	0.35	0.71	1.17	7.45	0.82	3.21	1.3	7.01
ILMN_1672536	FBLN1	NM_006486.2	0.83	8.2	1.87	22.19	-0.05	0.06	1.22	16.17	1.27	15.51	1.04	9.82
ILMN_1728799	FBP1	NM_000507.2	0.99	8.35	2.5	25.12	0.26	0.65	1.09	9.87	0.83	5.16	1.52	13.32
ILMN_1652280	FBXO32	NM_058229.2	-1.06	8.62	1.16	9.88	-2.53	19.79	-0.46	0.77	2.07	12.42	2.22	19.52
ILMN_1703955	FBXO32	NM_148177.1	-0.64	3.52	1.74	15.67	-2.77	18.41	-0.16	0.1	2.61	14.62	2.38	19.96
ILMN_1656691	FBXO4	NM_012176.2	1.02	9.33	0.37	1.63	1.34	14.17	0.36	1.11	-0.97	6.87	-0.66	3.77
ILMN_1741096	FDFT1	NM_004462.3	-0.67	7.69	0.33	2.37	-0.4	2.23	0.68	6.06	1.08	12.2	1.01	11.96
ILMN_2206722	FER1L4	NR_001442.2	-0.16	0.22	1.04	6.41	0.09	0.08	1.26	8.72	1.17	6.7	1.2	6.73
ILMN_1772686	FGD3	NM_033086.2	-1.01	8.79	0.99	8.59	-3.11	34.95	-1.37	9.79	1.73	13.09	2	18.64
ILMN_1764619	FLJ45244	NM_207443.1	0.62	3.66	1.22	10.66	-0.48	1.2	0.55	1.49	1.03	4.49	0.6	2.63
ILMN_1737343	FNIP1	NM_001008738.2	-0.57	3.42	0.43	2.07	-0.85	3.61	0.27	0.35	1.12	5.01	1.01	7.34
ILMN_1766650	FOXA1	NM_004496.2	0.93	9.16	2.86	30.88	-0.84	6.26	0.19	0.33	1.03	7.63	1.93	20.18
ILMN_1808707	FSCN1	NM_003088.2	1.04	12.72	0.27	1.37	1.11	20.41	0.19	0.7	-0.92	13.18	-0.77	7.27
ILMN_1658575	GALNT5	NM_014568.1	0.86	10.66	1.86	25.24	-0.08	0.1	0.96	9.61	1.03	9.77	0.99	11.35
ILMN_2211003	GAS2L3	NM_174942.1	-1.08	7.66	0.14	0.19	-1.24	8.27	-0.13	0.12	1.11	5.59	1.21	7.9
ILMN_1699631	GATS	NM_178831.4	-0.64	3.64	0.55	2.79	-1.79	15.22	-0.4	0.83	1.39	8.37	1.2	8.45
ILMN_1774077	GBP2	NM_004120.3	0.61	2	3.62	25.28	-1.84	9.87	1.24	4.53	3.08	19.86	3.01	19.41
ILMN_1756953	GBP6	NM_198460.1	1.65	15.46	3.09	28.89	0.78	4.34	1.38	11.81	0.61	2.05	1.44	11.75
ILMN_1653496	GLUL	NM_001033056.1	-0.11	0.17	0.99	8.41	-0.34	1.4	0.81	7.26	1.15	12.01	1.1	8.36
ILMN_1729417	GNE	NM_005476.3	-0.59	4.32	1.55	17.24	-1.32	15.56	0.03	0.03	1.35	13.93	2.14	21.94
ILMN_2407389	GPNMB	NM_002510.2	0.49	1.96	2.05	17.67	-1.14	6.89	0.97	4.87	2.11	17.04	1.56	11.23
ILMN_1801205	GPNMB	NM_001005340.1	0.99	8.25	2.52	25.09	-0.94	4.47	0.91	3.95	1.85	12.83	1.54	13.38
ILMN_1750497	GPR109A	NM_177551.3	-0.42	1.68	0.83	5.7	-1.48	11.06	0.08	0.07	1.56	10.37	1.25	9.11
ILMN_1744299	GPR110	NM_153840.2	0.89	7.65	1.36	13.77	0.19	0.33	1.19	10.69	1	6.87	0.47	1.89
ILMN_1787932	GPR110	NM_025048.2	0.9	7.79	1.55	16.23	-0.01	0	1.14	6.16	1.15	5.38	0.65	3.75
ILMN_1787648	GPR125	XM_944791.1	-0.19	0.4	0.86	6.2	-1.2	4.45	0.17	0.11	1.37	4.65	1.06	7.15
ILMN_1782944	GPR37L1	NM_004767.3	-0.06	0.09	2.07	23.09	-0.69	3.07	1.14	7.53	1.83	15.25	2.13	21.81
ILMN_1740762	GRB7	NM_005310.2	-0.16	0.33	1.47	14.01	-1.01	6.04	0.27	0.41	1.27	7.76	1.64	14.19
ILMN_2405254	GRB7	NM_001030002.1	-0.12	0.2	1.67	16.39	-0.98	4.45	0.39	0.64	1.37	6.87	1.79	15.95
ILMN_1658464	GTF3A	NM_002097.1	1.17	13.33	0.96	10.48	1.32	15.6	0.84	6.88	-0.48	1.85	-0.21	0.33
ILMN_1685415	HBP1	NM_012257.3	0.22	0.51	1.21	10.04	-0.51	1.61	0.53	1.62	1.05	5.35	0.98	6.24
ILMN_1797728	HMGCS1	NM_002130.6	-0.82	4.33	0.31	0.69	0.01	0.01	1.3	6.83	1.29	5.81	1.12	6.13
ILMN_1668179	HNRNPF	NM_001098204.1	1.31	10.61	0.89	6.13	1.12	9.88	0.64	3.35	-0.48	1.44	-0.41	0.99
ILMN_1657505	HR	NM_018411.3	1.1	8.14	0.43	1.6	1.23	11.75	0.42	1.41	-0.81	4.56	-0.67	2.84
ILMN_1840316			-0.19	0.62	1.37	16.76	-1.22	12.03	-0.07	0.08	1.15	9.21	1.55	17.46
ILMN_1865735			-1.41	14.9	-0.36	1.74	-1.86	10.75	-0.28	0.27	1.58	6.72	1.04	8.91
ILMN_1898059			0.71	4.98	1.97	19.95	-0.58	0.96	0.83	1.86	1.42	4.58	1.26	10.34
ILMN_1879480			0.57	3.63	1.82	19.05	-0.83	4.66	0.66	2.82	1.49	11.42	1.25	10.75
ILMN_1856480			-0.03	0.03	1.06	9.76	-1.27	9.79	-0.25	0.4	1.02	5.42	1.09	8.69
ILMN_1863962			0.6	3.58	3.22	30.38	-0.04	0.02	2.15	14.4	2.19	13.29	2.63	23.71
ILMN_1913701			0.46	2.41	1.32	12.98	0.46	2.1	1.61	19.31	1.15	9.77	0.86	5.94
ILMN_1915076			1.55	18.57	0.7	6.63	1.24	11.51	0.28	0.61	-0.96	6.04	-0.85	7.58
ILMN_1836776			1.64	14.43	0.78	5.05	1.52	12.8	1.17	7.81	-0.35	0.51	-0.86	4.77
ILMN_1823803			0.51	2.42	1.56	14.12	0.32	0.84	1.59	17.14	1.28	10.58	1.05	7.02
ILMN_1904628			1.14	10.17	1.75	17.41	0.47	1.9	1.34	12.89	0.87	5.14	0.6	2.88
ILMN_1881081			-0.98	7.2	0.47	2.04	-1.8	12.08	-0.47	0.84	1.33	5.89	1.45	11.08

ILMN_1724686	CLDN1	NM_021101.3	1.53	12.84	1.77	15.54	-1.25	8.92	-0.71	2.94	0.54	1.29	0.24	0.21
ILMN_1740418	CYP27B1	NM_000785.3	1.46	12.86	0.51	2.46	0.87	8.95	-0.16	0.32	-1.03	10.28	-0.95	5.96
ILMN_2394276	DST	NM_183380.1	1.08	4.36	2.07	11.8	-1.04	3.43	-0.17	0.12	0.87	1.82	0.99	2.91
ILMN_1779711	DTL	NM_016448.1	1.5	11.33	0.67	3.19	1.73	14.13	0.37	0.7	-1.35	7.85	-0.84	3.75
ILMN_2167011	ECHDC1	NM_018479.2	1.01	7.36	0.97	6.99	-0.96	4.91	-1.17	6.76	-0.21	0.17	-0.04	0
ILMN_1687538	ETS1	NM_005238.2	1.09	4.46	1.05	4.23	-1.32	5.39	-0.75	1.66	0.57	0.7	-0.04	0
ILMN_1673721	EXO1	NM_006027.3	1.46	15.11	0.9	8.06	1.39	13.79	0.16	0.22	-1.23	9.58	-0.56	2.83
ILMN_1699623	FAM81A	NM_152450.2	1.08	10.78	0.67	5.35	1.49	15.9	0.31	0.77	-1.18	9.13	-0.41	1.57
ILMN_2376050	FXYD3	NM_005971.2	0.53	2.33	1.32	10.53	-1.22	4.97	-0.76	1.83	0.46	0.48	0.79	3.82
ILMN_1730575	GCLC	NM_001498.2	1.16	12.02	0.61	4.6	0.25	0.5	-0.84	5.06	-1.09	7.27	-0.55	3.04
ILMN_1804357	GNG4	NM_001098721.1	2.01	16.94	1.32	10.22	3.07	40.41	1.88	20.65	-1.19	8.48	-0.68	2.75
ILMN_1802690	GULP1	NM_016315.2	0.38	1.66	1.69	17.3	-1.08	6.24	-0.35	0.61	0.73	2.29	1.31	11.25
ILMN_1694268	HES6	NM_018645.3	-2.29	19.98	-0.85	5.45	-1.37	16.3	-0.69	4.65	0.68	3.84	1.44	10.34
ILMN_2379788	HIF1A	NM_181054.1	0.45	2.72	1.22	13.05	-1.08	7.19	-0.84	4.2	0.25	0.26	0.77	5.64
ILMN_2409220	HMMR	NM_012484.1	-0.57	2.91	0.6	3.29	-1.25	11.14	-0.37	0.97	0.88	4.91	1.17	8.19
ILMN_1875342			1.21	2.7	1.39	3.48	-1.31	2.26	-0.84	0.85	0.47	0.2	0.18	0.01
ILMN_1864166			1.23	10.39	0.26	0.7	1.56	17.17	0.47	1.76	-1.09	8	-0.97	6.22
ILMN_1819608			0.04	0.04	1.04	8.66	-1.56	17.37	-0.62	3.13	0.94	6.16	1	6.93
ILMN_2228035	HS3ST1	NM_005114.2	1.3	12.86	0.78	6.35	2.07	27.77	0.88	6.72	-1.19	10.12	-0.51	2.33
ILMN_1676629	INSIG2	NM_016133.2	0.72	4	1.17	8.67	-1.6	8.89	-0.65	1.48	0.94	2.6	0.44	1.07
ILMN_1742332	KCTD12	NM_138444.3	1.5	9.75	0.9	4.45	-0.73	1.85	-1.22	4.79	-0.49	0.55	-0.6	1.51
ILMN_1739325	LOC284023	XM_941810.2	1.3	17.01	0.73	8.23	1.18	17.91	0.16	0.39	-1.02	12.22	-0.57	4.45
ILMN_1811117	LOC400986	XM_001126815.1	0.91	4.48	1.05	5.78	-1.71	6.96	-1.77	7.08	-0.06	0.02	0.14	0.02
ILMN_1893633	LOC439949	XM_001129241.1	1.13	10.45	0.35	1.48	1.34	10.9	0.29	0.52	-1.05	5.86	-0.77	4.96
ILMN_1718934	LOC728499	XM_001127568.1	0.47	1.07	1.1	5.14	-1.09	3.9	-0.3	0.28	0.79	1.56	0.63	1.29
ILMN_1764362	LYAR	NM_017816.1	1.05	12.59	0.29	1.51	1.66	15.94	0.57	2.1	-1.09	6.49	-0.76	6.94
ILMN_2413898	MCM10	NM_018518.3	1.65	14.13	0.71	4.14	2.14	19.3	0.39	0.74	-1.75	12.11	-0.94	5.38
ILMN_1663575	MGC87042	XM_001128032.1	1.45	15.81	0.59	4.34	1.92	21.02	0.41	1.15	-1.51	12.44	-0.87	6.86
ILMN_1673522	MOCOS	NM_017947.1	1.15	10.82	0.32	1.24	1.21	13.96	-0.05	0.05	-1.25	12.82	-0.83	5.66
ILMN_2049766	NFE2L3	NM_004289.5	1.44	12.93	0.52	2.66	0.88	5.9	-0.21	0.32	-1.09	7.26	-0.92	5.79
ILMN_1773154	NFKBIA	NM_020529.1	-0.45	2	0.82	5.92	-1.21	10.49	-0.35	0.87	0.86	4.65	1.27	9.76
ILMN_1807529	PADI4	NM_012387.1	1.2	12.79	0.43	2.58	1.06	13.37	-0.02	0.02	-1.08	11.91	-0.77	5.75
ILMN_1773760	PAICS	NM_006452.3	1.22	13.8	0.37	2.14	1.65	23.24	0.53	3.06	-1.12	10.83	-0.85	7.27
ILMN_2392546	PAICS	NM_006452.3	1.17	13.44	0.51	3.97	1.53	20.82	0.48	2.45	-1.06	9.77	-0.66	4.88
ILMN_1703324	PDSS1	NM_014317.3	1.06	10.18	0.14	0.29	1.71	15.97	0.35	0.73	-1.36	9.29	-0.92	7.13
ILMN_1693192	P13	NM_002638.2	1	6.07	0.15	0.19	4.35	44.74	0.91	3.6	-3.44	30.73	-0.85	3.74
ILMN_1798841	PLCXD3	NM_001005473.1	0.36	1.4	1.55	15.14	-1.6	14.25	-0.83	4.16	0.77	3.04	1.19	9.43
ILMN_1774336	POLE2	NM_002692.2	1.1	12.92	0.43	3.11	1.64	24.34	0.4	1.86	-1.24	13.84	-0.66	5.28
ILMN_2218780	PPM2C	NM_018444.2	1.17	9.15	1.89	17.11	-1.33	8.65	-0.61	1.79	0.72	2.11	0.72	3.43
ILMN_1660732	PPP2R2B	NM_181676.1	0.15	0.46	1.14	14.4	-1.31	16.68	-0.55	3.38	0.76	5.37	0.99	10.53
ILMN_1801105	PRKCD	NM_006254.3	-0.36	1.59	0.83	6.92	-1.08	8.23	-0.1	0.1	0.99	5.74	1.2	10.14
ILMN_1776105	PSPH	NM_004577.3	1.58	16.44	0.72	5.65	1.72	21.22	0.51	2.29	-1.21	10.35	-0.86	6.22
ILMN_1674135	RALGPS1	NM_014636.1	-0.63	3.62	0.83	5.87	-1.17	10.09	-0.54	2.16	0.64	2.53	1.46	11.62
ILMN_1814305	SAMD9	NM_017654.2	-0.43	2.7	0.64	5.44	-1.75	17.69	-0.82	4.51	0.93	4.86	1.07	10.17
ILMN_2317493	SEMA6D	NM_024966.2	-0.05	0.05	1.23	9.63	-1.6	13.03	-0.78	3.32	0.82	3.08	1.27	8.72
ILMN_1787813	SLC5A3	NM_006933.4	-0.12	0.15	0.91	5.69	-1.86	22.41	-1.12	9.5	0.74	3.75	1.03	5.71
ILMN_1775380	SMOX	NM_175842.1	1.52	21.72	0.85	11.29	2.91	41.84	1.67	19.48	-1.24	10.71	-0.67	6.92
ILMN_2367258	SMOX	NM_175840.1	2.18	19.05	1.35	10.92	3.82	47.02	2.61	29.46	-1.21	7.65	-0.83	4.21
ILMN_1892403	SNORD13	NR_003041.1	0.56	1.53	1.9	11.74	-1.43	6.74	-0.97	2.99	0.47	0.5	1.34	5.93
ILMN_1690017	SPINK5	NM_006846.2	1.17	12.65	0.26	0.96	1.71	19.89	0.4	1.25	-1.31	11.26	-0.92	7.86
ILMN_1761939	TIPIN	NM_017858.1	1.15	9.63	0.8	5.62	1.21	6.93	0.11	0.09	-1.1	4.72	-0.35	0.75
ILMN_2375879	VEGFA	NM_003376.4	1.17	10.98	0.72	5.35	1.64	21.33	0.39	1.45	-1.25	12.03	-0.45	1.69
ILMN_2174729	ZNF318	NM_014345.2	1.04	8.11	1.17	9.76	-1.1	7.89	-0.46	1.33	0.64	2.19	0.13	0.04