

ILMN Identifier		HGNC gene symbol	Refseq ID	1.27	12.97	1.25	13.18	1.58	28.07	1.6	28.33	0.02	0	-0.02	0.01
ILMN_1677814	ABCC3	NM_003786.2	B15_LR(0.08uM+TGFb/UT+DMSO+TGFb)												
ILMN_2330267	ABCE1	NM_001040876.1	1.07	3.8	0.94	3.17	1.22	5.98	1	4.01	-0.21	0.04	-0.13	0.04	
ILMN_1769694	ACCN2	NM_001095.2	0.37	0.97	1.63	12.41	1.18	9.83	1.15	9.2	-0.03	0	1.25	7.63	
ILMN_1683883	ACY1	NM_000666.1	0.92	3.12	1.26	5.71	1.19	9.87	1.09	8.23	-0.1	0.02	0.35	0.26	
ILMN_1708348	ADAM8	NM_001109.3	0.89	4.57	1.88	13.84	1.11	11	1.18	12.01	0.07	0.01	0.99	4.65	
ILMN_2333107	AES	NM_198970.1	0.87	3.01	1.51	7.88	0.99	6.96	1.03	7.27	0.04	0	0.64	1.2	
ILMN_1655645	AK2	NM_013411.3	1.04	6.11	0.74	3.6	1.22	10.63	1	7.35	-0.22	0.07	-0.3	0.35	
ILMN_2096372	ALDH1A1	NM_000689.3	0.58	1.34	1.84	10.21	1.64	8.82	1.52	7.48	-0.13	0.01	1.26	4.79	
ILMN_1702503	ALDH3A1	NM_000691.3	0.68	3.27	2.1	17.2	0.7	4.24	1.07	9.21	0.38	0.4	1.42	9.57	
ILMN_1728662	ALDH3B1	NM_000694.2	0.53	1.62	2.23	16.01	0.55	1.69	1.1	6.52	0.55	0.66	1.7	10.28	
ILMN_1763666	ALDH3B2	NM_001031615.1	0.9	3.37	2.67	16.71	1.44	10.65	1.62	12.79	0.18	0.04	1.77	8.93	
ILMN_1690252	ALKBH2	NM_001001655.1	0.88	4.36	1.72	12.23	1.01	9.51	0.9	7.65	-0.11	0.02	0.84	3.36	
ILMN_1699421	ANXA10	NM_007193.3	1.06	5.09	1.31	7.41	1.16	10.51	1	8.02	-0.15	0.04	0.25	0.17	
ILMN_1720360	BAD	NM_004322.2	0.73	3.03	1.12	6.55	0.95	6.32	1.02	7.01	0.07	0.01	0.39	0.55	
ILMN_1771652	BAIAP2L2	NM_025045.4	1.41	10.46	2.2	18.35	0.92	10.8	1.39	21.01	0.48	1.71	0.8	3.74	
ILMN_1766169	BCAT1	NM_005504.4	1.33	5.74	1.74	8.99	1.32	5.57	1.53	7.1	0.2	0.03	0.41	0.37	
ILMN_2206746	BGN	NM_001711.3	1.48	7.87	0.64	1.96	0.98	7.05	1.07	8.16	0.09	0.01	-0.84	2.53	
ILMN_1790100	C11orf82	NM_145018.2	0.56	0.81	1.05	2.9	1.67	12.09	0.88	3.56	-0.8	1.61	0.48	0.35	
ILMN_1658494	C13orf15	NM_014059.2	2.1	18.01	2.81	24.27	1.59	17.54	1.96	23.9	0.37	0.36	0.72	3.44	
ILMN_1730523	C16orf14	NM_138418.2	-0.03	0.02	0.98	5.4	1.23	11.95	1.04	8.68	-0.19	0.07	1.01	4.61	
ILMN_1656452	C16orf59	NM_025108.2	0.38	0.73	1.33	7.63	1.51	14.93	0.75	4.14	-0.76	2.67	0.95	3.54	
ILMN_1790537	C16orf75	NM_152308.1	0.54	1.84	1.44	9.99	1.45	18.11	0.68	4.68	-0.77	4.24	0.9	3.92	
ILMN_1740170	C22orf16	NM_213720.1	0.66	4.81	1.24	12.42	1.25	21.04	0.97	13.81	-0.28	0.47	0.58	3.15	
ILMN_1761138	C9orf142	NM_183241.1	0.67	2.65	1.02	5.72	1	9.14	0.6	3.34	-0.41	0.6	0.34	0.42	
ILMN_1810992	CAD	NM_004341.3	1.02	3.41	1.27	5.19	1.32	11.65	1.09	8.13	-0.23	0.08	0.24	0.11	
ILMN_1686920	CCDC58	NM_001017928.2	1.12	2.29	1.89	6.07	1.33	7.41	1.24	6.32	-0.09	0.01	0.77	0.73	
ILMN_2412384	CCNE2	NM_057735.1	1.12	3.45	1.9	8.55	1.38	9.54	0.6	1.85	-0.78	1.76	0.78	1.29	
ILMN_2115633	CD320	NM_016579.2	0.79	3.54	1.26	7.99	1.06	8.3	0.77	4.43	-0.29	0.16	0.47	0.91	
ILMN_1711005	CDC25A	NM_001789.2	1.04	4.82	1.45	8.42	1.34	7.7	0.54	1.24	-0.8	1.46	0.41	0.5	
ILMN_1774982	CDC42EP5	NM_145057.2	0.67	2.86	1.11	7.17	0.77	7.08	1.15	13.88	0.38	0.67	0.45	0.9	
ILMN_1769891	CDH4	NM_001794.2	0.78	1.32	1.42	4.29	1.37	11.93	1.09	7.8	-0.28	0.13	0.64	0.55	
ILMN_1689001	CDK4	NM_000075.2	0.59	4.44	1.3	14.19	1.23	15.07	1.15	13.17	-0.08	0.02	0.71	5.13	
ILMN_1712522	CEACAM6	NM_002483.3	1.29	8.41	1.56	11.38	1.04	6.62	0.83	4.15	-0.21	0.06	0.27	0.27	
ILMN_1782050	CEBPD	NM_005195.3	0.81	2.93	1.5	8.5	1.14	12.71	1.14	12.38	0	0	0.69	1.64	
ILMN_1737195	CENPK	NM_022145.3	0.88	1.46	1.46	4.03	1.24	6.65	0.69	2.02	-0.55	0.47	0.58	0.38	
ILMN_2368718	CENPM	NM_001002876.1	1.06	5.96	1.61	11.28	1.75	24.79	0.86	7.64	-0.89	6.09	0.55	1.34	
ILMN_2368721	CENPM	NM_001002876.1	0.66	3.52	1.05	7.92	1.03	9.16	0.26	0.63	-0.76	3.48	0.39	0.89	
ILMN_1668814	CENPM	NM_024053.3	0.77	3.41	1.27	8.18	1.44	15.56	0.73	4.56	-0.7	2.67	0.5	1.09	
ILMN_1669842	CHAF1A	NM_005483.2	0.32	0.81	1.05	7.57	1.08	8.72	0.59	2.66	-0.49	0.77	0.73	3.35	
ILMN_1664630	CHEK1	NM_001274.3	0.91	2.6	1.3	5.23	1.2	6.73	0.49	1.1	-0.71	1.13	0.4	0.29	
ILMN_1668910	CIDEB	NM_014430.2	0.62	3.24	1.23	10.15	1.01	9.5	0.91	7.73	-0.1	0.02	0.61	2.59	
ILMN_1787691	CITED4	NM_133467.2	1.09	5.1	0.53	1.45	0.85	6.89	1.03	9.57	0.18	0.07	-0.56	1.04	
ILMN_1803236	CLCA2	NM_006536.4	0.95	3.75	2.2	13.5	0.58	2.57	1.11	8.6	0.53	0.9	1.25	5.13	
ILMN_1679176	CLCA4	NM_012128.2	0.22	0.14	1.73	6.06	1.62	6.67	1.9	8.73	0.28	0.04	1.52	3.91	
ILMN_2072568	CLDN8	NM_199328.1	2.13	13.13	3.81	24.34	1.21	9.78	1.19	9.29	-0.02	0	1.68	8.81	
ILMN_1746676	CLDN8	NM_199328.1	0.82	2.55	2.33	13.47	1.38	12.95	1.12	8.87	-0.26	0.12	1.51	6.44	
ILMN_1784608	CLEC2B	NM_005127.2	1.21	4.49	1.87	9.29	0.88	4.94	1.04	6.63	0.16	0.03	0.67	1.05	
ILMN_2403247	CMTM7	NM_181472.1	1.08	5.31	0.48	1.26	1.47	9.72	1.06	5.22	-0.41	0.21	-0.6	1.34	
ILMN_1674038	CTSD	NM_001909.3	0.53	1.77	1.01	5.99	1.02	8.44	0.97	7.6	-0.05	0.01	0.48	1.04	
ILMN_1712305	CYBRD1	NM_024843.2	0.98	2.3	2.01	8.21	0.6	1.28	1.03	3.71	0.43	0.16	1.03	1.99	

ILMN_2093748	ZNF669	NM_024804.1	-1.36	6.78	-1.35	6.98	-1.19	9.03	-0.67	2.97	0.52	0.71	0.02	0
ILMN_1689431	APCDD1L	NM_153360.1	1.72	10.78	0.16	0.19	1.31	10.27	0.84	4.39	-0.47	0.51	-1.56	8.68
ILMN_2188264	CYR61	NM_001554.3	1.32	2.54	-0.41	0.3	1.23	4.83	0.81	2.04	-0.42	0.13	-1.72	3.47
ILMN_1677466	DUSP6	NM_001946.2	1.98	7.57	0.59	0.9	1.28	6.63	0.99	4.01	-0.28	0.07	-1.4	3.64
ILMN_1739222	ETV5	NM_004454.1	2.06	7.9	0.9	2.04	1.05	4.1	0.58	1.19	-0.47	0.22	-1.16	2.48
ILMN_2146761	FABP5	NM_001444.1	1.73	7.36	-1.08	3.6	1.64	14.83	0.69	2.94	-0.95	3.8	-2.81	13.03
ILMN_1769388	GJB2	NM_004004.4	3.44	17.3	1.88	8.6	1.48	7.86	0.84	2.57	-0.64	0.55	-1.55	5.35
ILMN_1864372			1.72	9.36	-0.49	1.15	1.17	10.58	0.7	3.97	-0.47	0.75	-2.21	12.13
ILMN_1682717	IER3	NM_003897.3	1.34	3.6	-0.4	0.4	1.09	4.04	0.41	0.56	-0.68	0.59	-1.74	4.85
ILMN_1717046	MOBKL2B	NM_024761.3	0.96	2.88	-0.3	0.35	1.04	5.91	0.76	3.13	-0.28	0.09	-1.26	3.96
ILMN_1721770	PAPPA	NM_002581.3	0.91	3.62	-0.56	1.55	1.13	8.26	0.34	0.74	-0.8	2.6	-1.47	7.06
ILMN_1811470	PLEK2	NM_016445.1	1.43	8.61	-0.71	2.94	1.34	13.47	0.73	4.32	-0.62	1.73	-2.14	13.53
ILMN_1797776	PRSS23	NM_007173.4	0.57	2.57	-1.12	8.42	1.28	10.83	1.17	9.14	-0.1	0.02	-1.69	12.76
ILMN_1730794	SERTAD4	NM_019605.2	0.35	0.37	-0.98	2.83	1.07	7.53	0.58	2.2	-0.49	0.63	-1.33	3.88
ILMN_1720513	SETBP1	NM_015559.1	0.45	2.17	-0.61	4.1	1.01	14.4	1.05	15.03	0.04	0.01	-1.06	8.33
ILMN_1806607	SFN	NM_006142.3	1.13	8.9	-0.12	0.19	1.15	13.41	0.86	8.04	-0.28	0.26	-1.25	9.46
ILMN_1758146	SIRPA	NM_001040023.1	-0.05	0.05	-1.85	17.41	1.36	16.47	0.82	6.71	-0.54	1.64	-1.8	15.36
ILMN_1695058	SLC38A5	NM_033518.1	0.79	1.87	-0.22	0.19	1.09	5.64	0.76	2.7	-0.33	0.12	-1	2.38
ILMN_1716591	SPRR1A	NM_005987.2	2.24	11.32	1.12	4.22	1.34	11.56	1.23	9.7	-0.11	0.02	-1.12	3.36
ILMN_1778668	TAGLN	NM_003186.3	1.39	9.53	-0.17	0.24	0.52	2.86	1.16	12.37	0.64	2.59	-1.56	10.24
ILMN_2138801	TP73L	NM_003722.3	1.15	2.01	-0.33	0.22	1.03	2.31	0.16	0.09	-0.87	0.62	-1.47	2.61
ILMN_1712708	TRIM47	NM_033452.2	0.49	0.97	-0.64	1.73	1.03	5.8	0.47	1.15	-0.56	0.69	-1.13	3.93
ILMN_2048700	ATAD2	NM_014109.2	0.86	1.59	1.54	4.94	1.45	7.13	0.3	0.33	-1.16	2.92	0.68	0.65
ILMN_1709484	BLM	NM_000057.2	0.54	1.63	1.08	6.03	1.67	14.38	0.44	1.08	-1.23	6.38	0.54	1.18
ILMN_1670238	CDC45L	NM_003504.3	1.29	7.34	2.21	15.35	1.92	25.94	0.79	5.75	-1.13	8.71	0.92	3.54
ILMN_1788131	CYP26A1	NM_057157.1	-0.1	0.09	1.71	11.49	-1.2	9.36	-0.43	1.23	0.77	2.46	1.81	10.93
ILMN_1651699	EPS8	NM_004447.4	-0.98	3.49	0.07	0.05	-1.37	9.21	-0.71	2.55	0.66	1.03	1.05	3.3
ILMN_1673721	EXO1	NM_006027.3	1.24	10.33	2.05	19.08	1.92	22.02	0.49	1.77	-1.43	10.89	0.8	4.69
ILMN_1755281	FBXO15	NM_152676.1	-0.29	0.62	0.74	3.91	-1.13	11.4	-0.91	7.59	0.22	0.11	1.03	5.65
ILMN_1809590	GINS2	NM_016095.1	1.44	10.74	2.44	20.21	2.35	34.14	1.33	14.59	-1.03	7.29	1	5.63
ILMN_1838942			0.91	3.94	1.29	7.29	-0.92	6.04	-1.01	7.05	-0.09	0.01	0.38	0.43
ILMN_1703692	LOC647000	XM_929980.2	1.01	3.85	0.58	1.44	1.15	7	0.14	0.13	-1.01	3.69	-0.43	0.46
ILMN_2413898	MCM10	NM_018518.3	1.07	4.02	2.35	13.28	2.38	21.83	1.2	6.85	-1.18	4.72	1.28	4.64
ILMN_1731184	MELK	NM_014791.2	1.01	5.59	0.72	3.32	1.48	15.66	0.46	1.66	-1.03	6.25	-0.29	0.29
ILMN_2148796	MND1	NM_032117.2	0.52	1.64	1.05	6.25	1.48	15.88	0.4	1.3	-1.08	7.03	0.54	1.31
ILMN_1658143	RFC3	NM_002915.3	0.65	1.02	1.1	3.02	0.93	5.18	-0.07	0.06	-1	4.05	0.45	0.27
ILMN_1678669	RRM2	NM_001034.1	1.35	4.9	1.3	4.87	1.6	9.58	0.03	0.02	-1.57	7	-0.04	0.01
ILMN_1806040	TYMS	NM_001071.1	1.02	3.92	1.28	6.11	1.7	20.31	0.5	2.14	-1.2	9.01	0.27	0.16