

ATR data for Transcript SNP IL13_rs20541

CEPH line	CEPH Family	sex	Resting cell lines			Activated cell lines		
			Replicate 1 (SE)	Replicate 2 (SE)	MEAN	Replicate 1 (SE)	Replicate 2 (SE)	MEAN
GM06985	1341-14	m	0.97 (0.04)		0.97	0.83 (0.02)		0.83
GM06994	1340-9	f	1.17 (0.03)	1.12 (0.03)	1.15	1.06 (0.02)	1.1 (0.03)	1.08
GM07022	1340-11	f	0.83 (0.06)	0.95 (0.03)	0.89	0.97 (0.03)	0.96 (0.01)	0.96
GM07034	1341-11	f		1.11 (0.04)	1.11	0.73 (0.05)	0.81 (0.01)	0.77
GM07055	1341-12	f		1.03 (0.03)	1.03	1.04 (0.02)	1.14 (0.03)	1.09
GM10830	1408-1	m	0.79 (0.04)		0.79	0.78 (0.03)		0.78
GM10851	1344-1	m	1.33 (0.09)		1.33	1.17 (0.05)		1.17
GM11840	1349-14	m	0.75 (0.04)	0.77 (0.03)	0.76	0.72 (0.02)	0.93 (0.06)	0.84
GM11992	1362-13	f	0.72 (0.05)	1.16 (0.04)	0.94	0.8 (0.03)	1.32 (0.04)	1.06
GM11993	1362-14	m	0.95 (0.07)		0.95	0.59 (0.02)	0.82 (0.04)	0.70
GM12004	1420-10	f				1.30 (0.02)		1.30
GM12005	1420-11	f	0.83 (0.06)		0.83	0.78 (0.02)	0.82 (0.03)	0.80
GM12043	1346-11	f				0.96 (0.12)	0.98 (0.03)	0.97
GM12144	1334-10	f	0.84 (0.04)	0.72 (0.01)	0.78	0.73 (0.02)	0.85 (0.01)	0.79
GM12146	1334-12	f		0.99 (0.11)	0.99	1.05 (0.02)	0.97 (0.05)	1.01
GM12248	1416-11	f		0.77 (0.02)	0.77	0.89 (0.03)	0.76 (0.03)	0.82
GM12249	1416-12	m	0.88 (0.03)	0.85 (0.03)	0.86	1.14 (0.03)	1.29 (0.03)	1.21
GM12707	1358-1	m	1.18 (0.09)	0.98 (0.06)	1.08	1.52 (0.04)	1.5 (0.03)	1.51
GM12760	1447-9	f		0.82 (0.03)	0.82		0.83 (0.04)	0.83
GM12814	1454-14	f		0.77 (0.03)	0.77	0.74 (0.03)	1.08 (0.02)	0.91
GM12815	1454-15	m	1.11 (0.07)	1.08 (0.04)	1.10	0.95 (0.04)	0.91 (0.03)	0.93
GM12874	1459-11	f	0.74 (0.05)	0.5 (0.02)	0.62	0.59 (0.04)	0.33 (0.02)	0.46
GM13044	13293-13	m	1.0 (0.1)		1.02	0.89 (0.03)		0.89
GM13045	13294-9	f	0.84 (0.04)	0.9 (0.03)	0.87	0.85 (0.04)	0.68 (0.02)	0.77
GM13046	13294-10	m	0.77 (0.09)		0.77	0.92 (0.03)		0.92
GM13048	13048-10	f	0.98 (0.01)	1.35 (0.07)	1.17	0.97 (0.03)	0.88 (0.01)	0.93