

Table S2. qPCR data (Ct values) of 56 genes for 21 cell lines and 3 culture conditions

Two batches of cDNA were used, as indicated between brackets following each gene name. RPLP0 was used as a reference gene. RPLP0 (1) was used for normalization of batch 1 and RPLP0 (2) was used for normalization of batch 2.

See Table S1 for gene symbols. NS = normal skin, PS = psoriasis, AD = atopic dermatitis

n/a = not analyzed

DIAG	STIM	S100A8 (f)	S100A9 (f)	SLPI (f)	SPRR2C (2)	TGFA (f)	TGM1 (f)	TLR2 (f)	TLR3 (f)	TNC (2)	TNF (f)	TNFRSF6 (f)	TP73L (f)
NS1	KGM	23.24	22.61	17.75	21.39	27.89	26.50	31.24	32.47	32.41	34.88	30.36	23.75
NS1	TH1	19.97	19.64	17.75	19.95	28.65	25.05	32.38	33.44	32.29	33.01	31.09	24.17
NS1	TH2	24.05	22.07	19.34	21.26	26.93	26.93	31.32	29.61	28.35	33.95	27.93	22.86
NS2	KGM	23.05	21.85	20.91	23.91	27.96	27.38	32.73	31.98	32.73	34.14	29.87	23.23
NS2	TH1	17.65	18.00	17.87	20.89	27.81	25.28	29.24	31.53	31.57	28.99	29.41	23.08
NS2	TH2	24.52	25.43	20.74	23.73	31.60	30.41	31.00	32.94	27.71	n/a	29.79	25.45
NS3	KGM	21.93	20.61	16.87	18.88	26.28	24.62	32.78	32.97	30.65	32.76	30.80	23.28
NS3	TH1	17.33	17.79	16.63	18.94	26.59	24.55	30.10	31.61	32.23	30.16	29.16	24.06
NS3	TH2	22.86	22.86	17.74	20.70	26.30	26.30	30.21	31.62	32.20	32.20	29.21	23.09
NS4	KGM	23.13	23.02	17.43	23.74	30.10	27.74	31.26	34.31	32.70	35.04	31.91	24.80
NS4	TH1	18.24	18.57	18.85	20.44	27.17	25.34	28.85	31.58	32.70	29.43	30.51	24.47
NS4	TH2	21.76	22.03	17.86	23.83	28.42	27.31	29.48	31.81	29.72	32.21	29.94	22.71
NS5	TH1	18.10	17.89	17.29	19.15	24.62	23.63	29.25	29.90	28.28	28.97	29.15	24.01
NS5	TH2	22.59	22.44	17.90	21.30	23.48	25.62	29.74	28.17	26.64	30.18	28.92	22.76
NS6	KGM	19.26	19.97	18.98	24.22	25.15	25.90	28.48	29.20	28.81	28.14	29.35	24.00
NS6	TH1	17.29	17.68	18.78	22.38	26.34	26.16	26.83	30.47	26.36	28.17	29.94	23.41
NS6	TH2	21.20	22.60	20.17	26.67	27.43	27.17	28.60	29.83	24.43	32.59	29.39	23.49
NS7	KGM	20.20	21.39	19.87	25.08	26.60	26.47	30.64	29.80	29.63	32.34	30.30	24.67
NS7	TH1	17.59	18.01	18.74	20.58	26.51	25.12	28.79	30.01	29.63	27.73	30.28	23.49
NS7	TH2	20.50	22.47	20.56	25.28	26.85	26.41	28.77	29.78	22.47	29.88	29.04	24.66
PS1	KGM	22.03	21.08	17.01	19.90	24.87	25.01	31.90	34.69	28.59	31.05	29.15	23.09
PS1	TH1	18.14	18.59	16.10	18.42	24.15	23.90	28.70	29.74	25.78	28.56	29.15	23.46
PS1	TH2	22.19	22.18	16.96	20.65	25.57	26.08	29.20	30.62	25.66	32.01	28.91	23.31
PS2	KGM	22.55	22.05	16.71	19.84	25.63	24.43	31.91	31.64	26.01	29.86	29.26	22.67
PS2	TH1	19.64	19.10	16.62	18.47	25.54	23.85	31.49	30.64	26.12	29.96	29.59	23.30
PS2	TH2	23.63	23.49	17.30	20.14	25.54	25.54	30.59	29.68	23.12	31.36	29.42	22.30
PS3	KGM	19.25	19.79	16.41	20.31	25.59	25.01	30.32	29.81	29.81	29.92	28.80	23.20
PS3	TH1	17.53	17.15	15.74	19.27	26.30	24.81	27.63	28.42	26.70	27.13	29.00	23.20
PS3	TH2	22.92	22.99	16.71	21.97	26.35	25.90	28.03	29.75	25.92	29.33	27.67	22.63
PS4	KGM	22.70	22.37	17.99	20.78	25.53	26.27	30.05	30.73	23.36	30.41	29.45	22.90
PS4	TH1	17.94	18.34	17.08	19.12	25.38	25.15	27.46	31.59	23.90	26.36	30.10	24.29
PS4	TH2	23.86	24.24	19.09	21.68	28.29	26.94	30.30	31.10	25.58	32.20	30.11	23.17
PS5	KGM	21.61	22.18	18.07	20.45	27.21	26.44	30.99	31.58	28.80	32.07	29.49	23.46
PS5	TH1	17.38	17.83	16.59	19.34	25.70	24.92	28.00	29.70	28.89	25.94	29.26	23.38
PS5	TH2	21.79	23.38	18.20	21.43	27.00	26.68	29.51	31.78	24.93	29.60	n/a	23.58
PS6	KGM	21.80	21.62	18.64	21.01	26.09	26.85	29.02	32.70	22.30	29.75	29.74	24.23
PS6	TH1	18.61	19.18	18.47	20.63	26.51	26.41	27.14	29.79	22.06	27.74	29.34	23.37
PS6	TH2	21.27	22.41	19.52	22.17	27.60	27.31	28.25	31.88	24.18	31.32	28.99	23.25
PS7	KGM	18.97	19.83	18.17	20.83	27.14	26.25	30.22	31.15	26.02	30.11	29.22	23.32
PS7	TH1	17.59	18.01	17.27	19.58	26.95	25.72	28.66	30.86	25.92	27.31	29.73	24.63
PS7	TH2	20.37	21.67	18.96	22.52	28.32	26.64	30.30	32.25	25.57	29.62	29.71	24.25
AD1	KGM	23.12	24.43	19.05	22.96	26.88	27.46	29.14	31.91	29.03	31.33	30.15	23.40
AD1	TH1	18.24	19.62	18.23	21.82	26.73	26.84	26.88	30.50	26.09	27.35	29.31	24.01
AD1	TH2	23.70	24.35	19.30	23.58	27.40	28.09	27.07	29.77	26.11	30.92	29.27	22.82
AD2	KGM	21.84	22.82	17.66	21.16	27.77	25.68	31.84	33.43	25.36	30.37	29.31	23.00
AD2	TH1	18.05	18.84	17.77	20.37	25.64	24.67	29.16	30.22	25.64	27.20	29.08	23.58
AD2	TH2	23.29	23.89	16.60	21.48	27.49	25.90	28.06	30.87	23.54	29.75	29.67	22.92
AD3	KGM	23.13	24.07	18.78	23.03	26.35	26.33	27.74	32.22	27.88	30.44	29.19	22.95
AD3	TH1	16.82	17.80	16.81	22.32	26.81	26.35	25.93	29.59	23.21	25.98	29.48	23.57
AD3	TH2	23.34	25.22	19.27	23.50	27.88	27.15	26.98	30.65	23.59	28.83	29.90	23.50
AD4	KGM	23.20	23.18	18.22	21.69	26.48	26.85	29.88	32.61	24.54	27.78	29.51	23.85
AD4	TH1	18.88	20.44	18.10	21.40	26.75	26.79	28.16	28.85	24.67	27.35	29.42	23.99
AD4	TH2	23.28	24.85	19.53	23.01	29.42	28.20	30.68	33.11	26.14	30.69	30.07	22.86
AD5	KGM	22.90	24.59	18.71	22.13	28.97	27.90	31.87	33.15	30.02	33.03	33.58	23.92
AD5	TH1	17.31	17.97	17.42	21.07	26.45	24.80	28.60	29.10	28.56	26.81	28.60	22.95
AD5	TH2	23.98	24.84	19.05	22.51	27.04	27.72	28.43	30.67	27.30	30.50	28.96	22.66
AD6	KGM	21.13	22.70	19.18	23.02	28.08	26.98	27.93	31.53	25.64	31.09	29.53	23.30
AD6	TH1	17.96	18.45	18.78	21.91	26.90	26.98	27.10	29.62	26.28	27.10	29.53	25.26
AD6	TH2	22.38	24.76	20.28	23.46	27.68	28.19	28.02	31.29	25.65	29.79	29.21	23.37
AD7	KGM	22.40	23.66	18.84	22.44	27.73	27.03	31.64	30.20	27.59	30.64	29.74	24.52
AD7	TH1	16.84	18.29	17.68	20.27	26.68	24.86	27.78	29.50	27.78	30.19	29.08	23.59
AD7	TH2	23.82	25.06	19.60	24.16	27.73	28.10	29.19	30.63	28.03	26.94	28.82	23.50