



Applied Taqman Assay ID	GENE SMI/50L	Screening set (n=23)				Complementary set (n=44)				Whole RCHOP group (n=67)				Screening set (n=23)				Complementary set (n=44)				Whole RCHOP group (n=67)				Cox models for OS MONI data set**	
		coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value	coefficient	p value
H800159383_m1	MARCKS	0.77	0.028	-0.59	0.138	0.19	0.473	0.11	0.74	0.038	-0.48	0.143	0.10	0.687	0.012	0.33	201670 s.at	0.21	0.1958								
H800733982_m1	MIG27165	0.15	0.316	0.01	0.894	0.05	0.525	0.432	0.21	0.162	-0.02	0.836	0.03	0.646	0.181	0.93	217022 s.at	0.03	0.3533								
H800381676_m1	MILF1IP	-0.87	0.015	0.28	0.237	-0.14	0.439	0.012	-0.94	0.011	-0.13	0.905	-0.30	0.057	0.067	0.95	218983 s.at	-0.17	0.2899								
H800135510_m1	MMIE	-0.31	0.009	-0.14	0.224	-0.22	0.004	0.331	-0.22	0.031	-0.04	0.660	-0.12	0.070	0.210	0.90	203434 s.at	-0.20	0.0009								
H800153510_m1	MMIE	-0.30	0.005	-0.14	0.224	-0.19	0.050	0.331	-0.22	0.031	-0.04	0.660	-0.12	0.070	0.210	0.87	203435 s.at	-0.28	0.0029								
H800277143_m1	MYBL1	-0.30	0.041	-0.10	0.434	-0.19	0.050	0.326	-0.18	0.198	-0.06	0.803	-0.10	0.225	0.484	0.93	213906 at	-0.19	0.0029								
H800192322_m1	MYO1E	-0.18	0.357	0.12	0.604	-0.05	0.728	0.318	-0.59	0.007	0.30	0.325	-0.14	0.311	0.008	0.96	203072 at	-0.32	0.0101								
H800172851_m1	MYO1E	-0.45	0.179	-0.07	0.859	-0.28	0.261	0.448	-0.27	0.388	0.30	0.325	-0.14	0.311	0.008	0.89	203675 at	0.26	0.1045								
H800377283_m1	PDOCD4	0.87	0.012	0.11	0.696	0.39	0.057	0.088	0.68	0.050	0.06	0.778	0.23	0.190	0.138	0.92	202731 at	0.11	0.5092								
H800277080_m1	PDE4B	0.99	0.031	0.37	0.303	0.59	0.030	0.310	0.10	0.778	0.29	0.315	0.22	0.333	0.695	0.86	203708 at	0.23	0.0409								
H800277080_m1	PDE4B	0.99	0.031	0.37	0.303	0.59	0.030	0.310	0.10	0.778	0.29	0.315	0.22	0.333	0.695	0.86	211302 s.at	0.13	0.4366								
H800211916_m1	PDGFC	-1.50	0.001	-0.03	0.908	-0.48	0.060	0.068	-0.90	0.023	0.03	0.892	-0.21	0.309	0.042	0.88	218178 at	0.26	0.1355								
H800171473_m1	PRM1	0.34	0.147	-0.07	0.778	0.15	0.370	0.241	0.14	0.540	-0.19	0.351	-0.04	0.789	0.279	0.95	209193 at	0.00	0.9982								
H800170182_m1	PLAU	-0.63	0.026	-0.18	0.527	-0.40	0.047	0.240	-0.27	0.371	-0.24	0.282	-0.25	0.163	0.838	0.88	206479 s.at	-0.15	0.3624								
H800170182_m1	PLAU	-0.63	0.026	-0.18	0.527	-0.40	0.047	0.240	-0.27	0.371	-0.24	0.282	-0.25	0.163	0.838	0.88	211688 s.at	-0.24	0.1214								
H800169473_m1	PRF1	0.51	0.040	-0.31	0.194	0.09	0.585	0.18	0.47	0.047	-0.27	0.161	0.04	0.815	0.015	0.98	214617 at	0.23	0.0332								
H800176988_m1	PRKCB1	0.17	0.596	-0.33	0.430	-0.01	0.957	0.343	0.13	0.667	-0.14	0.669	0.00	0.983	0.544	0.96	206865 s.at	-0.03	0.8616								
H800160482_m1	PRRS4P2	-0.79	0.031	-0.14	0.716	-0.50	0.066	0.239	-0.76	0.036	-0.08	0.785	-0.38	0.112	0.166	0.90	203537 at	-0.44	0.0061								
H800182280_m1	PTRN1	0.25	0.322	-0.17	0.540	0.06	0.740	0.265	0.21	0.436	-0.34	0.130	-0.11	0.530	0.113	0.96	202716 at	0.00	0.9999								
H800191243_m1	RAB33A	-0.52	0.027	-0.21	0.324	-0.35	0.027	0.330	-0.57	0.017	-0.23	0.183	-0.34	0.013	0.249	0.94	206039 at	0.10	0.4895								
H800412094_m1	RAFTLN	-0.75	0.017	-0.34	0.204	-0.50	0.012	0.311	-0.73	0.019	-0.10	0.660	-0.30	0.089	0.092	0.97	212646 at	-0.40	0.0112								
H800748952_s1	SCD	-0.41	0.255	0.24	0.503	-0.09	0.716	0.204	-0.16	0.623	-0.13	0.665	-0.14	0.515	0.942	0.95	200832 s.at	0.14	0.2018								
H800183137_m1	STEBP5	0.46	0.083	0.10	0.695	0.27	0.136	0.331	0.15	0.544	-0.12	0.443	-0.01	0.932	0.392	0.97	201810 s.at	0.27	0.0537								
H800266342_m1	SORL1	-0.24	0.191	-0.05	0.797	-0.15	0.261	0.475	-0.42	0.027	0.07	0.647	-0.42	0.288	0.046	0.98	203509 at	-0.48	0.0010								
H800266342_s1	SOX4	-1.33	0.003	-0.23	0.310	-0.43	0.022	0.068	-0.57	0.116	-0.18	0.289	-0.25	0.103	0.344	0.81	201417 at	-0.26	0.0069								
H800327839_m1	SFRP2	-0.46	0.053	-0.01	0.871	-0.26	0.137	0.215	-0.43	0.080	0.08	0.721	-0.14	0.359	0.125	0.92	212458 at	-0.33	0.0076								
H800230918_m1	SJLF1	-0.40	0.038	0.00	0.894	-0.14	0.216	0.097	-0.12	0.510	0.01	0.939	-0.03	0.761	0.556	0.91	212354 at	-0.16	0.0684								
H800198961_m1	SVFL	-0.69	0.049	-0.39	0.279	-0.55	0.030	0.668	-0.88	0.014	-0.54	0.070	-0.68	0.003	0.498	0.83	201259 s.at	-0.52	0.0011								
H800712040_m1	TCL1A	-0.10	0.297	0.03	0.739	-0.04	0.604	0.335	-0.17	0.061	0.07	0.402	-0.03	0.590	0.048	0.96	208995 s.at	0.04	0.5929								
H800206231_m1	TM4SF13	-0.35	0.155	-0.20	0.490	-0.27	0.113	0.049	-0.26	0.276	-0.12	0.530	-0.17	0.242	0.647	0.94	217979 at	0.10	0.4197								
H800186671_m1	TNKS	-0.54	0.208	-0.32	0.490	-0.43	0.163	0.729	-1.31	0.008	0.05	0.895	-0.43	0.107	0.034	0.92	202561 at	-0.70	0.0002								
H800135770_m1	TOSO	0.50	0.006	0.00	0.996	0.22	0.069	0.041	0.37	0.044	-0.06	0.637	0.08	0.434	0.055	0.98	221601 s.at	0.10	0.3626								
H800180361_m1	TPOD2	-0.38	0.319	-0.14	0.669	-0.24	0.330	0.635	-0.51	0.185	0.11	0.677	-0.09	0.674	0.185	0.85	201690 s.at	0.06	0.7058								
H800377979_m1	TRAF6	-0.10	0.728	0.58	0.090	0.19	0.389	0.133	-0.68	0.051	0.28	0.289	-0.08	0.692	0.029	0.93	208129 at	0.14	0.2074								
H800215487_m1	TSC	0.90	0.003	0.07	0.797	0.44	0.030	0.039	0.52	0.083	-0.07	0.728	0.11	0.501	0.102	0.97	218872 at	0.01	0.9231								
H800190581_m1	VNN2	-0.15	0.553	-0.91	0.005	-0.42	0.027	0.075	-0.07	0.801	-0.68	0.004	-0.38	0.023	0.082	0.95	206922 at	-0.06	0.6010								
H800200695_m1	VSGC4	0.08	0.355	0.09	0.411	0.09	0.216	0.955	0.13	0.172	-0.02	0.822	0.05	0.428	0.257	0.90	204787 at	0.20	0.0190								
H800245183_m1	ZFP361.1	0.63	0.183	0.02	0.943	0.22	0.428	0.301	0.12	0.809	0.24	0.372	0.21	0.370	0.824	0.68	211965 at	-0.11	0.3522								

\* The results are showed for the probeset with the highest correlation with quantitative RT-PCR data, except for APOBEC3G, FDXP1, MME, PDSE1B and PLAU, where the results are showed for the 2 probesets indicated in Table 1 of the manuscript.

\*\* Monli et al. Blood 2005;105:1851-61.