

SUPPLEMENTAL TABLE

Affymatrix ID	Gene Symbol	Fold change	Signal intensity					
		RST2 vs. Daudi	RST2_1	RST2_2	RST2_3	Daudi_1	Daudi_2	Daudi_3
<b><i>Genes downregulated in RST2 cells</i></b>								
219577_s_at	ABCA7	-2.2	81	51	75	148	145	154
212286_at	ANKRD12	-3.1	15	23	37	77	68	86
212280_x_at	Apg4B	-2.9	93	55	38	176	186	171
221653_x_at	APOL2	-4.3	21	27	25	102	130	80
207722_s_at	BTBD2	-3.0	65	85	26	178	149	211
211922_s_at	CAT	-2.0	181	209	218	371	438	425
202770_s_at	CCNG2	-2.0	156	240	251	330	603	393
203522_at	CCS	-2.6	35	57	18	136	93	63
203493_s_at	CEP57	-2.5	55	144	84	240	280	174
218446_s_at	CGI-148	-2.4	62	39	58	125	148	101
203104_at	CSF1R	-2.8	136	169	72	440	433	192
204971_at	CSTA	-4.8	92	107	21	626	208	217
201487_at	CTSC	-2.0	656	384	497	745	1383	895
202969_at	DYRK2	-2.0	566	298	444	757	1136	760
201135_at	ECHS1	-2.7	416	341	385	1007	918	1115
203646_at	FDX1	-2.1	52	97	79	132	213	134
218298_s_at	FLJ20950	-2.1	186	180	138	301	440	320
211829_s_at	GPR30	-2.3	34	27	43	69	110	62
212211_at	GTAR	-2.0	387	396	439	668	954	849
201833_at	HDAC2	-2.0	247	402	341	768	785	470
211530_x_at	HLA-G	-2.3	829	937	585	1877	2287	1253
221480_at	HNRPD	-2.0	171	112	163	238	415	234
208966_x_at	IFI16	-2.1	1480	1340	1333	2882	3341	2303
214453_s_at	IFI44	-10.0	137	93	47	1169	1167	427
204439_at	IFI44L	-24.5	103	11	5	1475	968	469
204415_at	IFI6	-7.1	10	39	9	197	125	83
203153_at	IFIT1	-39.3	3	13	6	487	287	115
204747_at	IFIT4	-2.5	152	116	136	511	276	219
205483_s_at	ISG15	-11.1	415	396	188	5808	3602	1715
205885_s_at	ITGA4	-2.5	83	71	156	209	328	231
212475_at	KIAA0241	-3.0	28	96	27	121	140	185
212428_at	KIAA0368	-2.5	100	138	175	204	396	420
212311_at	KIAA0746	-2.2	554	598	667	1330	1267	1332
217933_s_at	LAP3	-2.3	907	695	1218	1710	2994	1680

35160_at	LDB1	-2.1	122	96	67	208	243	134
201186_at	LRPAP1	-3.2	47	28	82	120	205	184
200796_s_at	MCL1	-2.3	107	160	145	240	338	368
36830_at	MIPEP	-3.8	17	22	39	95	102	102
205395_s_at	MRE11A	-2.3	146	112	131	267	356	256
203061_s_at	NFBD1	-3.2	52	44	57	142	163	188
201502_s_at	NFKBIA	-2.0	270	557	244	999	535	602
203964_at	NMI	-2.7	115	106	142	385	372	224
206023_at	NMU	-2.5	368	278	428	1133	622	888
202869_at	OAS1	-3.1	268	314	188	1099	839	432
204972_at	OAS2	-4.4	288	371	295	1846	1686	712
204300_at	PET112L	-2.0	71	70	78	165	156	106
201120_s_at	PGRMC1	-2.0	180	173	245	329	490	381
213227_at	PGRMC2	-2.3	74	38	83	152	185	120
202392_s_at	PISD	-2.1	104	76	126	191	261	189
201191_at	PITPN	-2.1	108	73	141	219	265	203
217677_at	PLEKHA2	-2.0	88	149	93	224	230	204
211012_s_at	PML	-16.5	26	16	24	741	222	115
204211_x_at	PRKR	-3.5	226	289	122	1264	385	608
210251_s_at	RIPX	-2.1	51	47	69	118	131	104
218209_s_at	RPRD1A	-2.0	130	111	166	236	292	289
208456_s_at	RRAS2	-2.0	294	361	378	712	779	556
216913_s_at	RRP12	-4.1	23	17	64	69	182	176
209486_at	SAS10	-2.6	380	230	561	1037	1197	843
213988_s_at	SAT	-3.7	113	50	72	493	209	172
212417_at	SCAMP1	-3.0	68	68	142	156	416	252
212902_at	SEC24A	-2.0	84	120	128	188	238	221
203155_at	SETDB1	-2.7	121	159	125	559	272	247
202111_at	SLC4A2	-3.3	31	50	23	100	144	101
212147_at	SMG5	-2.1	69	42	80	99	148	162
202864_s_at	SP100	-7.2	46	24	11	247	224	110
208012_x_at	SP110	-3.3	255	305	262	1116	1079	497
216202_s_at	SPTLC2	-2.1	95	122	95	207	248	207
213562_s_at	SQLE	-2.3	159	113	236	406	453	319
208803_s_at	SRP72	-2.2	162	152	250	458	346	440
200887_s_at	STAT1	-4.2	299	289	191	1538	1226	509
201079_at	SYNGR2	-2.0	876	785	829	2045	1344	1477
210875_s_at	TCF8	-3.4	22	36	38	79	142	105

202688_at	TNFSF10	-2.2	55	63	42	159	104	94
205875_s_at	TREX1	-6.9	14	23	31	180	167	125
220998_s_at	UNC93B1	-4.1	28	35	35	150	101	146
201797_s_at	VAR52	-2.0	213	269	196	402	544	424
209053_s_at	WHSC1	-2.3	310	645	339	949	1117	887
204139_x_at	ZNF42	-2.5	45	37	44	88	87	144
<b>Genes upregulated in RST2 cells</b>								
205839_s_at	BZRAP1	3.7	101	157	81	18	31	44
219012_s_at	C11orf30	4.7	93	70	53	8	20	18
205529_s_at	CBFA2T1	3.0	112	171	97	47	36	45
220227_at	CDH4	2.4	120	170	180	64	43	86
215999_at	CDRT1	2.5	163	115	87	42	41	61
214030_at	DKFZp667G2110	2.5	252	303	237	90	92	138
216835_s_at	DOK1	2.5	108	125	86	37	39	51
208138_at	GAS	2.5	212	178	172	96	69	61
220491_at	HAMP	2.9	134	95	94	33	45	34
211406_at	IER3IP1	2.4	480	413	402	153	157	229
215379_x_at	IGLJ3	2.4	1104	1005	1349	630	354	448
205889_s_at	JAKMIP2	2.5	71	88	76	38	22	35
213281_at	JUN	3.4	85	99	99	37	19	28
207723_s_at	KLRC3	2.6	220	281	208	91	88	92
215930_s_at	MGEA6	2.5	54	66	50	30	20	18
89476_r_at	NPEPL1	2.9	147	138	119	54	26	57
205549_at	PCP4	6.0	118	167	276	13	36	46
207537_at	PFKFB1	2.6	152	135	142	39	58	66
210758_at	PSIP1	3.5	181	197	189	63	55	44
221960_s_at	RAB2	2.4	258	198	247	81	108	102
206035_at	REL	2.9	339	390	224	89	69	167
205596_s_at	SMURF2	2.5	442	376	385	118	198	173
219677_at	SSB1	2.7	116	155	73	38	55	34
209228_x_at	TUSC3	2.7	203	291	234	99	67	99