

SSP	Annotation	Spot densities and ratios compared to UT _{t1} (relative t ₀)							
		UT _{t1}	Ratio	T _{t1}	Ratio	T _{t2}	Ratio	T _{t3}	Ratio
213		2869318	1	4508792	1.57	704448.6	0.25	1620015	0.56
215		2190291	1	1211539	0.55	2648142	1.21	6412805	2.93
1102	Falcipain-2 (PF11_0165)	894643.1	1	867554.4	0.97	1054486	1.18	2591236	2.9
1707		1336849	1	1334699	1	2516244	1.88	3913316	2.93
1722		523218.4	1	759024.9	1.45	1902331	3.64	1688286	3.23
1828	Human erythroid alpha-spectrin	3680704	1	6395287	1.74	17274236	4.69	6387526	1.74
2006		2906660	1	2399245	0.83	703373.1	0.24	1071395	0.37
2105		68055.3	1	259310	3.81	206534.4	3.03	117665.2	1.73
2407	Human beta-actin (fragment)	3364660	1	4195220	1.25	4597818	1.37	9063727	2.69
2903	Human erythroid alpha-spectrin	4843321	1	4921287	1.02	14176775	2.93	16156811	3.34
2904	Human erythroid alpha-spectrin	1050788	1	6207556	5.91	3713321	3.53	4896287	4.66
2908	Human erythroid alpha-spectrin	2718422	1	2733009	1.01	7295902	2.68	4799249	1.77
4102		708179.6	1	1431409	2.02	212995.4	0.3	353271.8	0.5
4109		192948.1	1	248409.8	1.29	399896.7	2.07	509413.5	2.64
4110		87676.6	1	73404	0.84	228144	2.6	335853.4	3.83
4302		498470.5	1	604199.6	1.21	198448.3	0.4	258172.3	0.52
5106		1254317	1	1421460	1.13	843056.3	0.67	472447.8	0.38
5201		837381.3	1	281030.6	0.34	204343.8	0.24	274797	0.33
5207		1261177	1	336372.6	0.27	435374.7	0.35	482240.5	0.38
5314		396261.8	1	419522.3	1.06	1267443	3.2	1447255	3.65
6005		527602.3	1	200273.7	0.38	147416.9	0.28	161523	0.31
6104		770410.6	1	1071500	1.39	234813.4	0.3	187354.2	0.24
6105		2009352	1	2198885	1.09	666891.9	0.33		0
6505	S-adenosylmethionine synthetase (PF11090w) ^a	2509872	1	2592248	1.03	1781118	0.71	1200317	0.48
6506	Ornithine aminotransferase (PFF0435w) ^b	1042238	1	1420503	1.36	1489907	1.43	2313307	2.22
7103		6916921	1	2988248	0.43	4419540	0.64	1842549	0.27
7104		2328269	1	877185.6	0.38	272572	0.12	358543.8	0.15
7106		1721574	1	2139107	1.24	416634.8	0.24	844223.1	0.49
7718		2544799	1	588543.2	0.23	1293771	0.51	1031303	0.41
7817	Elongation factor 2 (PF14_0486)	572182.3	1	594160.1	1.04	1607578	2.81	873754.8	1.53
8001		1304672	1	978171.5	0.75	447955.2	0.34	522991.3	0.4
8002		3035502	1	980220.3	0.32	843949.5	0.28	898411.9	0.3
8004		897505.3	1	1179157	1.31	132369.3	0.15	362044.2	0.4
8018		4875407	1	812069.3	0.17	1234645	0.25	442642.3	0.09

8102		38253012	1	10144632	0.27	9096348	0.24	8096272	0.21
8115		1058422	1	991311.4	0.94	58656.1	0.06	143936.8	0.14
8201	Pyridoxal-5-phosphate synthase pdx1 (PFF1025c)	948551.1	1	943220.3	0.99	1506122	1.59	2393723	2.52
8203		4472542	1	6114925	1.37	1879056	0.42	1532572	0.34
8207	L-lactate dehydrogenase (PF13_0141)	4834714	1	13598061	2.81	7629889	1.58	5320832	1.1
8301		423341	1	516469.9	1.22	173934.7	0.41	109692.3	0.26
8705		1807778	1	254621.6	0.14	848141.9	0.47	556224.1	0.31

a. Spot saturated on high intensity scan set, thus SSP number, spot densities and ratios provided obtained from medium intensity scan set.

b. Spot saturated on high intensity scan set, thus SSP number, spot densities and ratios provided obtained from medium intensity scan set.