

TreeBase ID	Sequences	Codons	Mean divergence	Rates of false positives / replicate, %		
				Mean	Range	95% Confidence Interval
51737	96	209	0.0404	0.191	0 – 0.478	2.39 – 8.13
47706	97	298	0.0148	0.268	0 – 0.671	2.68 – 7.72
26258	158	78	0.0139	0.385	0 – 2.56	1.28 – 10.3
27576	155	218	0.00874	0.459	0 – 1.83	2.29 – 8.26
50407	112	233	0.027	0.515	0 – 1.29	2.58 – 8.15
50250	147	912	0.0271	0.537	0.219 – 1.43	3.62 – 6.47
47670	97	231	0.0392	0.693	0 – 1.3	2.6 – 8.23
47707	131	222	0.0416	0.991	0.45 – 1.35	2.25 – 8.11
47708	133	176	0.0503	1.65	0 – 3.98	2.27 – 8.52
43786	190	467	0.109	1.99	1.07 – 2.57	3.21 – 7.07
51735	102	217	0.181	2.03	0.922 – 3.69	2.3 – 8.29
26268	101	216	0.0637	2.13	0.463 – 3.24	2.31 – 8.33
20793	139	155	0.0552	2.39	0.645 – 4.52	1.94 – 8.39
43799	109	216	0.124	2.92	1.39 – 4.63	2.31 – 7.87
50066	191	279	0.0868	3.33	1.08 – 5.73	2.51 – 7.53
31473	177	467	0.113	4.07	2.36 – 5.78	3 – 7.07
49597	191	210	0.155	4.33	2.38 – 6.19	2.38 – 8.1
50488	100	157	0.158	4.39	2.55 – 6.37	1.91 – 8.92
47638	110	233	0.286	4.68	2.15 – 6.87	2.58 – 8.15
48737	146	193	0.0777	4.72	3.11 – 6.74	2.07 – 8.29
31472	160	217	0.114	4.93	3.23 – 7.37	2.3 – 8.29
31078	156	213	0.195	5.02	2.82 – 6.57	2.35 – 7.98
50259	167	216	0.509	5.05	1.85 – 7.87	2.31 – 7.87
49966	153	215	0.14	5.12	4.19 – 6.98	2.33 – 8.37
7471	103	569	0.459	5.29	2.99 – 6.33	3.34 – 6.85
48654	169	351	0.402	5.3	2.85 – 7.98	2.85 – 7.41
44942	190	203	0.351	5.32	2.46 – 8.87	2.46 – 8.37
50998	108	234	0.215	5.34	2.56 – 8.12	2.56 – 8.12
26494	121	199	0.753	5.38	2.01 – 10.1	2.01 – 8.04
25455	137	299	0.277	5.52	3.34 – 8.03	2.68 – 7.36
45979	142	208	0.185	5.53	1.92 – 9.13	2.4 – 8.17
9873	171	469	0.298	5.76	4.69 – 7.25	3.2 – 7.04
31295	104	231	0.406	5.8	3.9 – 7.36	2.16 – 7.79
29855	160	210	0.359	5.81	4.76 – 8.57	2.38 – 8.1
50442	104	545	0.41	5.87	4.59 – 9.36	3.3 – 6.79
48655	169	221	0.637	6.43	2.71 – 10.4	2.26 – 8.14