

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Pr
Metaseiulus_occidentalis	CTA	False	False	19	1.458616	8	15	False	1	1	0.0
Paracentrotus	CTT	False	False	46	1.278771	11	41	False	1	1	0.0
Aphrocallistes_vastus	CTC	False	False	20	1.230964	9	16	False	1	1	0.0
	TTA	False	False	74	1.152219	12	61	False	1	1	0.0
Schistosoma	CTA	False	False	12	1.237687	6	9	False	1	1	0.0
	TTA	False	False	105	0.818449	12	64	False	3	4	0.0
Fasciola	TTG	False	False	50	0.962090	12	36	False	3	3	0.0
	TTG	False	False	127	0.898428	12	86	False	2	3	0.0
Trichobilharzia_regenti	TTA	False	False	91	0.726560	12	58	False	1	2	0.0
	TTG	False	False	87	0.899438	11	58	False	2	2	0.0
Versteria_mustelae	CTT	False	False	14	1.016484	6	9	False	1	1	0.0
	TTG	False	False	107	0.964065	12	77	False	2	3	0.0
Euseius_nicholsi	TTG	False	False	25	1.349794	6	22	False	1	1	0.0
Haplorchis_taichui	CTA	False	False	8	1.150641	8	6	False	1	1	0.0
	CTT	False	False	48	0.645281	11	26	False	2	2	0.0
	TTA	False	False	52	0.852435	10	36	False	1	1	0.0
Halocynthia	CTT	False	False	14	1.036496	7	10	False	1	1	0.0
	TTA	False	False	110	0.974651	11	92	False	3	3	0.0
	TTG	False	False	71	1.064260	12	62	False	1	1	0.0
Echinococcus_equinus	CTT	False	False	11	1.255869	5	8	False	1	1	0.0
	TTA	False	False	60	0.933870	11	43	False	1	1	0.0
	TTG	False	False	123	0.885560	12	88	False	1	2	0.0
Taenia_crassiceps	CTT	False	False	14	0.666055	7	10	False	1	1	0.0
	TTA	False	False	120	0.883627	12	81	False	2	3	0.0
Bfloridae	TTG	False	False	63	1.018315	11	47	False	1	1	0.0
	CTT	False	False	21	1.158845	6	18	False	1	1	0.0
Katharina	TTG	False	False	30	1.293299	7	28	False	1	1	0.0
Tetrancistrum_nebulosi	CTA	False	False	23	0.965556	9	16	False	1	1	0.0
	TTA	False	False	91	0.872743	12	64	False	2	2	0.0
Trichinella	CTA	False	False	64	0.988682	8	49	False	2	3	0.0
	CTT	False	False	17	0.665663	9	10	False	1	1	0.0
Balanoglossus	CTC	False	False	68	1.193028	11	61	False	1	1	0.0
Pseudogarypus_banksi	CTC	False	False	7	1.227106	5	4	False	1	1	0.0
	TTA	False	False	122	0.853521	12	100	False	2	2	0.0
Benedenia_hoshinai	TTA	False	False	124	0.943869	12	86	False	3	4	0.0
	CTA	False	False	43	1.046045	10	37	False	1	1	0.0
Myxine	CTC	False	False	38	1.046622	10	32	False	1	1	0.0
	CTT	False	False	45	1.137536	11	43	False	1	1	0.0

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Ciona	TTG	False	False	8	1.412903	6	7	False	1	1	0.0
	CTA	False	False	28	0.970136	9	24	False	1	1	0.0
	TTA	False	False	116	0.964877	12	100	False	1	1	0.0
Epigonichthys	TTG	False	False	19	0.917232	6	15	False	1	1	0.0
	CTT	False	False	23	1.079734	11	22	False	1	1	0.0
Caenorhabditis	CTA	False	False	15	0.455959	9	8	False	1	1	0.0
	TTA	False	False	121	0.985560	12	96	False	6	6	0.0
Homo	TTG	False	False	34	1.297134	11	30	False	1	1	0.0
	CTC	False	False	53	1.030318	11	49	False	1	1	0.0
Ascidella aspersa	CTC	False	False	17	1.216541	10	16	False	1	1	0.0
	TTG	False	False	60	1.120903	11	53	False	1	1	0.0
Blanceolatum	CTT	False	False	13	1.218324	6	12	False	1	1	0.0
Saccoglossus	TTA	False	False	63	1.130257	10	61	False	1	1	0.0