

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Schistosoma	ATG	False	False	39	0.108339	9	11	False	5	5
Aphrocallistes_vastus	ATG	False	False	16	0.218650	5	11	False	1	1
Fasciola	ATG	False	False	38	-0.097015	8	11	False	2	2
Trichobilharzia_regenti	ATG	False	False	40	-0.010911	10	11	False	2	2
Versteria_mustelae	ATG	False	False	32	-0.226506	9	10	False	1	1
Haplorchis_taichui	ATG	False	False	40	-0.069166	9	11	False	1	2
Limulus	ATG	False	False	6	1.137500	2	0	False	2	2
Echinococcus_equinus	ATG	False	False	36	-0.018545	11	10	False	2	2
Taenia_crassiceps	ATG	False	False	34	-0.070348	9	10	False	2	2
Axinella	ATG	False	False	19	0.035326	7	9	False	1	1
Suberites_domuncula	ATG	False	False	19	0.190217	6	11	False	1	1
Geodia	ATG	False	False	16	0.229402	6	10	False	1	1
Metridium	ATG	False	False	21	0.196078	6	11	False	1	1
Tetrancistrum_nebulosi	ATG	False	False	34	-0.012830	10	9	False	1	2
Trichinella	ATG	False	False	19	0.439575	5	2	False	2	2
Amphimedon_queenslandica	ATG	False	False	24	0.155172	9	11	False	1	1
Acropora	ATG	False	False	20	0.217784	6	11	False	1	1
Pseudogarypus_banksi	ATG	False	False	2	3.375000	2	0	False	2	2
Benedenia_hoshinai	ATG	False	False	29	-0.066071	8	11	False	1	1
Ascidella_aspersa	ATG	False	False	18	-0.197122	8	6	False	1	1
Blanceolatum	ATG	False	False	4	1.118750	3	0	False	1	1
Epigonichthys	ATG	False	False	8	0.397436	5	5	False	1	1