

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	AAT	False	False	30	0.595218	8	21	False	1	1	0.
Halocynthia	AAT	False	False	31	0.909693	9	28	False	2	2	0.
Schistosoma	AAC	False	False	2	2.105263	2	1	False	1	1	0.
Aphrocallistes_vastus	AAC	False	False	20	0.948914	8	15	False	1	1	0.
Pseudogarypus_banksi	AAC	False	False	14	0.660584	3	10	False	1	1	0.
	AAT	False	False	29	0.456965	10	21	False	1	1	0.
Echinococcus_equinus	AAT	False	False	36	0.708155	7	15	False	3	5	0.
Taenia_crassiceps	AAT	False	False	23	0.698985	6	10	False	3	3	0.
Benedenia_hoshinai	AAT	False	False	24	0.244111	8	9	False	2	2	0.
Ciona	AAT	False	False	41	0.946115	9	32	False	3	3	0.
Saccoglossus	AAC	False	False	16	0.918919	7	15	False	1	1	0.
Fasciola	AAT	False	False	27	0.668582	8	13	False	3	3	0.
Trichobilharzia_regenti	AAC	False	False	2	0.937500	1	0	False	1	1	0.
	AAT	False	False	22	0.580683	7	9	False	2	2	0.
Tetrancistrum_nebulosi	AAT	False	False	18	0.691429	8	8	False	3	3	0.
Versteria_mustelae	AAT	False	False	24	0.779333	6	9	False	3	4	0.
Ascidella_aspersa	AAT	False	False	30	0.799142	8	26	False	1	1	0.
Trichinella	AAT	False	False	17	0.489362	8	10	False	1	1	0.
Haplorchis_taichui	AAT	False	False	12	0.995690	5	8	False	1	1	0.
Lumbricus	AAT	False	False	22	0.848946	8	18	False	1	1	0.