

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	TGT	False	False	6	-0.033898	3	3	False	1	1	0
Daphnia	TGT	False	False	5	0.203046	4	4	False	1	1	0
Schistosoma	TGC	False	False	6	-0.311688	4	2	False	1	1	0
	TGT	False	False	29	0.027507	9	1	False	5	6	0
Trichobilharzia_regenti	TGT	False	False	32	-0.209302	9	3	False	5	5	0
Taenia_crassiceps	TGT	False	False	33	0.184723	9	3	False	5	7	0
Benedenia_hoshinai	TGC	False	False	8	-0.083067	5	2	False	1	1	0
	TGT	False	False	24	-0.027987	9	1	False	4	5	0
Echinococcus_equinus	TGT	False	False	31	0.275459	10	3	False	5	7	0
Halocynthia	TGT	False	False	13	0.196429	7	4	False	2	2	0
Fasciola	TGC	False	False	2	0.525641	2	0	False	1	1	0
	TGT	False	False	36	-0.155050	10	3	False	3	5	0
Tetrancistrum_nebulosi	TGC	False	False	12	-0.751082	6	1	False	1	1	0
	TGT	False	False	21	0.100122	9	2	False	3	4	0
Versteria_mustelae	TGC	False	False	2	2.820513	2	0	False	2	2	0
	TGT	False	False	32	-0.134196	9	3	False	4	5	0
Drosophila	TGT	False	False	8	0.114286	6	6	False	1	1	0
Strigamia_maritima	TGT	False	False	6	0.797468	6	4	False	2	2	0
Caenorhabditis	TGT	False	False	19	-0.270636	9	4	False	2	2	0
Haplorchis_taichui	TGT	False	False	33	0.070543	10	3	False	4	6	0
Lumbricus	TGT	False	False	4	0.518987	4	2	False	1	1	0