CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Metaseiulus_occidentalis	TTA	False	False	101	-0.943786	10	92	False	1	1
	TTG	False	False	22	-0.910242	7	18	False	1	1
Pseudogarypus_banksi	CTA	False	False	23	-0.682594	9	16	False	2	2
	TTA	False	False	122	-0.983445	12	100	False	1	1
Schistosoma	TTA	False	False	105	-0.835332	12	64	False	3	5
	TTG	False	False	50	-0.858607	12	36	False	1	1
Tetrancistrum_nebulosi	CTT	False	False	28	-0.930083	8	20	False	1	1
	TTA	False	False	91	-1.043172	12	64	False	1	1
	CTA	False	False	5	-0.367347	5	4	False	1	1
Taenia_crassiceps	TTA	False	False	120	-0.906774	12	81	False	2	2
	TTG	False	False	63	-0.976394	11	47	False	1	1
Benedenia_hoshinai	TTA	False	False	124	-0.976388	12	86	False	3	3
Echinococcus_equinus	TTG	False	False	123	-0.845402	12	88	False	5	5
Limulus	TTA	False	False	71	-0.956257	11	67	False	1	1
Ciona	TTA	False	False	116	-0.954495	12	100	False	1	1
Fasciola	TTA	False	False	43	-0.871352	10	32	False	1	1
	TTG	False	False	127	-0.988311	12	86	False	1	1
Terebratulina	CTT	False	False	52	-0.922963	10	49	False	1	1
Amphimedon_queenslandica	TTA	False	False	92	-0.962726	12	84	False	1	1
Trichobilharzia_regenti	TTA	False	False	91	-0.951022	12	58	False	1	1
	TTG	False	False	87	-0.984324	11	58	False	1	1
Euseius_nicholsi	TTA	False	False	113	-0.876046	11	98	False	3	3
Versteria_mustelae	TTA	False	False	73	-1.006314	10	54	False	1	1
	TTG	False	False	107	-0.878153	12	77	False	3	3
Trichinella	CTA	False	False	64	-0.947858	8	49	False	1	1
	CTC	False	False	20	-0.875000	8	14	False	1	1
	CTT	False	False	17	-0.634036	9	10	False	2	2
Caenorhabditis	TTG	False	False	34	-0.850679	11	30	False	1	1
Katharina	TTA	False	False	108	-0.951046	12	102	False	1	1
Haplorchis_taichui	CTT	False	False	48	-1.008484	11	26	False	2	2
Lumbricus	TTG	False	False	9	-0.682720	6	8	False	1	1