

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	TTT	False	False	104	-1.576280	10	75	False	1	1	0
Schistosoma	TTC	False	False	8	-0.993631	4	4	False	1	1	0
Tetrancistrum_nebulosi	TTT	False	False	108	-1.688878	12	61	False	1	1	0
Echinococcus_equinus	TTT	False	False	133	-1.696800	12	72	False	1	1	0
Taenia_crassiceps	TTT	False	False	132	-1.676048	12	68	False	1	1	0
Benedenia_hoshinai	TTC	False	False	10	-1.132316	5	6	False	1	1	0
Ciona	TTT	False	False	119	-1.589351	11	89	False	1	2	0
Fasciola	TTT	False	False	130	-1.722889	11	67	False	1	1	0
Trichobilharzia_regenti	TTT	False	False	116	-1.707947	12	65	False	1	1	0
Suberites_domuncula	TTT	False	False	117	-1.699978	11	92	False	1	1	0
Versteria_mustelae	TTT	False	False	122	-1.694345	11	67	False	1	1	0
Ascidiella_aspersa	TTC	False	False	32	-1.508380	7	24	False	1	1	0
Caenorhabditis	TTT	False	False	140	-1.664471	12	88	False	1	1	0
Haplorchis_taichui	TTT	False	False	103	-1.681943	11	57	False	1	1	0
Lumbricus	TTT	False	False	85	-1.672097	12	75	False	1	1	0