

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	-1.612943	10	75	False	1	1	0
Halocynthia	TTT	False	False	137	-1.582257	11	100	False	2	2	0
Schistosoma	TTT	False	False	107	-1.500360	11	62	False	2	2	0
Pseudogarypus_banksi	TTT	False	False	106	-1.642320	11	85	False	1	1	0
Echinococcus_equinus	TTT	False	False	133	-1.557247	12	72	False	1	1	0
Taenia_crassiceps	TTT	False	False	132	-1.512616	12	68	False	1	1	0
Benedenia_hoshinai	TTT	False	False	110	-1.639814	12	63	False	1	1	0
Limulus	TTT	False	False	83	-1.678925	10	73	False	1	1	0
Ciona	TTT	False	False	119	-1.655314	11	89	False	1	1	0
Fasciola	TTT	False	False	130	-1.574690	11	67	False	1	1	0
Trichobilharzia_regenti	TTT	False	False	116	-1.535320	12	65	False	2	2	0
Tetrancistrum_nebulosi	TTT	False	False	108	-1.623192	12	61	False	1	1	0
Versteria_mustelae	TTT	False	False	122	-1.552029	11	67	False	1	1	0
Ascidrella_aspersa	TTT	False	False	104	-1.612043	11	80	False	1	1	0
Haplorchis_taichui	TTT	False	False	103	-1.553674	11	57	False	2	2	0