

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	TTC	False	False	31	-0.464672	7	24	False	1	1	0.
Schistosoma	TTT	False	False	107	-0.449484	11	62	False	3	3	0.
Trichobilharzia_regenti	TTT	False	False	116	-0.484106	12	65	False	3	4	0.
Taenia_crassiceps	TTT	False	False	132	-0.513199	12	68	False	2	4	0.
Benedenia_hoshinai	TTT	False	False	110	-0.593481	12	63	False	2	2	0.
Myxine	TTT	False	False	72	-0.593214	11	65	False	1	1	0.
Echinococcus_equinus	TTT	False	False	133	-0.492483	12	72	False	2	4	0.
Fasciola	TTT	False	False	130	-0.510726	11	67	False	2	3	0.
Tetrancistrum_nebulosi	TTC	False	False	16	-0.738019	7	9	False	1	1	0.
	TTT	False	False	108	-0.504624	12	61	False	2	2	0.
Versteria_mustelae	TTT	False	False	122	-0.521547	11	67	False	2	4	0.
Caenorhabditis	TTT	False	False	140	-0.512075	12	88	False	1	1	0.
Haplorchis_taichui	TTC	False	False	13	-0.221135	7	7	False	2	2	0.
	TTT	False	False	103	-0.538979	11	57	False	1	1	0.