CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Metaseiulus_occidentalis	TTT	False	False	104	0.312008	10	75	False	2	2
Paracentrotus	TTT	False	False	86	0.532696	12	77	False	1	1
Aphrocallistes_vastus	TTC	False	False	37	0.847232	9	34	False	1	1
	TTT	False	False	82	0.493870	11	65	False	2	2
Schistosoma	TTT	False	False	107	0.206144	11	62	False	3	4
Fasciola	TTT	False	False	130	0.044676	11	67	False	2	2
Trichobilharzia_regenti	TTT	False	False	116	0.134216	12	65	False	3	4
Versteria_mustelae	TTT	False	False	122	0.076519	11	67	False	3	3
Euseius_nicholsi	TTT	False	False	123	0.384712	11	92	False	2	2
Patiria	TTT	False	False	74	0.498435	11	65	False	1	1
Haplorchis_taichui	TTT	False	False	103	0.271233	11	57	False	3	3
Echinococcus_equinus	TTT	False	False	133	0.112567	12	72	False	4	4
Taenia_crassiceps	TTT	False	False	132	0.078610	12	68	False	4	4
Axinella	TTT	False	False	121	0.670059	11	102	False	4	4
Suberites_domuncula	TTT	False	False	117	0.610147	11	92	False	4	4
Geodia	TTC	False	False	19	1.017639	7	17	False	1	1
	TTT	False	False	113	0.417084	12	91	False	2	2
Metridium	TTT	False	False	101	0.621889	11	85	False	3	3
Tetrancistrum_nebulosi	TTC	False	False	16	1.108626	7	9	False	2	2
	TTT	False	False	108	0.267489	12	61	False	1	1
Trichinella	TTC	False	False	56	0.384332	7	44	False	1	1
Amphimedon_queenslandica	TTC	False	False	29	1.331541	9	24	False	3	3
	TTT	False	False	98	0.364872	12	84	False	1	1
Acropora	TTT	False	False	126	0.548308	12	106	False	3	3
Pseudogarypus_banksi	TTT	False	False	106	0.299927	11	85	False	1	1
Benedenia_hoshinai	TTT	False	False	110	0.399534	12	63	False	4	4
Ciona	TTT	False	False	119	0.316448	11	89	False	2	2
Caenorhabditis	TTT	False	False	140	0.268752	12	88	False	4	5
Ascidiella_aspersa	TTT	False	False	104	0.205331	11	80	False	1	1