CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	
Metaseiulus_occidentalis	**AAA**	False	False	28	-0.054695	8	21	True	1	1	0
Halocynthia	**AAA**	False	False	13	0.358974	7	10	True	1	1	0
Schistosoma	**AAA**	True	True	11	0.939252	7	1	True	3	3	0
	AAG	False	False	27	0.011396	8	22	False	1	1	0
Pseudogarypus_banksi	**AAA**	False	False	29	0.089631	8	24	True	1	1	0
Echinococcus_equinus	**AAG**	False	False	27	0.016008	8	22	False	1	1	0
Taenia_crassiceps	**AAA**	True	True	13	1.601190	6	1	True	2	5	0
	AAG	False	False	25	0.114837	7	22	False	1	1	0
Benedenia_hoshinai	**AAA**	True	True	14	1.836066	9	1	True	4	5	0
Paracentrotus	**AAA**	True	True	13	4.186770	6	0	True	6	13	0
Fasciola	**AAA**	True	True	5	1.855670	4	1	True	1	2	0
	AAG	False	False	28	-0.008167	8	23	False	1	1	0
Trichobilharzia_regenti	**AAA**	True	True	10	2.389744	5	1	True	3	4	0
	AAG	False	False	26	0.118164	7	22	False	1	1	0
Tetrancistrum_nebulosi	**AAA**	True	True	12	2.431624	6	0	True	4	6	0
Versteria_mustelae	**AAA**	True	True	13	1.677800	5	0	True	2	5	0
	AAG	False	False	26	0.024462	8	22	False	1	1	0
Euseius_nicholsi	**AAA**	False	False	28	-0.073370	8	23	True	1	1	0
Patiria	**AAA**	True	True	15	4.357143	5	0	True	5	15	0
Trichinella	**AAA**	False	False	17	0.147368	5	14	True	1	1	0
Caenorhabditis	**AAA**	False	False	30	-0.190355	10	19	True	1	1	0
Haplorchis_taichui	**AAA**	True	True	14	2.210238	5	1	True	3	6	0
	AAG	False	False	29	0.024540	8	24	False	1	1	0