CoReTracker report

		Clad. Val	Reassigned	Tot.	Telford	Tot. Genes	Use.	Trans. Val	Rea. Genes	Rea. count
Metaseiulus_occidentalis	GAC	False	False	12	2.045455	4	11	False	1	1
	GAT	False	False	24	2.042872	8	21	False	3	3
Aphrocallistes_vastus	GAT	False	False	21	1.871571	6	20	False	1	1
Fasciola	GAT	False	False	32	1.826861	8	26	False	2	2
Trichobilharzia_regenti	GAT	False	False	30	1.510363	8	22	False	1	1
Versteria_mustelae	GAT	False	False	36	1.759885	8	31	False	1	1
Euseius_nicholsi	GAT	False	False	35	1.973432	7	30	False	4	4
Haplorchis_taichui	GAT	False	False	28	1.521140	7	22	False	1	1
Lumbricus	GAC	False	False	16	1.955056	6	15	False	1	1
Halocynthia	GAT	False	False	30	1.773471	8	27	False	1	1
Limulus	GAT	False	False	23	1.931767	8	22	False	1	1
Echinococcus_equinus	GAT	False	False	39	1.675066	8	30	False	2	2
Taenia_crassiceps	GAT	False	False	42	1.679803	8	32	False	3	3
Axinella	GAT	False	False	31	1.868114	8	30	False	1	1
Terebratulina	GAC	False	False	27	1.901639	8	26	False	1	1
Suberites_domuncula	GAT	False	False	30	1.836788	6	28	False	1	1
Katharina	GAT	False	False	28	2.023234	7	26	False	2	2
Geodia	GAT	False	False	35	1.840118	8	34	False	1	1
Metridium	GAT	False	False	29	1.931373	7	27	False	2	2
Tetrancistrum_nebulosi	GAC	False	False	16	1.977346	7	14	False	1	1
	GAT	False	False	17	1.938066	7	13	False	1	1
Strigamia_maritima	GAC	False	False	24	1.908108	6	22	False	2	2
Trichinella	GAC	False	False	16	1.720661	5	14	False	1	1
	GAT	False	False	18	2.072443	7	14	False	2	2
Amphimedon_queenslandica	GAC	False	False	16	1.907618	7	15	False	1	1
	GAT	False	False	27	1.820439	8	25	False	1	1
Acropora	GAC	False	False	10	2.093750	6	9	False	1	1
	GAT	False	False	31	1.792154	7	29	False	1	1
Benedenia_hoshinai	GAT	False	False	26	1.800995	8	21	False	1	1
Myxine	GAC	False	False	23	1.704289	7	20	False	1	1
	GAT	False	False	18	1.952924	6	17	False	1	1
Drosophila	GAT	False	False	36	1.915412	8	34	False	2	2
Ciona	GAT	False	False	29	1.917260	8	28	False	1	1
Daphnia	GAC	False	False	18	1.879360	7	17	False	1	1
	GAT	False	False	23	1.930804	7	22	False	1	1
Caenorhabditis	GAT	False	False	34	1.769522	8	31	False	1	1
Homo	GAC	False	False	31	1.934167	7	28	False	2	2
Ascidiella_aspersa	GAT	False	False	32	2.004847	8	28	False	3	3

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea.
Epigonichthys	GAT	False	False	29	1.847534	8	28	False	1	1