CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Pr
Paracentrotus	GCC	False	False	32	-0.202220	8	22	False	1	2	0.0
Aphrocallistes_vastus	GCA	False	False	40	-0.421323	9	29	False	1	1	0.0
	GCT	False	False	22	-0.443931	8	14	False	1	1	0.0
Schistosoma	GCT	False	False	28	-0.157895	7	13	False	3	4	0.0
Fasciola	GCT	False	False	40	-0.607692	9	20	False	2	2	0.0
Trichobilharzia_regenti	GCC	False	False	7	-0.073529	4	2	False	1	1	0.0
	GCG	False	False	6	0.207627	4	3	False	1	1	0.0
	GCT	False	False	36	-0.390780	9	20	False	2	3	0.0
Versteria_mustelae	GCT	False	False	35	-0.402930	9	17	False	2	2	0.0
Euseius_nicholsi	GCT	False	False	33	-0.214011	7	30	False	1	1	0.0
	GCC	False	False	6	0.186441	5	4	False	1	1	0.0
Haplorchis_taichui	GCG	False	False	16	0.200957	9	8	False	3	3	0.0
	GCT	False	False	31	-0.649587	7	14	False	1	1	0.0
Halocynthia	GCG	False	False	6	-0.160338	4	3	False	1	1	0.0
	GCT	False	False	37	-0.070491	9	29	False	1	1	0.0
Limulus	GCC	False	False	14	-0.060998	6	13	False	1	1	0.0
Echinococcus_equinus	GCT	False	False	32	-0.495994	8	16	False	2	2	0.0
Taenia_crassiceps	GCT	False	False	34	-0.378480	8	17	False	2	2	0.0
Terebratulina	GCT	False	False	20	-0.057252	8	18	False	1	1	0.0
Metridium	GCT	False	False	38	-0.290604	9	33	False	1	1	0.0
Tetrancistrum_nebulosi	GCT	False	False	36	-0.680912	9	13	False	2	3	0.0
Trichinella	GCA	False	False	19	0.083447	5	10	False	2	3	0.0
	GCT	False	False	28	-0.346399	5	20	False	1	2	0.0
Balanoglossus	GCC	False	False	32	-0.262530	10	25	False	1	1	0.0
Acropora	GCT	False	False	30	-0.222504	9	24	False	1	1	0.0
Pseudogarypus_banksi	GCA	False	False	20	-0.256410	8	17	False	1	1	0.0
	GCT	False	False	24	-0.334755	9	16	False	2	2	0.0
Benedenia_hoshinai	GCT	False	False	33	-0.589506	7	15	False	3	3	0.0
Myxine	GCA	False	False	24	-0.286788	7	18	False	2	2	0.0
Ascidiella_aspersa	GCG	False	False	7	-0.126812	5	4	False	1	1	0.0
	GCT	False	False	31	0.051937	8	28	False	1	1	0.0