## CoReTracker report

		Clad. Val	Reassigned	Tot.	Telford	Tot. Genes	Use.	Trans. Val	Rea. Genes	Rea.
Schistosoma	GCT	False	False	28	-0.796733	7	13	False	2	2
Fasciola	GCT	False	False	40	-1.294231	9	20	False	2	2
	GCC	False	False	7	-0.014706	4	2	False	1	2
Trichobilharzia_regenti	GCG	False	False	6	-0.470339	4	3	False	1	1
	GCT	False	False	36	-1.231915	9	20	False	1	1
Versteria_mustelae	GCA	False	False	5	-0.654639	5	2	False	1	1
	GCT	False	False	35	-1.010256	9	17	False	2	2
Haplorchis_taichui	GCT	False	False	31	-1.041322	7	14	False	2	2
Lumbricus	GCG	False	False	4	0.569620	2	2	False	1	1
Halocynthia	GCT	False	False	37	-0.757429	9	29	False	2	2
Echinococcus_equinus	GCA	False	False	5	-0.553299	4	3	False	1	1
	GCT	False	False	32	-1.110577	8	16	False	3	3
Taenia_crassiceps	GCG	False	False	3	0.433333	3	1	False	1	1
	GCT	False	False	34	-1.005267	8	17	False	3	3
Terebratulina	GCA	False	False	21	-1.062802	7	15	False	1	1
	GCC	False	False	38	-0.916611	8	30	False	2	2
Bfloridae	GCG	False	False	9	-0.717087	6	6	False	1	1
Katharina	GCA	False	False	19	-1.177333	6	14	False	1	1
	GCC	False	False	12	-1.056250	4	10	False	1	1
Amphimedon_queenslandica	GCG	False	False	29	-0.906690	9	21	False	1	1
Tetrancistrum_nebulosi	GCA	False	False	11	0.043478	5	7	False	3	3
	GCC	False	False	6	-1.176471	5	3	False	1	1
	GCT	False	False	36	-1.240741	9	13	False	2	2
Trichinella	GCC	False	False	13	-0.821218	5	9	False	2	2
Balanoglossus	GCC	False	False	32	-1.012729	10	25	False	1	1
Drosophila	GCA	False	False	13	-0.747036	6	12	False	1	1
Pseudogarypus_banksi	GCT	False	False	24	-1.221748	9	16	False	1	1
Benedenia_hoshinai	GCA	False	False	8	-1.208861	4	4	False	1	1
	GCT	False	False	33	-1.307870	7	15	False	1	1
Saccoglossus	GCA	False	False	35	-0.943755	7	30	False	2	2
Strigamia_maritima	GCT	False	False	23	-1.019956	8	20	False	1	1
Caenorhabditis	GCA	False	False	20	-0.638817	6	13	False	1	2
	GCC	False	False	6	-0.718487	5	4	False	1	1
	GCG	False	False	2	1.350000	2	0	False	1	1
	GCT	False	False	36	-1.201418	7	25	False	1	1
Homo	GCA	False	False	26	-0.945098	8	20	False	1	1
Ascidiella_aspersa	GCT	False	False	31	-0.766694	8	28	False	1	1
Blanceolatum	GCA	False	False	10	-0.635000	5	8	False	1	1

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea.
	GCC	False	False	8	-0.987500	4	5	False	1	1
	GCT	False	False	23	-0.740741	6	19	False	1	1
Epigonichthys	GCT	False	False	25	-1.046559	7	21	False	1	1