## CoReTracker report

		Clad. Val	Reassigned	Tot.	Telford	Tot. Genes	Use.	Trans. Val	Rea. Genes	Rea.
Metaseiulus_occidentalis	GCT	False	False	31	1.018062	8	29	False	2	2
Paracentrotus	GCC	False	False	32	0.768438	8	22	False	3	4
Aphrocallistes_vastus	GCA	False	False	40	0.933205	9	29	False	3	6
	GCT	False	False	22	0.717919	8	14	False	2	2
Schistosoma	GCG	False	False	4	0.736842	2	1	False	1	1
	GCT	False	False	28	0.459165	7	13	False	3	4
	GCA	False	False	2	1.846154	2	1	False	1	1
Fasciola	GCG	False	False	7	0.985560	6	5	False	1	1
	GCT	False	False	40	0.687179	9	20	False	2	4
Trichobilharzia_regenti	GCG	False	False	6	0.423729	4	3	False	1	1
	GCT	False	False	36	0.472340	9	20	False	2	2
Versteria_mustelae	GCT	False	False	35	0.542125	9	17	False	4	4
Euseius_nicholsi	GCT	False	False	33	0.799076	7	30	False	1	1
Patiria	GCT	False	False	16	0.748808	6	14	False	1	1
Haplorchis_taichui	GCG	False	False	16	0.500797	9	8	False	2	2
	GCT	False	False	31	0.294215	7	14	False	1	2
	GCA	False	False	28	0.804209	9	25	False	1	1
Lumbricus	GCC	False	False	23	0.952062	8	17	False	2	2
	GCT	False	False	25	0.871457	8	23	False	2	2
Halocynthia	GCA	False	False	16	0.833866	9	12	False	1	1
	GCG	False	False	6	0.603376	4	3	False	1	1
	GCT	False	False	37	0.517623	9	29	False	1	1
Limulus	GCA	False	False	26	0.900198	8	23	False	1	1
	GCT	False	False	35	0.933962	8	33	False	1	2
Echinococcus_equinus	GCT	False	False	32	0.610577	8	16	False	3	3
Taenia_crassiceps	GCA	False	False	6	0.904762	4	4	False	1	1
	GCT	False	False	34	0.306998	8	17	False	1	1
Axinella	GCA	False	False	21	0.761847	7	16	False	3	3
	GCG	False	False	14	0.996350	7	12	False	2	2
Terebratulina	GCA	False	False	21	0.919082	7	15	False	3	3
	GCC	False	False	38	0.933423	8	30	False	3	5
Suberites_domuncula	GCG	False	False	14	0.891501	8	10	False	1	1
	GCT	False	False	34	0.842461	7	28	False	2	3
Bfloridae	GCG	False	False	9	0.694678	6	6	False	1	1
	GCT	False	False	24	0.885297	5	23	False	1	1
	GCA	False	False	19	0.950667	6	14	False	3	3
Katharina	GCC	False	False	12	0.879167	4	10	False	1	1
	GCG	False	False	7	1.496296	4	5	False	1	1

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Geodia	GCT	False	False	38	0.777478	10	35	False	1	1
	GCA	False	False	18	0.570822	7	14	False	1	1
	GCC	False	False	8	0.909677	6	6	False	1	1
	GCG	False	False	10	1.230964	6	9	False	1	1
	GCT	False	False	43	0.669226	10	37	False	1	1
	GCA	False	False	12	0.915433	5	8	False	1	1
Metridium	GCG	False	False	10	1.110256	6	8	False	1	2
	GCT	False	False	38	0.744295	9	33	False	1	1
Balanoglossus	GCT	False	False	15	0.868376	7	13	False	1	1
Tetrancistrum_nebulosi	GCG	False	False	5	1.035714	4	3	False	1	1
	GCT	False	False	36	0.069088	9	13	False	2	2
Strigamia_maritima	GCA	False	False	27	0.997124	6	22	False	2	2
	GCT	False	False	23	0.950111	8	20	False	1	2
Trichinella	GCT	False	False	28	0.773017	5	20	False	2	3
	GCA	False	False	27	1.038023	8	24	False	2	2
Amphimedon_queenslandica	GCC	False	False	16	0.781847	8	14	False	1	1
Acropora	GCT	False	False	11	1.094037	7	9	False	2	2
	GCG	False	False	24	1.066028	10	19	False	3	4
	GCT	False	False	30	0.675127	9	24	False	1	1
	GCA	False	False	20	0.742308	8	17	False	1	1
Pseudogarypus_banksi	GCC	False	False	4	1.101911	2	3	False	1	1
	GCT	False	False	24	0.439232	9	16	False	1	1
Benedenia_hoshinai	GCT	False	False	33	0.315586	7	15	False	2	2
Myxine	GCA	False	False	24	0.777658	7	18	False	2	2
Drosophila	GCT	False	False	55	0.898887	8	52	False	3	3
Ciona	GCT	False	False	40	0.776072	9	36	False	1	1
	GCA	False	False	26	0.900493	7	24	False	1	1
Epigonichthys	GCG	False	False	19	0.837366	8	16	False	1	1
Daphnia	GCT	False	False	25	0.799595	7	21	False	2	2
	GCA	False	False	13	1.387226	6	9	False	2	3
	GCT	False	False	25	0.847321	7	24	False	1	1
Caenorhabditis	GCC	False	False	6	0.638655	5	4	False	1	1
	GCT	False	False	36	0.587943	7	25	False	3	3
Homo	GCA	False	False	26	0.716667	8	20	False	1	1
Ascidiella_aspersa	GCA	False	False	7	1.218978	4	5	False	1	1
	GCT	False	False	31	0.589448	8	28	False	1	1
Blanceolatum	GCA	False	False	10	0.752500	5	8	False	1	1
	GCT	False	False	23	0.550109	6	19	False	1	1
Saccoglossus	GCA	False	False	35	0.756026	7	30	False	1	1
	GCC	False	False	16	0.998420	8	14	False	1	1
	GCT	False	False	27	0.769668	9	25	False	1	1