CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Paracentrotus	**GCC**	False	False	32	-0.346550	8	22	False	2	2
	GCT	False	False	16	-0.030351	8	14	False	1	1
	GCA	False	False	40	-0.213873	9	29	False	1	1
Aphrocallistes_vastus	**GCC**	False	False	16	0.629393	9	13	False	2	3
	GCT	False	False	22	-0.026590	8	14	False	2	3
Schistosoma	**GCG**	False	False	4	0.335526	2	1	False	1	1
	GCT	False	False	28	-0.754991	7	13	False	1	1
Fasciola	**GCG**	False	False	7	0.231047	6	5	False	1	1
	GCT	False	False	40	0.101923	9	20	False	5	5
Trichobilharzia_regenti	**GCT**	False	False	36	-0.002837	9	20	False	3	5
Versteria_mustelae	**GCG**	False	False	4	1.363636	2	1	False	2	2
	GCT	False	False	35	-0.336996	9	17	False	3	3
Euseius_nicholsi	**GCT**	False	False	33	-0.030023	7	30	False	1	1
Patiria	**GCA**	False	False	22	0.086247	6	20	False	1	1
	GCT	False	False	16	0.025437	6	14	False	1	1
Haplorchis_taichui	**GCG**	False	False	16	-0.272727	9	8	False	2	2
	GCT	False	False	31	-0.586777	7	14	False	2	2
Lumbricus	**GCA**	False	False	28	0.298262	9	25	False	2	2
	GCC	False	False	23	-0.257525	8	17	False	1	1
Halocynthia	**GCA**	False	False	16	0.041534	9	12	False	1	1
	GCG	False	False	6	-0.320675	4	3	False	1	1
Limulus	**GCA**	False	False	26	0.131423	8	23	False	2	2
Echinococcus_equinus	**GCG**	False	False	7	-0.315217	4	3	False	1	1
	GCT	False	False	32	-0.185096	8	16	False	3	3
Caenorhabditis	**GCA**	False	False	20	-0.638817	6	13	False	1	1
	GCT	False	False	36	-0.086525	7	25	False	3	3
Axinella	**GCT**	False	False	24	0.120507	9	20	False	2	2
Terebratulina	**GCA**	False	False	21	-0.157005	7	15	False	2	2
	GCT	False	False	20	-0.199746	8	18	False	1	1
Suberites_domuncula	**GCG**	False	False	14	-0.079566	8	10	False	1	1
	GCT	False	False	34	-0.258065	7	28	False	1	1
Katharina	**GCA**	False	False	19	0.117333	6	14	False	1	1
	GCG	False	False	7	0.640741	4	5	False	1	1
Geodia	**GCT**	False	False	43	-0.346722	10	37	False	2	2
Metridium	**GCA**	False	False	12	0.704017	5	8	False	2	3
	GCT	False	False	38	-0.185906	9	33	False	1	1
Amphimedon_queenslandica	**GCA**	False	False	27	0.067490	8	24	False	1	1
	GCC	False	False	16	-0.171975	8	14	False	1	1

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Tetrancistrum_nebulosi	**GCG**	False	False	29	-0.501761	9	21	False	1	1
	GCC	False	False	6	0.361345	5	3	False	1	1
	GCT	False	False	36	-0.533476	9	13	False	3	4
	GCA	False	False	19	-0.391245	5	10	False	3	3
Trichinella	**GCC**	False	False	13	-0.383104	5	9	False	1	1
	GCG	False	False	2	2.900000	1	1	False	1	1
	GCT	False	False	28	-0.240656	5	20	False	1	1
Balanoglossus	**GCC**	False	False	32	-0.037391	10	25	False	3	3
Acropora	**GCG**	False	False	24	0.195953	10	19	False	1	1
	GCT	False	False	30	-0.254653	9	24	False	1	1
Pseudogarypus_banksi	**GCA**	False	False	20	0.417949	8	17	False	1	1
	GCT	False	False	24	0.162047	9	16	False	2	3
Benedenia_hoshinai	**GCG**	False	False	1	3.225000	1	0	False	1	1
	GCT	False	False	33	-0.347222	7	15	False	2	3
	GCA	False	False	24	0.082707	7	18	False	2	2
Myxine	**GCC**	False	False	17	0.061194	6	15	False	1	1
	GCT	False	False	28	-0.085688	8	27	False	1	1
Ciona	**GCA**	False	False	10	0.267176	6	8	False	1	1
	GCT	False	False	40	-0.214971	9	36	False	1	1
Epigonichthys	**GCT**	False	False	25	-0.092105	7	21	False	1	1
Strigamia_maritima	**GCA**	False	False	27	0.051774	6	22	False	2	2
	GCC	False	False	16	-0.092504	8	14	False	1	1
Taenia_crassiceps	**GCT**	False	False	34	-0.395034	8	17	False	3	3
	GCA	False	False	26	-0.318627	8	20	False	1	1
Homo	**GCC**	False	False	39	-0.199089	9	38	False	1	1
	GCT	False	False	11	0.482353	4	9	False	2	2
Ascidiella_aspersa	**GCA**	False	False	7	0.529197	4	5	False	1	1
	GCG	False	False	7	-0.083333	5	4	False	1	1
Saccoglossus	**GCA**	False	False	35	-0.248356	7	30	False	1	1
	GCT	False	False	27	-0.266351	9	25	False	1	1