CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	
Metaseiulus_occidentalis	GTT	False	False	26	-0.316733	8	17	False	4	4	0
	GTA	False	False	35	-0.127365	10	9	False	3	3	0
Schistosoma	GTG	False	False	24	-0.514286	10	8	False	1	1	0
	GTT	False	False	65	0.092058	12	18	False	4	6	0
Aphrocallistes_vastus	GTA	False	False	35	-0.774003	8	29	False	1	1	0
Fasciola	GTT	False	False	98	-0.004195	11	31	False	6	11	0
Trichobilharzia_regenti	GTG	False	False	30	-0.093562	9	8	False	2	4	0
	GTT	False	False	74	-0.438062	12	22	False	2	5	0
	GTA	False	False	15	-0.239316	8	5	False	1	1	0
Versteria_mustelae	GTG	False	False	26	-0.410256	11	7	False	2	2	0
	GTT	False	False	64	-0.554038	11	23	False	2	4	0
Euseius_nicholsi	GTT	False	False	26	-0.422732	9	15	False	1	1	0
Patiria	GTC	False	False	8	-0.142857	4	6	False	1	1	0
	GTC	False	False	13	-0.337945	6	5	False	1	1	0
Haplorchis_taichui	GTG	False	False	45	-0.283352	10	12	False	3	4	0
	GTT	False	False	53	-0.301211	11	16	False	3	4	0
Lumbricus	GTG	False	False	16	-0.346405	6	12	False	1	1	0
Limulus	GTC	False	False	11	-0.213115	4	7	False	1	1	0
	GTA	False	False	12	0.807611	7	2	False	4	4	0
Echinococcus_equinus	GTG	False	False	36	-0.588953	11	7	False	1	1	0
	GTT	False	False	74	-0.495119	11	31	False	3	4	0
	GTA	False	False	24	-0.421505	8	10	False	2	2	0
Taenia_crassiceps	GTG	False	False	12	0.919149	7	4	False	3	3	0
	GTT	False	False	70	-0.327694	11	20	False	4	6	0
Suberites_domuncula	GTT	False	False	25	-0.497921	8	20	False	1	1	0
Bfloridae	GTA	False	False	18	-0.269177	7	9	False	1	1	0
	GTG	False	False	10	-0.277078	6	5	False	1	1	0
Tetrancistrum_nebulosi	GTA	False	False	50	-0.483854	10	17	False	3	3	0
	GTT	False	False	39	-0.507561	10	12	False	3	4	0
	GTA	False	False	34	-0.722222	8	17	False	1	1	0
Trichinella	GTG	False	False	12	-0.130252	5	4	False	2	2	0
	GTT	False	False	22	-0.060606	7	9	False	2	2	0
Drosophila	GTG	False	False	4	0.267123	4	3	False	1	1	0
Pseudogarypus_banksi	GTG	False	False	8	0.176849	5	5	False	1	1	0
	GTA	False	False	32	-0.205457	9	11	False	3	3	0
Benedenia_hoshinai	GTG	False	False	6	0.021277	5	2	False	1	1	0
	GTT	False	False	60	-0.459494	11	19	False	3	4	0
Ciona	GTA	False	False	35	-0.586716	9	26	False	1	1	0

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	
Daphnia	**GTA**	False	False	35	-0.587583	8	26	False	1	1	0
Caenorhabditis	**GTA**	False	False	30	-0.621972	9	18	False	1	1	0
	GTT	False	False	46	-0.764771	11	23	False	1	1	0
Ascidiella_aspersa	**GTT**	False	False	50	-0.616255	12	30	False	1	1	0
Blanceolatum	**GTG**	False	False	8	-0.239748	7	5	False	1	1	0
Saccoglossus	**GTT**	False	False	42	-0.542679	10	37	False	1	1	0