CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	ı
Metaseiulus_occidentalis	GTA	False	False	26	-0.135644	9	18	False	2	2	0
	GTA	False	False	35	-0.280932	10	9	False	3	4	0
Schistosoma	GTG	False	False	24	-0.095238	10	8	False	4	4	0
	GTT	False	False	65	-0.732912	12	18	False	2	2	0
Aphrocallistes_vastus	GTA	False	False	35	0.064993	8	29	False	2	3	0
- -asciola	GTA	False	False	13	-0.162427	8	1	False	2	2	0
Fasciola	GTT	False	False	98	-0.512061	11	31	False	5	7	0
	GTA	False	False	17	-0.183333	8	5	False	1	2	0
Trichobilharzia_regenti	GTG	False	False	30	-0.349356	9	8	False	3	3	0
	GTT	False	False	74	-0.504498	12	22	False	3	3	0
Versteria_mustelae	GTT	False	False	64	-0.241864	11	23	False	5	7	0
Euseius_nicholsi	GTA	False	False	30	-0.230241	10	23	False	1	1	0
Euseius_IIICIIOISI	GTG	False	False	11	-0.271028	5	6	False	1	1	0
Haplorchis_taichui	GTA	False	False	20	-0.304236	10	7	False	2	2	0
	GTG	False	False	45	-0.444698	10	12	False	2	3	0
	GTT	False	False	53	-0.422276	11	16	False	4	4	0 0 0
Lumbricus	GTA	False	False	33	-0.081187	9	29	False	1	1	0
I la la aventeia	GTA	False	False	43	-0.272455	11	22	False	2	2	0
Halocynthia	GTG	False	False	17	-0.391172	7	8	False	1	1	0
Limulus	GTA	False	False	35	-0.186567	11	26	False	1	1	0
	GTA	False	False	12	-0.520085	7	2	False	1	1	
Echinococcus oquinus	GTC	False	False	3	0.372881	3	1	False	1	1	0
Echinococcus_equinus	GTG	False	False	36	-0.616212	11	7	False	1	1	0
	GTT	False	False	74	-0.365063	11	31	False	4	5	0
Taenia_crassiceps	GTA	False	False	24	-0.377419	8	10	False	1	1	0
raema_crassiceps	GTT	False	False	70	-0.325855	11	20	False	5	7	
Suberites_domuncula	GTA	False	False	35	-0.061074	7	28	False	1	1	0
Acropora	GTT	False	False	35	-0.160180	9	27	False	1	1	0
Katharina	GTG	False	False	9	-0.011662	4	6	False	1	1	0
Geodia	GTT	False	False	38	-0.114519	8	30	False	1	1	0
Metridium	GTG	False	False	17	-0.196672	8	13	False	1	1	0
	GTA	False	False	50	-0.413634	10	17	False	2	3	0
Tetrancistrum_nebulosi	GTC	False	False	9	-0.233429	5	3	False	1	1	0
	GTT	False	False	39	-0.384615	10	12	False	2	2	0
Strigamia_maritima	GTT	False	False	31	-0.150204	8	21	False	1	1	0
**************************************	GTA	False	False	34	-0.019787	8	17	False	3	4	0
Trichinella	GTG	False	False	12	-0.302521	5	4	False	1	1	0
Balanoglossus	GTC	False	False	37	-0.145455	10	30	False	1	1	0

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	
Drosophila	GTA	False	False	33	-0.282592	10	24	False	1	1	(
Pseudogarypus_banksi	GTA	False	False	36	-0.087518	9	21	False	2	3	1
Benedenia_hoshinai	GTA	False	False	32	-0.458266	9	11	False	2	2	
	GTT	False	False	60	-0.395199	11	19	False	3	3	
/lyxine	GTC	False	False	13	-0.209486	7	8	False	1	1	
Ciona	GTA	False	False	35	-0.194096	9	26	False	1	1	
	GTT	False	False	36	-0.105797	9	28	False	1	1	
Paphnia	GTA	False	False	35	-0.198817	8	26	False	1	1	
Caenorhabditis	GTA	False	False	30	-0.305363	9	18	False	1	1	
	GTT	False	False	46	-0.107023	11	23	False	4	5	
scidiella_aspersa	GTG	False	False	36	-0.229885	10	21	False	2	2	
Blanceolatum	GTA	False	False	17	0.028358	7	12	False	1	1	
Epigonichthys	GTA	False	False	28	-0.108635	7	23	False	1	1	