

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Metaseiulus_occidentalis	TCA	False	False	17	0.739394	6	12	False	1	1
	TCG	False	False	4	1.205128	3	3	False	1	1
	TCT	False	False	39	1.009843	9	29	False	4	5
	AGT	False	False	32	0.149325	10	13	False	2	2
Schistosoma	TCA	False	False	17	0.521148	8	9	False	1	1
	TCG	False	False	6	0.575107	3	2	False	1	1
	TCT	False	False	38	0.361486	8	14	False	2	3
Aphrocallistes_vastus	AGT	False	False	5	0.924242	3	2	False	1	1
	TCA	False	False	28	0.800731	8	19	False	4	4
	TCC	False	False	14	0.747715	8	11	False	1	1
Paracentrotus	TCG	False	False	4	0.868750	3	2	False	1	1
	AGT	False	False	36	0.288682	10	20	False	2	2
	TCT	False	False	52	0.517174	10	20	False	4	6
Fasciola	AGT	False	False	45	0.056561	9	15	False	3	3
	TCC	False	False	2	1.300000	1	0	False	1	1
	TCG	False	False	6	0.926407	2	2	False	1	2
Trichobilharzia_regenti	TCT	False	False	37	0.472222	8	21	False	1	1
	AGT	False	False	37	0.088599	9	15	False	2	3
	TCA	False	False	7	0.637363	7	2	False	1	1
Versteria_mustelae	TCG	False	False	7	0.505455	6	2	False	1	1
	TCT	False	False	53	0.400000	11	20	False	3	6
	TCA	False	False	17	1.029096	7	14	False	2	2
Euseius_nicholsi	TCG	False	False	3	1.862069	3	2	False	1	1
	TCT	False	False	38	0.717172	8	28	False	2	2
	AGC	False	False	7	0.793358	5	5	False	1	1
Patiria	TCT	False	False	19	0.877333	6	16	False	2	2
	AGT	False	False	21	0.369645	9	10	False	1	1
	TCG	False	False	10	0.212821	6	3	False	1	1
Haplorchis_taichui	TCT	False	False	43	0.530113	8	15	False	4	6
	TCT	False	False	15	0.874355	9	13	False	1	1
	AGT	False	False	32	0.433841	10	21	False	3	3
Halocynthia	TCA	False	False	8	1.100000	6	6	False	1	1
	TCG	False	False	6	1.240506	5	4	False	2	2
	TCT	False	False	39	0.653997	10	29	False	2	2
Limulus	TCT	False	False	29	0.874126	8	25	False	2	2
	AGT	False	False	23	-0.015521	7	12	False	1	1
	TCA	False	False	10	0.491094	8	3	False	1	2
Echinococcus_equinus	TCT	False	False	41	0.625079	10	18	False	2	4

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Taenia_crassiceps	AGT	False	False	36	0.241501	9	17	False	2	3
	TCA	False	False	15	0.752137	4	10	False	1	1
	TCG	False	False	7	0.380435	6	2	False	1	1
Terebratulina	TCT	False	False	44	0.405279	11	14	False	3	6
	TCC	False	False	30	0.734468	10	27	False	1	1
	TCT	False	False	24	0.517609	10	18	False	1	1
Suberites_domuncula	TCC	False	False	10	0.989848	5	8	False	1	1
	TCT	False	False	26	0.642012	8	23	False	1	1
Bfloridae	TCA	False	False	12	0.809224	6	11	False	1	1
	TCG	False	False	7	0.884892	4	5	False	1	1
Katharina	TCT	False	False	42	0.866176	9	41	False	1	1
Geodia	TCT	False	False	34	0.768018	9	32	False	1	1
Amphimedon_queenslandica	TCC	False	False	9	0.865169	5	7	False	1	1
Tetrancistrum_nebulosi	AGT	False	False	25	-0.034729	9	7	False	2	2
	TCA	False	False	10	0.639386	7	5	False	1	1
	TCC	False	False	8	0.704180	5	2	False	2	2
Strigamia_maritima	TCT	False	False	38	0.608901	8	16	False	2	5
	TCT	False	False	29	0.862762	8	25	False	2	2
Trichinella	AGC	False	False	10	0.143959	6	3	False	1	1
	TCA	False	False	22	0.201632	9	13	False	1	1
	TCC	False	False	16	0.650160	6	8	False	2	3
Balanoglossus	TCT	False	False	19	0.796496	6	11	False	3	4
	TCG	False	False	5	0.944444	4	4	False	1	1
Acropora	TCT	False	False	36	0.743369	8	32	False	1	1
Pseudogarypus_banksi	AGT	False	False	7	0.848921	3	2	False	2	2
	TCA	False	False	30	0.696658	10	16	False	3	4
	TCC	False	False	6	1.587719	2	3	False	2	3
	TCG	False	False	3	1.508475	3	2	False	1	1
Benedenia_hoshinai	TCT	False	False	43	0.966071	11	28	False	4	7
	AGC	False	False	12	0.404255	6	6	False	1	1
	AGT	False	False	33	0.320155	9	14	False	3	3
	TCC	False	False	7	0.898551	6	1	False	2	3
	TCT	False	False	53	0.340887	10	23	False	4	4
Myxine	AGC	False	False	11	0.572770	5	7	False	1	1
	TCA	False	False	39	0.743775	10	33	False	1	1
	TCC	False	False	19	0.636608	7	14	False	1	1
Drosophila	TCA	False	False	30	0.926976	8	28	False	2	2
	TCT	False	False	31	0.724816	9	26	False	1	1
Ciona	AGC	False	False	3	1.211864	2	2	False	1	1
	AGT	False	False	27	0.452133	8	21	False	1	1
	TCA	False	False	26	0.813793	9	17	False	2	4

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Daphnia	TCC	False	False	5	1.576531	3	4	False	1	1
	TCT	False	False	33	0.985981	10	21	False	4	5
	AGT	False	False	9	0.752137	6	5	False	2	2
	TCG	False	False	10	0.913043	8	9	False	1	1
	TCT	False	False	28	0.906703	8	24	False	1	3
Caenorhabditis	AGT	False	False	13	0.926441	7	8	False	2	2
	TCC	False	False	2	2.113924	2	1	False	1	1
Homo	TCT	False	False	23	0.577053	9	12	False	2	3
	TCC	False	False	25	0.838611	8	24	False	1	1
Ascidella_aspersa	TCA	False	False	8	0.936709	5	6	False	1	1
	TCC	False	False	9	0.634561	5	6	False	1	1
Blanceolatum	TCT	False	False	36	1.097718	9	24	False	5	7
	TCC	False	False	6	0.929167	3	5	False	1	1
	TCT	False	False	23	0.723497	6	22	False	1	1
Epigonichthys	AGT	False	False	5	1.269841	4	4	False	1	1
	TCT	False	False	41	0.882390	8	37	False	2	3