

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Metaseiulus_occidentalis	ACA	False	False	19	0.083333	5	17	False	1	1
Aphrocallistes_vastus	ACA	False	False	47	-0.014795	9	31	False	2	2
	ACT	False	False	15	0.132759	6	14	False	1	1
Fasciola	ACG	False	False	4	1.012987	4	1	False	1	1
	ACT	False	False	34	0.259063	10	24	False	3	3
Trichobilharzia_regenti	ACA	False	False	4	0.551282	4	3	False	1	1
	ACT	False	False	28	0.041705	6	19	False	1	1
Versteria_mustelae	ACT	False	False	45	0.202286	10	29	False	3	3
Daphnia	ACT	False	False	24	0.011853	8	22	False	1	1
Haplorchis_taichui	ACG	False	False	9	0.258621	5	5	False	1	1
	ACT	False	False	27	0.401896	10	19	False	3	3
Halocynthia	ACT	False	False	31	0.023372	8	27	False	1	1
Echinococcus_equinus	ACG	False	False	11	0.209302	7	6	False	1	1
	ACT	False	False	41	0.174623	8	27	False	3	3
Taenia_crassiceps	ACA	False	False	10	0.536458	4	5	False	2	2
	ACT	False	False	38	0.212508	9	24	False	3	3
Axinella	ACA	False	False	25	0.023810	6	22	False	1	1
Suberites_domuncula	ACA	False	False	30	0.013841	9	27	False	1	1
Geodia	ACA	False	False	14	-0.025974	5	12	False	1	1
	ACC	False	False	8	0.207120	6	5	False	1	1
Tetrancistrum_nebulosi	ACG	False	False	8	0.288462	5	5	False	1	1
	ACT	False	False	27	0.036190	9	16	False	2	2
Trichinella	ACT	False	False	13	-0.125000	7	5	False	1	1
Amphimedon_queenslandica	ACA	False	False	22	-0.017689	7	21	False	1	1
Pseudogarypus_banksi	ACA	False	False	25	0.188272	7	17	False	1	3
	ACC	False	False	4	0.794872	3	2	False	1	1
Benedenia_hoshinai	ACA	False	False	14	0.412963	7	6	False	2	2
	ACC	False	False	5	0.510417	3	3	False	1	1
Myxine	ACT	False	False	33	-0.038401	7	24	False	1	1
	ACA	False	False	25	-0.169948	8	19	False	1	1
Ciona	ACT	False	False	20	0.086452	8	17	False	1	1
Strigamia_maritima	ACT	False	False	36	-0.063446	10	26	False	1	1
Caenorhabditis	ACA	False	False	28	0.008333	7	24	False	1	1
Homo	ACA	False	False	38	-0.123474	9	21	False	1	1
Ascidella_aspersa	ACA	False	False	30	0.228769	9	27	False	2	2
	ACT	False	False	21	0.215686	7	16	False	2	2