

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	AGA	False	False	21	-0.547648	9	0	False	1	1	0
Pseudogarypus_banksi	AGA	False	False	30	-0.691327	9	0	False	1	1	0
Schistosoma	AGG	False	False	17	-0.571856	9	0	False	1	1	0
	CGT	False	False	20	-1.714469	7	17	False	1	1	0
Taenia_crassiceps	CGT	False	False	26	-1.737769	7	22	False	1	1	0
Trichobilharzia_regenti	AGA	False	False	10	-0.221662	5	0	False	1	1	0
	CGA	False	False	4	-0.512821	4	3	False	1	1	0
Caenorhabditis	AGG	False	False	15	-0.443686	7	0	False	1	1	0
Benedenia_hoshinai	CGT	False	False	22	-1.659328	7	17	False	1	1	0
Echinococcus_equinus	CGT	False	False	23	-1.710468	8	19	False	1	1	0
Ciona	CGC	False	False	2	1.346154	2	1	False	1	1	0
Fasciola	AGG	False	False	12	-0.527660	7	0	False	1	1	0
	CGT	False	False	22	-1.667820	7	17	False	1	1	0
	AGA	False	False	20	-0.423077	6	0	False	1	1	0
Tetrancistrum_nebulosi	AGG	False	False	26	-0.808824	9	1	False	2	2	0
	CGT	False	False	14	-1.553734	7	13	False	1	1	0
Versteria_mustelae	CGT	False	False	20	-1.613027	6	17	False	1	1	0
Trichinella	AGA	False	False	32	-0.889871	7	0	False	1	1	0
Haplorchis_taichui	AGG	False	False	22	-0.505814	8	1	False	1	1	0
	CGA	False	False	4	-0.329032	3	2	False	1	1	0