

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	AGG	False	False	5	-0.281915	3	0	False	1	1	0
Halocynthia	AGG	False	False	31	-1.729218	9	0	False	1	1	0
	CGA	False	False	13	-1.309665	8	12	False	1	1	0
Schistosoma	AGG	False	False	17	-1.047904	9	0	False	1	1	0
Echinococcus_equinus	AGG	False	False	16	-1.097946	9	0	False	1	1	0
Caenorhabditis	AGA	False	False	45	-0.745888	9	0	False	4	5	0
	AGG	False	False	15	-0.390785	7	0	False	2	2	0
Benedenia_hoshinai	AGA	False	False	7	-0.510870	5	0	False	1	1	0
	AGG	False	False	4	0.335443	3	0	False	1	1	0
Ciona	AGA	False	False	78	-1.761810	11	0	False	3	3	0
Fasciola	AGG	False	False	12	-0.914894	7	0	False	1	1	0
	CGT	False	False	22	-1.726644	7	17	False	1	1	0
Trichobilharzia_regenti	AGG	False	False	4	0.392405	3	0	False	1	1	0
Tetrancistrum_nebulosi	AGA	False	False	20	-0.644872	6	0	False	2	2	0
	AGG	False	False	26	-0.989216	9	1	False	3	3	0
Versteria_mustelae	AGG	False	False	7	-0.715328	4	0	False	1	1	0
Trichinella	AGA	False	False	32	-1.201768	7	0	False	2	2	0
	CGT	False	False	5	-0.459184	3	4	False	1	1	0
Haplorchis_taichui	AGG	False	False	22	-0.608140	8	1	False	3	3	0
	CGG	False	False	12	-1.588110	7	8	False	1	1	0