CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	ı
Metaseiulus_occidentalis	AGA	False	False	21	0.290712	9	0	False	1	1	0
Halocynthia	AGA	False	False	29	0.211879	10	0	False	3	3	0
	AGG	False	False	31	0.104527	9	0	False	2	2	0
Schistosoma	AGA	False	False	18	0.384615	6	0	False	3	3	0
	AGG	False	False	17	0.497006	9	0	False	3	3	0
Aphrocallistes_vastus	AGA	False	False	32	0.509193	8	0	False	3	4	0
Pseudogarypus_banksi	AGA	False	False	30	0.353741	9	0	False	3	3	0
Echinococcus_equinus	AGA	False	False	12	-0.006369	6	3	False	1	1	0
	AGG	False	False	16	0.611374	9	0	False	2	3	0
Taenia_crassiceps	AGG	False	False	9	0.817416	6	0	False	2	2	0
Benedenia_hoshinai	AGA	False	False	7	0.692029	5	0	False	1	1	0
	AGG	False	False	4	0.607595	3	0	False	1	1	0
Balanoglossus	AGA	False	False	4	0.906250	3	0	False	1	1	0
Ciona	AGA	False	False	78	0.041295	11	0	False	4	6	0
Fasciola	AGG	False	False	12	0.400000	7	0	False	2	2	0
Terebratulina	AGG	False	False	4	0.406250	4	0	False	1	1	0
Tetrancistrum_nebulosi	AGA	False	False	20	0.341026	6	0	False	3	3	0
	AGG	False	False	26	-0.091176	9	1	False	2	2	0
Trichobilharzia_regenti	AGA	False	False	10	0.513854	5	0	False	1	2	0
Ascidiella_aspersa	AGA	False	False	6	0.218884	5	0	False	1	1	0
	AGG	False	False	26	0.379310	8	0	False	3	4	0
Versteria_mustelae	AGA	False	False	13	0.888454	8	0	False	3	4	0
Trichinella	AGA	False	False	32	0.030547	7	0	False	1	1	0
	CGA	False	False	24	-0.822757	8	23	False	1	1	0
Caenorhabditis	AGA	False	False	45	0.409529	9	0	False	4	5	0
	AGG	False	False	15	0.392491	7	0	False	1	1	0
Katharina	AGA	False	False	7	0.399281	5	0	False	1	1	0
	AGG	False	False	4	0.987013	3	0	False	1	1	0
Haplorchis_taichui	AGG	False	False	22	0.261628	8	1	False	4	4	0
Euseius_nicholsi	AGA	False	False	26	0.389831	9	0	False	1	2	0