CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	
Metaseiulus_occidentalis	AGG	False	False	5	0.037234	3	0	False	1	1	0
Pseudogarypus_banksi	AGA	False	False	30	0.162415	9	0	False	2	2	0
Echinococcus_equinus	AGG	False	False	16	-0.157978	9	0	False	1	1	0
Taenia_crassiceps	AGG	False	False	9	0.466292	6	0	False	1	1	0
Benedenia_hoshinai	AGG	False	False	4	1.031646	3	0	False	1	1	0
Drosophila	AGA	False	False	9	0.412607	6	0	False	1	1	0
Strigamia_maritima	AGA	False	False	12	1.036717	8	0	False	2	2	0
Caenorhabditis	AGA	False	False	45	0.086784	9	0	False	1	1	0
	AGG	False	False	15	0.612628	7	0	False	2	2	0
Tetrancistrum_nebulosi	AGA	False	False	20	0.496154	6	0	False	2	2	0
Versteria_mustelae	AGA	False	False	13	0.123288	8	0	False	1	1	0
Euseius_nicholsi	AGA	False	False	26	0.393819	9	0	False	3	4	0
Trichinella	AGA	False	False	32	-0.124598	7	0	False	1	1	0
Katharina	AGA	False	False	7	0.075540	5	0	False	1	1	0
	AGG	False	False	4	1.668831	3	0	False	1	1	0
Lumbricus	AGA	False	False	11	0.502315	7	0	False	1	1	0