## CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea.	
Metaseiulus_occidentalis	AGA	True	True	21	1.487334	9	0	True	9	13	(
	AGG	True	True	5	0.430851	3	0	True	1	1	(
Paracentrotus	AGA	True	True	7	2.386029	4	0	True	4	7	C
	AGG	True	True	2	2.600000	2	0	True	2	2	C
Aphrocallistes_vastus	AGA	True	True	32	1.310951	8	0	True	6	13	C
Schistosoma	AGA	True	True	18	1.015670	6	0	True	4	6	C
	AGG	True	True	17	1.360778	9	0	True	6	8	C
Fasciola	AGG	True	True	12	1.306383	7	0	True	5	5	C
Trichobilharzia_regenti	AGA	True	True	10	1.715365	5	0	True	3	5	C
	AGG	True	True	4	0.582278	3	0	True	2	2	C
Versteria_mustelae	AGA	True	True	13	1.091977	8	0	True	3	4	C
	AGG	True	True	7	1.248175	4	0	True	3	4	C
Euseius_nicholsi	AGA	True	True	26	1.478564	9	0	True	6	12	C
	AGG	True	True	5	1.440415	3	0	True	3	3	C
Patiria	AGA	True	True	4	2.506494	3	0	True	3	4	C
	AGG	True	True	1	2.425000	1	0	True	1	1	C
Haplorchis_taichui	AGA	True	True	3	2.741379	2	0	True	2	3	C
	AGG	True	True	22	1.295349	8	1	True	6	9	C
Lumbricus	AGA	True	True	11	2.194444	7	0	True	6	9	1
Halocynthia	AGG	False	False	31	0.162140	9	0	True	2	2	C
Limulus	AGA	True	True	7	1.764493	4	0	True	3	6	C
Echinococcus_equinus	AGA	True	True	12	0.815287	6	3	True	3	4	C
	AGG	True	True	16	1.146919	9	0	True	4	5	C
Taenia_crassiceps	AGA	False	False	12	0.466951	7	2	True	3	3	C
	AGG	True	True	9	1.814607	6	0	True	4	5	(
Terebratulina	AGA	True	True	8	2.159236	5	0	True	5	7	1
	AGG	True	True	4	0.312500	4	0	True	1	1	(
Katharina	AGA	True	True	7	1.190647	5	0	True	3	3	C
	AGG	True	True	4	1.668831	3	0	True	2	2	C
Tetrancistrum_nebulosi	AGA	True	True	20	1.567949	6	0	True	5	9	C
	AGG	True	True	26	0.686275	9	1	True	6	8	(
Strigamia_maritima	AGA	True	True	12	2.222462	8	0	True	6	10	1
Trichinella	AGA	True	True	32	0.995177	7	0	True	6	14	(
	AGG	True	True	5	2.318182	3	0	True	2	4	(
Balanoglossus	AGA	True	True	4	2.168750	3	0	True	2	3	(
Drosophila	AGA	True	True	9	1.722063	6	0	True	5	7	(
Pseudogarypus_banksi	AGA	True	True	30	1.359694	9	0	True	6	15	(
Benedenia_hoshinai	AGA	False	False	7	1.775362	5	0	True	4	4	(

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	
	AGG	False	False	4	1.272152	3	0	True	1	1	0
Ciona	AGG	False	False	25	0.080943	10	0	True	1	1	O
Saccoglossus	AGA	True	True	1	2.425000	1	0	True	1	1	0
Daphnia	AGA	True	True	7	1.599265	3	0	True	2	6	1
Caenorhabditis	AGA	True	True	45	1.581963	9	0	True	8	24	0
	AGG	True	True	15	1.740614	7	0	True	4	8	O
Ascidiella_aspersa	AGG	False	False	26	0.268966	8	0	True	2	2	0
	CGA	False	False	6	-0.101266	5	5	False	1	1	O
Blanceolatum	AGA	True	True	1	3.100000	1	0	True	1	1	0
Epigonichthys	AGA	True	True	2	2.537500	2	0	True	2	2	0