

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	P
Metaseiulus_occidentalis	CAA	False	False	21	0.671698	8	18	False	1	1	0.
Paracentrotus	CAA	False	False	19	0.844444	10	18	False	1	1	0.
Aphrocallistes_vastus	CAA	False	False	19	0.720708	9	16	False	1	1	0.
Drosophila	CAA	False	False	22	0.877088	10	21	False	1	1	0.
Taenia_crassiceps	CAA	False	False	11	1.048837	6	8	False	1	1	0.
Echinococcus_equinus	CAG	False	False	10	0.982143	7	6	False	1	1	0.
Balanoglossus	CAA	False	False	19	0.882759	9	18	False	1	1	0.
Ciona	CAA	False	False	14	0.970037	9	13	False	1	1	0.
Fasciola	CAG	False	False	8	1.137821	5	6	False	1	1	0.
Acropora	CAA	False	False	16	0.805324	9	13	False	1	1	0.
Homo	CAA	False	False	24	0.940152	10	21	False	2	2	0.
Versteria_mustelae	CAG	False	False	8	1.098726	5	4	False	1	1	0.
Euseius_nicholsi	CAG	False	False	5	1.654639	4	3	False	2	2	0.
Patiria	CAA	False	False	17	0.936434	8	16	False	1	1	0.
Caenorhabditis	CAA	False	False	16	0.598413	10	13	False	1	1	0.
Katharina	CAA	False	False	17	0.937690	9	16	False	1	1	0.
Saccoglossus	CAA	False	False	21	0.926799	9	20	False	1	1	0.
Terebratulina	CAG	False	False	4	1.297468	4	3	False	1	1	0.
Lumbricus	CAA	False	False	20	0.808042	9	18	False	1	1	0.