

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
Halocynthia	AGA	True	True	23	3.709797	5	0	True	5	18	0.995
	AGG	True	True	19	4.328859	4	0	True	4	17	0.995
Schistosoma	AGA	False	False	13	-0.160131	3	1	True	2	2	0.273
	CGC	False	False	1	3.583333	1	0	False	1	1	0.384
Caenorhabditis	AGA	False	False	32	-0.362184	5	1	True	1	2	0.176
	AGG	False	False	10	0.034783	4	0	True	1	2	0.204
Ciona	AGA	True	True	56	4.332055	5	0	True	5	49	0.991
	AGG	True	True	14	4.647416	5	0	True	5	13	0.995
Fasciola	AGG	False	False	5	1.049587	2	0	True	2	2	0.289
	CGG	False	False	5	-0.606838	4	3	False	1	1	0.150
Terebratulina	AGA	False	False	5	0.393162	2	0	True	1	1	0.114
Daphnia	AGA	False	False	7	0.000000	2	0	True	1	1	0.186
Trichinella	AGA	False	False	25	-0.172650	4	0	True	2	4	0.218