CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
Halocynthia	**AGG**	False	False	19	0.055928	4	0	True	1	1	0.175
Schistosoma	**AGA**	False	False	13	0.787582	3	1	True	2	4	0.316
	AGG	True	True	6	2.133803	3	0	True	2	4	0.504
Metridium	**CGA**	False	False	6	-0.390071	3	4	False	1	1	0.009
Geodia	**CGA**	False	False	8	-0.576087	4	7	False	1	1	0.005
Caenorhabditis	**AGA**	True	True	32	1.392810	5	1	True	4	15	0.989
	AGG	True	True	10	1.178261	4	0	True	2	4	0.993
Paracentrotus	**AGA**	True	True	7	2.506329	4	0	True	4	7	0.995
Drosophila	**AGA**	True	True	5	2.614035	4	0	True	4	5	0.822
Axinella	**CGG**	False	False	3	-0.072464	3	2	False	1	1	0.066
Ciona	**AGG**	False	False	14	0.115502	5	0	True	1	1	0.179
Fasciola	**AGG**	False	False	5	0.239669	2	0	True	1	1	0.203
Terebratulina	**AGA**	True	True	5	2.111111	2	0	True	2	4	0.661
Patiria	**AGA**	True	True	5	2.194690	3	0	True	3	4	0.595
Daphnia	**AGA**	True	True	7	1.798742	2	0	True	1	5	0.821
Limulus	**AGA**	True	True	6	2.544118	3	0	True	3	6	0.927
Katharina	**AGA**	True	True	4	2.173913	3	0	True	3	3	0.820
	AGG	True	True	5	1.547009	3	0	True	2	2	0.765
Trichinella	**AGA**	True	True	25	1.088889	4	0	True	4	10	0.988
Lumbricus	**AGA**	True	True	10	2.323404	5	0	True	5	7	0.998