

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Halocynthia	GAT	False	False	30	1.049096	8	27	False	2	2
Schistosoma	GAT	False	False	30	0.830596	8	25	False	1	1
Tetrancistrum_nebulosi	GAT	False	False	17	1.297583	7	13	False	2	2
Echinococcus_equinus	GAT	False	False	39	1.037798	8	30	False	2	2
Taenia_crassiceps	GAC	False	False	1	4.657895	1	0	False	1	1
	GAT	False	False	42	0.820197	8	32	False	1	1
Benedenia_hoshinai	GAT	False	False	26	1.127363	8	21	False	2	2
Myxine	GAC	False	False	23	1.079007	7	20	False	2	2
Amphimedon_queenslandica	GAT	False	False	27	0.968481	8	25	False	1	1
Fasciola	GAT	False	False	32	1.070388	8	26	False	2	2
Trichobilharzia_regenti	GAT	False	False	30	0.954231	8	22	False	2	2
Suberites_domuncula	GAT	False	False	30	0.976684	6	28	False	1	1
Homo	GAC	False	False	31	0.993333	7	28	False	1	1
	GAC	False	False	3	2.094828	3	2	False	1	1
Versteria_mustelae	GAT	False	False	36	0.989216	8	31	False	1	1
Ascidella_aspersa	GAT	False	False	32	0.926494	8	28	False	1	1
Trichinella	GAT	False	False	18	1.024148	7	14	False	1	1
	GAC	False	False	4	1.288462	4	2	False	1	1
Caenorhabditis	GAT	False	False	34	0.923427	8	31	False	1	1
	GAT	False	False	28	1.061581	7	22	False	3	3