CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Pı
Schistosoma	GGA	False	False	16	-0.183280	5	9	False	1	1	0.0
	GGG	False	False	24	0.102345	8	18	False	1	1	0.0
	GGT	False	False	63	-0.133361	10	49	False	1	1	0.0
Caenorhabditis	GGG	False	False	19	0.170501	6	17	False	1	1	0.0
Tetrancistrum_nebulosi	GGT	False	False	39	0.160560	9	28	False	3	3	0.0
Pseudogarypus_banksi	GGG	False	False	45	-0.103645	10	42	False	1	1	0.0
Benedenia_hoshinai	GGC	False	False	12	0.170940	8	9	False	1	1	0.0
	GGT	False	False	56	-0.072278	10	43	False	2	2	0.0
Taenia_crassiceps	GGA	False	False	21	0.238329	7	15	False	2	2	0.0
Echinococcus_equinus	GGG	False	False	24	-0.117457	9	18	False	1	1	0.0
	GGT	False	False	63	-0.176495	10	52	False	1	1	0.0
Epigonichthys	GGC	False	False	20	-0.039846	10	18	False	1	1	0.0
Fasciola	GGT	False	False	61	-0.076891	11	46	False	3	3	0.0
Trichobilharzia_regenti	GGA	False	False	8	0.570064	4	7	False	1	1	0.0
	GGG	False	False	35	-0.140950	10	24	False	1	1	0.0
	GGT	False	False	74	-0.285863	10	54	False	1	1	0.0
Suberites_domuncula	GGC	False	False	10	0.328982	7	8	False	1	1	0.0
Versteria_mustelae	GGG	False	False	15	0.025997	7	12	False	1	1	0.0
	GGT	False	False	77	-0.107955	11	61	False	1	1	0.0
Ascidiella_aspersa	GGT	False	False	42	0.161449	11	34	False	3	3	0.0
Acropora	GGG	False	False	68	-0.162580	12	60	False	1	1	0.0
Blanceolatum	GGT	False	False	23	-0.083969	7	20	False	1	1	0.1
	GGA	False	False	58	-0.061470	9	50	False	2	2	0.0
Trichinella	GGC	False	False	16	0.234910	5	14	False	1	1	0.0
	GGT	False	False	26	0.000000	7	22	False	1	1	0.0
Bfloridae	GGG	False	False	31	-0.072535	9	28	False	1	1	0.1
Haplorchis_taichui	GGA	False	False	14	0.354244	7	11	False	1	1	0.0
	GGG	False	False	57	-0.296346	10	39	False	1	1	0.0