

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Metaseiulus_occidentalis	GTA	False	False	26	0.879208	9	18	False	2	2
	GTC	False	False	4	1.526316	2	3	False	1	1
	GTT	False	False	26	0.925299	8	17	False	2	2
Paracentrotus	GTA	False	False	41	1.248731	10	32	False	3	5
	GTC	False	False	11	0.697674	7	9	False	1	1
	GTG	False	False	7	1.207885	4	6	False	1	1
Aphrocallistes_vastus	GTT	False	False	17	1.202144	7	15	False	1	1
	GTA	False	False	35	0.948301	8	29	False	1	1
	GTT	False	False	10	1.226562	5	8	False	1	1
Schistosoma	GTA	False	False	35	0.946143	10	9	False	5	10
	GTC	False	False	3	0.681034	2	0	False	1	1
	GTG	False	False	24	0.753439	10	8	False	3	3
Fasciola	GTT	False	False	65	1.165152	12	18	False	10	17
	GTA	False	False	13	0.369863	8	1	False	2	4
	GTG	False	False	23	0.896552	12	9	False	5	5
Trichobilharzia_regenti	GTT	False	False	98	1.029628	11	31	False	9	21
	GTA	False	False	17	1.701515	8	5	False	5	5
	GTG	False	False	30	0.583691	9	8	False	3	4
Versteria_mustelae	GTT	False	False	74	1.065744	12	22	False	11	22
	GTA	False	False	15	1.261538	8	5	False	4	4
	GTG	False	False	26	1.054241	11	7	False	5	6
Euseius_nicholsi	GTT	False	False	64	0.646846	11	23	False	8	12
	GTA	False	False	30	1.006873	10	23	False	2	2
	GTG	False	False	11	1.476636	5	6	False	2	3
Patiria	GTT	False	False	26	1.516451	9	15	False	4	6
	GTA	False	False	37	0.929371	9	34	False	2	2
	GTT	False	False	22	1.081049	8	16	False	1	1
Haplorchis_taichui	GTA	False	False	20	1.015404	10	7	False	5	6
	GTC	False	False	13	1.162055	6	5	False	3	3
	GTG	False	False	45	0.803307	10	12	False	6	9
Lumbricus	GTT	False	False	53	1.097821	11	16	False	9	14
	GTA	False	False	33	0.971116	9	29	False	2	2
	GTC	False	False	13	1.635644	7	6	False	2	3
Halocynthia	GTG	False	False	16	1.238562	6	12	False	1	2
	GTT	False	False	17	1.371166	8	13	False	2	2
	GTA	False	False	43	1.261078	11	22	False	8	11
Halocynthia	GTC	False	False	6	1.052863	3	5	False	1	1
	GTG	False	False	17	0.998478	7	8	False	3	3

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Limulus	GTT	False	False	43	1.353011	10	25	False	6	9
	GTA	False	False	35	1.120149	11	26	False	3	3
	GTC	False	False	11	1.330211	4	7	False	2	2
	GTT	False	False	23	1.455866	8	18	False	2	4
Echinococcus_equinus	GTA	False	False	12	0.657505	7	2	False	3	3
	GTC	False	False	3	1.194915	3	1	False	1	1
	GTG	False	False	36	1.133429	11	7	False	8	12
Taenia_crassiceps	GTT	False	False	74	0.761158	11	31	False	7	10
	GTA	False	False	24	1.097849	8	10	False	4	5
	GTG	False	False	12	0.974468	7	4	False	1	1
Axinella	GTT	False	False	70	0.902538	11	20	False	10	16
	GTA	False	False	23	1.168350	9	20	False	3	3
	GTT	False	False	32	1.052377	9	26	False	3	3
Terebratulina	GTC	False	False	23	1.343017	10	13	False	2	3
	GTT	False	False	14	1.455390	8	10	False	2	2
Suberites_domuncula	GTA	False	False	35	1.049301	7	28	False	2	2
	GTG	False	False	20	1.042691	7	15	False	2	2
	GTT	False	False	25	1.250520	8	20	False	2	2
Acropora	GTG	False	False	30	1.228326	8	25	False	3	3
	GTT	False	False	35	1.060629	9	27	False	2	3
Bfloridae	GTA	False	False	18	1.567643	7	9	False	4	6
	GTC	False	False	6	1.527426	6	5	False	1	1
	GTG	False	False	10	1.161209	6	5	False	3	3
Katharina	GTT	False	False	15	0.918197	5	11	False	1	1
	GTA	False	False	32	1.212783	11	21	False	4	5
	GTT	False	False	40	1.192506	10	34	False	2	3
Geodia	GTA	False	False	32	1.312500	8	23	False	3	4
	GTT	False	False	38	0.986367	8	30	False	3	3
Metridium	GTA	False	False	27	1.161041	8	22	False	2	3
	GTG	False	False	17	1.012103	8	13	False	1	1
Balanoglossus	GTT	False	False	30	1.064212	7	24	False	1	2
	GTA	False	False	22	1.021102	8	17	False	2	2
	GTC	False	False	37	1.119580	10	30	False	2	2
	GTG	False	False	6	1.453390	5	5	False	1	1
	GTT	False	False	16	1.252874	6	14	False	2	2
Tetrancistrum_nebulosi	GTA	False	False	50	0.718093	10	17	False	7	12
	GTC	False	False	9	1.420749	5	3	False	1	2
	GTG	False	False	27	1.697143	9	9	False	7	9
	GTT	False	False	39	0.652202	10	12	False	3	3
Strigamia_maritima	GTC	False	False	5	2.158730	5	2	False	2	2
	GTG	False	False	14	1.679035	7	7	False	4	4

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
Trichinella	GTT	False	False	31	1.142857	8	21	False	2	4
	GTA	False	False	34	0.741248	8	17	False	1	2
	GTG	False	False	12	0.886555	5	4	False	1	2
Amphimedon_queenslandica	GTT	False	False	22	1.625874	7	9	False	5	7
	GTA	False	False	26	1.162722	8	19	False	4	4
	GTG	False	False	49	1.114407	8	38	False	3	4
Drosophila	GTA	False	False	33	1.267760	10	24	False	4	4
	GTT	False	False	39	1.109861	9	34	False	3	3
	GTA	False	False	36	1.117647	9	21	False	5	7
Pseudogarypus_banksi	GTG	False	False	8	1.331190	5	5	False	1	2
	GTT	False	False	20	1.520513	7	14	False	3	3
	GTA	False	False	32	0.926164	9	11	False	4	8
Benedenia_hoshinai	GTC	False	False	6	0.519149	6	2	False	1	1
	GTG	False	False	6	1.178723	5	2	False	1	1
	GTT	False	False	60	0.768538	11	19	False	7	9
Myxine	GTA	False	False	30	1.084120	9	26	False	1	1
	GTC	False	False	13	1.442688	7	8	False	2	2
	GTT	False	False	19	1.216366	8	15	False	3	3
Ciona	GTA	False	False	35	1.086347	9	26	False	4	5
	GTG	False	False	5	1.680203	4	1	False	1	2
	GTT	False	False	36	1.129710	9	28	False	3	3
Saccoglossus	GTA	False	False	19	1.156757	8	15	False	2	2
	GTG	False	False	6	1.587983	3	4	False	2	2
	GTT	False	False	42	1.095327	10	37	False	1	1
Daphnia	GTA	False	False	35	1.124908	8	26	False	2	3
	GTC	False	False	7	1.566667	5	5	False	1	1
	GTG	False	False	8	1.571885	4	5	False	2	2
Caenorhabditis	GTT	False	False	33	1.223612	10	26	False	3	4
	GTA	False	False	30	0.681661	9	18	False	3	3
	GTC	False	False	6	1.017167	5	3	False	1	1
Homo	GTT	False	False	46	0.866221	11	23	False	4	4
	GTA	False	False	24	1.094565	7	20	False	2	2
	GTC	False	False	24	1.194805	10	21	False	1	1
Ascidiaella_aspersa	GTA	False	False	7	1.812500	5	2	False	3	3
	GTC	False	False	10	1.519182	7	7	False	2	3
	GTG	False	False	36	0.738506	10	21	False	4	5
Blanceolatum	GTT	False	False	50	1.290638	12	30	False	6	10
	GTA	False	False	17	1.188060	7	12	False	4	4
	GTT	False	False	17	1.186391	6	10	False	4	4
Epigonichthys	GTA	False	False	28	0.922934	7	23	False	1	1
	GTC	False	False	14	1.174954	11	10	False	2	2

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count
	GTT	False	False	25	1.304840	10	16	False	4	4