

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Pr
Metaseiulus_occidentalis	ATA	True	True	44	1.707189	10	7	True	4	10	0.9
Versteria_mustelae	ATT	False	False	43	1.022156	10	23	False	1	1	0.0
Euseius_nicholsi	ATA	True	True	36	1.587896	10	3	True	4	7	0.9
Patiria	ATA	False	True	50	1.243590	11	31	True	1	1	0.5
Lumbricus	ATA	True	True	29	1.801406	10	1	True	2	8	0.9
Halocynthia	ATA	False	False	27	1.531754	6	7	True	3	5	0.2
	ATT	False	False	56	0.935780	10	39	False	1	1	0.0
Limulus	ATA	True	True	28	2.083640	8	3	True	4	11	0.9
Caenorhabditis	ATA	True	True	33	1.448518	11	2	True	4	7	0.8
	ATT	False	False	76	1.080553	12	36	False	1	1	0.4
Axinella	ATA	False	True	41	1.233022	10	26	True	1	1	0.5
Terebratulina	ATA	True	True	27	1.766541	9	2	True	4	8	0.9
Bfloridae	ATA	True	True	9	2.030726	4	1	True	1	2	0.6
Katharina	ATA	True	True	22	2.060185	9	4	True	4	7	0.9
Tetrancistrum_nebulosi	ATT	False	False	29	1.154804	10	11	False	2	2	0.0
Strigamia_maritima	ATA	True	True	31	2.004184	8	4	True	4	10	0.9
Trichinella	ATA	True	True	56	1.031293	11	7	True	3	9	0.7
Drosophila	ATA	True	True	25	2.399384	8	3	True	4	11	0.9
Pseudogarypus_banksi	ATA	True	True	51	1.455321	11	7	True	4	10	0.9
Myxine	ATA	True	True	37	1.772884	8	3	True	4	11	0.9
Ciona	ATA	True	True	49	1.464995	11	5	True	3	8	0.9
Daphnia	ATA	True	True	11	2.424883	7	2	True	4	6	0.9
Homo	ATA	True	True	26	2.213861	9	2	True	4	9	0.9
Ascidella_aspersa	ATA	True	True	29	1.543517	11	2	True	4	5	0.9
Blanceolatum	ATA	True	True	10	2.272727	5	1	True	2	3	0.6
Epigonichthys	ATA	True	True	21	1.998761	7	1	True	4	6	0.9