

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	CAA	False	False	44	-1.007085	12	35	False	1	1	0
CfrijA	CAA	False	False	37	-0.853436	12	36	False	1	1	0
CgallA	CAA	False	False	41	-0.973195	12	36	False	1	1	0
CviswA	CAA	False	False	37	-0.853436	11	36	False	1	1	0
CdefoA	CAA	False	False	42	-0.996648	12	36	False	1	1	0
MstarA	CAA	False	False	46	-1.042184	12	35	False	1	1	0
NbaciA	CAA	False	False	26	-1.012286	7	21	False	1	1	0
Ctetra	CAG	False	False	6	0.023091	4	5	False	1	1	0
MmagnB	CAG	False	False	7	0.028912	7	6	False	1	1	0
MmagnA	CAA	False	False	39	-1.016626	11	29	False	1	1	0
LstarA	CAA	False	False	33	-1.096210	8	23	False	1	1	0
Mtetra	CAA	False	False	24	-0.889641	9	18	False	1	1	0
CtropB	CAG	False	False	19	-0.710333	9	18	False	1	1	0
CtropA	CAG	False	False	19	-0.710333	9	18	False	1	1	0
CphanA	CAA	False	False	42	-0.996648	12	36	False	1	1	0
CsojaA	CAA	False	False	21	-0.706897	7	20	False	1	1	0
BcustA	CAA	False	False	37	-0.956919	12	34	False	1	1	0
AgossA	CAA	False	False	27	-0.918033	7	21	False	1	1	0
AgossB	CAA	False	False	28	-0.921569	7	22	False	1	1	0
CglabA	CAA	False	False	28	-0.910003	7	25	False	1	1	0
NdelpA	CAA	False	False	28	-0.910003	7	25	False	1	1	0
ScryoA	CAA	False	False	25	-1.189179	7	16	False	1	1	0
CneerA	CAA	False	False	37	-0.853479	12	36	False	1	1	0
HuvarA	CAA	False	False	13	-0.543575	5	10	False	1	1	0
CalimA	CAA	False	False	41	-0.996556	12	35	False	1	1	0
CmaltA	CAA	False	False	36	-0.909893	12	32	False	1	1	0
SoctoA	CAA	False	False	24	-1.198802	7	15	False	1	1	0