

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MmagnB	CAA	False	False	38	-0.780488	11	29	False	1	1	0
CphanA	CAA	False	False	42	-0.804796	12	36	False	1	1	0
Cgalla	CAA	False	False	41	-0.822452	12	36	False	1	1	0
MmagnA	CAA	False	False	39	-0.767512	11	29	False	1	1	0
CdefoA	CAA	False	False	42	-0.804796	12	36	False	1	1	0
MingeA	CAA	False	False	44	-0.812118	12	35	False	1	1	0
MstarA	CAA	False	False	46	-0.801500	12	35	False	1	1	0
YlipoB	CAA	False	False	41	-0.802384	12	36	False	1	1	0
YlipoA	CAA	False	False	41	-0.802384	12	36	False	1	1	0
CsantA	CAA	False	False	24	-0.915018	10	18	False	1	1	0
CalimA	CAA	False	False	41	-0.799470	12	35	False	1	1	0
SjapoA	CAA	False	False	26	-0.600595	7	15	False	3	3	0
NbaciA	CAA	False	False	26	-0.813477	7	21	False	1	1	0
CcastA	CAA	False	False	24	-0.793998	7	22	False	1	1	0