

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	CAA	False	False	44	-1.759345	12	35	False	1	1	0.000
MmagnB	CAA	False	False	38	-1.730866	11	29	False	1	1	0.000
CgigaA	CAA	False	False	16	-1.638298	10	14	False	1	1	0.000
MmagnA	CAA	False	False	39	-1.733442	11	29	False	1	1	0.000
LstarA	CAA	False	False	33	-1.675737	8	23	False	1	1	0.001
MstarA	CAA	False	False	46	-1.743379	12	35	False	1	1	0.000
Csalma	CAA	False	False	40	-1.872811	11	30	False	1	1	0.000
MtetrA	CAA	False	False	24	-1.585624	9	18	False	1	1	0.000
SjapoA	CAA	False	False	26	-1.813313	7	15	False	1	1	0.000
SpombA	CAA	False	False	28	-1.902364	7	17	False	1	1	0.000
CsubhB	CAG	False	False	22	-1.655299	9	21	False	1	1	0.000
CsubhA	CAG	False	False	22	-1.655299	9	21	False	1	1	0.000