CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	TTA	False	False	335	-1.881042	13	283	False	1	1	0.000
CgallA	CTA	False	False	12	-1.474164	9	10	False	1	1	0.000
	TTA	False	False	312	-1.919277	13	263	False	1	1	0.000
CdefoA	TTA	False	False	318	-1.914385	13	268	False	1	1	0.000
PpastA	TTA	False	False	339	-1.898758	13	303	False	1	1	0.000
MstarA	TTA	False	False	333	-1.890846	13	285	False	1	1	0.000
NbaciA	CTA	False	False	8	0.544686	4	0	False	1	1	0.001
MtetrA	CTG	False	False	124	-1.887704	12	104	False	1	1	0.000
MmagnB	CTG	False	False	62	-1.820974	12	52	False	1	1	0.000
MmagnA	CTG	False	False	63	-1.824713	13	53	False	1	1	0.000
SpastA	TTA	False	False	195	-1.887587	7	175	False	1	1	0.000
CsantA	CTA	False	False	191	-1.792455	13	152	False	2	2	0.000
ScereC	TTA	False	False	194	-1.898061	7	176	False	1	1	0.000
ScereA	TTA	False	False	193	-1.898285	7	176	False	1	1	0.000
ScereD	CTT	False	False	2	0.334951	1	0	False	1	1	0.050
	TTA	False	False	194	-1.898061	7	176	False	1	1	0.000
ScereE	CTT	False	False	2	0.334951	1	0	False	1	1	0.061
	TTA	False	False	195	-1.898568	7	177	False	1	1	0.001
CphanA	TTA	False	False	333	-1.918903	13	281	False	1	1	0.000
YlipoB	TTA	False	False	317	-1.914377	13	266	False	1	1	0.000
YlipoA	TTA	False	False	317	-1.914377	13	266	False	1	1	0.000
CsalmA	TTA	False	False	294	-1.825959	12	228	False	1	1	0.000
AgossA	CTA	False	False	24	0.433858	5	2	False	1	1	0.000
AgossB	СТА	False	False	24	0.433858	5	2	False	1	1	0.000
SarboA	CTT	False	False	2	0.334951	1	0	False	1	1	0.061
	TTA	False	False	192	-1.896641	7	174	False	1	1	0.001
HuvarA	CTG	False	False	3	1.528481	2	0	False	1	1	0.072
	TTA	False	False	166	-1.817462	6	122	False	1	1	0.055
ScastA	CTA	False	False	5	0.644612	3	0	False	1	1	0.003
CalimA	TTA	False	False	324	-1.915927	13	275	False	1	1	0.000