

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LkluyF	TGT	False	False	13	-1.099180	5	9	False	1	1	0.000
MingeA	TGT	False	False	18	-0.963190	10	9	False	2	2	0.036
MmagnB	TGT	False	False	19	-1.004703	10	10	False	2	2	0.034
CphanA	TGT	False	False	16	-1.257566	8	8	False	1	1	0.001
CgallA	TGT	False	False	16	-1.257566	8	8	False	1	1	0.001
MmagnA	TGT	False	False	19	-1.004703	10	10	False	2	2	0.034
CdefoA	TGT	False	False	16	-1.257566	8	8	False	1	1	0.001
HuvarA	TGT	False	False	12	-1.134522	4	5	False	1	1	0.027
MstarA	TGT	False	False	19	-1.020196	10	9	False	2	2	0.037
YlipoB	TGT	False	False	16	-1.257566	8	8	False	1	1	0.001
YlipoA	TGT	False	False	16	-1.257566	8	8	False	1	1	0.001
CalimA	TGT	False	False	16	-1.257566	8	8	False	1	1	0.001
CsantA	TGT	False	False	21	-1.388651	10	6	False	1	1	0.000
MtetrA	TGT	False	False	18	-1.316020	10	10	False	1	1	0.000
SjapoA	TGT	False	False	9	-0.419424	5	3	False	1	1	0.008
LkluyD	TGT	False	False	12	-1.027375	5	8	False	1	1	0.000
LkluyC	TGT	False	False	13	-1.099180	5	9	False	1	1	0.000
CsalmA	TGT	False	False	24	-0.695592	10	11	False	3	3	0.135
LkluyA	TGT	False	False	13	-1.099180	5	9	False	1	1	0.000
LkluyE	TGT	False	False	13	-1.099180	5	9	False	1	1	0.000