CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
PpastA	**ATG**	False	False	58	-1.103973	11	48	False	1	1	0.000
MmagnB	**ATG**	False	False	57	-1.148708	11	39	False	1	1	0.000
MmagnA	**ATG**	False	False	57	-1.148708	11	39	False	1	1	0.000
LstarA	**ATG**	False	False	63	-1.185393	9	40	False	1	1	0.000
MstarA	**ATG**	False	False	56	-1.049539	10	39	False	2	2	0.001
BcustA	**ATG**	False	False	62	-1.143677	11	43	False	1	1	0.000
CcastA	**ATG**	False	False	41	-1.072787	6	30	False	1	1	0.000
CsantA	**ATG**	False	False	58	-1.145517	11	37	False	1	1	0.000
MtetrA	**ATG**	False	False	58	-1.184765	10	38	False	1	1	0.000
NbaciA	**ATG**	False	False	53	-1.112441	6	34	False	1	1	0.000
PkluyA	**ATG**	False	False	60	-1.127824	11	45	False	1	1	0.000
OanguA	**ATG**	False	False	54	-1.104483	11	45	False	1	1	0.000
MingeA	**ATG**	False	False	55	-1.142912	10	39	False	1	1	0.000
ScryoA	**ATG**	False	False	40	-1.165440	6	24	False	1	1	0.000