CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
HuvarA	TGA	False	False	30	-1.443378	4	24	False	2	2	0.106
MstarA	TGA	False	False	48	-1.704004	9	43	False	2	2	0.013
SjapoA	TGG	False	False	31	-1.516179	4	27	False	2	2	0.064
SpombA	TGG	False	False	31	-1.511864	5	26	False	2	2	0.069
ClusiA	TGA	False	False	46	-1.786370	9	45	False	1	1	0.000
MmagnB	TGA	False	False	45	-1.718802	9	42	False	2	2	0.012
MmagnA	TGA	False	False	45	-1.718802	9	42	False	2	2	0.012
LstarA	TGG	False	False	37	-1.821119	7	36	False	1	1	0.000
MguilA	TGA	False	False	45	-1.782015	9	44	False	1	1	0.000
CsantA	TGA	False	False	27	-1.406760	7	24	False	2	3	0.000
MtetrA	TGA	False	False	36	-1.796024	9	32	False	1	1	0.000
	TGG	False	False	9	-1.367898	6	8	False	1	1	0.000
MfariA	TGA	False	False	44	-1.775151	9	43	False	1	1	0.000
LelonA	TGA	False	False	47	-1.735687	9	45	False	2	2	0.015
PcanaA	TGA	False	False	47	-1.855917	9	43	False	1	1	0.000
DhansA	TGA	False	False	45	-1.802255	9	44	False	1	1	0.000
CsalmA	TGA	False	False	45	-1.427999	8	39	False	3	4	0.000
CsubhA	TGA	False	False	38	-1.713808	9	36	False	2	2	0.000
CsubhB	TGA	False	False	38	-1.713808	9	36	False	2	2	0.000
ScryoA	TGG	False	False	30	-1.529506	5	26	False	2	2	0.065
CsakeA	TGA	False	False	46	-1.786370	9	45	False	1	1	0.000
MingeA	TGA	False	False	46	-1.797686	9	42	False	1	1	0.000
CzempA	TGA	False	False	31	-1.416460	5	26	False	2	2	0.069
SoctoA	TGG	False	False	30	-1.529506	5	26	False	2	2	0.065