CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CgallA	AGT	False	False	42	-0.122409	12	35	False	1	1	0.000
	TCA	False	False	81	-0.305271	12	63	False	1	1	0.000
HuvarA	AGT	False	False	29	-0.210664	4	22	False	1	1	0.000
CphanA	TCA	False	False	85	-0.348453	12	63	False	1	1	0.000
	TCT	False	False	34	-0.118179	10	28	False	1	1	0.000
CgigaA	AGC	False	False	46	-0.093391	9	43	False	1	1	0.000
CbuenA	TCA	False	False	47	-0.108935	10	41	False	1	1	0.000
CdefoA	AGT	False	False	44	-0.178881	12	37	False	1	1	0.000
	TCA	False	False	80	-0.278348	12	62	False	1	1	0.000
LstarA	AGT	False	False	49	-0.283552	8	34	False	1	1	0.000
YlipoB	AGT	False	False	43	-0.177288	12	35	False	1	1	0.000
	TCA	False	False	75	-0.291795	12	57	False	1	1	0.000
YlipoA	AGT	False	False	43	-0.177288	12	35	False	1	1	0.000
	TCA	False	False	75	-0.291795	12	57	False	1	1	0.000
CalimA	TCA	False	False	82	-0.310010	12	63	False	1	1	0.000
	TCT	False	False	37	-0.285065	9	26	False	1	1	0.000
SjapoA	TCC	False	False	3	1.137821	3	0	False	1	1	0.009
	TCT	False	False	61	-0.453127	7	38	False	1	1	0.000
CsaraA	AGT	False	False	65	-0.145053	12	59	False	1	1	0.000
CtropB	TCG	False	False	9	0.433376	6	7	False	1	1	0.061
CzempA	TCA	False	False	38	-0.404893	6	20	False	1	1	0.000
CtropA	TCG	False	False	9	0.433376	6	7	False	1	1	0.061
SpombA	TCA	False	False	41	-0.375319	6	22	False	1	1	0.000
PkluyA	TCA	False	False	122	-0.278643	13	99	False	1	1	0.000
CtetrA	TCC	False	False	2	1.994253	2	1	False	1	1	0.115
CalaiA	TCA	False	False	61	-0.123775	11	55	False	1	1	0.000
ScryoA	TCC	False	False	3	0.668790	3	0	False	1	1	0.082