CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CsakeA	GCT	False	False	73	-1.946229	11	68	False	1	1	0.000
CphanA	GCT	False	False	108	-1.890099	12	71	False	1	1	0.000
CgallA	GCT	False	False	86	-1.874190	11	56	False	1	1	0.000
ClusiA	GCT	False	False	83	-1.897362	11	78	False	1	1	0.000
CcoryA	GCT	False	False	95	-1.920706	13	83	False	1	1	0.046
CdefoA	GCT	False	False	88	-1.891462	11	58	False	1	1	0.000
YlipoB	GCT	False	False	98	-1.852838	12	63	False	1	1	0.000
YlipoA	GCT	False	False	98	-1.852838	12	63	False	1	1	0.000
SoctoA	GCG	False	False	1	3.520000	1	0	False	1	1	0.082
	GCT	False	False	68	-1.486330	6	37	False	2	4	0.000
CalimA	GCT	False	False	109	-1.921324	12	69	False	1	1	0.000
SjapoA	GCT	False	False	61	-1.896001	6	37	False	1	1	0.000
CchauA	GCA	False	False	27	-1.739164	10	24	False	1	1	0.060
CsubhA	GCA	False	False	29	-1.731321	9	25	False	1	1	0.000
CsubhB	GCA	False	False	29	-1.731321	9	25	False	1	1	0.000
SpombA	GCA	False	False	22	-1.406655	6	10	False	2	2	0.000
	GCT	False	False	68	-1.534969	6	40	False	2	4	0.000
PkluyA	GCA	False	False	76	-1.919263	9	58	False	1	1	0.051
OanguA	GCT	False	False	82	-1.934116	12	70	False	1	1	0.046
ScryoA	GCA	False	False	32	-1.416642	6	20	False	2	2	0.000
	GCT	False	False	54	-1.451653	6	27	False	2	4	0.000