CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CneerA	GCA	False	False	20	-1.781926	6	18	False	1	1	0.000
LstarA	GCT	False	False	71	-1.867062	10	46	False	1	1	0.000
MmagnB	GCA	False	False	53	-1.709841	9	35	False	1	1	0.000
CfrijA	GCT	False	False	95	-1.932432	13	89	False	1	1	0.000
CgigaA	GCC	False	False	36	-1.737255	11	31	False	1	1	0.000
MmagnA	GCA	False	False	51	-1.690167	9	33	False	1	1	0.000
CsojaA	GCA	False	False	28	-1.835966	6	25	False	1	1	0.056
HuvarA	GCA	False	False	26	-1.656193	5	18	False	1	1	0.000
	GCT	False	False	59	-1.848563	6	39	False	2	2	0.000
CsalmA	GCC	False	False	7	0.156969	4	1	False	2	2	0.005
	GCT	False	False	86	-1.798188	11	63	False	1	1	0.000
AgossA	GCA	False	False	12	-1.290584	5	10	False	1	1	0.000
MtetrA	GCA	False	False	32	-1.729064	8	26	False	1	1	0.000
KlactA	GCT	False	False	54	-1.852842	6	42	False	1	1	0.000
AgossB	GCA	False	False	11	-1.230292	5	9	False	1	1	0.000
SpombA	GCC	False	False	5	-0.774131	4	3	False	1	1	0.000
CzempA	GCT	False	False	45	-1.962916	6	26	False	1	1	0.000
PpastA	GCT	False	False	54	-1.840871	12	48	False	1	1	0.000
SoctoA	GCT	False	False	68	-1.605468	6	37	False	1	1	0.000
MfariB	GCT	False	False	83	-1.973179	12	78	False	1	1	0.000
ScryoA	GCT	False	False	54	-1.640597	6	27	False	1	1	0.000