CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	**GCT**	False	False	61	-0.889619	6	51	False	1	1	0.000
CgallA	**GCA**	False	False	58	-0.894523	13	38	False	1	1	0.000
CdefoA	**GCA**	False	False	61	-0.880142	13	38	False	1	1	0.000
LdasiA	**GCT**	False	False	57	-0.851270	6	47	False	1	1	0.000
LkluyF	**GCT**	False	False	47	-0.924524	7	38	False	1	1	0.000
LkluyE	**GCT**	False	False	48	-0.947869	7	37	False	1	1	0.000
LkluyD	**GCT**	False	False	48	-0.934390	7	38	False	1	1	0.000
LkluyC	**GCT**	False	False	47	-0.924524	7	38	False	1	1	0.000
LkluyA	**GCT**	False	False	47	-0.924524	7	38	False	1	1	0.000
ClusiA	**GCA**	False	False	46	-0.935454	11	43	False	1	1	0.000
KtherA	**GCT**	False	False	58	-0.916749	6	51	False	1	1	0.000
MfariB	**GCA**	False	False	49	-0.871127	10	40	False	1	1	0.055
LanspB	**GCT**	False	False	58	-0.893847	6	49	False	1	1	0.000
CphanA	**GCA**	False	False	49	-0.860623	12	32	False	1	1	0.000
LanspA	**GCA**	False	False	35	-0.818896	5	27	False	1	1	0.000
YlipoB	**GCA**	False	False	54	-0.905955	11	37	False	1	1	0.000
YlipoA	**GCA**	False	False	54	-0.905955	11	37	False	1	1	0.000
LnothA	**GCT**	False	False	62	-0.892424	6	50	False	1	1	0.000
BcustA	**GCT**	False	False	66	-1.082525	11	51	False	1	1	0.000
OanguA	**GCA**	False	False	49	-0.809262	9	41	False	1	1	0.000
LelonA	**GCT**	False	False	93	-0.965006	13	88	False	1	1	0.000
HuvarA	**GCT**	False	False	59	-1.044075	6	39	False	1	1	0.046
CalimA	**GCC**	False	False	8	-0.088571	6	7	False	1	1	0.000
KlactA	**GCT**	False	False	54	-0.889853	6	42	False	1	1	0.000