CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
PpastA	**TGA**	False	False	46	-1.982403	9	42	False	1	1	0
LdasiA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
LkluyF	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
LkluyE	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
LkluyD	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
NbaciA	**TGA**	False	False	33	-1.917322	5	30	False	1	1	0
LkluyA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
SpastA	**TGA**	False	False	34	-1.942906	5	30	False	1	1	0
ScereC	**TGA**	False	False	33	-1.917322	5	30	False	1	1	0
ScereA	**TGA**	False	False	33	-1.917322	5	30	False	1	1	0
ScereD	**TGA**	False	False	34	-1.942906	5	30	False	1	1	0
ScereE	**TGA**	False	False	34	-1.942906	5	30	False	1	1	0
CcastA	**TGA**	False	False	34	-1.944272	5	30	False	1	1	0
PcanaA	**TGA**	False	False	47	-1.888889	9	43	False	1	1	0
LanspB	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
LanspA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
LnothA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
AgossB	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
CglabA	**TGA**	False	False	33	-1.923233	5	30	False	1	1	0
NdelpA	**TGA**	False	False	33	-1.923233	5	30	False	1	1	0
AgossA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
SarboA	**TGA**	False	False	34	-1.942906	5	30	False	1	1	0
ScastA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
KtherA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
KlactA	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0
LkluyC	**TGA**	False	False	35	-1.949739	5	31	False	1	1	0