CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
BarxA	**CAA**	False	False	8	-1.095455	1	4	False	1	1	0.262
MmagnB	**CAA**	False	False	38	-1.788618	11	29	False	2	2	0.000
MmagnA	**CAA**	False	False	39	-1.813573	11	29	False	2	2	0.000
CbuenA	**CAA**	False	False	36	-1.801157	11	33	False	1	1	0.000
LstarA	**CAA**	False	False	33	-1.822157	8	23	False	1	1	0.000
MstarA	**CAA**	False	False	46	-1.841575	12	35	False	2	2	0.086
CsalmA	**CAA**	False	False	40	-1.608461	11	30	False	2	2	0.092
BcustA	**CAA**	False	False	37	-1.859546	12	34	False	1	1	0.000
CsantA	**CAA**	False	False	24	-1.698741	10	18	False	2	2	0.000
MtetrA	**CAA**	False	False	24	-1.972939	9	18	False	1	1	0.000
	CAG	False	False	19	-1.610202	9	17	False	1	1	0.000
SjapoA	**CAA**	False	False	26	-1.491633	7	15	False	2	2	0.000
NbaciA	**CAA**	False	False	26	-1.761355	7	21	False	1	1	0.000
SoctoA	**CAA**	False	False	24	-1.216367	7	15	False	3	3	0.000
SpombA	**CAA**	False	False	28	-1.356629	7	17	False	3	3	0.042
CalaiA	**CAA**	False	False	33	-1.773427	10	31	False	1	1	0.000
MingeA	**CAA**	False	False	44	-1.870511	12	35	False	2	2	0.082
ScryoA	**CAA**	False	False	25	-1.245203	7	16	False	3	3	0.000