CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
AgossA	**TAT**	False	False	72	-1.977515	6	67	False	1	1	0.00
LstarA	**TAC**	False	False	14	-1.559219	3	12	False	1	1	0.06
MmagnB	**TAT**	False	False	33	-1.898332	9	28	False	1	1	0.00
CphanA	**TAT**	False	False	124	-2.050436	12	109	False	1	1	0.00
CgallA	**TAT**	False	False	112	-2.028099	13	100	False	1	1	0.00
MmagnA	**TAT**	False	False	29	-1.880697	9	24	False	1	1	0.00
CdefoA	**TAT**	False	False	114	-1.972038	13	101	False	2	2	0.00
MingeA	**TAT**	False	False	142	-2.050310	13	124	False	1	1	0.00
MstarA	**TAT**	False	False	145	-2.016426	13	129	False	1	1	0.00
YlipoB	**TAT**	False	False	114	-2.020209	13	102	False	1	1	0.00
WmucoA	**TAT**	False	False	120	-2.021091	12	104	False	1	1	0.00
CsalmA	**TAT**	False	False	133	-1.885140	12	112	False	2	2	0.00
CsantA	**TAT**	False	False	120	-1.927210	13	100	False	2	2	0.00
CalimA	**TAC**	False	False	25	-1.954230	10	21	False	1	1	0.00
	TAT	False	False	117	-2.014816	12	106	False	1	1	0.00
AgossB	**TAT**	False	False	72	-1.977515	6	67	False	1	1	0.00
SpombA	**TAT**	False	False	55	-1.973006	6	41	False	1	1	0.00
YlipoA	**TAT**	False	False	114	-2.020209	13	102	False	1	1	0.00
BcaliA	**TAT**	False	False	143	-2.034874	13	125	False	1	1	0.00
SoctoA	**TAT**	False	False	56	-1.886030	7	42	False	2	2	0.00
MtetrA	**TAC**	False	False	136	-1.972368	13	115	False	2	2	0.00
ScryoA	**TAT**	False	False	46	-1.968350	7	36	False	1	1	0.00