CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CglabA	**AGT**	False	False	12	-0.945643	5	10	False	1	1	0
SarboA	**TCT**	False	False	33	-0.861829	5	25	False	1	1	0
CsubhB	**AGC**	False	False	62	-0.919490	11	55	False	1	1	0
ScastA	**AGT**	False	False	17	-0.745753	6	13	False	1	1	0
SpastA	**TCA**	False	False	59	-1.008068	7	51	False	1	1	0
ScereC	**TCT**	False	False	28	-0.842862	5	23	False	1	1	0
ScereA	**TCT**	False	False	28	-0.842862	5	23	False	1	1	0
NbaciA	**TCT**	False	False	38	-0.976773	6	33	False	1	1	0
ScereD	**TCT**	False	False	27	-0.841546	5	21	False	1	1	0
ScereE	**TCT**	False	False	28	-0.847075	5	22	False	1	1	0
CsubhA	**AGC**	False	False	62	-0.919490	11	55	False	1	1	0
CcastA	**TCT**	False	False	22	-0.957667	5	16	False	1	1	0
NdelpA	**AGT**	False	False	11	-0.940818	5	9	False	1	1	0