

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
NbaciA	TCA	False	False	50	-1.654667	6	40	False	1	1	0.000
LstarA	TCT	False	False	53	-1.468257	7	37	False	1	1	0.000
MmagnB	TCA	False	False	52	-1.412589	11	34	False	1	1	0.000
CgigaA	TCA	False	False	39	-1.581651	10	34	False	1	1	0.000
MmagnA	TCA	False	False	52	-1.412589	11	34	False	1	1	0.000
CjiufA	AGT	False	False	50	-1.661748	10	47	False	1	1	0.056
HuvarA	TCA	False	False	50	-1.398281	6	31	False	1	1	0.000
	TCT	False	False	26	-1.235055	6	16	False	1	1	0.000
MstarA	AGT	False	False	74	-1.652220	12	56	False	1	1	0.000
MtetrA	TCA	False	False	37	-1.246392	9	26	False	1	1	0.000
BarxA	TCA	False	False	21	-1.069727	1	12	False	1	1	0.064
SpombA	AGT	False	False	31	-1.222973	6	17	False	1	1	0.000
	TCA	False	False	41	-1.133612	6	22	False	1	1	0.000
CzempA	TCA	False	False	38	-1.178982	6	20	False	1	1	0.000
	TCT	False	False	34	-1.383389	6	20	False	1	1	0.000
SoctoA	TCT	False	False	35	-1.244652	5	25	False	2	2	0.000
PkluyA	TCA	False	False	122	-1.648854	13	99	False	1	1	0.000
MingeA	AGT	False	False	73	-1.707850	12	56	False	1	1	0.000
ScryoA	TCT	False	False	29	-1.326309	6	20	False	1	1	0.000