

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	TCA	False	False	73	-0.316767	12	53	False	1	1	0.000
MmagnB	TCA	False	False	52	-0.313184	11	34	False	1	1	0.000
MmagnA	TCA	False	False	52	-0.313184	11	34	False	1	1	0.000
CparaE	TCA	False	False	52	-0.009352	10	48	False	1	1	0.056
CparaC	TCA	False	False	52	-0.009352	10	48	False	1	1	0.056
CparaB	TCA	False	False	52	-0.009352	10	48	False	1	1	0.056
MstarA	TCT	False	False	40	-0.290175	11	29	False	1	1	0.000
CparaD	TCA	False	False	52	-0.009352	10	48	False	1	1	0.056
BcustA	AGT	False	False	47	-0.329879	11	32	False	1	1	0.000
	TCT	False	False	27	-0.146151	9	22	False	1	1	0.000
CparaF	TCA	False	False	52	-0.009352	10	48	False	1	1	0.056
CsantA	AGT	False	False	78	-0.322656	11	54	False	1	1	0.000
MtetrA	TCT	False	False	25	0.022191	9	21	False	1	1	0.000
SjapoA	TCT	False	False	61	-0.408205	7	38	False	1	1	0.000
SpombA	AGT	False	False	31	-0.354115	6	17	False	1	1	0.000
	TCA	False	False	41	-0.339828	6	22	False	1	1	0.000
CzempA	TCA	False	False	38	-0.332501	6	20	False	1	1	0.000
SoctoA	TCA	False	False	35	-0.266504	6	21	False	1	1	0.000
PkluyA	AGT	False	False	46	-0.274011	11	33	False	1	1	0.000
OanguA	AGT	False	False	40	-0.271336	10	28	False	1	1	0.000
ScryoA	AGT	False	False	31	-0.368083	6	14	False	1	1	0.000
	TCA	False	False	37	-0.254968	6	24	False	1	1	0.000