

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	TCA	False	False	73	-1.870894	12	53	False	1	1	0.000
MmagnB	TCA	False	False	52	-1.886010	11	34	False	1	1	0.000
CphanA	TCA	False	False	85	-1.982851	12	63	False	1	1	0.000
CgallA	TCA	False	False	81	-1.973774	12	63	False	1	1	0.000
MmagnA	TCA	False	False	52	-1.886010	11	34	False	1	1	0.000
CdefoA	TCT	False	False	38	-1.831835	10	28	False	1	1	0.000
HuvarA	TCA	False	False	50	-1.977268	6	31	False	1	1	0.000
YlipoB	TCT	False	False	41	-1.855044	9	31	False	1	1	0.000
YlipoA	TCT	False	False	41	-1.855044	9	31	False	1	1	0.000
CsantA	AGC	False	False	24	-1.803476	10	16	False	1	1	0.000
CalimA	TCT	False	False	37	-1.846713	9	26	False	1	1	0.000
SjapoA	TCT	False	False	61	-1.809546	7	38	False	1	1	0.000
NbaciA	TCT	False	False	38	-1.908841	6	33	False	1	1	0.000
CzempA	AGT	False	False	38	-1.745601	6	21	False	2	2	0.000
	TCA	False	False	38	-1.832751	6	20	False	1	1	0.000
CsubhB	AGC	False	False	62	-1.827454	11	55	False	1	1	0.000
MfariA	AGC	False	False	3	0.407767	1	2	False	1	1	0.063
CsubhA	AGC	False	False	62	-1.827454	11	55	False	1	1	0.000