

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	TTT	False	False	125	-0.728641	13	95	False	1	1	0.000
CgallA	TTT	False	False	70	-0.736745	13	50	False	1	1	0.000
CdefoA	TTT	False	False	69	-0.795143	12	51	False	1	1	0.000
MstarA	TTC	False	False	40	-0.595162	8	31	False	2	2	0.000
SjapoA	TTC	False	False	75	-0.794443	6	51	False	1	1	0.000
	TTT	False	False	58	-0.301812	6	32	False	1	1	0.000
SpombA	TTC	False	False	76	-0.660083	7	52	False	1	1	0.000
	TTT	False	False	43	-0.582684	6	28	False	1	1	0.000
AacerA	TTT	False	False	22	-0.461957	2	19	False	1	1	0.116
MmagnB	TTC	False	False	157	-0.716816	13	116	False	1	1	0.000
MmagnA	TTC	False	False	157	-0.728525	13	117	False	1	1	0.000
LstarA	TTT	False	False	92	-0.593692	9	59	False	1	1	0.000
CsantA	TTT	False	False	109	-0.765216	13	88	False	2	2	0.000
CphanA	TTT	False	False	77	-0.751893	13	59	False	1	1	0.000
YlipoB	TTT	False	False	68	-0.751287	12	49	False	1	1	0.000
BcustA	TTC	False	False	71	-0.621571	12	55	False	1	1	0.000
	TTT	False	False	102	-0.599666	13	76	False	1	1	0.000
AgossA	TTT	False	False	78	-0.769910	7	68	False	1	1	0.051
AgossB	TTT	False	False	75	-0.761856	7	66	False	1	1	0.051
YlipoA	TTT	False	False	68	-0.751287	12	49	False	1	1	0.000
ScryoA	TTC	False	False	78	-0.663730	6	55	False	2	2	0.000
HuvarA	TTC	False	False	57	-0.700034	6	45	False	1	1	0.000
CalimA	TTT	False	False	57	-0.602487	12	42	False	1	1	0.000
CzempA	TTC	False	False	39	-0.812105	7	28	False	1	1	0.000
	TTT	False	False	81	-0.617340	7	59	False	1	1	0.000
SoctoA	TTC	False	False	68	-0.599327	6	47	False	2	2	0.000