## CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	AAA	False	False	40	2.104284	10	38	False	2	2	0.194
LmeyeA	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
CgallA	AAA	False	False	39	2.084355	11	33	False	3	3	0.333
CviswA	AAA	False	False	39	2.067975	10	37	False	1	1	0.194
CdefoA	AAA	False	False	39	2.084355	11	33	False	3	3	0.333
MstarA	AAA	False	False	42	1.849738	11	38	False	1	1	0.194
ScastA	AAA	False	False	20	1.907425	7	18	False	1	1	0.286
AgossA	AAA	False	False	21	1.971954	6	19	False	1	1	0.286
LkluyF	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
SjapoA	AAA	False	False	20	0.494881	6	9	False	1	1	0.267
LkluyD	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
NbaciA	AAA	False	False	22	2.059211	6	19	False	2	2	0.455
CvartA	AAA	False	False	43	2.188638	11	40	False	3	3	0.408
LkluyA	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
MmagnB	AAA	False	False	30	1.907038	11	27	False	1	1	0.194
ScereC	AAA	False	False	19	1.995947	7	17	False	1	1	0.194
MmagnA	AAA	False	False	30	1.907038	11	27	False	1	1	0.194
WmucoA	AAA	False	False	45	2.191671	11	42	False	3	3	0.410
CalimA	AAA	False	False	41	2.080751	11	35	False	3	3	0.420
CsantA	AAA	False	False	34	1.523982	12	26	False	2	2	0.174
MtetrA	AAA	False	False	24	2.169829	11	23	False	1	1	0.194
ScereA	AAA	False	False	19	1.995947	7	17	False	1	1	0.194
CsuavA	AAA	False	False	41	2.198957	11	38	False	3	3	0.333
ScereD	AAA	False	False	20	1.961409	7	18	False	1	1	0.286
ScereE	AAA	False	False	20	1.961409	7	18	False	1	1	0.286
MfariA	AAA	False	False	45	1.843574	11	42	False	1	1	0.194
PkluyA	AAA	False	False	45	1.795755	13	39	False	2	2	0.291
CcastA	AAA	False	False	21	1.908674	6	18	False	1	1	0.286
PcanaA	AAA	False	False	43	2.032516	11	40	False	2	2	0.194
WpijpA	AAA	False	False	20	2.181159	6	19	False	1	1	0.194
LanspB	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
CphanA	AAA	False	False	40	2.082705	11	34	False	3	3	0.333
LanspA	AAA	False	False	19	2.145473	6	17	False	2	2	0.357
LdasiA	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
CsalmA	AAA	False	False	39	1.853165	11	33	False	2	2	0.292
YlipoB	AAA	False	False	40	2.042547	11	32	False	4	4	0.385
YlipoA	AAA	False	False	40	2.042547	11	32	False	4	4	0.385
SoctoA	AAA	False	False	21	0.424270	6	10	False	1	1	0.267

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CjadiA	AAA	False	False	45	2.126119	11	41	False	3	3	0.411
AgossB	AAA	False	False	21	1.971954	6	19	False	1	1	0.286
CsubhA	AAG	False	False	40	1.944461	12	38	False	1	1	0.000
BcustA	AAA	False	False	47	2.044070	12	40	False	4	5	0.195
CglabA	AAA	False	False	19	1.995947	7	17	False	1	1	0.194
CsubhB	AAG	False	False	40	1.944461	12	38	False	1	1	0.000
LkluyC	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
OanguA	AAA	False	False	45	1.750789	13	39	False	2	2	0.194
NdelpA	AAA	False	False	19	1.995947	7	17	False	1	1	0.286
ScryoA	AAA	False	False	20	0.399029	6	10	False	1	1	0.267
LnothA	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
SarboA	AAA	False	False	21	1.878509	7	18	False	1	1	0.286
MingeA	AAA	False	False	40	1.841357	11	36	False	1	1	0.194
LkluyE	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
CalaiA	AAA	False	False	41	1.990368	9	38	False	2	2	0.194
KtherA	AAA	False	False	19	2.063862	6	18	False	1	1	0.286
KlactA	AAA	False	False	20	2.168593	6	18	False	2	2	0.457
BcaliA	AAA	False	False	43	2.167714	11	40	False	3	3	0.333
SpombA	AAA	False	False	16	0.751364	6	9	False	1	1	0.116
SpastA	AAA	False	False	20	1.961409	7	18	False	1	1	0.286