

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	TGA	False	False	44	2.148192	9	42	False	1	1	0.000
LmeyerA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
CgallA	TGA	False	False	46	2.290174	9	42	False	2	2	0.000
CdefoA	TGA	False	False	45	2.296043	9	41	False	2	2	0.000
PpastA	TGA	False	False	46	2.154202	9	42	False	2	2	0.000
MstarA	TGA	False	False	48	2.183536	9	43	False	3	3	0.002
AgossA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
LkluyF	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
LkluyE	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
LkluyD	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
NbaciA	TGA	False	False	33	2.054148	5	30	False	1	1	0.000
CvartA	TGA	False	False	43	2.135493	9	42	False	1	1	0.000
LkluyA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
SpombA	TGG	False	False	31	2.207088	5	26	False	1	1	0.059
AacerA	TGA	False	False	15	2.207477	2	14	False	1	1	0.084
MmagnB	TGA	False	False	45	2.068503	9	42	False	1	1	0.000
ScereC	TGA	False	False	33	2.054148	5	30	False	1	1	0.000
MmagnA	TGA	False	False	45	2.068503	9	42	False	1	1	0.000
WmucoA	TGA	False	False	43	2.135493	9	42	False	1	1	0.000
CalimA	TGA	False	False	46	2.290174	9	42	False	2	2	0.000
MtetrA	TGA	False	False	36	1.934025	9	32	False	1	1	0.000
ScereA	TGA	False	False	33	2.054148	5	30	False	1	1	0.000
CsuavA	TGA	False	False	43	2.135493	9	42	False	1	1	0.000
ScereD	TGA	False	False	34	2.075749	5	30	False	1	1	0.000
ScereE	TGA	False	False	34	2.075749	5	30	False	1	1	0.000
PkluyA	TGA	False	False	40	2.262552	9	38	False	2	2	0.000
MfariB	TGA	False	False	46	2.159080	9	45	False	1	1	0.000
CcastA	TGA	False	False	34	2.168034	5	30	False	2	2	0.000
PcanaA	TGA	False	False	47	1.868112	9	43	False	1	1	0.000
WpijpA	TGA	False	False	31	2.173172	5	30	False	1	1	0.000
BcaliA	TGA	False	False	44	2.148192	9	42	False	1	1	0.000
LanspB	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
CphanA	TGA	False	False	46	2.290174	9	42	False	2	2	0.000
LanspA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
LdasiA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
YlipoB	TGA	False	False	45	2.297947	9	41	False	2	2	0.000
YlipoA	TGA	False	False	46	2.181984	9	41	False	2	2	0.000
Csalma	TGA	False	False	45	1.991987	8	39	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CjadiA	TGA	False	False	43	2.135493	9	42	False	1	1	0.000
AgossB	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
CsubhA	TGG	False	False	9	2.401451	6	8	False	1	1	0.000
BcustA	TGA	False	False	42	2.334575	9	38	False	3	3	0.003
CglabA	TGA	False	False	33	2.079093	5	30	False	1	1	0.000
CsubhB	TGG	False	False	9	2.401451	6	8	False	1	1	0.000
OanguA	TGA	False	False	42	2.269773	9	39	False	2	2	0.000
NdelpA	TGA	False	False	33	2.079093	5	30	False	1	1	0.000
MingeA	TGA	False	False	46	2.241974	9	42	False	3	3	0.003
LnothA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
SarboA	TGA	False	False	34	2.075749	5	30	False	1	1	0.000
HuvarA	TGA	False	False	30	1.793026	4	24	False	1	1	0.000
ScastA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
KtherA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
KlactA	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
CblacA	TGA	False	False	11	2.201337	1	10	False	1	1	0.172
CzempA	TGA	False	False	31	2.020730	5	26	False	1	1	0.000
LkluyC	TGA	False	False	35	2.096402	5	31	False	1	1	0.000
SpastA	TGA	False	False	34	2.075749	5	30	False	1	1	0.000