

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	TGA	False	False	44	1.166627	9	42	False	1	1	0.000
LmeyeA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
CfrijA	TGA	False	False	42	1.168349	10	41	False	1	1	0.000
CgallA	TGA	False	False	46	1.294307	9	42	False	2	2	0.000
CbuenA	TGA	False	False	44	1.168222	10	43	False	1	1	0.000
CviswA	TGA	False	False	45	1.165412	10	44	False	1	1	0.000
CdefoA	TGA	False	False	45	1.299105	9	41	False	2	2	0.000
PpastA	TGA	False	False	46	1.152582	9	42	False	1	1	0.000
LdasiA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
ScastA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
LkluyF	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
LkluyE	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
CchauA	TGA	False	False	45	1.134112	9	43	False	1	1	0.000
NbaciA	TGA	False	False	33	1.112082	5	30	False	1	1	0.000
SpombA	TGG	False	False	31	1.343297	5	26	False	2	2	0.003
LkluyA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
CdublA	TGA	False	False	38	1.196338	9	37	False	1	1	0.000
CalbiE	TGA	False	False	21	1.282787	5	20	False	1	1	0.000
CtetrA	TGA	False	False	44	1.173524	10	43	False	1	1	0.000
CalbiC	TGA	False	False	46	1.164828	10	45	False	1	1	0.000
CalbiB	TGA	False	False	46	1.164828	10	45	False	1	1	0.000
CalbiA	TGA	False	False	38	1.196338	9	37	False	1	1	0.000
CalbiD	TGA	False	False	38	1.196338	9	37	False	1	1	0.000
CgigaA	TGG	False	False	10	1.491474	6	9	False	1	1	0.000
CalimA	TGA	False	False	46	1.294307	9	42	False	2	2	0.000
CcoryA	TGA	False	False	45	1.183919	9	44	False	1	1	0.000
CbohiA	TGA	False	False	44	1.188525	9	43	False	1	1	0.000
SpastA	TGA	False	False	34	1.224703	5	30	False	2	2	0.004
ScereC	TGA	False	False	33	1.112082	5	30	False	1	1	0.000
	TGG	False	False	1	4.980583	1	0	False	1	1	0.190
CsaraA	TGA	False	False	43	1.168690	10	42	False	1	1	0.000
CtropB	TGA	False	False	44	1.168222	10	43	False	1	1	0.000
BcaliA	TGA	False	False	44	1.166627	9	42	False	1	1	0.000
ScereD	TGA	False	False	34	1.224703	5	30	False	2	2	0.004
CtropA	TGA	False	False	44	1.168222	10	43	False	1	1	0.000
SjapoA	TGG	False	False	31	1.341757	4	27	False	2	2	0.002
CcastA	TGA	False	False	34	1.158699	5	30	False	1	1	0.000
LanspB	TGA	False	False	35	1.231255	5	31	False	2	2	0.001

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CphanA	TGA	False	False	46	1.294307	9	42	False	2	2	0.000
LanspA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
CsojaA	TGA	False	False	32	1.159160	7	31	False	1	1	0.000
ScereE	TGA	False	False	34	1.224703	5	30	False	2	2	0.004
YlipoB	TGA	False	False	45	1.300980	9	41	False	2	2	0.000
YlipoA	TGA	False	False	46	1.210419	9	41	False	2	2	0.000
LnothA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
AgossB	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
BcustA	TGA	False	False	42	1.285892	9	38	False	1	1	0.000
CglabA	TGA	False	False	33	1.125327	5	30	False	1	1	0.000
SoctoA	TGG	False	False	30	1.288995	5	26	False	2	2	0.003
OanguA	TGA	False	False	42	1.263980	9	39	False	1	1	0.000
NdelpA	TGA	False	False	33	1.125327	5	30	False	1	1	0.000
ScryoA	TGG	False	False	30	1.288995	5	26	False	2	2	0.003
CneerA	TGA	False	False	45	1.165412	10	44	False	1	1	0.000
AgossA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
SarboA	TGA	False	False	34	1.224703	5	30	False	2	2	0.004
HuvarA	TGA	False	False	30	1.046065	4	24	False	1	1	0.000
ScereA	TGA	False	False	33	1.112082	5	30	False	1	1	0.000
	TGG	False	False	1	4.980583	1	0	False	1	1	0.190
KtherA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
KlactA	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
LkluyD	TGA	False	False	35	1.231255	5	31	False	2	2	0.001
CzempA	TGA	False	False	31	1.194307	5	26	False	1	1	0.000
CmaltA	TGA	False	False	45	1.170133	10	44	False	1	1	0.000
LkluyC	TGA	False	False	35	1.231255	5	31	False	2	2	0.001