

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CfrijA	CGT	False	False	22	2.083709	10	21	False	1	1	0.059
CgallA	AGA	False	False	51	1.994023	12	46	False	3	3	0.199
CbuenA	AGA	False	False	32	2.068896	10	30	False	1	1	0.004
CviswA	CGT	False	False	25	2.085281	11	24	False	1	1	0.059
CdefoA	AGA	False	False	51	1.994023	12	46	False	3	3	0.199
PpastA	CGT	False	False	8	2.219945	5	7	False	1	1	0.000
MstarA	CGT	False	False	17	2.128892	9	16	False	1	1	0.000
ScastA	AGA	False	False	33	1.962737	7	31	False	1	1	0.005
AgossA	AGA	False	False	32	1.915966	7	30	False	1	1	0.070
LkluyF	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
SjapoA	AGA	False	False	26	1.882873	7	21	False	1	2	0.005
	CGT	False	False	5	2.381783	4	4	False	1	1	0.000
CchauA	CGT	False	False	40	2.160441	11	36	False	2	3	0.000
SpombA	AGA	False	False	27	1.725726	7	23	False	1	1	0.005
CvartA	AGA	False	False	29	1.954907	10	27	False	1	1	0.006
	CGT	False	False	22	2.093296	9	21	False	1	1	0.000
LkluyA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CdublA	CGT	False	False	32	2.077181	10	31	False	1	1	0.000
CalbiE	CGT	False	False	17	2.147476	6	16	False	1	1	0.000
CalbiD	CGT	False	False	31	2.079944	10	30	False	1	1	0.000
CalbiC	CGT	False	False	33	2.062500	11	32	False	1	1	0.000
CalbiB	CGT	False	False	33	2.062500	11	32	False	1	1	0.000
CalbiA	CGT	False	False	31	2.079944	10	30	False	1	1	0.000
Ctetra	AGA	False	False	28	2.058891	10	26	False	1	1	0.067
	CGT	False	False	25	2.085509	11	24	False	1	1	0.055
MmagnB	AGA	False	False	33	2.007097	11	31	False	1	1	0.006
	CGT	False	False	19	2.115670	9	18	False	1	1	0.000
ScereC	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CgigaA	AGA	False	False	14	2.231360	7	13	False	1	1	0.006
MmagnA	AGA	False	False	33	2.007097	11	31	False	1	1	0.006
	CGT	False	False	19	2.115670	9	18	False	1	1	0.000
CcoryA	CGT	False	False	48	2.138988	11	44	False	2	3	0.000
CbohiA	CGA	False	False	2	3.209302	2	1	False	1	1	0.038
	CGT	False	False	42	2.114691	10	39	False	1	2	0.000
LstarA	AGA	False	False	33	2.256664	7	30	False	2	3	0.006
SpastA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CalimA	AGA	False	False	51	1.994023	12	46	False	3	3	0.199

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CsantA	AGA	False	False	19	2.163032	9	18	False	1	1	0.006
	AGG	False	False	10	2.000000	5	8	False	1	1	0.003
MtetrA	AGA	False	False	32	1.958027	11	29	False	1	1	0.006
	CGT	False	False	18	2.120546	9	17	False	1	1	0.000
LelonA	AGA	False	False	35	2.064858	11	34	False	1	1	0.067
	CGT	False	False	19	1.971248	10	17	False	1	1	0.055
CsaraA	CGT	False	False	23	2.116853	9	22	False	1	1	0.059
CtropB	CGT	False	False	28	2.080784	11	27	False	1	1	0.000
CsuavA	AGA	False	False	30	1.956712	10	28	False	1	1	0.006
	CGT	False	False	21	2.098369	9	20	False	1	1	0.000
ScereD	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CtropA	CGT	False	False	28	2.080784	11	27	False	1	1	0.000
PkluyA	AGA	False	False	41	1.966614	11	39	False	1	1	0.006
	CGT	False	False	8	2.220246	5	7	False	1	1	0.000
MfariB	CGT	False	False	16	2.172505	8	15	False	1	1	0.060
CcastA	AGA	False	False	34	1.999717	7	31	False	2	2	0.010
LanspB	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CphanA	AGA	False	False	51	1.994023	12	46	False	3	3	0.199
LanspA	AGA	False	False	33	1.993597	7	30	False	2	2	0.010
CsojaA	CGT	False	False	22	2.068663	7	21	False	1	1	0.000
ScereE	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
YlipoB	AGA	False	False	50	1.985479	12	45	False	3	3	0.197
YlipoA	AGA	False	False	50	1.985479	12	45	False	3	3	0.197
SoctoA	AGA	False	False	28	1.886573	7	23	False	1	2	0.005
Cjadia	AGA	False	False	31	1.957223	10	29	False	1	1	0.006
ScereA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
AgossB	AGA	False	False	32	1.915966	7	30	False	1	1	0.070
CsubhA	AGG	False	False	5	2.664474	4	4	False	1	1	0.004
BcustA	AGA	False	False	41	1.815761	11	37	False	1	1	0.006
CglabA	AGA	False	False	34	1.853507	7	31	False	1	1	0.005
CsubhB	AGG	False	False	5	2.664474	4	4	False	1	1	0.004
LkluyC	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
OanguA	CGT	False	False	15	2.098516	5	14	False	1	1	0.000
NdelpA	AGA	False	False	34	1.853507	7	31	False	1	1	0.005
ScryoA	AGA	False	False	24	1.866164	7	21	False	1	1	0.005
	CGT	False	False	1	3.912621	1	0	False	1	1	0.191
CneerA	CGT	False	False	20	2.094795	9	19	False	1	1	0.059
LnothA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
SarboA	AGA	False	False	33	2.062827	7	31	False	2	2	0.010
MingeA	AGA	False	False	37	2.005108	10	35	False	1	1	0.006
	CGT	False	False	15	2.149351	9	14	False	1	1	0.000

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LkluyE	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
KtherA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
KlactA	AGA	False	False	33	1.975837	7	30	False	2	2	0.011
LkluyD	AGA	False	False	34	1.984737	7	31	False	2	2	0.010
CzempA	AGA	False	False	21	1.824070	6	16	False	1	2	0.005
CmaltA	CGT	False	False	14	2.217593	9	13	False	1	1	0.000
NbaciA	AGA	False	False	31	2.034941	7	30	False	1	1	0.005
LdasiA	AGA	False	False	34	1.984737	7	31	False	2	2	0.010