

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	AGT	False	False	13	0.618943	6	12	False	1	1	0.000
	TCA	False	False	55	0.037647	7	51	False	1	1	0.000
	TCT	False	False	28	0.252295	5	25	False	2	2	0.000
SpombA	AGT	False	False	31	-0.249079	6	17	False	1	1	0.000
	TCT	False	False	29	-0.483871	6	17	False	1	1	0.000
SpastA	TCA	False	False	59	0.027110	7	51	False	1	1	0.000
	TCT	False	False	27	0.057133	6	23	False	1	1	0.000
AacerA	AGT	False	False	7	1.103356	2	5	False	1	1	0.023
	TCA	False	False	27	0.216281	2	23	False	1	1	0.009
	TCT	False	False	13	0.108108	2	9	False	1	1	0.013
CgigaA	AGC	False	False	46	0.048372	9	43	False	1	1	0.000
	AGT	False	False	23	0.138614	12	22	False	1	1	0.000
	TCA	False	False	39	-0.104063	10	34	False	1	1	0.000
CsantA	AGC	False	False	24	0.403743	10	16	False	3	3	0.000
	AGT	False	False	78	0.044756	11	54	False	4	4	0.000
	TCT	False	False	41	-0.209921	9	28	False	1	1	0.000
CtropB	TCA	False	False	60	-0.035475	11	57	False	1	1	0.000
CtropA	TCA	False	False	61	-0.035194	12	58	False	1	1	0.000
WpijpA	AGT	False	False	23	0.152829	4	19	False	1	1	0.000
	TCA	False	False	59	0.057791	6	51	False	2	2	0.000
	TCT	False	False	25	0.286920	6	21	False	2	2	0.000
BarxA	TCA	False	False	21	-0.227804	1	12	False	1	1	0.064
CoxycA	TCA	False	False	12	0.198027	1	9	False	1	1	0.037
	TCT	False	False	11	0.266942	1	8	False	1	1	0.040
CsubhA	TCA	False	False	43	0.216148	12	39	False	2	2	0.000
	TCG	False	False	22	0.129082	9	20	False	1	1	0.000
CsubhB	TCA	False	False	43	0.216148	12	39	False	2	2	0.000
	TCG	False	False	22	0.129082	9	20	False	1	1	0.000
OanguA	AGT	False	False	40	0.238670	10	28	False	3	4	0.000
	TCA	False	False	84	-0.123364	13	71	False	2	2	0.000
SarboA	TCA	False	False	57	0.084672	7	50	False	2	2	0.000
CalaiaA	AGT	False	False	54	-0.118935	10	47	False	1	1	0.000
	TCA	False	False	61	0.031650	11	55	False	2	2	0.000
KtherA	TCT	False	False	23	0.399421	5	19	False	2	3	0.000
CblacA	TCT	False	False	15	0.180000	1	14	False	1	1	0.061
CnorvA	AGT	False	False	48	0.273715	11	41	False	3	3	0.000
	TCA	False	False	96	0.133588	12	82	False	3	5	0.000
	TCT	False	False	26	0.225859	10	21	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CviswA	AGT	False	False	63	0.116729	13	57	False	2	2	0.000
	TCA	False	False	68	0.093610	13	65	False	1	2	0.000
MstarA	AGT	False	False	74	0.023095	12	56	False	2	2	0.000
	TCA	False	False	69	-0.135678	11	51	False	2	2	0.000
NbaciA	TCT	False	False	40	-0.372005	11	29	False	1	1	0.000
	TCA	False	False	50	0.044762	6	40	False	1	2	0.000
CalbiD	TCT	False	False	38	-0.011988	6	33	False	1	1	0.000
	TCA	False	False	55	0.061768	9	54	False	1	1	0.000
CbohiA	AGT	False	False	60	0.021012	11	52	False	2	2	0.000
	TCA	False	False	80	-0.011513	12	74	False	2	2	0.000
LstarA	AGT	False	False	49	0.021690	8	34	False	2	2	0.000
	TCT	False	False	53	-0.139327	7	37	False	2	2	0.000
MfariA	AGT	False	False	55	0.056145	11	51	False	1	1	0.000
	TCA	False	False	55	0.101046	9	52	False	2	2	0.000
MfariB	TCT	False	False	60	-0.065477	12	51	False	1	1	0.000
	TCA	False	False	79	-0.018576	12	72	False	2	2	0.000
CcastA	TCT	False	False	46	0.060079	11	43	False	1	1	0.000
	AGT	False	False	12	0.513578	5	11	False	1	1	0.000
LanspB	TCA	False	False	64	0.177599	6	58	False	2	2	0.000
	TCT	False	False	22	0.312311	5	16	False	3	3	0.000
CphanaA	AGT	False	False	14	0.525992	5	12	False	1	1	0.000
	TCA	False	False	61	0.028460	7	57	False	1	1	0.000
LanspA	TCT	False	False	23	0.322139	5	20	False	2	2	0.000
	TCA	False	False	85	-0.031813	12	63	False	3	3	0.000
CsojaA	TCT	False	False	34	-0.143957	10	28	False	1	1	0.000
	TCA	False	False	57	0.045569	6	55	False	1	1	0.000
CglabA	TCT	False	False	29	0.297599	6	25	False	2	2	0.000
	TCA	False	False	44	0.125773	7	41	False	1	1	0.000
ScryoA	AGT	False	False	12	0.503597	5	10	False	1	1	0.000
	TCT	False	False	27	0.379044	6	21	False	2	3	0.000
CneerA	AGT	False	False	31	-0.301437	6	14	False	1	1	0.000
	TCA	False	False	37	-0.125935	6	24	False	1	1	0.000
CsakeA	TCA	False	False	62	-0.147167	11	54	False	1	1	0.000
	AGT	False	False	50	0.015600	10	45	False	1	1	0.000
HuvarA	TCA	False	False	79	-0.005706	11	75	False	1	1	0.000
	TCT	False	False	41	0.087981	10	37	False	1	1	0.000
CjiufA	AGT	False	False	29	0.253517	4	22	False	2	3	0.000
	TCA	False	False	50	-0.400191	6	31	False	1	1	0.000
KlactA	TCT	False	False	62	0.042212	12	56	False	2	2	0.000
CzempA	TCA	False	False	69	0.100208	6	61	False	2	3	0.000
	AGT	False	False	38	0.299145	6	21	False	2	3	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CfrijA	TCA	False	False	66	-0.010486	12	56	False	1	2	0.000
CgallA	TCA	False	False	81	-0.095463	12	63	False	2	2	0.000
	TCT	False	False	39	0.075542	10	28	False	2	2	0.000
PpastA	AGT	False	False	58	0.345252	11	46	False	3	5	0.000
	TCA	False	False	113	-0.065966	13	91	False	3	4	0.000
SjapoA	AGT	False	False	26	0.336521	6	11	False	2	3	0.000
	TCT	False	False	61	-0.287787	7	38	False	2	2	0.000
CalbiE	TCA	False	False	39	0.080101	6	38	False	1	1	0.000
	AGC	False	False	6	0.566474	5	5	False	1	1	0.000
CtetrA	AGT	False	False	63	0.016586	11	57	False	1	1	0.000
	TCA	False	False	64	0.026245	12	61	False	1	1	0.000
	TCT	False	False	40	0.060506	11	37	False	1	1	0.000
CalbiC	TCA	False	False	59	0.051020	10	58	False	1	1	0.000
CalbiB	TCA	False	False	59	0.051020	10	58	False	1	1	0.000
CalbiA	TCA	False	False	54	0.063079	9	53	False	1	1	0.000
MmagnB	AGT	False	False	51	-0.007708	11	42	False	1	1	0.000
	TCA	False	False	52	-0.334677	11	34	False	1	1	0.000
MmagnA	AGT	False	False	53	0.008204	11	43	False	1	1	0.000
	TCA	False	False	52	-0.334677	11	34	False	1	1	0.000
CcoryA	TCA	False	False	60	0.081588	11	56	False	1	1	0.000
	TCT	False	False	46	-0.022426	11	43	False	1	1	0.000
CparaB	AGT	False	False	54	-0.023294	11	51	False	1	1	0.000
CparaD	AGT	False	False	54	-0.023294	11	51	False	1	1	0.000
MtetrA	AGT	False	False	33	-0.028618	10	26	False	1	1	0.000
	TCA	False	False	37	-0.250853	9	26	False	1	1	0.000
CsaraA	AGT	False	False	65	0.005008	12	59	False	1	1	0.000
	TCT	False	False	43	0.042564	9	41	False	1	1	0.000
	AGT	False	False	47	0.331986	10	38	False	4	4	0.000
CsuavA	TCA	False	False	85	0.018184	11	72	False	3	3	0.000
	TCC	False	False	2	1.576355	2	1	False	1	1	0.037
	TCT	False	False	39	0.089152	10	35	False	1	1	0.000
PkluyA	AGT	False	False	46	0.360041	11	33	False	3	5	0.000
	TCA	False	False	122	-0.043144	13	99	False	3	3	0.000
	AGT	False	False	51	0.400319	11	40	False	5	5	0.000
PcanaA	TCA	False	False	89	0.111611	12	78	False	4	4	0.000
	TCT	False	False	33	0.209150	8	27	False	2	2	0.000
YlipoB	TCA	False	False	75	-0.052824	12	57	False	2	2	0.000
	TCT	False	False	41	0.014412	9	31	False	2	2	0.000
YlipoA	TCA	False	False	75	-0.052824	12	57	False	2	2	0.000
	TCT	False	False	41	0.014412	9	31	False	2	2	0.000
CjadiA	AGT	False	False	47	0.361681	10	39	False	4	4	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
AgossB	TCA	False	False	101	0.106436	13	85	False	4	5	0.000
	AGT	False	False	16	0.915173	6	13	False	1	2	0.000
	TCA	False	False	57	0.228290	6	44	False	3	4	0.000
	TCT	False	False	32	0.054444	7	29	False	1	1	0.000
NdelpA	AGT	False	False	11	0.548303	5	9	False	1	1	0.000
	TCT	False	False	26	0.372790	5	18	False	2	3	0.000
LelonA	TCA	False	False	82	-0.081036	12	75	False	1	1	0.000
LkluyF	TCA	False	False	62	0.138265	6	58	False	2	2	0.000
	TCT	False	False	22	0.016052	5	19	False	1	1	0.000
CalimA	TCA	False	False	82	-0.018495	12	63	False	3	3	0.000
	TCT	False	False	37	-0.220539	9	26	False	1	1	0.000
CchauA	AGT	False	False	54	0.043972	12	53	False	1	1	0.000
	TCA	False	False	65	0.015392	11	59	False	1	1	0.000
BcaliA	AGT	False	False	47	0.387705	11	39	False	4	4	0.000
	TCA	False	False	88	0.088884	11	76	False	3	4	0.000
	TCC	False	False	1	3.436893	1	0	False	1	1	0.182
	TCT	False	False	35	0.031220	11	30	False	1	1	0.000
SoctoA	AGC	False	False	4	0.484412	3	0	False	1	1	0.000
	TCA	False	False	35	-0.101874	6	21	False	1	1	0.000
LkluyB	TCA	False	False	22	0.303845	1	20	False	1	1	0.061
CbuenA	AGT	False	False	67	0.065770	11	65	False	1	1	0.000
	TCT	False	False	49	0.009339	11	47	False	1	1	0.000
CdefoA	TCA	False	False	80	0.026835	12	62	False	3	3	0.000
	TCT	False	False	38	-0.167266	10	28	False	1	1	0.000
LdasiA	AGT	False	False	15	0.533844	5	12	False	1	1	0.000
	TCT	False	False	24	0.365834	5	19	False	2	3	0.000
CpseuA	TCT	False	False	14	0.327698	1	11	False	1	1	0.060
LkluyE	TCA	False	False	63	0.135503	6	59	False	2	2	0.000
	TCT	False	False	21	0.018619	5	18	False	1	1	0.000
LkluyD	TCA	False	False	60	0.217696	6	55	False	3	3	0.000
	TCT	False	False	23	0.037267	5	20	False	1	1	0.000
LkluyC	TCA	False	False	62	0.138265	6	58	False	2	2	0.000
	TCT	False	False	22	0.016052	5	19	False	1	1	0.000
CvartA	AGT	False	False	49	0.311106	11	40	False	4	4	0.000
	TCA	False	False	90	-0.041744	13	77	False	3	3	0.000
	TCT	False	False	36	0.311291	10	31	False	2	2	0.000
LkluyA	TCA	False	False	62	0.138265	6	58	False	2	2	0.000
	TCT	False	False	22	0.016052	5	19	False	1	1	0.000
CdublA	TCA	False	False	44	0.058463	10	43	False	1	1	0.000
	AGT	False	False	54	0.077413	11	48	False	1	1	0.000
ClusiA	TCA	False	False	72	0.011377	12	66	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MguilA	TCT	False	False	47	0.041314	11	46	False	1	1	0.000
	TCA	False	False	62	0.023418	11	60	False	1	1	0.000
	TCT	False	False	54	-0.028441	13	48	False	1	1	0.000
WmucoA	AGT	False	False	54	0.263809	12	43	False	4	4	0.000
	TCA	False	False	62	0.075844	10	52	False	3	3	0.000
ScereC	TCT	False	False	55	0.285538	12	49	False	2	3	0.000
	TCA	False	False	60	0.106555	7	53	False	2	2	0.000
ScereA	TCA	False	False	59	0.101031	7	53	False	2	2	0.000
ScereD	TCA	False	False	59	0.100210	7	53	False	2	2	0.000
ScereE	TCA	False	False	59	0.100210	7	53	False	2	2	0.000
DhansA	AGT	False	False	61	-0.041339	11	54	False	1	1	0.000
	TCA	False	False	74	0.010932	12	69	False	2	2	0.000
	TCT	False	False	41	0.049740	10	38	False	1	1	0.000
CsalmaA	AGT	False	False	79	-0.020374	11	54	False	4	5	0.000
	TCT	False	False	43	-0.097272	9	29	False	1	1	0.000
LnothA	AGT	False	False	15	0.513496	5	13	False	1	1	0.000
	TCA	False	False	61	0.087914	7	56	False	1	2	0.000
BcustA	TCT	False	False	23	0.166667	5	21	False	1	1	0.000
	AGT	False	False	47	0.613246	11	32	False	5	7	0.000
	TCA	False	False	97	-0.049834	12	76	False	3	3	0.000
AgossaA	AGT	False	False	16	0.915173	6	13	False	1	2	0.000
	TCA	False	False	56	0.228703	6	43	False	3	4	0.000
	TCT	False	False	33	0.059058	7	30	False	1	1	0.000
MingeA	AGT	False	False	73	0.144794	12	56	False	3	3	0.000
	TCA	False	False	73	-0.306912	12	53	False	1	1	0.000
ScastA	AGT	False	False	17	-0.058324	6	13	False	1	1	0.000
	TCA	False	False	60	0.132010	7	51	False	2	2	0.000
	TCT	False	False	23	0.076571	5	19	False	1	1	0.000
CmaltA	AGT	False	False	71	0.000635	11	65	False	2	2	0.000
KmarxA	TCA	False	False	24	0.211145	1	22	False	1	1	0.061