

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	AGC	False	False	3	1.377049	2	2	False	1	1	0.001
	AGT	False	False	31	0.937039	6	17	False	3	4	0.000
SpombA	TCA	False	False	41	0.480399	6	22	False	1	1	0.000
	TCT	False	False	29	1.082142	6	17	False	4	5	0.000
SpastA	TCA	False	False	59	0.880910	7	51	False	1	1	0.000
AacerA	TCT	False	False	13	0.889047	2	9	False	1	1	0.071
	AGT	False	False	51	1.081477	11	49	False	2	2	0.000
CmetaD	TCA	False	False	60	1.014524	11	59	False	1	1	0.000
	TCT	False	False	58	1.003062	10	54	False	1	1	0.000
CgigaA	AGC	False	False	46	0.892241	9	43	False	1	1	0.000
	AGC	False	False	24	0.842692	10	16	False	2	3	0.000
CsantA	AGT	False	False	78	0.582565	11	54	False	1	1	0.000
	TCT	False	False	41	0.762209	9	28	False	2	2	0.000
CtropB	AGC	False	False	24	1.065014	9	22	False	1	1	0.000
	AGT	False	False	40	0.988372	10	38	False	1	1	0.000
CtropA	AGT	False	False	40	1.094477	10	37	False	1	2	0.000
	AGT	False	False	23	0.875671	4	19	False	1	1	0.000
WpijpA	TCA	False	False	59	0.965712	6	51	False	1	3	0.000
	TCT	False	False	25	0.893748	6	21	False	1	1	0.000
BarxA	AGT	False	False	13	1.046919	1	11	False	1	1	0.036
	TCA	False	False	21	0.978779	1	12	False	1	3	0.044
CoxycA	AGT	False	False	14	1.156697	1	12	False	1	2	0.046
	AGC	False	False	62	0.841949	11	55	False	1	1	0.000
CsubhA	AGT	False	False	11	1.017154	6	9	False	1	1	0.000
	TCG	False	False	22	0.986735	9	20	False	1	1	0.000
	AGC	False	False	62	0.841949	11	55	False	1	1	0.000
CsubhB	AGT	False	False	11	1.017154	6	9	False	1	1	0.000
	TCG	False	False	22	0.986735	9	20	False	1	1	0.000
OanguA	AGT	False	False	40	0.678046	10	28	False	2	2	0.000
	TCA	False	False	84	0.964559	13	71	False	3	4	0.000
SarboA	TCA	False	False	57	0.848394	7	50	False	1	1	0.000
CcastA	TCA	False	False	64	0.885170	6	58	False	1	1	0.000
KtherA	AGC	False	False	1	2.610000	1	0	False	1	1	0.183
	TCA	False	False	62	0.979234	7	58	False	1	1	0.001
CnorvA	TCA	False	False	96	0.925909	12	82	False	2	4	0.000
CviswA	AGT	False	False	63	0.835216	13	57	False	1	1	0.000
	AGT	False	False	74	0.717906	12	56	False	2	2	0.000
MstarA	TCA	False	False	69	0.695980	11	51	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
NbaciA	TCT	False	False	40	0.745895	11	29	False	1	1	0.000
	AGT	False	False	13	0.792160	5	11	False	1	1	0.000
	TCA	False	False	50	0.855048	6	40	False	1	1	0.000
CalbiD	AGT	False	False	58	0.941906	11	55	False	1	1	0.000
CbohiA	AGT	False	False	60	0.849443	11	52	False	1	1	0.000
	TCA	False	False	80	0.971354	12	74	False	2	2	0.000
LstarA	AGT	False	False	49	0.725712	8	34	False	1	2	0.000
	TCA	False	False	27	0.802103	8	22	False	1	1	0.000
MfariA	TCT	False	False	53	0.833404	7	37	False	3	3	0.000
	AGT	False	False	55	0.947734	11	51	False	1	1	0.000
MfariB	TCT	False	False	60	0.976090	12	51	False	2	3	0.000
	AGT	False	False	51	0.891593	11	46	False	1	1	0.000
	TCA	False	False	79	0.936678	12	72	False	1	1	0.000
CalaiA	AGC	True	True	1	2.981818	1	0	False	1	1	0.574
	AGT	False	False	54	0.953335	10	47	False	1	3	0.284
	TCC	False	False	1	4.714286	1	0	False	1	1	0.500
LanspB	TCT	False	False	40	0.911153	8	37	False	1	1	0.284
	AGT	False	False	14	0.784542	5	12	False	1	1	0.000
CphanA	TCA	False	False	85	0.838449	12	63	False	5	5	0.000
CsojaA	AGT	False	False	42	0.963849	7	39	False	1	1	0.000
CglabA	TCA	False	False	60	0.919892	7	54	False	1	1	0.000
	AGC	False	False	6	1.652104	3	3	False	2	2	0.004
ScryoA	AGT	False	False	31	0.446041	6	14	False	1	1	0.000
	TCA	False	False	37	0.687226	6	24	False	1	2	0.000
	TCC	False	False	3	0.863057	3	0	False	1	1	0.009
CneerA	TCT	False	False	29	0.926243	6	20	False	2	2	0.000
	AGT	False	False	63	0.970648	12	61	False	1	1	0.000
	TCA	False	False	62	1.060854	11	54	False	3	4	0.000
CsakeA	TCT	False	False	43	1.025093	12	39	False	1	1	0.000
	AGT	False	False	50	0.908433	10	45	False	1	1	0.000
	TCA	False	False	79	0.993320	11	75	False	1	1	0.000
HuvarA	TCT	False	False	41	0.872742	10	37	False	1	1	0.000
	AGT	False	False	29	0.873405	4	22	False	1	2	0.000
	TCA	False	False	50	0.421203	6	31	False	1	1	0.000
CjiufA	TCT	False	False	26	0.885609	6	16	False	3	3	0.000
	AGT	False	False	50	1.068785	10	47	False	2	2	0.000
	TCA	False	False	52	1.062020	10	49	False	2	2	0.000
CzempA	TCA	False	False	38	1.244134	6	20	False	4	8	0.000
	TCT	False	False	34	0.673658	6	20	False	2	3	0.000
CmetaB	AGT	False	False	51	1.081477	11	49	False	2	2	0.000
	TCA	False	False	60	1.014524	11	59	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaC	TCT	False	False	58	1.003062	10	54	False	1	1	0.000
	AGT	False	False	51	1.081477	11	49	False	2	2	0.000
	TCA	False	False	60	1.014524	11	59	False	1	1	0.000
CmetaA	TCT	False	False	58	1.003062	10	54	False	1	1	0.000
	AGT	False	False	51	1.081477	11	49	False	2	2	0.000
	TCA	False	False	60	1.014524	11	59	False	1	1	0.000
CmetaF	TCT	False	False	58	1.003062	10	54	False	1	1	0.000
	AGT	False	False	51	1.081477	11	49	False	2	2	0.000
	TCA	False	False	60	1.014524	11	59	False	1	1	0.000
CfrijA	TCT	False	False	58	1.003062	10	54	False	1	1	0.000
	AGT	False	False	65	0.963353	12	62	False	1	1	0.000
	TCA	False	False	66	0.984998	12	56	False	3	3	0.000
CgallA	TCT	False	False	43	0.973534	13	40	False	1	1	0.000
	AGT	False	False	42	0.860504	12	35	False	1	1	0.000
	TCA	False	False	81	0.843037	12	63	False	4	4	0.000
CmetaE	TCC	False	False	1	4.218182	1	0	False	1	1	0.139
	TCT	False	False	39	0.689282	10	28	False	1	1	0.000
	AGT	False	False	51	1.081477	11	49	False	2	2	0.000
PpastA	TCA	False	False	60	1.014524	11	59	False	1	1	0.000
	TCT	False	False	58	1.003062	10	54	False	1	1	0.000
	AGT	False	False	58	0.664168	11	46	False	1	1	0.000
SjapoA	TCA	False	False	113	0.935577	13	91	False	3	5	0.000
	TCT	False	False	17	1.086231	9	15	False	1	1	0.000
	AGT	False	False	26	0.321074	6	11	False	1	1	0.000
CalbiE	TCT	False	False	61	0.579473	7	38	False	1	3	0.000
Ctetra	AGT	False	False	31	0.939305	6	28	False	1	1	0.000
CalbiC	AGT	False	False	63	0.894898	11	57	False	1	1	0.000
CalbiB	AGT	False	False	62	0.947697	12	59	False	1	1	0.000
CalbiA	AGT	False	False	62	0.947697	12	59	False	1	1	0.000
MmagnB	AGT	False	False	59	0.942707	11	56	False	1	1	0.000
	AGC	False	False	22	0.615721	10	15	False	1	1	0.000
	AGT	False	False	51	0.847390	11	42	False	1	1	0.000
MmagnA	TCA	False	False	52	0.770294	11	34	False	2	3	0.000
	AGT	False	False	53	0.884098	11	43	False	2	2	0.000
	TCA	False	False	52	0.770294	11	34	False	2	3	0.000
CparaA	AGT	False	False	43	1.063789	10	42	False	1	1	0.000
	TCA	False	False	35	1.162080	9	33	False	2	2	0.000
CcoryA	AGT	False	False	59	1.068449	12	56	False	2	2	0.000
CparaC	AGT	False	False	53	1.075910	11	51	False	2	2	0.000
	TCA	False	False	52	1.035121	10	48	False	2	2	0.000
CparaB	AGT	False	False	54	1.039373	11	51	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CparaE	TCA	False	False	52	1.035121	10	48	False	2	2	0.000
	AGT	False	False	53	1.075910	11	51	False	2	2	0.000
CparaD	TCA	False	False	52	1.035121	10	48	False	2	2	0.000
	AGT	False	False	54	1.039373	11	51	False	2	2	0.000
CparaF	TCA	False	False	52	1.035121	10	48	False	2	2	0.000
	AGT	False	False	53	1.075910	11	51	False	2	2	0.000
MtetrA	TCA	False	False	52	1.035121	10	48	False	2	2	0.000
	AGC	False	False	39	0.714038	12	31	False	1	1	0.000
	AGT	False	False	33	0.810526	10	26	False	1	1	0.000
CsaraA	TCC	False	False	19	0.962633	9	14	False	1	1	0.000
	TCG	False	False	17	1.203730	10	15	False	1	1	0.000
	AGT	False	False	65	0.879814	12	59	False	1	1	0.000
CsuavA	AGT	False	False	47	0.862557	10	38	False	2	2	0.000
	TCA	False	False	85	0.906868	11	72	False	2	3	0.000
PkluyA	TCT	False	False	39	0.954703	10	35	False	1	1	0.000
	AGT	False	False	46	0.494864	11	33	False	1	1	0.000
	TCA	False	False	122	0.864035	13	99	False	2	3	0.000
PcanaA	AGT	False	False	51	0.779221	11	40	False	2	2	0.000
	TCA	False	False	89	0.908057	12	78	False	1	3	0.000
	TCT	False	False	33	0.859649	8	27	False	1	1	0.000
YlipoB	AGT	False	False	43	0.824346	12	35	False	1	1	0.000
	TCA	False	False	75	0.923012	12	57	False	4	5	0.000
	TCT	False	False	41	0.658259	9	31	False	1	1	0.000
YlipoA	AGT	False	False	43	0.824346	12	35	False	1	1	0.000
	TCA	False	False	75	0.923012	12	57	False	4	5	0.000
	TCT	False	False	41	0.658259	9	31	False	1	1	0.000
CjadiA	AGT	False	False	47	0.797899	10	39	False	1	1	0.000
	TCA	False	False	101	0.864469	13	85	False	2	3	0.000
	TCT	False	False	26	1.027682	10	23	False	1	1	0.000
AgossB	TCA	False	False	57	0.698397	6	44	False	1	1	0.000
NdelpA	TCA	False	False	63	0.935794	7	58	False	1	1	0.000
	AGT	False	False	54	0.998134	10	51	False	2	2	0.000
LelonA	TCA	False	False	82	1.084660	12	75	False	3	4	0.000
	TCT	False	False	37	1.022975	10	35	False	1	1	0.000
LkluyF	AGT	False	False	15	0.788732	5	12	False	1	1	0.000
	AGT	False	False	45	0.858466	12	38	False	1	1	0.000
CalimA	TCA	False	False	82	0.901061	12	63	False	5	5	0.000
	TCT	False	False	37	0.595274	9	26	False	1	1	0.000
CchauA	AGC	False	False	9	1.218506	5	8	False	1	1	0.000
BcaliA	TCA	False	False	88	0.837686	11	76	False	1	2	0.000
	TCT	False	False	35	1.091057	11	30	False	2	2	0.000

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SoctoA	AGT	False	False	32	0.748884	6	15	False	3	3	0.000
	TCA	False	False	35	0.740288	6	21	False	2	3	0.000
	TCT	False	False	35	0.945145	5	25	False	2	2	0.000
CbuenA	AGT	False	False	67	0.934903	11	65	False	1	1	0.000
	TCA	False	False	47	1.033986	10	41	False	2	2	0.000
	TCT	False	False	49	1.016173	11	47	False	1	1	0.000
CdefoA	TCA	False	False	80	0.967245	12	62	False	5	6	0.000
LdasiA	AGT	False	False	15	0.731801	5	12	False	1	1	0.000
CpseuA	AGT	False	False	9	1.130567	1	8	False	1	1	0.070
LkluyE	AGT	False	False	15	0.788732	5	12	False	1	1	0.000
LkluyD	AGT	False	False	15	0.788732	5	12	False	1	1	0.000
LkluyC	AGT	False	False	15	0.788732	5	12	False	1	1	0.000
CvartA	AGT	False	False	49	0.871432	11	40	False	2	2	0.000
	TCA	False	False	90	1.021628	13	77	False	3	5	0.000
LkluyA	AGT	False	False	15	0.788732	5	12	False	1	1	0.000
CdublA	AGT	False	False	58	0.949814	11	55	False	1	1	0.000
CorthB	AGT	False	False	52	1.078402	11	50	False	2	2	0.000
	TCA	False	False	49	1.087140	10	47	False	2	2	0.000
CorthC	AGT	False	False	51	1.080508	11	49	False	2	2	0.000
	TCA	False	False	49	1.085733	10	47	False	2	2	0.000
CorthA	AGT	False	False	52	1.078402	11	50	False	2	2	0.000
	TCA	False	False	48	1.089829	10	46	False	2	2	0.000
ClusiA	TCA	False	False	72	0.982635	12	66	False	1	2	0.000
MguilA	AGT	False	False	51	0.921388	11	47	False	1	1	0.000
	TCT	False	False	54	1.066224	13	48	False	2	3	0.000
WmucoA	AGT	False	False	54	0.819791	12	43	False	2	2	0.000
	TCA	False	False	62	0.955563	10	52	False	1	3	0.000
	TCT	False	False	55	0.862564	12	49	False	1	1	0.000
ScereC	TCA	False	False	60	0.905795	7	53	False	1	1	0.000
ScereA	TCA	False	False	59	0.925556	7	53	False	1	1	0.000
ScereD	TCA	False	False	59	0.924802	7	53	False	1	1	0.000
ScereE	TCA	False	False	59	0.924802	7	53	False	1	1	0.000
DhansA	AGT	False	False	61	0.963135	11	54	False	2	2	0.000
	TCA	False	False	74	0.957710	12	69	False	2	2	0.000
	AGT	False	False	79	0.779514	11	54	False	5	6	0.000
CsalmaA	TCA	False	False	36	0.893516	8	27	False	2	2	0.000
	TCT	False	False	43	0.776377	9	29	False	1	1	0.000
LnothA	AGT	False	False	15	0.771851	5	13	False	1	1	0.000
BcustA	AGT	False	False	47	0.416582	11	32	False	1	1	0.000
	TCA	False	False	97	0.880688	12	76	False	2	4	0.000
CtheaA	AGT	False	False	51	1.081258	11	49	False	2	2	0.000

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AgossaA	TCA	False	False	63	1.003153	12	61	False	1	1	0.000
	TCT	False	False	55	1.027927	12	53	False	1	1	0.000
	TCA	False	False	56	0.707533	6	43	False	1	1	0.000
	AGT	False	False	73	0.631912	12	56	False	1	1	0.000
MingeA	TCA	False	False	73	0.712511	12	53	False	2	2	0.000
	TCT	False	False	36	0.792308	9	26	False	1	1	0.000
ScastA	TCA	False	False	60	0.836855	7	51	False	1	1	0.000
CmaltA	AGC	False	False	12	1.172117	6	10	False	1	1	0.000
	AGT	False	False	71	0.937143	11	65	False	2	2	0.000