

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	TCA	False	False	55	1.007287	7	51	False	2	2	0.000
	AGC	False	False	3	-0.015873	3	0	False	1	1	0.007
SpombA	AGT	False	False	31	0.939189	6	17	False	3	4	0.000
	TCA	False	False	41	1.118998	6	22	False	3	8	0.002
	TCT	False	False	29	0.700698	6	17	False	2	2	0.000
SpastA	AGT	False	False	12	0.876997	5	10	False	1	1	0.000
	TCA	False	False	59	1.057931	7	51	False	3	4	0.000
AacerA	TCT	False	False	27	1.092619	6	23	False	1	2	0.000
	TCA	False	False	27	1.303815	2	23	False	1	3	0.011
CmetaD	TCT	False	False	13	1.020626	2	9	False	1	1	0.013
	TCT	False	False	58	0.924608	10	54	False	1	1	0.000
CgigaA	TCA	False	False	39	1.061861	10	34	False	2	2	0.000
	AGC	False	False	24	0.757576	10	16	False	1	1	0.000
CsantA	AGT	False	False	78	1.056604	11	54	False	6	11	0.000
	TCA	False	False	31	1.250424	8	23	False	3	5	0.000
	TCT	False	False	41	1.161595	9	28	False	4	6	0.000
CtropB	AGT	False	False	40	1.043605	10	38	False	1	1	0.000
	TCA	False	False	60	1.075362	11	57	False	1	1	0.000
	TCC	False	False	8	1.188119	6	7	False	1	1	0.000
CtropA	AGT	False	False	40	1.006105	10	37	False	1	1	0.000
	TCA	False	False	61	1.096741	12	58	False	1	1	0.000
	TCC	False	False	8	1.188119	6	7	False	1	1	0.000
WpijpA	AGT	False	False	23	1.126394	4	19	False	2	2	0.000
	TCA	False	False	59	0.988731	6	51	False	2	2	0.000
	TCT	False	False	25	0.995781	6	21	False	1	1	0.000
BarxA	AGT	False	False	13	1.108543	1	11	False	1	1	0.036
	TCA	False	False	21	1.034647	1	12	False	1	3	0.044
CoxycA	TCT	False	False	11	1.079339	1	8	False	1	1	0.064
	AGC	False	False	62	0.849109	11	55	False	1	1	0.051
CsubhA	TCA	False	False	43	0.936978	12	39	False	1	1	0.056
	TCC	False	False	30	1.096404	11	27	False	1	2	0.056
	AGC	False	False	62	0.849109	11	55	False	1	1	0.051
CsubhB	TCA	False	False	43	0.936978	12	39	False	1	1	0.056
	TCC	False	False	30	1.096404	11	27	False	1	2	0.056
	AGT	False	False	40	0.755150	10	28	False	2	2	0.000
OanguA	TCA	False	False	84	0.945172	13	71	False	2	2	0.000
	TCT	False	False	46	1.083421	11	42	False	1	2	0.000
SarboA	TCA	False	False	57	0.991466	7	50	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CalaiA	TCT	False	False	33	1.167779	5	25	False	3	5	0.000
	AGT	False	False	54	0.946934	10	47	False	1	1	0.051
	TCA	False	False	61	0.946307	11	55	False	1	1	0.051
KtherA	AGT	False	False	14	1.064164	5	13	False	1	1	0.056
	TCA	False	False	62	1.005076	7	58	False	2	2	0.051
CblacA	AGT	False	False	12	1.006829	1	10	False	1	1	0.061
CnorvA	AGT	False	False	48	1.145788	11	41	False	4	4	0.000
	TCA	False	False	96	0.981365	12	82	False	3	4	0.000
CviswA	TCT	False	False	26	1.072422	10	21	False	2	2	0.000
	AGC	False	False	6	1.484848	4	5	False	1	1	0.000
	AGT	False	False	63	1.039627	13	57	False	2	2	0.000
MstarA	TCT	False	False	35	1.172550	11	32	False	1	2	0.000
	AGT	False	False	74	1.059538	12	56	False	5	8	0.000
	TCA	False	False	69	1.056062	11	51	False	6	9	0.000
NbaciA	TCT	False	False	40	0.941857	11	29	False	2	3	0.000
	TCA	False	False	50	1.088381	6	40	False	2	5	0.000
CbohiA	AGT	False	False	60	0.880687	11	52	False	1	1	0.055
	TCT	False	False	34	1.039080	9	32	False	1	1	0.060
LstarA	AGC	False	False	6	0.790660	4	4	False	1	1	0.000
	AGT	False	False	49	0.857659	8	34	False	3	5	0.000
	TCA	False	False	27	0.914007	8	22	False	1	1	0.000
MfariA	TCT	False	False	53	1.109715	7	37	False	4	6	0.000
	AGT	False	False	55	0.977338	11	51	False	1	1	0.000
	TCA	False	False	55	1.077166	9	52	False	1	1	0.000
MfariB	TCT	False	False	60	0.918153	12	51	False	2	2	0.000
	AGT	False	False	51	1.125664	11	46	False	3	3	0.000
	TCA	False	False	79	1.047186	12	72	False	2	3	0.000
CcastA	TCT	False	False	46	1.077706	11	43	False	1	1	0.000
	TCA	False	False	64	1.031236	6	58	False	2	3	0.000
	TCT	False	False	22	0.893737	5	16	False	1	1	0.000
LanspB	TCA	False	False	61	1.015637	7	57	False	2	2	0.000
	AGT	False	False	45	1.007916	12	38	False	3	4	0.000
CphanA	TCA	False	False	85	1.027091	12	63	False	6	8	0.000
	TCT	False	False	34	0.792434	10	28	False	1	1	0.000
LanspA	TCA	False	False	57	1.013235	6	55	False	1	1	0.051
	TCT	False	False	29	1.049655	6	25	False	1	1	0.052
CsojaA	AGT	False	False	42	1.028181	7	39	False	1	1	0.000
	TCA	False	False	44	1.098669	7	41	False	1	1	0.000
CglabA	TCA	False	False	60	1.065775	7	54	False	3	3	0.000
	TCT	False	False	27	1.113924	6	21	False	2	2	0.000
ScryoA	AGT	False	False	31	0.967594	6	14	False	4	7	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CneerA	TCA	False	False	37	0.990452	6	24	False	4	5	0.000
	TCC	False	False	3	0.898089	3	0	False	1	1	0.007
	TCT	False	False	29	1.047744	6	20	False	2	3	0.000
	AGT	False	False	63	1.061273	12	61	False	1	1	0.000
	TCT	False	False	43	1.216149	12	39	False	2	3	0.000
CsakeA	AGT	False	False	50	1.142211	10	45	False	3	3	0.000
	TCA	False	False	79	0.980935	11	75	False	1	1	0.000
HuvarA	TCT	False	False	41	1.102121	10	37	False	1	1	0.000
	TCA	False	False	50	0.841070	6	31	False	3	6	0.000
CjiufA	TCT	False	False	26	0.759410	6	16	False	2	2	0.000
	TCT	False	False	62	0.966267	12	56	False	2	2	0.051
KlactA	AGT	False	False	13	1.360267	5	11	False	1	2	0.000
	TCA	False	False	69	0.981592	6	61	False	2	3	0.000
	TCT	False	False	18	1.313386	4	15	False	2	3	0.000
CzempA	AGT	False	False	38	1.083459	6	21	False	4	9	0.001
	TCA	False	False	38	0.655267	6	20	False	1	3	0.000
	TCT	False	False	34	0.656879	6	20	False	2	2	0.000
CmetaB	TCT	False	False	58	0.924608	10	54	False	1	1	0.000
CmetaC	TCT	False	False	58	0.924608	10	54	False	1	1	0.000
CmetaA	TCT	False	False	58	0.924608	10	54	False	1	1	0.000
CmetaF	TCT	False	False	58	0.924608	10	54	False	1	1	0.000
CfrijA	AGT	False	False	65	1.027003	12	62	False	1	1	0.000
	TCA	False	False	66	1.096467	12	56	False	2	4	0.000
	TCT	False	False	43	1.007265	13	40	False	1	1	0.000
CgallA	AGT	False	False	42	1.097759	12	35	False	3	4	0.000
	TCA	False	False	81	0.974561	12	63	False	2	5	0.000
	TCT	False	False	39	1.157355	10	28	False	6	6	0.000
CmetaE	TCT	False	False	58	0.924608	10	54	False	1	1	0.000
PpastA	AGT	False	False	58	0.915994	11	46	False	3	3	0.000
	TCA	False	False	113	1.064037	13	91	False	5	8	0.000
	AGC	False	False	4	1.000000	3	2	False	1	1	0.000
SjapoA	AGT	False	False	26	1.319603	6	11	False	5	8	0.000
	TCA	False	False	10	1.423041	4	9	False	1	1	0.000
	TCC	False	False	3	0.532051	3	0	False	1	1	0.010
Ctetra	TCT	False	False	61	0.909998	7	38	False	4	7	0.000
	AGT	False	False	63	1.013521	11	57	False	1	2	0.000
	TCA	False	False	64	1.044132	12	61	False	1	1	0.000
	TCT	False	False	40	1.015677	11	37	False	1	1	0.000
	AGC	False	False	22	0.673459	10	15	False	2	2	0.000
MmagnB	AGT	False	False	51	1.257652	11	42	False	4	6	0.000
	TCA	False	False	52	0.979850	11	34	False	4	6	0.000

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MmagnA	TCT	False	False	46	1.141383	11	40	False	3	4	0.000
	AGC	False	False	20	0.717532	10	14	False	2	2	0.000
	AGT	False	False	53	1.214346	11	43	False	4	6	0.000
	TCA	False	False	52	0.979850	11	34	False	4	6	0.000
CcoryA	TCT	False	False	46	1.141383	11	40	False	3	4	0.000
	AGT	False	False	59	1.079826	12	56	False	1	1	0.055
	TCG	False	False	1	2.238532	1	0	False	1	1	0.389
	TCT	False	False	46	0.966241	11	43	False	1	1	0.056
MtetrA	AGC	False	False	39	1.019353	12	31	False	3	4	0.000
	AGT	False	False	33	1.232566	10	26	False	3	4	0.000
	TCA	False	False	37	1.298609	9	26	False	3	6	0.000
	TCC	False	False	19	0.705813	9	14	False	1	1	0.000
CsaraA	TCG	False	False	17	1.034433	10	15	False	1	1	0.000
	TCT	False	False	25	0.985668	9	21	False	2	2	0.000
	AGT	False	False	65	1.083060	12	59	False	2	3	0.000
	TCA	False	False	56	1.190549	12	53	False	2	2	0.000
CsuavA	AGT	False	False	47	1.018949	10	38	False	3	3	0.000
	TCA	False	False	85	1.075316	11	72	False	4	5	0.000
PkluyA	TCT	False	False	39	1.075591	10	35	False	2	2	0.000
	AGT	False	False	46	0.813559	11	33	False	2	3	0.000
	TCA	False	False	122	1.054869	13	99	False	3	9	0.000
PcanaA	AGT	False	False	51	1.128047	11	40	False	4	4	0.000
	TCA	False	False	89	0.972867	12	78	False	2	2	0.000
	TCT	False	False	33	0.952528	8	27	False	1	2	0.000
YlipoB	AGT	False	False	43	1.048747	12	35	False	3	4	0.000
	TCA	False	False	75	1.034560	12	57	False	4	6	0.000
	TCT	False	False	41	0.966186	9	31	False	4	4	0.000
YlipoA	AGT	False	False	43	1.048747	12	35	False	3	4	0.000
	TCA	False	False	75	1.034560	12	57	False	4	6	0.000
	TCT	False	False	41	0.966186	9	31	False	4	4	0.000
CjadiA	AGT	False	False	47	1.035268	10	39	False	3	3	0.000
	TCA	False	False	101	1.078702	13	85	False	4	6	0.000
	TCT	False	False	26	0.941609	10	23	False	1	1	0.000
AgossB	TCA	False	False	57	1.170341	6	44	False	3	6	0.000
	TCT	False	False	32	1.074456	7	29	False	1	1	0.000
	TCA	False	False	63	1.104316	7	58	False	3	3	0.000
NdelpA	TCT	False	False	26	1.118369	5	18	False	2	3	0.000
	TCA	False	False	82	0.956375	12	75	False	1	1	0.000
LkluyF	AGT	False	False	15	0.839949	5	12	False	1	1	0.000
	TCA	False	False	62	1.026146	6	58	False	2	2	0.000
CalimA	AGT	False	False	45	1.062790	12	38	False	3	4	0.000

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LkluyD	TCA	False	False	82	1.012545	12	63	False	5	6	0.000
	TCT	False	False	37	0.895789	9	26	False	4	4	0.000
	AGT	False	False	15	0.839949	5	12	False	1	1	0.000
	TCA	False	False	60	1.025780	6	55	False	2	2	0.000
	AGT	False	False	47	1.148240	11	39	False	4	4	0.000
BcaliA	TCA	False	False	88	0.948011	11	76	False	2	3	0.000
	TCT	False	False	35	1.085203	11	30	False	2	2	0.000
SoctoA	AGT	False	False	32	0.872954	6	15	False	4	6	0.000
	TCA	False	False	35	0.991578	6	21	False	3	5	0.000
	TCT	False	False	35	1.107241	5	25	False	3	4	0.000
LkluyB	TCA	False	False	22	1.059529	1	20	False	1	1	0.061
CbuenA	TCA	False	False	47	0.981319	10	41	False	2	2	0.000
	AGT	False	False	44	1.063567	12	37	False	3	4	0.000
CdefoA	TCA	False	False	80	0.989740	12	62	False	3	5	0.000
	TCT	False	False	38	1.057554	10	28	False	5	5	0.000
LdasiA	AGT	False	False	15	0.961047	5	12	False	1	1	0.000
	TCA	False	False	61	1.045426	7	58	False	2	2	0.000
	TCT	False	False	24	0.969877	5	19	False	1	1	0.000
CpseuA	TCA	False	False	13	1.143457	1	11	False	1	1	0.095
	TCT	False	False	14	1.184655	1	11	False	1	1	0.094
LkluyE	AGT	False	False	15	0.839949	5	12	False	1	1	0.000
	TCA	False	False	63	1.023770	6	59	False	2	2	0.000
CchauA	TCA	False	False	65	0.956637	11	59	False	2	2	0.051
	TCT	False	False	40	1.007127	10	38	False	1	1	0.056
LkluyC	AGT	False	False	15	0.839949	5	12	False	1	1	0.000
	TCA	False	False	62	1.026146	6	58	False	2	2	0.000
CvartA	AGT	False	False	49	1.029743	11	40	False	3	3	0.000
	TCA	False	False	90	0.978140	13	77	False	3	3	0.000
	TCT	False	False	36	1.176544	10	31	False	3	3	0.000
LkluyA	AGT	False	False	15	0.839949	5	12	False	1	1	0.056
	TCA	False	False	62	1.026146	6	58	False	2	2	0.051
ClusiA	AGT	False	False	54	1.068378	11	48	False	3	3	0.000
	TCA	False	False	72	1.033383	12	66	False	2	2	0.000
MguilA	AGT	False	False	51	0.975626	11	47	False	1	1	0.000
	TCA	False	False	62	1.066585	11	60	False	1	1	0.000
	TCT	False	False	54	0.981109	13	48	False	1	1	0.000
WmucoA	AGT	False	False	54	1.061207	12	43	False	4	4	0.000
	TCA	False	False	62	1.033077	10	52	False	3	3	0.000
	TCT	False	False	55	0.970872	12	49	False	2	2	0.000
ScereC	TCA	False	False	60	1.031824	7	53	False	2	3	0.000
	TCT	False	False	28	1.013694	5	23	False	2	2	0.000

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ScereA	TCA	False	False	59	0.993232	7	53	False	2	2	0.000
	TCT	False	False	28	1.013694	5	23	False	2	2	0.000
ScereD	TCA	False	False	59	0.992577	7	53	False	2	2	0.000
	TCT	False	False	27	1.087912	5	21	False	2	3	0.000
ScereE	TCA	False	False	59	0.992577	7	53	False	2	2	0.000
	TCT	False	False	28	1.083818	5	22	False	2	3	0.000
DhansA	AGT	False	False	61	0.973157	11	54	False	2	2	0.000
	TCT	False	False	41	1.029786	10	38	False	1	1	0.000
CsalmaA	AGT	False	False	79	0.890874	11	54	False	5	7	0.000
	TCA	False	False	36	1.039687	8	27	False	3	4	0.000
	TCC	False	False	1	3.957746	1	0	False	1	1	0.136
	TCT	False	False	43	1.276891	9	29	False	4	8	0.000
LnothA	TCA	False	False	61	1.007495	7	56	False	2	2	0.051
	AGT	False	False	47	0.781345	11	32	False	2	3	0.000
BcustA	TCA	False	False	97	1.130411	12	76	False	4	9	0.000
	TCT	False	False	27	1.151370	9	22	False	3	3	0.000
AgossaA	TCA	False	False	56	1.140622	6	43	False	3	6	0.000
	TCT	False	False	33	1.127247	7	30	False	1	1	0.000
MingeA	AGT	False	False	73	1.044900	12	56	False	5	7	0.000
	TCA	False	False	73	1.068123	12	53	False	4	9	0.000
	TCT	False	False	36	1.044551	9	26	False	4	5	0.000
	AGT	False	False	17	0.966025	6	13	False	1	1	0.000
ScastA	TCA	False	False	60	1.117712	7	51	False	4	4	0.000
	TCT	False	False	23	1.215980	5	19	False	3	3	0.000
KmarxA	TCT	False	False	6	1.457576	1	5	False	1	1	0.075