

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	TCT	False	False	28	0.917035	5	25	False	1	1	0.000
SpombA	AGT	False	False	31	0.225430	6	17	False	1	1	0.000
	TCA	False	False	41	0.120158	6	22	False	3	3	0.000
AacerA	AGT	False	False	7	1.422819	2	5	False	1	1	0.155
CgallA	AGT	False	False	42	0.689916	12	35	False	1	1	0.000
CgigaA	TCT	False	False	18	1.129714	7	16	False	1	2	0.000
	AGT	False	False	78	0.381308	11	54	False	1	1	0.000
CsantA	TCA	False	False	31	0.529691	8	23	False	1	1	0.000
	TCT	False	False	41	0.196114	9	28	False	1	1	0.000
CtropB	AGC	False	False	24	0.841441	9	22	False	1	1	0.000
	TCG	False	False	9	1.206986	6	7	False	1	1	0.000
CtropA	AGC	False	False	24	0.885875	9	23	False	1	1	0.000
	TCG	False	False	9	1.206986	6	7	False	1	1	0.000
CoxycA	TCA	False	False	12	0.671472	1	9	False	1	1	0.037
	TCT	False	False	11	0.848760	1	8	False	1	1	0.040
CsubhA	AGC	False	False	62	0.608802	11	55	False	1	1	0.000
	TCC	False	False	30	0.808340	11	27	False	1	1	0.000
CsubhB	AGC	False	False	62	0.608802	11	55	False	1	1	0.000
	TCC	False	False	30	0.808340	11	27	False	1	1	0.000
OanguA	AGT	False	False	40	0.684226	10	28	False	1	1	0.000
	TCA	False	False	84	0.564213	13	71	False	1	1	0.000
KtherA	TCT	False	False	23	0.872930	5	19	False	1	1	0.000
CblacA	AGT	False	False	12	0.783005	1	10	False	1	1	0.245
	TCG	False	False	1	3.660550	1	0	False	1	1	0.357
CviswA	TCT	False	False	35	0.695480	11	32	False	1	1	0.000
MstarA	AGT	False	False	74	0.526995	12	56	False	2	2	0.000
	TCA	False	False	69	0.440955	11	51	False	1	2	0.000
NbaciA	TCT	False	False	38	0.756494	6	33	False	1	2	0.000
CalbiD	AGT	False	False	58	0.883025	11	55	False	1	2	0.000
CbohiA	AGT	False	False	60	0.892746	11	52	False	3	3	0.000
	TCA	False	False	80	0.857593	12	74	False	2	2	0.000
	AGC	False	False	6	1.046699	4	4	False	1	1	0.000
LstarA	AGT	False	False	49	0.692951	8	34	False	2	3	0.000
	TCA	False	False	27	0.615471	8	22	False	1	1	0.000
	TCT	False	False	53	0.430550	7	37	False	2	2	0.000
MfariA	AGT	False	False	55	0.809514	11	51	False	1	1	0.000
	TCT	False	False	60	0.781865	12	51	False	2	2	0.000
MfariB	TCT	False	False	46	0.764647	11	43	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CalaiA	TCA	False	False	61	0.774868	11	55	False	1	1	0.000
LanspB	TCT	False	False	23	0.928275	5	20	False	1	1	0.000
CphanA	AGT	False	False	45	0.842186	12	38	False	2	2	0.000
LanspA	TCT	False	False	29	0.737915	6	25	False	1	1	0.000
CsojaA	AGT	False	False	42	0.796186	7	39	False	1	1	0.000
	TCA	False	False	44	0.792677	7	41	False	1	1	0.000
CglabA	TCT	False	False	27	0.608650	6	21	False	1	1	0.000
	AGT	False	False	31	0.327117	6	14	False	3	3	0.000
ScryoA	TCA	False	False	37	0.401548	6	24	False	1	1	0.000
	TCT	False	False	29	0.360882	6	20	False	1	1	0.000
CneerA	TCA	False	False	62	0.791630	11	54	False	1	1	0.000
CsakeA	TCA	False	False	79	0.860145	11	75	False	1	1	0.000
HuvarA	TCT	False	False	26	0.426937	6	16	False	1	1	0.000
CjiufA	TCA	False	False	52	0.891111	10	49	False	1	1	0.000
	TCT	False	False	62	0.745069	12	56	False	1	1	0.000
	AGT	False	False	38	0.159879	6	21	False	1	1	0.000
CzempA	TCA	False	False	38	0.167000	6	20	False	1	1	0.000
	TCT	False	False	34	0.135906	6	20	False	1	1	0.000
CmetaB	TCT	False	False	58	0.908534	10	54	False	1	2	0.000
CmetaC	TCT	False	False	58	0.908534	10	54	False	1	2	0.000
CmetaA	TCT	False	False	58	0.908534	10	54	False	1	2	0.000
CmetaF	TCT	False	False	58	0.908534	10	54	False	1	2	0.000
CfrijA	AGT	False	False	65	0.811853	12	62	False	1	1	0.000
	TCT	False	False	43	0.791386	13	40	False	1	1	0.000
CmetaD	TCT	False	False	58	0.908534	10	54	False	1	2	0.000
CmetaE	TCT	False	False	58	0.908534	10	54	False	1	2	0.000
PpastA	AGT	False	False	58	0.753749	11	46	False	2	2	0.000
SjapoA	AGT	False	False	26	-0.091578	6	11	False	1	1	0.000
	TCT	False	False	61	0.188894	7	38	False	2	2	0.000
CalbiE	AGT	False	False	31	0.893541	6	28	False	1	2	0.000
Ctetra	AGT	False	False	63	0.816297	11	57	False	2	2	0.000
	TCT	False	False	40	0.814081	11	37	False	1	1	0.000
CalbiC	AGT	False	False	62	0.864562	12	59	False	1	2	0.000
CalbiB	AGT	False	False	62	0.864562	12	59	False	1	2	0.000
CalbiA	AGT	False	False	59	0.884638	11	56	False	1	2	0.000
MmagnB	AGC	False	False	22	0.866084	10	15	False	2	2	0.000
	TCA	False	False	52	0.345231	11	34	False	1	2	0.000
MmagnA	AGC	False	False	20	0.960498	10	14	False	2	2	0.000
	TCA	False	False	52	0.345231	11	34	False	1	2	0.000
CcoryA	TCA	False	False	60	1.091240	11	56	False	3	3	0.000
	TCT	False	False	46	0.843019	11	43	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CparaC	TCA	False	False	52	0.870740	10	48	False	1	1	0.000
	TCT	False	False	67	0.913982	11	66	False	1	1	0.000
CparaB	TCA	False	False	52	0.870740	10	48	False	1	1	0.000
	TCT	False	False	67	0.913982	11	66	False	1	1	0.000
CparaE	TCA	False	False	52	0.870740	10	48	False	1	1	0.000
	TCT	False	False	67	0.913982	11	66	False	1	1	0.000
CparaD	TCA	False	False	52	0.870740	10	48	False	1	1	0.000
	TCT	False	False	67	0.913982	11	66	False	1	1	0.000
CparaF	TCA	False	False	52	0.870740	10	48	False	1	1	0.000
	TCT	False	False	67	0.913982	11	66	False	1	1	0.000
MtetrA	AGC	False	False	39	0.590121	12	31	False	1	1	0.000
	TCC	False	False	19	0.916370	9	14	False	1	2	0.000
CsaraA	TCT	False	False	43	0.939194	9	41	False	1	1	0.000
PkluyA	TCA	False	False	122	0.548067	13	99	False	2	2	0.000
YlipoB	AGT	False	False	43	0.609205	12	35	False	1	1	0.000
YlipoA	AGT	False	False	43	0.609205	12	35	False	1	1	0.000
AgossA	AGT	False	False	16	0.994026	6	13	False	1	1	0.000
	TCA	False	False	56	0.517599	6	43	False	1	1	0.000
AgossB	AGT	False	False	16	0.994026	6	13	False	1	1	0.000
	TCA	False	False	57	0.490982	6	44	False	1	1	0.000
NdelpA	TCT	False	False	26	0.496211	5	18	False	1	1	0.000
	TCA	False	False	82	0.842460	12	75	False	1	1	0.000
LelonA	TCG	False	False	1	3.660550	1	0	False	1	1	0.146
	TCT	False	False	37	0.821947	10	35	False	1	1	0.000
LkluyF	TCT	False	False	22	0.842950	5	19	False	1	1	0.000
CalimA	AGT	False	False	45	0.734174	12	38	False	1	1	0.000
CchauA	TCA	False	False	65	0.967064	11	59	False	3	3	0.000
	TCT	False	False	40	0.852052	10	38	False	1	1	0.000
	AGC	False	False	4	1.652278	3	0	False	2	2	0.037
SoctoA	AGT	False	False	32	0.246058	6	15	False	2	2	0.001
	TCA	False	False	35	0.378158	6	21	False	2	2	0.001
CbuenA	TCA	False	False	47	0.755796	10	41	False	1	1	0.000
CdefoA	AGT	False	False	44	0.604150	12	37	False	1	1	0.000
LdasiA	TCT	False	False	24	0.773286	5	19	False	1	1	0.000
CpseuA	TCA	False	False	13	0.885934	1	11	False	1	1	0.037
	TCT	False	False	14	0.621586	1	11	False	1	1	0.036
LkluyE	TCT	False	False	21	0.837875	5	18	False	1	1	0.000
LkluyD	TCT	False	False	23	0.831056	5	20	False	1	1	0.000
LkluyC	TCT	False	False	22	0.842950	5	19	False	1	1	0.000
LkluyA	TCT	False	False	22	0.842950	5	19	False	1	1	0.000
CdublA	AGT	False	False	58	0.865456	11	55	False	1	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CorthB	TCT	False	False	67	0.958326	12	65	False	1	2	0.000
CorthC	TCT	False	False	69	0.951001	12	67	False	1	2	0.000
CorthA	TCT	False	False	68	0.950267	12	66	False	1	2	0.000
ClusiA	AGT	False	False	54	0.704928	11	48	False	1	1	0.000
	TCA	False	False	72	0.782784	12	66	False	1	1	0.000
MguilA	TCT	False	False	54	0.792610	13	48	False	1	1	0.000
SpastA	AGT	False	False	12	1.027955	5	10	False	1	1	0.000
DhansA	AGT	False	False	61	0.816034	11	54	False	2	2	0.000
	TCT	False	False	41	0.796992	10	38	False	1	1	0.000
	AGT	False	False	79	0.466927	11	54	False	2	2	0.000
Csalma	TCA	False	False	36	0.643376	8	27	False	2	2	0.000
	TCT	False	False	43	0.428976	9	29	False	2	2	0.000
LnothA	TCT	False	False	23	0.956882	5	21	False	1	1	0.000
BcustA	AGT	False	False	47	0.653438	11	32	False	1	1	0.000
	TCA	False	False	97	0.447170	12	76	False	1	1	0.000
CtheaA	TCA	False	False	63	0.950604	12	61	False	1	1	0.000
	TCT	False	False	55	0.848594	12	53	False	1	1	0.000
	AGT	False	False	73	0.535935	12	56	False	2	2	0.000
MingeA	TCA	False	False	73	0.278492	12	53	False	1	1	0.000
	TCT	False	False	36	0.492628	9	26	False	1	1	0.000
ScastA	TCA	False	False	60	0.655441	7	51	False	1	1	0.000
CmaltA	AGT	False	False	71	0.774921	11	65	False	1	1	0.000
	TCA	False	False	48	1.049258	11	46	False	1	2	0.000