

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	GCT	False	False	75	-0.117117	12	61	False	3	4	0.000
LmeyerA	GCT	False	False	61	-0.175109	6	51	False	1	1	0.000
CfrijA	GCA	False	False	27	-0.086904	9	26	False	1	1	0.000
	GCT	False	False	95	-0.166667	13	89	False	1	1	0.000
CgallA	GCA	False	False	58	-0.346040	13	38	False	2	2	0.000
	GCT	False	False	86	-0.163680	11	56	False	2	3	0.000
CviswA	GCA	False	False	44	-0.139474	9	39	False	1	1	0.055
CdefoA	GCA	False	False	61	-0.315248	13	38	False	1	2	0.000
	GCT	False	False	88	-0.213379	11	58	False	2	3	0.000
PpastA	GCA	False	False	78	-0.214286	10	64	False	1	1	0.000
	GCT	False	False	54	-0.182365	12	48	False	1	1	0.000
MstarA	GCA	False	False	41	-0.208411	8	34	False	1	1	0.000
	GCT	False	False	84	-0.293820	12	62	False	2	2	0.000
ScastA	GCT	False	False	53	-0.242181	6	46	False	1	1	0.000
AgossaA	GCT	False	False	42	-0.242942	6	36	False	1	1	0.046
LkluyF	GCA	False	False	51	-0.228780	5	42	False	1	1	0.000
	GCT	False	False	47	-0.298462	7	38	False	1	1	0.000
LkluyE	GCA	False	False	51	-0.195172	5	43	False	1	1	0.000
	GCT	False	False	48	-0.356789	7	37	False	1	1	0.000
LkluyD	GCA	False	False	50	-0.202520	5	42	False	1	1	0.000
	GCT	False	False	48	-0.331417	7	38	False	1	1	0.000
SpombA	GCA	False	False	22	-0.341400	6	10	False	2	2	0.000
	GCC	False	False	5	0.252896	4	3	False	1	1	0.000
	GCG	False	False	4	-0.648193	4	0	False	1	1	0.000
	GCT	False	False	68	-0.505499	6	40	False	2	2	0.000
CvartA	GCA	False	False	51	-0.122213	10	46	False	1	1	0.000
	GCC	False	False	3	0.892508	3	2	False	1	1	0.020
LkluyA	GCT	False	False	80	-0.400767	11	64	False	1	2	0.000
	GCA	False	False	51	-0.228780	5	42	False	1	1	0.000
ClusiA	GCT	False	False	47	-0.298462	7	38	False	1	1	0.000
	GCT	False	False	83	-0.117524	11	78	False	2	2	0.046
CoxycA	GCT	False	False	15	0.184242	1	13	False	1	1	0.051
NbaciA	GCT	False	False	70	-0.291934	6	56	False	2	2	0.000
AacerA	GCT	False	False	16	-0.253480	2	10	False	1	1	0.015
	GCA	False	False	53	-0.210229	9	35	False	1	1	0.000
MmagnB	GCC	False	False	31	-0.224719	9	25	False	1	1	0.000
	GCT	False	False	33	-0.239818	8	26	False	2	2	0.000
ScereC	GCT	False	False	53	-0.354798	6	41	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CgigaA	GCT	False	False	64	-0.207880	10	54	False	2	2	0.000
	GCA	False	False	51	-0.189760	9	33	False	1	1	0.000
MmagnA	GCC	False	False	31	-0.224719	9	25	False	1	1	0.000
	GCT	False	False	34	-0.270882	8	27	False	2	2	0.000
CcoryA	GCA	False	False	42	-0.262466	10	35	False	1	1	0.056
KmarxA	GCT	False	False	25	-0.276100	1	21	False	1	1	0.066
LstarA	GCA	False	False	37	-0.232419	7	25	False	1	2	0.000
	GCT	False	False	71	-0.230769	10	46	False	3	4	0.000
WmucoA	GCA	False	False	47	-0.307538	11	40	False	1	1	0.000
	GCT	False	False	85	-0.184110	12	69	False	3	4	0.000
	GCA	False	False	37	-0.239222	9	25	False	1	1	0.000
CsantA	GCC	False	False	21	-0.161503	8	18	False	1	1	0.000
	GCT	False	False	52	-0.492327	10	35	False	1	1	0.000
MtetrA	GCC	False	False	60	-0.211730	10	43	False	1	1	0.000
	GCT	False	False	15	-0.243598	6	9	False	2	2	0.000
CsaraA	GCA	False	False	40	-0.168700	10	37	False	1	1	0.055
	GCA	False	False	57	-0.176848	11	52	False	1	1	0.000
CsuavA	GCC	False	False	7	0.302452	4	6	False	1	1	0.000
	GCT	False	False	71	-0.434756	11	53	False	1	2	0.000
ScereD	GCT	False	False	51	-0.351275	6	40	False	1	1	0.000
ScereE	GCT	False	False	51	-0.349841	6	40	False	1	1	0.000
MfariA	GCT	False	False	68	-0.148373	12	64	False	1	1	0.046
PkluyA	GCA	False	False	76	-0.359645	9	58	False	1	1	0.000
	GCT	False	False	39	-0.186813	11	34	False	1	1	0.000
MfariB	GCA	False	False	49	-0.194277	10	40	False	1	1	0.000
CcastA	GCT	False	False	57	-0.309301	6	46	False	1	1	0.000
PcanaA	GCA	False	False	46	-0.155814	10	39	False	1	1	0.000
	GCT	False	False	89	-0.299129	11	71	False	3	4	0.000
DhansA	GCA	False	False	41	-0.066310	9	37	False	1	1	0.052
	GCT	False	False	90	-0.314382	12	80	False	1	1	0.046
BcaliA	GCT	False	False	77	-0.248649	12	59	False	2	3	0.000
BarxA	GCT	False	False	15	-0.489684	1	10	False	1	1	0.052
LanspB	GCT	False	False	58	-0.200164	6	49	False	1	1	0.000
CphanaA	GCA	False	False	49	-0.294557	12	32	False	2	2	0.000
	GCT	False	False	108	-0.240683	12	71	False	2	3	0.000
LanspA	GCC	False	False	5	0.685934	3	4	False	1	1	0.000
LdasiA	GCT	False	False	57	-0.159425	6	47	False	1	1	0.000
YlipoB	GCA	False	False	54	-0.280329	11	37	False	1	1	0.000
	GCT	False	False	98	-0.220250	12	63	False	2	4	0.000
YlipoA	GCA	False	False	54	-0.280329	11	37	False	1	1	0.000
	GCT	False	False	98	-0.220250	12	63	False	2	4	0.000

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CsalmA	GCT	False	False	86	-0.224711	11	63	False	2	3	0.000
CjadiA	GCT	False	False	82	-0.272054	12	62	False	3	4	0.050
ScereA	GCT	False	False	52	-0.314097	6	41	False	1	1	0.000
AgossB	GCT	False	False	43	-0.232703	6	37	False	1	1	0.046
CglabA	GCT	False	False	50	-0.273694	6	41	False	1	1	0.000
LkluyC	GCA	False	False	51	-0.228780	5	42	False	1	1	0.000
	GCT	False	False	47	-0.298462	7	38	False	1	1	0.000
OanguA	GCA	False	False	49	-0.106999	9	41	False	1	1	0.000
	GCT	False	False	82	-0.218917	12	70	False	2	2	0.000
NdelpA	GCT	False	False	47	-0.331523	6	38	False	1	1	0.000
	GCA	False	False	32	-0.115050	6	20	False	1	1	0.000
ScryoA	GCC	False	False	8	0.650602	5	5	False	2	2	0.001
	GCT	False	False	54	-0.731426	6	27	False	1	1	0.000
CneerA	GCT	False	False	97	-0.163681	13	93	False	1	1	0.000
LnothA	GCT	False	False	62	-0.227755	6	50	False	1	1	0.000
CsakeA	GCA	False	False	49	-0.082361	9	45	False	1	1	0.000
SarboA	GCT	False	False	50	-0.290582	6	40	False	1	1	0.000
KtherA	GCT	False	False	58	-0.179639	6	51	False	1	1	0.000
HuvarA	GCT	False	False	59	-0.515660	6	39	False	1	1	0.000
SjapoA	GCC	False	False	4	0.305882	3	2	False	1	1	0.000
	GCT	False	False	61	-0.474474	6	37	False	2	2	0.000
CalimA	GCA	False	False	42	-0.477938	9	27	False	2	2	0.000
	GCT	False	False	109	-0.276917	12	69	False	2	2	0.000
KlactA	GCT	False	False	54	-0.241190	6	42	False	1	1	0.000
CchauA	GCG	False	False	9	0.248563	5	8	False	1	1	0.061
CzempA	GCA	False	False	35	-0.157007	6	27	False	1	2	0.000
	GCT	False	False	45	-0.716677	6	26	False	1	1	0.000
	GCA	False	False	23	-0.530705	5	12	False	1	1	0.000
SoctoA	GCC	False	False	6	0.646605	2	4	False	1	1	0.000
	GCT	False	False	68	-0.428389	6	37	False	2	2	0.000
SpastA	GCT	False	False	47	-0.366395	6	35	False	1	1	0.000
WpijpA	GCT	False	False	49	-0.062853	6	41	False	2	3	0.000
MingeA	GCT	False	False	88	-0.185464	12	63	False	3	4	0.000
LkluyB	GCT	False	False	17	-0.337968	1	12	False	1	1	0.067