

# CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	GCA	False	False	54	-0.910588	11	47	False	1	1	0.000
	GCT	False	False	75	-1.004927	12	61	False	2	2	0.000
LmeyeA	GCA	False	False	37	-1.010518	6	30	False	1	1	0.000
CorthA	GCT	False	False	116	-1.036347	13	110	False	1	1	0.000
CgallA	GCA	False	False	58	-1.089933	13	38	False	2	2	0.000
	GCT	False	False	86	-1.083492	11	56	False	2	2	0.000
CbuenA	GCA	False	False	30	-0.782117	8	27	False	1	1	0.000
	GCT	False	False	104	-1.047518	13	94	False	1	1	0.000
CdefoA	GCA	False	False	61	-1.137943	13	38	False	2	2	0.000
	GCT	False	False	88	-1.001232	11	58	False	2	3	0.000
PpastA	GCT	False	False	54	-1.031120	12	48	False	1	1	0.000
MstarA	GCA	False	False	41	-0.893532	8	34	False	1	1	0.000
	GCT	False	False	84	-0.857396	12	62	False	3	4	0.000
AgossA	GCT	False	False	42	-0.907104	6	36	False	1	1	0.000
LkluyF	GCA	False	False	51	-1.052277	5	42	False	1	1	0.000
LkluyE	GCA	False	False	51	-1.040606	5	43	False	1	1	0.000
LkluyD	GCA	False	False	50	-1.058217	5	42	False	1	1	0.000
LkluyC	GCA	False	False	51	-1.052277	5	42	False	1	1	0.000
CvartA	GCA	False	False	51	-0.933522	10	46	False	1	1	0.000
	GCT	False	False	80	-1.090125	11	64	False	1	1	0.000
LkluyA	GCA	False	False	51	-1.052277	5	42	False	1	1	0.000
CorthB	GCT	False	False	115	-1.036398	13	109	False	1	1	0.000
CorthC	GCT	False	False	116	-1.036210	13	110	False	1	1	0.000
CparaF	GCT	False	False	105	-1.020829	12	100	False	1	1	0.000
DhansA	GCT	False	False	90	-0.958516	12	80	False	2	2	0.000
SpombA	GCG	False	False	4	0.262651	4	0	False	1	1	0.003
AacerA	GCT	False	False	16	-0.714037	2	10	False	1	1	0.016
	GCA	False	False	53	-1.170362	9	35	False	1	1	0.000
MmagnB	GCC	False	False	31	-0.860300	9	25	False	1	1	0.000
	GCG	False	False	14	-0.377701	9	11	False	2	2	0.000
ScereC	GCT	False	False	53	-1.069805	6	41	False	1	1	0.000
CgigaA	GCT	False	False	64	-1.020340	10	54	False	1	1	0.000
MmagnA	GCA	False	False	51	-1.167005	9	33	False	1	1	0.000
	GCC	False	False	31	-0.860300	9	25	False	1	1	0.000
	GCG	False	False	15	-0.433307	9	12	False	2	2	0.000
CparaA	GCT	False	False	84	-1.013041	11	81	False	1	1	0.000
CcoryA	GCA	False	False	42	-0.947656	10	35	False	1	1	0.000
KmarxA	GCA	False	False	8	-0.455581	1	6	False	1	1	0.129

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LstarA	GCA	False	False	37	-0.982276	7	25	False	1	1	0.000
	GCT	False	False	71	-1.168045	10	46	False	1	1	0.000
CparaE	GCT	False	False	105	-1.020829	12	100	False	1	1	0.000
CparaD	GCT	False	False	105	-1.020829	12	100	False	1	1	0.000
WmucoA	GCA	False	False	47	-0.957850	11	40	False	1	1	0.000
	GCT	False	False	85	-0.994805	12	69	False	2	2	0.000
CalimA	GCA	False	False	42	-1.128656	9	27	False	1	1	0.000
	GCT	False	False	109	-1.060982	12	69	False	2	3	0.000
CsantA	GCC	False	False	21	-0.737430	8	18	False	1	1	0.000
	GCT	False	False	52	-1.086556	10	35	False	1	1	0.000
	GCC	False	False	60	-1.076171	10	43	False	1	1	0.000
	GCG	False	False	27	-0.747656	10	21	False	2	2	0.000
MtetrA	GCT	False	False	15	-0.702561	6	9	False	1	1	0.000
	GCT	False	False	52	-1.054953	6	41	False	1	1	0.000
ScereA	GCT	False	False	52	-1.054953	6	41	False	1	1	0.000
CsuavA	GCT	False	False	71	-0.880952	11	53	False	3	3	0.000
ScereD	GCT	False	False	51	-1.069580	6	40	False	1	1	0.000
ScereE	GCT	False	False	51	-1.071094	6	40	False	1	1	0.000
MfariA	GCA	False	False	51	-0.905153	9	49	False	1	1	0.000
PkluyA	GCA	False	False	76	-1.074782	9	58	False	1	1	0.000
	GCT	False	False	39	-1.045604	11	34	False	1	1	0.000
MfariB	GCA	False	False	49	-0.995052	10	40	False	1	1	0.000
	GCT	False	False	83	-1.031028	12	78	False	1	1	0.000
CcastA	GCA	False	False	35	-0.932245	7	27	False	1	1	0.000
	GCT	False	False	57	-1.073436	6	46	False	1	1	0.000
PcanaA	GCA	False	False	46	-0.827442	10	39	False	2	2	0.000
	GCT	False	False	89	-1.089965	11	71	False	1	1	0.000
WpijpA	GCA	False	False	36	-0.746829	7	32	False	2	2	0.000
BcaliA	GCT	False	False	77	-0.883917	12	59	False	3	3	0.000
BarxA	GCT	False	False	15	-0.874393	1	10	False	1	1	0.052
LanspB	GCA	False	False	36	-0.989153	5	28	False	1	1	0.000
CphanaA	GCA	False	False	49	-1.138060	12	32	False	1	1	0.000
	GCT	False	False	108	-1.081027	12	71	False	2	3	0.000
LanspA	GCA	False	False	35	-1.011911	5	27	False	1	1	0.000
LdasiA	GCA	False	False	38	-0.951299	6	30	False	1	1	0.000
YlipoB	GCA	False	False	54	-1.111891	11	37	False	1	1	0.000
	GCT	False	False	98	-1.084265	12	63	False	2	3	0.000
YlipoA	GCA	False	False	54	-1.111891	11	37	False	1	1	0.000
	GCT	False	False	98	-1.084265	12	63	False	2	3	0.000
CsalmA	GCC	False	False	7	-1.368065	4	1	False	1	1	0.001
LnothA	GCA	False	False	36	-0.920307	6	29	False	1	1	0.000
	GCT	False	False	62	-1.092208	6	50	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
AgossB	GCT	False	False	43	-0.910790	6	37	False	1	1	0.000
	GCA	False	False	29	-0.956226	9	25	False	1	1	0.000
CsubhA	GCC	False	False	80	-1.019221	12	70	False	1	1	0.000
	GCG	False	False	14	-0.876763	8	11	False	1	1	0.000
BcustA	GCA	False	False	51	-0.840448	10	39	False	2	2	0.000
	GCT	False	False	66	-1.041427	11	51	False	1	1	0.000
CglabA	GCT	False	False	50	-0.902754	6	41	False	2	2	0.000
	GCA	False	False	29	-0.956226	9	25	False	1	1	0.000
CsubhB	GCC	False	False	80	-1.019221	12	70	False	1	1	0.000
	GCG	False	False	14	-0.876763	8	11	False	1	1	0.000
OanguA	GCT	False	False	82	-1.088845	12	70	False	1	1	0.000
NdelpA	GCT	False	False	47	-0.842486	6	38	False	2	2	0.000
MingeA	GCA	False	False	37	-1.000000	9	29	False	1	1	0.000
	GCT	False	False	88	-0.952986	12	63	False	2	3	0.000
CjadiA	GCT	False	False	82	-0.918000	12	62	False	3	3	0.000
CsakeA	GCT	False	False	73	-0.967438	11	68	False	1	1	0.000
SarboA	GCT	False	False	50	-1.045176	6	40	False	1	1	0.000
CparaB	GCT	False	False	105	-1.020829	12	100	False	1	1	0.000
	GCA	False	False	26	-1.105225	5	18	False	1	1	0.000
HuvarA	GCC	False	False	2	2.394366	2	0	False	1	1	0.023
	GCT	False	False	59	-1.180497	6	39	False	1	1	0.000
CjiufA	GCT	False	False	98	-1.051083	13	92	False	1	1	0.000
ScastA	GCA	False	False	26	-0.837243	6	22	False	1	1	0.000
CalaiA	GCT	False	False	100	-1.006767	10	88	False	1	1	0.000
	GCA	False	False	39	-1.016858	6	28	False	1	1	0.000
KtherA	GCT	False	False	58	-1.043021	6	51	False	1	1	0.000
	GCT	False	False	54	-1.133345	6	42	False	1	1	0.000
CchauA	GCT	False	False	98	-1.041354	13	83	False	1	1	0.000
CblacA	GCC	False	False	1	5.709091	1	0	False	1	1	0.347
	GCT	False	False	26	-0.756123	1	22	False	1	1	0.245
CzempA	GCT	False	False	45	-1.154058	6	26	False	1	1	0.000
NbaciA	GCT	False	False	70	-0.999591	6	56	False	1	2	0.000
SpastA	GCT	False	False	47	-0.977597	6	35	False	2	2	0.000
SjapoA	GCT	False	False	61	-0.913387	6	37	False	2	3	0.000
CparaC	GCT	False	False	105	-1.020829	12	100	False	1	1	0.000
LkluyB	GCA	False	False	18	-0.782499	1	17	False	1	1	0.060