CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	ACA	False	False	56	-0.806888	11	45	False	1	1	0.000
LmeyeA	ACA	False	False	48	-0.868563	7	44	False	1	1	0.000
CfrijA	ACT	False	False	56	-0.894576	10	52	False	1	1	0.000
CgallA	ACA	False	False	73	-0.736207	12	47	False	4	5	0.000
CyaliA	ACT	False	False	44	-0.910202	11	32	False	1	1	0.000
CbuenA	ACT	False	False	77	-0.854285	11	71	False	1	1	0.000
CviswA	ACT	False	False	53	-0.832296	11	48	False	1	1	0.000
CdefoA	ACA	False	False	78	-0.818613	12	52	False	3	4	0.000
	ACT	False	False	39	-0.771412	11	26	False	2	2	0.000
PpastA	ACA	False	False	87	-0.701352	12	73	False	1	1	0.000
MstarA	ACA	False	False	55	-1.014020	10	41	False	1	1	0.000
MStarA	ACT	False	False	55	-0.730691	10	44	False	3	3	0.000
LkluyF	ACA	False	False	41	-0.745898	7	36	False	2	2	0.000
LkluyE	ACA	False	False	41	-0.745898	7	36	False	2	2	0.000
LkluyD	ACA	False	False	41	-0.731130	7	36	False	2	2	0.000
SpombA	ACA	False	False	23	-0.068623	6	13	False	3	3	0.000
	ACT	False	False	36	-0.814904	6	25	False	1	1	0.000
CvartA	ACT	False	False	53	-0.792073	10	41	False	2	2	0.000
LkluyA	ACA	False	False	41	-0.745898	7	36	False	2	2	0.000
CdublA	ACA	False	False	27	-0.695321	7	26	False	1	1	0.000
CalbiE	ACA	False	False	20	-0.723734	4	18	False	1	1	0.000
CalbiD	ACA	False	False	35	-0.791517	8	33	False	1	1	0.000
CalbiC	ACA	False	False	41	-0.862462	9	39	False	1	1	0.000
CalbiB	ACA	False	False	41	-0.862462	9	39	False	1	1	0.000
CalbiA	ACA	False	False	35	-0.791517	8	33	False	1	1	0.000
CphanA	ACA	False	False	81	-0.810313	13	55	False	2	2	0.000
СрпапА	ACT	False	False	35	-0.916273	9	24	False	2	3	0.000
CtetrA	ACT	False	False	47	-0.786319	11	45	False	1	1	0.000
MmagnR	ACA	False	False	89	-0.872169	11	65	False	3	3	0.000
MmagnB	ACT	False	False	26	-0.727938	10	18	False	1	1	0.000
CgigaA	ACC	False	False	43	-0.668520	9	37	False	1	2	0.000
MmagnA	ACA	False	False	90	-0.877527	11	65	False	3	3	0.000
	ACT	False	False	26	-0.712010	10	19	False	1	1	0.000
CcoryA	ACA	False	False	47	-0.864551	10	44	False	1	1	0.000
CbohiA	ACT	False	False	45	-0.778922	10	42	False	1	1	0.000
	ACA	False	False	37	-0.746219	8	25	False	3	3	0.000
LstarA	ACG	False	False	3	0.904762	3	2	False	1	1	0.017
	ACT	False	False	54	-0.806426	8	42	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
WmucoA	ACA	False	False	51	-0.847191	11	43	False	1	1	0.000
	ACT	False	False	47	-0.967973	11	38	False	1	1	0.000
CsantA	ACA	False	False	56	-0.680194	13	35	False	4	5	0.000
	ACT	False	False	54	-0.563109	11	37	False	3	3	0.000
MtetrA	ACA	False	False	63	-0.796372	11	41	False	3	3	0.000
	ACT	False	False	29	-0.874497	9	24	False	1	1	0.000
CsaraA	ACA	False	False	56	-0.921887	11	52	False	1	1	0.000
CtropB	ACT	False	False	42	-0.883786	10	37	False	1	1	0.000
CsuavA	ACA	False	False	64	-0.855311	13	54	False	1	1	0.000
	ACT	False	False	35	-1.033987	9	29	False	1	1	0.000
CtropA	ACT	False	False	44	-0.878303	10	39	False	1	1	0.000
PkluyA	ACA	False	False	99	-0.693076	11	80	False	2	2	0.000
CcastA	ACA	False	False	26	-0.860294	6	22	False	1	1	0.000
	ACT	False	False	35	-0.688797	6	29	False	1	1	0.000
PcanaA	ACA	False	False	59	-0.871140	12	51	False	1	1	0.000
	ACT	False	False	41	-0.909240	10	32	False	1	1	0.000
WpijpA	ACA	False	False	49	-0.719250	7	42	False	1	2	0.000
BcaliA	ACA	False	False	50	-0.827867	10	41	False	1	1	0.000
BarxA	ACA	False	False	15	-0.717576	1	8	False	1	1	0.044
	ACT	False	False	10	0.132969	1	7	False	1	2	0.053
LanspB	ACA	False	False	45	-0.861995	7	41	False	1	1	0.000
CoxycA	ACA	False	False	10	-0.322404	1	8	False	1	1	0.036
LdasiA	ACA	False	False	43	-0.894666	7	39	False	1	1	0.000
CsojaA	ACT	False	False	39	-0.842987	8	37	False	1	1	0.000
YlipoB	ACA	False	False	70	-0.899529	12	46	False	2	3	0.000
	ACT	False	False	47	-0.802800	11	32	False	2	2	0.000
YlipoA	ACA	False	False	70	-0.899529	12	46	False	2	3	0.000
	ACT	False	False	47	-0.802800	11	32	False	2	2	0.000
CsalmA	ACA	False	False	37	-0.989344	10	24	False	1	1	0.000
	ACT	False	False	70	-0.824511	10	44	False	2	2	0.000
LnothA	ACA	False	False	42	-0.884818	7	38	False	1	1	0.000
Coulbb A	ACT	False	False	22	-0.688546	5	21	False	1	1	0.000
CsubhA	ACC	False	False	47	-0.917258	9	39	False	1	1	0.000
BcustA	ACA	False	False	61	-0.685729	11	43	False	2	2	0.000
CglabA	ACC	False	False	32	-0.631595	5	30	False	1	1	0.000
CsubhB	ACC	False	False	47	-0.917258	9	39	False	1	1	0.000
LkluyC	ACA	False	False	41	-0.745898	7	36	False False	2	2	0.000
NdelpA	ACA	False	False	34	-0.892290	7	27	False	1	1	0.000
ScryoA	ACA	False	False False	29 37	-0.613681	7 6	18 23		1	1	0.000
0	ACT	False			-0.272563			False	3	5	0.000
CneerA	ACT	False	False	53	-0.816083	11	50	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CjadiA	ACT	False	False	35	-0.692333	10	26	False	2	2	0.000
KtherA	ACT	False	False	21	-0.756682	5	19	False	1	1	0.000
HuvarA	ACA	False	False	21	-0.399907	5	11	False	1	1	0.000
	ACT	False	False	35	-0.266339	6	17	False	3	4	0.000
ScastA	ACA	False	False	38	-0.787732	7	30	False	1	1	0.000
CalimA	ACA	False	False	84	-0.898770	12	58	False	3	4	0.000
	ACT	False	False	31	-0.652883	10	20	False	1	1	0.000
KlactA	ACA	False	False	46	-0.766716	6	41	False	1	1	0.000
CchauA	ACT	False	False	42	-0.837449	10	41	False	1	1	0.000
CzempA	ACA	False	False	32	-0.543046	6	21	False	2	2	0.000
	ACT	False	False	33	-0.708065	5	21	False	1	1	0.000
SoctoA	ACA	False	False	26	-0.524723	7	18	False	2	2	0.000
	ACT	False	False	39	-0.443193	6	24	False	3	4	0.000
SjapoA	ACT	False	False	58	-0.637842	6	32	False	3	3	0.000
MingeA	ACA	False	False	67	-0.789880	10	50	False	2	3	0.000
	ACT	False	False	44	-0.935501	10	34	False	2	2	0.000