CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	ACT	False	False	50	-0.755951	12	49	False	1	1	0.000
CnorvA	ACA	False	False	56	-0.783595	11	45	False	1	1	0.000
MingeA	ACA	False	False	67	-0.786097	10	50	False	2	2	0.000
CmetaF	ACT	False	False	50	-0.755951	12	49	False	1	1	0.000
CfrijA	ACA	False	False	45	-0.776190	11	43	False	1	1	0.000
CgallA	ACA	False	False	73	-0.692960	12	47	False	2	2	0.000
	ACT	False	False	44	-0.890276	11	32	False	2	2	0.000
CmetaC	ACT	False	False	50	-0.755951	12	49	False	1	1	0.000
CmetaE	ACT	False	False	50	-0.755951	12	49	False	1	1	0.000
CdefoA	ACA	False	False	78	-0.760955	12	52	False	2	2	0.000
	ACT	False	False	39	-0.783945	11	26	False	2	2	0.000
PpastA	ACA	False	False	87	-0.559604	12	73	False	3	4	0.000
MstarA	ACA	False	False	55	-0.966199	10	41	False	2	2	0.000
LkluyF	ACT	False	False	24	-0.748888	5	23	False	1	1	0.000
SjapoA	ACC	False	False	5	-0.384470	2	3	False	1	1	0.000
CchauA	ACA	False	False	58	-0.877021	10	54	False	1	1	0.000
SpombA	ACA	False	False	23	-0.024804	6	13	False	3	3	0.000
	ACT	False	False	36	-0.793382	6	25	False	1	1	0.000
CvartA	ACT	False	False	53	-0.760946	10	41	False	1	1	0.000
LkluyA	ACT	False	False	24	-0.748888	5	23	False	1	1	0.000
CbuenA	ACT	False	False	77	-0.865820	11	71	False	1	1	0.000
CorthB	ACA	False	False	47	-0.731279	9	44	False	1	1	0.000
CorthC	ACA	False	False	48	-0.714477	9	45	False	1	1	0.000
CorthA	ACA	False	False	47	-0.731279	9	44	False	1	1	0.000
ClusiA	ACA	False	False	42	-0.730929	9	41	False	1	1	0.000
LkluyE	ACT	False	False	24	-0.748888	5	23	False	1	1	0.000
NbaciA	ACT	False	False	40	-0.638942	6	35	False	1	1	0.000
MmagnB	ACA	False	False	89	-0.823133	11	65	False	2	2	0.000
CmetaD	ACT	False	False	50	-0.755951	12	49	False	1	1	0.000
CgigaA	ACC	False	False	43	-0.708656	9	37	False	1	1	0.000
MmagnA	ACA	False	False	90	-0.829845	11	65	False	2	2	0.000
CcoryA	ACT	False	False	53	-0.825822	8	51	False	1	1	0.000
CparaC	ACA	False	False	45	-0.841446	11	43	False	1	1	0.000
CparaB	ACA	False	False	45	-0.841446	11	43	False	1	1	0.000
CparaE	ACA	False	False	45	-0.841446	11	43	False	1	1	0.000
CparaD	ACA	False	False	45	-0.841446	11	43	False	1	1	0.000
WmucoA	ACA	False	False	51	-0.834910	11	43	False	1	1	0.000
CparaF	ACA	False	False	45	-0.841446	11	43	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CsantA	ACA	False	False	56	-0.782136	13	35	False	1	1	0.000
	ACT	False	False	54	-0.486425	11	37	False	2	2	0.000
MtetrA	ACA	False	False	63	-0.738846	11	41	False	1	1	0.000
	ACG	False	False	17	-1.053501	7	11	False	1	1	0.000
CsuavA	ACA	False	False	64	-0.841664	13	54	False	1	1	0.000
PkluyA	ACA	False	False	99	-0.605166	11	80	False	4	5	0.000
LkluyD	ACT	False	False	24	-0.762232	5	23	False	1	1	0.000
CalaiA	ACT	False	False	53	-0.863880	10	47	False	1	1	0.000
PcanaA	ACA	False	False	59	-0.871497	12	51	False	1	1	0.000
WpijpA	ACA	False	False	49	-0.735834	7	42	False	1	1	0.000
BcaliA	ACA	False	False	50	-0.797577	10	41	False	1	1	0.000
BarxA	ACA	False	False	15	-0.620000	1	8	False	1	1	0.048
CphanA	ACA	False	False	81	-0.685248	13	55	False	4	4	0.000
CmetaA	ACT	False	False	50	-0.755951	12	49	False	1	1	0.000
YlipoB	ACA	False	False	70	-0.829153	12	46	False	2	2	0.000
	ACT	False	False	47	-0.780835	11	32	False	2	2	0.000
YlipoA	ACA	False	False	70	-0.829153	12	46	False	2	2	0.000
	ACT	False	False	47	-0.780835	11	32	False	2	2	0.000
CsalmA	ACA	False	False	37	-0.968033	10	24	False	1	1	0.000
	ACT	False	False	70	-0.798442	10	44	False	2	2	0.000
CjadiA	ACT	False	False	35	-0.643393	10	26	False	1	1	0.000
AgossB	ACT	False	False	48	-0.865764	7	41	False	1	1	0.000
CsubhA	ACG	False	False	21	-0.600660	11	17	False	1	1	0.000
BcustA	ACA	False	False	61	-0.610277	11	43	False	4	4	0.000
	ACT	False	False	58	-0.638700	9	44	False	1	2	0.000
CglabA	ACT	False	False	32	-0.625899	5	30	False	1	1	0.000
CsubhB	ACG	False	False	21	-0.600660	11	17	False	1	1	0.000
LkluyC	ACT	False	False	24	-0.748888	5	23	False	1	1	0.000
CtheaA	ACT	False	False	62	-0.802804	12	60	False	1	1	0.000
OanguA	ACA	False	False	51	-0.640725	10	38	False	3	3	0.000
	ACC	False	False	6	-0.283228	3	4	False	1	1	0.000
	ACT	False	False	60	-0.916956	11	46	False	1	1	0.000
NdelpA	ACT	False	False	35	-0.883618	6	31	False	1	1	0.000
ScryoA	ACA	False	False	29	-0.499009	7	18	False	3	3	0.000
	ACT	False	False	37	-0.351435	6	23	False	1	1	0.000
AgossA	ACT	False	False	49	-0.851259	7	42	False	1	1	0.000
CsakeA	ACT	False	False	44	-0.822481	10	41	False	1	1	0.000
HuvarA	ACA	False	False	21	-0.207617	5	11	False	3	4	0.000
	ACT	False	False	35	-0.293957	6	17	False	2	2	0.000
ScastA	ACA	False	False	38	-0.825910	7	30	False	1	1	0.000
CalimA	ACA	False	False	84	-0.876189	12	58	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
	ACT	False	False	31	-0.510622	10	20	False	3	3	0.000
LstarA	ACT	False	False	54	-0.801778	8	42	False	1	1	0.000
CblacA	ACT	False	False	15	-0.708914	1	13	False	1	1	0.040
CzempA	ACA	False	False	32	-0.600542	6	21	False	2	2	0.000
SoctoA	ACA	False	False	26	-0.596679	7	18	False	1	1	0.000
	ACT	False	False	39	-0.451028	6	24	False	2	2	0.000
CbohiA	ACT	False	False	45	-0.782108	10	42	False	1	1	0.000