CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	ACA	False	False	48	0.932278	7	44	False	1	1	0.000
	ACA	False	False	23	0.272427	6	13	False	2	2	0.000
SpombA	ACT	False	False	36	0.920635	6	25	False	4	4	0.000
SpastA	ACA	False	False	33	0.948487	6	29	False	1	1	0.000
AacerA	ACT	False	False	18	1.224120	2	15	False	1	2	0.011
CgigaA	ACA	False	False	5	1.424569	4	4	False	1	1	0.001
Cylyan	ACC	False	False	43	0.747340	9	37	False	1	1	0.000
	ACA	False	False	56	0.621553	13	35	False	5	5	0.000
CsantA	ACG	False	False	5	2.009634	3	0	False	2	3	0.238
	ACT	False	False	54	0.500311	11	37	False	2	2	0.000
CtropB	ACT	False	False	42	1.056331	10	37	False	3	4	0.000
CtropA	ACT	False	False	44	1.043636	10	39	False	3	4	0.000
WpijpA	ACA	False	False	49	0.776308	7	42	False	1	1	0.000
[][ACT	False	False	19	1.223911	6	16	False	1	2	0.000
	ACA	False	False	34	0.926421	11	32	False	1	1	0.000
CsubhA	ACC	False	False	47	0.991726	9	39	False	2	2	0.000
	ACG	False	False	21	0.887789	11	17	False	2	2	0.000
	ACA	False	False	34	0.926421	11	32	False	1	1	0.000
CsubhB	ACC	False	False	47	0.991726	9	39	False	2	2	0.000
	ACG	False	False	21	0.887789	11	17	False	2	2	0.000
	ACA	False	False	51	0.687256	10	38	False	2	2	0.000
OanguA	ACC	False	False	6	0.753165	3	4	False	1	1	0.000
	ACT	False	False	60	0.972963	11	46	False	6	6	0.000
SarboA	ACA	False	False	36	0.957600	7	32	False	1	1	0.000
CcastA	ACT	False	False	35	0.756293	6	29	False	2	2	0.000
KtherA	ACT	False	False	21	0.895392	5	19	False	1	1	0.000
CblacA	ACT	False	False	15	0.927229	1	13	False	1	1	0.040
CnorvA	ACA	False	False	56	0.898865	11	45	False	4	5	0.000
	ACT	False	False	44	1.178459	10	34	False	2	4	0.000
CviswA	ACA	False	False	49	0.948081	10	47	False	1	1	0.000
	ACT	False	False	53 EE	0.939948	11	48	False	1	2	0.000
MstarA	ACA ACT	False False	False False	55 55	0.925101 0.808943	10	41 44	False False	3	5 3	0.000
NbaciA	ACA	False	False	25	1.333463	7	18	False	4	5	0.006
NDACIA	ACA	False	False	35	0.919355	8	33	False	1	1	0.000
CalbiD	ACT	False	False	53	0.919333	10	50	False	1	2	0.000
CbohiA	ACT	False	False	45	0.867157	10	42	False	1	1	0.000
ODOINA	ACA	False	False	37	0.763541	8	25	False	1	3	0.000
LstarA	AUA	i aise	1 disc	01	0.700041	U	20	i disc	_	J	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
	ACT	False	False	54	0.761972	8	42	False	3	3	0.000
MfariA	ACA	False	False	45	1.012522	9	44	False	1	1	0.000
	ACT	False	False	59	1.027571	12	55	False	2	3	0.000
MfariB	ACT	False	False	42	0.994256	9	40	False	1	1	0.000
CalaiA	ACA	False	False	45	1.048159	9	44	False	1	1	0.000
	ACT	False	False	53	0.958986	10	47	False	3	3	0.000
LanspB	ACA	False	False	45	0.931639	7	41	False	1	1	0.000
O I A	ACA	False	False	81	0.627546	13	55	False	4	6	0.000
CphanA	ACT	False	False	35	0.895341	9	24	False	4	4	0.000
LanspA	ACA	False	False	40	0.946939	7	38	False	1	1	0.000
CsojaA	ACA	False	False	38	1.046955	6	36	False	2	2	0.000
Osojan	ACT	False	False	39	0.971479	8	37	False	1	1	0.000
CglabA	ACA	False	False	37	1.183935	7	30	False	3	4	0.000
ScryoA	ACA	False	False	29	0.846993	7	18	False	4	5	0.000
Coryon	ACT	False	False	37	0.400310	6	23	False	2	2	0.000
	ACA	False	False	45	0.946949	9	42	False	2	2	0.000
CneerA	ACC	False	False	4	1.616279	2	3	False	1	1	0.002
	ACT	False	False	53	0.904395	11	50	False	1	1	0.000
CsakeA	ACA	False	False	63	0.897762	10	58	False	2	2	0.000
HuvarA	ACA	False	False	21	0.525778	5	11	False	2	3	0.000
	ACT	False	False	35	0.413180	6	17	False	3	4	0.000
KlactA	ACA	False	False	46	0.871935	6	41	False	1	1	0.000
CzempA	ACA	False	False	32	0.586394	6	21	False	2	3	0.000
·	ACT	False	False	33	0.785798	5	21	False	3	3	0.000
CfrijA	ACA	False	False	45	0.901754	11	43	False	1	1	0.000
•	ACT	False	False	56	0.970779	10	52	False	1	2	0.000
CgallA	ACA	False	False	73	0.543103	12	47	False	4	5	0.000
	ACT	False	False	44	0.925877	11	32	False	5	5	0.000
PpastA	ACA	False	False	87	0.707864	12	73	False	2	3	0.000
SjapoA	ACA	False	False	8	1.153216	2	6	False	1	1	0.000
	ACT	False	False	58	0.809959	6	32	False	5	9	0.000
CalbiE	ACA	False	False	20	0.922139	4	18	False	1	1	0.000
	ACT	False	False	24	1.050040	5	22	False	1	2	0.000
CtetrA	ACA	False	False	52	1.029772	11	49	False	2	2	0.000
	ACT	False	False	47	0.891866	11	45	False	1	1	0.000
CalbiC	ACA	False	False	41	0.955167	9	39	False	1	1	0.000
	ACT	False	False	62	0.978902	11	59	False	1	2	0.000
CalbiB	ACA	False	False	41	0.955167	9	39	False	1	1	0.000
	ACT	False	False	62	0.978902	11	59	False	1	2	0.000
CalbiA	ACA	False	False	35	0.919355	8	33	False	1	1	0.000
	ACT	False	False	53	0.976241	10	50	False	1	2	0.000

		Clad. Val	Reassigned	Tot.	Telford	Tot. Genes	Use.	Trans. Val	Rea. Genes	Rea. count	Prob
MmagnB	ACA	False	False	89	0.954699	11	65	False	6	10	0.000
	ACT	False	False	26	0.395179	10	18	False	1	1	0.000
Managara	ACA	False	False	90	0.955648	11	65	False	6	10	0.000
MmagnA	ACT	False	False	26	0.388721	10	19	False	1	1	0.000
CparaA	ACC	False	False	2	2.215116	2	1	False	1	1	0.041
CcoryA	ACA	False	False	47	0.950983	10	44	False	1	2	0.000
CparaC	ACC	False	False	2	2.215116	2	1	False	1	1	0.055
CparaB	ACC	False	False	2	2.215116	2	1	False	1	1	0.055
CparaE	ACC	False	False	2	2.215116	2	1	False	1	1	0.055
CparaD	ACC	False	False	2	2.215116	2	1	False	1	1	0.055
CparaF	ACC	False	False	2	2.215116	2	1	False	1	1	0.055
	ACA	False	False	63	0.914365	11	41	False	6	10	0.000
MtetrA	ACG	False	False	17	1.140047	7	11	False	2	2	0.000
	ACT	False	False	29	0.762898	9	24	False	1	1	0.000
CsaraA	ACA	False	False	56	0.918794	11	52	False	1	1	0.000
Osuran	ACC	False	False	4	1.205195	4	3	False	1	1	0.001
CsuavA	ACA	False	False	64	0.872082	13	54	False	3	3	0.000
	ACT	False	False	35	1.168627	9	29	False	3	3	0.000
PkluyA	ACA	False	False	99	0.780334	11	80	False	4	5	0.000
PcanaA	ACA	False	False	59	0.961271	12	51	False	4	4	0.000
	ACT	False	False	41	1.025424	10	32	False	2	3	0.000
CsalmA	ACA	False	False	37	1.069672	10	24	False	5	7	0.000
	ACT	False	False	70	0.701160	10	44	False	3	5	0.000
YlipoA	ACA	False	False	70	0.784314	12	46	False	4	8	0.000
	ACT	False	False	47	0.746078	11	32	False	5	5	0.000
	ACA	False	False	65	1.060601	11	59	False	3	3	0.000
CjadiA	ACC	False	False	1	3.450704	1	0	False	1	1	0.185
	ACT	False	False	35	0.779119	10	26	False	3	3	0.000
AgossB	ACT	False	False	48	1.070504	7	41	False	4	5	0.000
NdelpA	ACA	False	False	34	1.085034	7	27	False	3	4	0.000
1 -1 0	ACT	False	False	35	0.974076	6	31	False	1	1	0.000
LelonA	ACA	False	False	51	0.874066	10	48	False	1	1	0.000
CalimA	ACA	False	False	84	0.673601	12	58	False	4	6	0.000
Lklast	ACT	False	False	31	0.762519	10	20	False	4	4	0.000
LkluyD	ACA	False	False	41	0.845992	7	36	False	1	1	0.000
BcaliA	ACA	False	False	50	0.947357	10	41	False	3	4	0.000
	ACT	False	False	50	1.094395	12	41	False	3	4	0.000
SoctoA	ACA	False	False	26	0.633948	7	18	False	2	2	0.000
	ACT	False	False	39	0.775955	6	24	False	1	6	0.000
Chuan A	ACA	False	False	30	0.942924	7	28	False	1	1	0.000
CbuenA	ACC	False	False	1	3.400000	1	0	False	1	1	0.182

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
	ACT	False	False	77	0.962622	11	71	False	3	3	0.000
CdefoA	ACA	False	False	78	0.625017	12	52	False	4	6	0.000
	ACT	False	False	39	0.854670	11	26	False	5	5	0.000
LdasiA	ACA	False	False	43	0.967195	7	39	False	1	1	0.000
LkluyF	ACA	False	False	41	0.852789	7	36	False	1	1	0.000
LkluyE	ACA	False	False	41	0.852789	7	36	False	1	1	0.000
CchauA	ACA	False	False	58	0.938056	10	54	False	1	2	0.000
LkluyC	ACA	False	False	41	0.852789	7	36	False	1	1	0.000
CvartA	ACA	False	False	48	1.098942	12	43	False	2	3	0.000
Ovartit	ACT	False	False	53	0.884208	10	41	False	4	4	0.000
LkluyA	ACA	False	False	41	0.852789	7	36	False	1	1	0.000
CdublA	ACT	False	False	61	0.974231	10	58	False	2	2	0.000
ClusiA	ACT	False	False	64	0.916752	11	59	False	1	1	0.000
MguilA	ACA	False	False	51	0.988828	10	49	False	1	1	0.000
WmucoA	ACA	False	False	51	0.924904	11	43	False	3	3	0.000
	ACT	False	False	47	1.054493	11	38	False	4	4	0.000
ScereC	ACA	False	False	38	0.917110	7	33	False	1	1	0.000
ScereA	ACA	False	False	38	0.917110	7	33	False	1	1	0.000
ScereD	ACA	False	False	37	0.979979	7	33	False	1	1	0.000
ScereE	ACT	False	False	25	0.921257	5	24	False	1	1	0.000
DhansA	ACT	False	False	53	0.907118	11	50	False	1	1	0.000
YlipoB	ACA	False	False	70	0.784314	12	46	False	4	8	0.000
	ACT	False	False	47	0.746078	11	32	False	5	5	0.000
LnothA	ACA	False	False	42	0.957179	7	38	False	1	1	0.000
BcustA	ACA	False	False	61	0.770570	11	43	False	4	5	0.000
Double	ACT	False	False	58	0.746350	9	44	False	3	4	0.000
AgossA	ACT	False	False	49	1.056217	7	42	False	4	5	0.000
MingeA	ACA	False	False	67	0.731873	10	50	False	3	5	0.000
90, .	ACT	False	False	44	0.909959	10	34	False	3	3	0.000
ScastA	ACA	False	False	38	1.045559	7	30	False	4	4	0.000
CmaltA	ACT	False	False	62	1.010493	10	59	False	1	1	0.000