## CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	ACA	False	False	51	-0.962924	8	50	False	1	1	0.000
CmetaC	ACA	False	False	51	-0.962924	8	50	False	1	1	0.000
CmetaA	ACA	False	False	51	-0.962924	8	50	False	1	1	0.000
CmetaF	ACA	False	False	51	-0.962924	8	50	False	1	1	0.000
CfrijA	ACT	False	False	56	-0.974790	10	52	False	1	1	0.051
CmetaD	ACA	False	False	51	-0.962924	8	50	False	1	1	0.000
CmetaE	ACA	False	False	51	-0.962924	8	50	False	1	1	0.000
CbuenA	ACA	False	False	30	-0.884846	7	28	False	1	1	0.000
CviswA	ACA	False	False	49	-0.931377	10	47	False	1	1	0.000
CdefoA	ACA	False	False	78	-1.003120	12	52	False	1	1	0.000
MguilA	ACA	False	False	51	-0.932530	10	49	False	1	1	0.000
CchauA	ACA	False	False	58	-0.968211	10	54	False	1	1	0.000
CvartA	ACT	False	False	53	-0.993152	10	41	False	1	1	0.000
CdubIA	ACT	False	False	61	-0.956618	10	58	False	1	1	0.000
CorthB	ACA	False	False	47	-1.023973	9	44	False	1	1	0.000
CorthC	ACA	False	False	48	-1.032791	9	45	False	1	1	0.000
CtetrA	ACA	False	False	52	-0.908034	11	49	False	1	1	0.000
CalbiC	ACT	False	False	62	-0.959950	11	59	False	1	1	0.000
CalbiB	ACT	False	False	62	-0.959950	11	59	False	1	1	0.000
CalbiA	ACT	False	False	53	-0.958422	10	50	False	1	1	0.000
CorthA	ACA	False	False	47	-1.023973	9	44	False	1	1	0.000
CgallA	ACA	False	False	73	-1.039368	12	47	False	1	1	0.000
CgigaA	ACT	False	False	34	-0.948371	10	33	False	1	1	0.000
CparaA	ACA	False	False	34	-0.938137	10	33	False	1	1	0.000
CcoryA	ACT	False	False	53	-0.996549	8	51	False	1	1	0.000
CbohiA	ACA	False	False	58	-0.979726	9	56	False	1	1	0.000
CparaB	ACA	False	False	45	-0.984819	11	43	False	1	1	0.000
CparaE	ACA	False	False	45	-0.984819	11	43	False	1	1	0.000
CparaD	ACA	False	False	45	-0.984819	11	43	False	1	1	0.000
WmucoA	ACT	False	False	47	-0.885994	11	38	False	1	1	0.000
CparaF	ACA	False	False	45	-0.984819	11	43	False	1	1	0.000
CsantA	ACA	False	False	56	-1.049126	13	35	False	1	1	0.000
CsaraA	ACA	False	False	56	-0.949729	11	52	False	1	1	0.000
CtropB	ACA	False	False	49	-0.984066	12	48	False	1	1	0.051
CsuavA	ACA	False	False	64	-1.011181	13	54	False	1	1	0.000
CtropA	ACA	False	False	48	-0.979487	12	47	False	1	1	0.051
MfariA	ACT	False	False	59	-0.967713	12	55	False	1	1	0.000
MfariB	ACA	False	False	62	-0.981769	11	61	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CalaiA	ACG	False	False	1	2.303922	1	0	False	1	1	0.278
	ACT	False	False	53	-0.976025	10	47	False	1	1	0.000
PcanaA	ACA	False	False	59	-0.956809	12	51	False	1	1	0.000
WpijpA	ACA	False	False	49	-1.035341	7	42	False	1	1	0.000
CphanA	ACA	False	False	81	-1.036031	13	55	False	1	1	0.000
CnorvA	ACT	False	False	44	-0.842792	10	34	False	1	1	0.000
YlipoB	ACA	False	False	70	-0.927345	12	46	False	1	1	0.000
YlipoA	ACA	False	False	70	-0.927345	12	46	False	1	1	0.000
CsalmA	ACT	False	False	70	-1.016373	10	44	False	1	1	0.000
CjadiA	ACA	False	False	65	-0.916290	11	59	False	1	1	0.000
CsubhA	ACA	False	False	34	-0.933718	11	32	False	1	1	0.000
BcustA	ACA	False	False	61	-1.053090	11	43	False	1	1	0.000
CsubhB	ACA	False	False	34	-0.933718	11	32	False	1	1	0.000
CalbiD	ACT	False	False	53	-0.958422	10	50	False	1	1	0.000
CtheaA	ACT	False	False	62	-1.018282	12	60	False	1	1	0.000
LelonA	ACA	False	False	51	-0.987033	10	48	False	1	1	0.000
CneerA	ACA	False	False	45	-0.916065	9	42	False	1	1	0.055
CsakeA	ACT	False	False	44	-1.011141	10	41	False	1	1	0.000
ClusiA	ACT	False	False	64	-0.949201	11	59	False	1	1	0.000
CjiufA	ACA	False	False	50	-0.990353	10	48	False	1	1	0.000
CalimA	ACA	False	False	84	-0.955854	12	58	False	1	1	0.000
LstarA	ACT	False	False	54	-1.052132	8	42	False	1	1	0.000
BcaliA	ACA	False	False	50	-0.960936	10	41	False	1	1	0.000
CmaltA	ACT	False	False	62	-0.924449	10	59	False	1	1	0.000
DhansA	ACT	False	False	53	-0.982422	11	50	False	1	1	0.000
CparaC	ACA	False	False	45	-0.984819	11	43	False	1	1	0.000