CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CmetaB	GCT	False	False	105	-1.070702	13	101	False	1	1	0.000
CmetaC	GCT	False	False	105	-1.070702	13	101	False	1	1	0.000
CparaB	GCA	False	False	30	-0.992566	6	28	False	1	1	0.000
CmetaF	GCT	False	False	105	-1.070702	13	101	False	1	1	0.000
CgallA	GCA	False	False	58	-1.083087	13	38	False	1	1	0.000
	GCT	False	False	86	-0.888169	11	56	False	4	5	0.000
CmetaE	GCT	False	False	105	-1.070702	13	101	False	1	1	0.000
CdefoA	GCA	False	False	61	-1.034574	13	38	False	2	2	0.000
	GCT	False	False	88	-0.946532	11	58	False	4	4	0.000
PpastA	GCA	False	False	78	-1.009917	10	64	False	1	1	0.000
MstarA	GCT	False	False	84	-1.128799	12	62	False	2	2	0.000
CpseuA	GCT	False	False	11	-0.767769	1	10	False	1	1	0.046
SjapoA	GCT	False	False	61	-1.205785	6	37	False	1	2	0.000
LmeyeA	GCT	False	False	61	-0.982645	6	51	False	1	1	0.000
SpombA	GCT	False	False	68	-1.177383	6	40	False	2	2	0.000
CorthB	GCT	False	False	115	-1.092227	13	109	False	1	1	0.000
CorthC	GCT	False	False	116	-1.090525	13	110	False	1	1	0.000
CorthA	GCT	False	False	116	-1.093032	13	110	False	1	1	0.000
CoxycA	GCT	False	False	15	-0.640606	1	13	False	1	1	0.051
NbaciA	GCT	False	False	70	-1.143033	6	56	False	1	1	0.000
MmagnB	GCA	False	False	53	-0.863866	9	35	False	2	3	0.000
CmetaD	GCT	False	False	105	-1.070702	13	101	False	1	1	0.000
CgigaA	GCT	False	False	64	-1.061339	10	54	False	1	1	0.000
MmagnA	GCA	False	False	51	-0.831166	9	33	False	2	3	0.000
CcoryA	GCA	False	False	42	-1.102208	10	35	False	1	1	0.000
	GCT	False	False	95	-1.040628	13	83	False	1	1	0.000
CparaC	GCA	False	False	30	-0.992566	6	28	False	1	1	0.000
LstarA	GCA	False	False	37	-1.020869	7	25	False	1	1	0.000
	GCT	False	False	71	-0.967234	10	46	False	3	3	0.000
CparaE	GCA	False	False	30	-0.992566	6	28	False	1	1	0.000
CparaD	GCA	False	False	30	-0.992566	6	28	False	1	1	0.000
SpastA	GCT	False	False	47	-1.170061	6	35	False	1	1	0.000
CparaF	GCA	False	False	30	-0.992566	6	28	False	1	1	0.000
CsantA	GCA	False	False	37	-0.985630	9	25	False	1	1	0.000
	GCT	False	False	52	-1.278494	10	35	False	1	1	0.000
MtetrA	GCA	False	False	32	-1.070505	8	26	False	1	1	0.000
	GCC	False	False	60	-0.887311	10	43	False	1	1	0.000
LelonA	GCA	False	False	40	-1.080166	10	36	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
ScereA	GCT	False	False	52	-1.112663	6	41	False	1	1	0.000
CalimA	GCT	False	False	109	-1.010397	12	69	False	5	6	0.000
ScereD	GCT	False	False	51	-1.162791	6	40	False	1	1	0.000
ScereE	GCT	False	False	51	-1.160195	6	40	False	1	1	0.000
MfariB	GCA	False	False	49	-0.994621	10	40	False	1	1	0.000
DhansA	GCA	False	False	41	-0.923974	9	37	False	1	1	0.000
LanspB	GCT	False	False	58	-1.004102	6	49	False	1	1	0.000
CphanA	GCA	False	False	49	-1.064311	12	32	False	1	1	0.000
	GCT	False	False	108	-0.963433	12	71	False	4	5	0.000
CmetaA	GCT	False	False	105	-1.070702	13	101	False	1	1	0.000
LdasiA	GCT	False	False	57	-0.953041	6	47	False	1	1	0.000
YlipoB	GCA	False	False	54	-1.088430	11	37	False	1	1	0.000
	GCT	False	False	98	-0.910256	12	63	False	4	5	0.000
YlipoA	GCA	False	False	54	-1.088430	11	37	False	1	1	0.000
	GCT	False	False	98	-0.910256	12	63	False	4	5	0.000
CsalmA	GCC	False	False	7	-0.368065	4	1	False	1	1	0.001
	GCT	False	False	86	-0.950679	11	63	False	3	3	0.000
LnothA	GCT	False	False	62	-1.018903	6	50	False	1	1	0.000
CsubhA	GCG	False	False	14	-1.110616	8	11	False	1	1	0.000
CsubhB	GCG	False	False	14	-1.110616	8	11	False	1	1	0.000
CtheaA	GCA	False	False	33	-0.962350	9	31	False	1	1	0.000
OanguA	GCA	False	False	49	-0.932755	9	41	False	1	1	0.000
ScryoA	GCA	False	False	32	-0.735056	6	20	False	2	3	0.000
	GCT	False	False	54	-1.394063	6	27	False	2	2	0.000
CsakeA	GCA	False	False	49	-0.933852	9	45	False	1	1	0.000
SarboA	GCT	False	False	50	-1.100115	6	40	False	1	1	0.000
HuvarA	GCA	False	False	26	-0.792839	5	18	False	2	2	0.000
	GCT	False	False	59	-1.262028	6	39	False	1	1	0.000
CjiufA	GCT	False	False	98	-1.066193	13	92	False	1	1	0.000
ScastA	GCT	False	False	53	-1.080965	6	46	False	1	1	0.000
KtherA	GCC	False	False	4	-0.227603	3	2	False	1	1	0.000
KlactA	GCA	False	False	34	-0.944831	7	29	False	1	1	0.000
	GCT	False	False	54	-1.017886	6	42	False	1	1	0.000
CchauA	GCT	False	False	98	-1.106528	13	83	False	2	2	0.000
CzempA	GCA	False	False	35	-0.934694	6	27	False	1	1	0.000
	GCT	False	False	45	-1.448824	6	26	False	1	1	0.000
ScereC	GCT	False	False	53	-1.156385	6	41	False	1	1	0.000
CmaltA	GCT	False	False	99	-1.119275	13	92	False	1	1	0.000
SoctoA	GCA	False	False	23	-1.248963	5	12	False	1	1	0.000
	GCT	False	False	68	-1.061623	6	37	False	3	5	0.000
CbohiA	GCT	False	False	108	-0.997591	13	93	False	2	2	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MingeA	**GCT**	False	False	88	-0.970385	12	63	False	2	2	0.000