CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	GCA	False	False	37	-1.142381	6	30	False	1	1	0
CgallA	GCA	False	False	58	-1.114360	13	38	False	1	1	0
	GCT	False	False	86	-1.019952	11	56	False	1	1	0
CbuenA	GCT	False	False	104	-1.128383	13	94	False	1	1	0
CdefoA	GCA	False	False	61	-1.106206	13	38	False	1	1	0
	GCT	False	False	88	-1.045337	11	58	False	1	1	0
PpastA	GCA	False	False	78	-1.026132	10	64	False	2	2	0
MstarA	GCA	False	False	41	-1.019164	8	34	False	1	1	0
LkluyF	GCA	False	False	51	-1.080382	5	42	False	1	1	0
SjapoA	GCA	False	False	12	-1.500000	3	7	False	1	1	0
	GCT	False	False	61	-1.285127	6	37	False	1	1	0
CchauA	GCA	False	False	27	-0.985943	10	24	False	1	1	0
SpombA	GCA	False	False	22	-1.133535	6	10	False	1	1	0
	GCT	False	False	68	-1.258319	6	40	False	1	1	0
LkluyA	GCA	False	False	51	-1.080382	5	42	False	1	1	0
CdublA	GCT	False	False	88	-1.089595	12	83	False	1	1	0
CalbiE	GCA	False	False	15	-0.952896	5	14	False	1	1	0
CalbiD	GCA	False	False	26	-0.979158	10	25	False	1	1	0
CalbiC	GCA	False	False	33	-1.008798	11	31	False	1	1	0
CalbiB	GCA	False	False	33	-1.008798	11	31	False	1	1	0
CalbiA	GCA	False	False	27	-0.990684	10	26	False	1	1	0
MmagnB	GCA	False	False	53	-0.912485	9	35	False	2	2	0
CgigaA	GCC	False	False	36	-0.784594	11	31	False	2	2	0
MmagnA	GCA	False	False	51	-0.883584	9	33	False	2	2	0
CcoryA	GCT	False	False	95	-1.032779	13	83	False	2	2	0
CbohiA	GCT	False	False	108	-0.990061	13	93	False	3	4	0
LstarA	GCA	False	False	37	-1.073185	7	25	False	1	1	0
CalimA	GCA	False	False	42	-1.207238	9	27	False	1	1	0
	GCT	False	False	109	-1.113366	12	69	False	1	1	0
CsantA	GCA	False	False	37	-1.070161	9	25	False	1	1	0
	GCT	False	False	52	-1.315736	10	35	False	1	1	0
MtetrA	GCC	False	False	60	-0.836961	10	43	False	3	4	0
	GCG	False	False	27	-1.066496	10	21	False	1	1	0
MfariA	GCA	False	False	51	-1.017450	9	49	False	1	1	0
PkluyA	GCA	False	False	76	-1.107326	9	58	False	1	2	0
LelonA	GCT	False	False	93	-1.039265	13	88	False	1	1	0
CcastA	GCA	False	False	35	-1.170612	7	27	False	1	1	0
	GCT	False	False	57	-1.110070	6	46	False	1	1	0

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LanspB	**GCA**	False	False	36	-1.127778	5	28	False	1	1	0
CphanA	**GCT**	False	False	108	-1.064042	12	71	False	2	2	0
YlipoB	**GCT**	False	False	98	-1.016108	12	63	False	2	2	0
LanspA	**GCA**	False	False	35	-0.952355	5	27	False	2	2	0
	GCT	False	False	58	-1.161370	6	51	False	1	1	0
LdasiA	**GCT**	False	False	57	-0.979612	6	47	False	1	1	0
CsalmA	**GCA**	False	False	25	-1.250530	8	18	False	1	1	0
	GCT	False	False	86	-1.007801	11	63	False	2	2	0
BcustA	**GCA**	False	False	51	-1.242437	10	39	False	1	1	0
SoctoA	**GCC**	False	False	6	-0.100309	2	4	False	1	1	0
LnothA	**GCT**	False	False	62	-1.037959	6	50	False	1	1	0
CsubhA	**GCA**	False	False	29	-0.811698	9	25	False	2	2	0
YlipoA	**GCT**	False	False	98	-1.016108	12	63	False	2	2	0
CsubhB	**GCA**	False	False	29	-0.811698	9	25	False	2	2	0
OanguA	**GCA**	False	False	49	-0.939099	9	41	False	1	1	0
	GCT	False	False	82	-1.110502	12	70	False	1	1	0
ScryoA	**GCA**	False	False	32	-0.833884	6	20	False	1	1	0
MingeA	**GCA**	False	False	37	-1.173876	9	29	False	1	1	0
	GCT	False	False	88	-1.051209	12	63	False	1	1	0
LkluyE	**GCA**	False	False	51	-1.048466	5	43	False	1	1	0
KtherA	**GCA**	False	False	39	-1.120938	6	28	False	2	2	0
KlactA	**GCA**	False	False	34	-0.947618	7	29	False	1	1	0
LkluyD	**GCA**	False	False	50	-1.050200	5	42	False	1	1	0
LkluyC	**GCA**	False	False	51	-1.080382	5	42	False	1	1	0