CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
LmeyeA	TAT	False	False	61	3.046427	6	58	False	2	2	0.000
SpombA	TAC	False	False	17	3.270787	5	15	False	1	2	0.000
	TAT	False	False	55	2.680806	6	41	False	4	5	0.000
SpastA	TAT	False	False	69	2.952282	7	64	False	1	2	0.000
AacerA	TAT	False	False	32	3.025095	2	29	False	2	2	0.037
CmetaD	TAC	False	False	25	3.302034	9	22	False	2	3	0.000
	TAT	False	False	131	3.040845	11	123	False	4	7	0.000
CgigaA	TAC	False	False	100	3.070620	12	96	False	2	3	0.000
	TAT	False	False	31	3.086144	10	28	False	1	2	0.000
CsantA	TAC	False	False	42	3.142820	11	34	False	4	5	0.000
	TAT	False	False	120	2.812119	13	100	False	6	11	0.000
CtropB	TAC	False	False	81	3.005386	11	80	False	1	1	0.000
	TAT	False	False	69	2.989551	12	64	False	2	3	0.000
CtropA	TAC	False	False	82	3.007298	11	81	False	1	1	0.000
	TAT	False	False	68	2.986683	12	63	False	2	3	0.000
WpijpA	TAC	False	False	48	3.125549	7	43	False	2	4	0.000
	TAT	False	False	32	2.868161	6	27	False	2	2	0.000
BarxA	TAT	False	False	21	3.120157	1	17	False	1	3	0.038
CoxycA	TAT	False	False	13	2.997179	1	9	False	1	3	0.127
CsubhA	TAC	False	False	132	3.176295	13	122	False	5	9	0.000
	TAT	False	False	11	3.030435	6	10	False	1	1	0.000
CsubhB	TAC	False	False	132	3.176295	13	122	False	5	9	0.000
	TAT	False	False	11	3.030435	6	10	False	1	1	0.000
OanguA	TAT	False	False	124	2.826192	12	112	False	2	2	0.000
SarboA	TAT	False	False	69	2.959850	7	64	False	1	2	0.000
CcastA	TAT	False	False	67	2.964772	7	62	False	2	2	0.000
KtherA	TAT	False	False	54	3.080880	6	50	False	2	3	0.000
CblacA	TAT	False	False	23	3.496835	1	19	False	1	4	0.043
CnorvA	TAT	False	False	149	2.995798	13	130	False	7	11	0.000
CviswA	TAT	False	False	138	3.063828	13	131	False	4	6	0.000
MstarA	TAT	False	False	145	2.952348	13	129	False	5	9	0.000
NbaciA	TAT	False	False	65	3.037288	7	59	False	2	3	0.000
CalbiD	TAC	False	False	28	3.223976	8	27	False	1	1	0.000
	TAT	False	False	111	3.071756	11	106	False	3	5	0.000
CbohiA	TAT	False	False	145	2.992465	13	138	False	3	5	0.000
LstarA	TAT	False	False	87	2.919492	10	74	False	5	7	0.000
MfariA	TAT	False	False	117	2.997536	12	113	False	2	3	0.000
MfariB	TAT	False	False	126	3.065121	12	121	False	3	5	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CalaiA	**TAC**	False	False	8	3.360636	4	7	False	1	1	0.000
	TAT	False	False	133	2.946806	11	128	False	2	3	0.000
LanspB	**TAT**	False	False	59	3.048908	6	56	False	2	2	0.000
CphanA	**TAT**	False	False	124	2.792567	12	109	False	3	4	0.000
LanspA	**TAT**	False	False	61	3.002674	6	58	False	2	2	0.000
CsojaA	**TAT**	False	False	90	2.992393	8	87	False	2	2	0.000
CglabA	**TAT**	False	False	71	3.085080	7	63	False	3	5	0.000
ScryoA	**TAC**	False	False	27	3.018008	4	19	False	3	5	0.000
	TAT	False	False	46	2.810732	7	36	False	3	4	0.000
CneerA	**TAT**	False	False	144	3.007623	13	137	False	5	5	0.000
CsakeA	**TAT**	False	False	131	3.028477	12	127	False	4	4	0.000
HuvarA	**TAC**	False	False	47	3.036766	6	39	False	1	4	0.000
	TAT	False	False	16	2.472057	6	12	False	2	2	0.000
CjiufA	**TAC**	False	False	25	3.263823	9	22	False	2	3	0.000
	TAT	False	False	126	2.982250	11	121	False	4	4	0.000
KlactA	**TAT**	False	False	63	3.084993	7	59	False	2	3	0.000
CzempA	**TAT**	False	False	82	2.830780	6	64	False	4	9	0.000
CmetaB	**TAC**	False	False	25	3.302034	9	22	False	2	3	0.000
	TAT	False	False	131	3.040845	11	123	False	4	7	0.000
CmetaC	**TAC**	False	False	25	3.302034	9	22	False	2	3	0.000
	TAT	False	False	131	3.040845	11	123	False	4	7	0.000
CmetaA	**TAC**	False	False	25	3.302034	9	22	False	2	3	0.000
	TAT	False	False	131	3.040845	11	123	False	4	7	0.000
CmetaF	**TAC**	False	False	25	3.302034	9	22	False	2	3	0.000
	TAT	False	False	131	3.040845	11	123	False	4	7	0.000
CfrijA	**TAT**	False	False	145	3.016873	13	138	False	5	5	0.000
CgallA	**TAT**	False	False	112	2.776573	13	100	False	2	3	0.000
CmetaE	**TAC**	False	False	25	3.302034	9	22	False	2	3	0.000
	TAT	False	False	131	3.040845	11	123	False	4	7	0.000
PpastA	**TAT**	False	False	133	2.862377	12	123	False	2	2	0.000
SjapoA	**TAT**	False	False	62	2.625948	6	45	False	4	7	0.000
CalbiE	**TAC**	False	False	21	3.220428	5	20	False	1	1	0.000
	TAT	False	False	51	3.154735	6	46	False	3	5	0.000
CtetrA	**TAC**	False	False	36	3.127875	11	34	False	2	2	0.000
	TAT	False	False	111	3.034317	13	108	False	3	3	0.000
CalbiC	**TAC**	False	False	29	3.207021	9	28	False	1	1	0.000
	TAT	False	False	123	3.038618	12	118	False	3	5	0.000
CalbiB	**TAC**	False	False	29	3.207021	9	28	False	1	1	0.000
	TAT	False	False	123	3.038618	12	118	False	3	5	0.000
CalbiA	**TAC**	False	False	28	3.223808	8	27	False	1	1	0.000
	TAT	False	False	111	3.071771	11	106	False	3	5	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
MmagnB	TAC	False	False	115	2.973658	13	104	False	3	6	0.000
	TAT	False	False	33	2.816549	9	28	False	3	3	0.000
MmagnA	TAC	False	False	119	2.970226	13	108	False	3	6	0.000
	TAT	False	False	29	2.805484	9	24	False	3	3	0.000
CparaA	TAC	False	False	14	3.140613	7	13	False	1	1	0.000
	TAT	False	False	116	3.023990	10	113	False	3	3	0.000
CcoryA	TAT	False	False	138	3.047805	13	127	False	5	9	0.000
CparaC	TAC	False	False	18	3.240991	8	16	False	2	2	0.000
	TAT	False	False	138	3.045364	11	130	False	4	7	0.000
CparaB	TAC	False	False	18	3.240991	8	16	False	2	2	0.000
	TAT	False	False	138	3.045364	11	130	False	4	7	0.000
CparaE	TAC	False	False	18	3.240991	8	16	False	2	2	0.000
	TAT	False	False	138	3.045364	11	130	False	4	7	0.000
CparaD	TAC	False	False	18	3.240991	8	16	False	2	2	0.000
	TAT	False	False	138	3.045364	11	130	False	4	7	0.000
CparaF	TAC	False	False	18	3.240991	8	16	False	2	2	0.000
	TAT	False	False	138	3.045364	11	130	False	4	7	0.000
MtetrA	TAC	False	False	136	2.775343	13	115	False	5	9	0.000
CsaraA	TAT	False	False	137	3.073685	13	130	False	4	6	0.000
CsuavA	TAC	False	False	31	3.122247	10	30	False	1	1	0.000
	TAT	False	False	120	2.912603	12	104	False	6	8	0.000
PkluyA	TAT	False	False	117	2.844674	12	106	False	3	3	0.000
PcanaA	TAT	False	False	140	2.975872	12	122	False	7	10	0.000
YlipoB	TAC	False	False	25	3.135081	11	23	False	1	1	0.000
	TAT	False	False	114	2.747642	13	102	False	1	2	0.000
YlipoA	TAC	False	False	25	3.135081	11	23	False	1	1	0.000
	TAT	False	False	114	2.747642	13	102	False	1	2	0.000
CjadiA	TAC	False	False	32	2.963426	9	29	False	2	2	0.000
	TAT	False	False	119	2.941770	12	104	False	5	7	0.000
AgossB	TAT	False	False	72	2.947046	6	67	False	3	3	0.000
NdelpA	TAT	False	False	74	3.040689	6	66	False	3	5	0.000
LelonA	TAC	False	False	30	3.100809	10	28	False	2	2	0.000
	TAT	False	False	122	3.043499	12	115	False	4	6	0.000
LkluyF	TAT	False	False	62	3.053640	5	58	False	2	3	0.000
CalimA	TAT	False	False	117	2.822717	12	106	False	2	3	0.000
CchauA	TAT	False	False	139	3.135802	12	125	False	5	12	0.000
BcaliA	TAC	False	False	12	3.167585	5	11	False	1	1	0.000
	TAT	False	False	143	2.927432	13	125	False	7	9	0.000
SoctoA	TAC	False	False	17	3.126940	5	13	False	1	3	0.000
	TAT	False	False	56	2.591780	7	42	False	4	4	0.000
LkluyB	TAT	False	False	18	3.177868	1	16	False	1	2	0.035

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CbuenA	TAT	False	False	141	3.013876	12	137	False	3	3	0.000
CdefoA	TAC	False	False	28	3.123369	12	25	False	1	1	0.000
	TAT	False	False	114	2.689002	13	101	False	1	2	0.000
LdasiA	TAT	False	False	57	3.005548	6	54	False	2	2	0.000
CpseuA	TAC	False	False	11	3.426799	1	9	False	1	2	0.036
	TAT	False	False	13	2.750353	1	11	False	1	1	0.039
LkluyE	TAT	False	False	62	3.053640	5	58	False	2	3	0.000
LkluyD	TAT	False	False	62	3.055650	5	58	False	2	3	0.000
LkluyC	TAT	False	False	62	3.053640	5	58	False	2	3	0.000
CvartA	TAC	False	False	23	3.102297	8	22	False	1	1	0.000
	TAT	False	False	127	2.936488	12	111	False	6	8	0.000
LkluyA	TAT	False	False	62	3.053640	5	58	False	2	3	0.000
CdublA	TAT	False	False	125	3.118695	12	118	False	3	7	0.000
CorthB	TAC	False	False	20	3.355069	9	17	False	2	3	0.000
	TAT	False	False	137	3.025002	11	130	False	3	6	0.000
CorthC	TAC	False	False	19	3.274584	8	17	False	1	2	0.000
	TAT	False	False	138	3.040263	11	130	False	4	7	0.000
CorthA	TAC	False	False	20	3.254582	8	18	False	1	2	0.000
	TAT	False	False	137	3.041534	11	129	False	4	7	0.000
ClusiA	TAC	False	False	15	3.235294	7	14	False	1	1	0.000
	TAT	False	False	130	3.019641	12	125	False	3	4	0.000
MguilA	TAT	False	False	125	3.093220	11	118	False	4	7	0.000
WmucoA	TAC	False	False	34	3.158925	7	30	False	3	3	0.000
	TAT	False	False	120	2.954974	12	104	False	7	9	0.000
ScereC	TAT	False	False	64	2.967127	7	59	False	1	2	0.000
ScereA	TAT	False	False	66	2.968536	7	61	False	1	2	0.000
ScereD	TAT	False	False	65	2.962826	7	60	False	1	2	0.000
ScereE	TAT	False	False	67	3.002292	7	61	False	1	3	0.000
DhansA	TAT	False	False	111	3.048658	11	106	False	3	5	0.000
CsalmA	TAT	False	False	133	2.727144	12	112	False	7	9	0.000
LnothA	TAT	False	False	58	3.009417	6	55	False	2	2	0.000
BcustA	TAT	False	False	139	2.922748	12	122	False	3	7	0.000
CtheaA	TAC	False	False	19	3.375594	8	16	False	2	3	0.000
	TAT	False	False	138	3.027692	11	131	False	4	6	0.000
AgossA	TAT	False	False	72	2.947046	6	67	False	3	3	0.000
MingeA	TAT	False	False	142	2.924932	13	124	False	5	9	0.000
ScastA	TAT	False	False	63	3.003952	6	60	False	2	2	0.000
CmaltA	TAC	False	False	19	3.141563	10	17	False	2	2	0.000
	TAT	False	False	128	3.071871	12	124	False	2	4	0.000
KmarxA	TAT	False	False	18	3.216271	1	15	False	1	3	0.035