

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	GAT	False	False	60	2.056407	9	58	False	2	2	0.000
CfrijA	GAT	False	False	57	2.075763	9	56	False	1	1	0.055
CgallA	GAT	False	False	53	2.054856	9	49	False	1	2	0.000
CviswA	GAT	False	False	60	2.112752	9	58	False	2	2	0.000
CdefoA	GAT	False	False	54	2.048188	9	50	False	1	2	0.000
PpastA	GAT	False	False	56	2.076864	9	53	False	2	2	0.000
CpseuA	GAT	False	False	10	2.039126	1	8	False	1	1	0.039
SjapoA	GAT	False	False	38	1.629191	5	27	False	3	3	0.000
CchauA	GAT	False	False	56	2.123703	9	54	False	2	2	0.000
SpombA	GAT	False	False	37	1.732435	5	27	False	3	3	0.000
CvartA	GAT	False	False	57	1.996563	9	54	False	2	2	0.000
CtetrA	GAT	False	False	51	1.997681	9	49	False	1	1	0.000
CoxycA	GAT	False	False	9	2.040445	1	7	False	1	1	0.041
MmagnB	GAC	False	False	36	2.018255	9	34	False	1	1	0.000
MmagnA	GAC	False	False	35	2.015054	9	33	False	1	1	0.000
CcoryA	GAT	False	False	57	2.051625	9	55	False	1	1	0.000
LstarA	GAT	False	False	40	1.841883	7	36	False	1	1	0.000
WmucoA	GAT	False	False	58	2.049963	9	56	False	2	2	0.000
CsantA	GAC	False	False	13	2.337748	5	12	False	1	1	0.000
	GAT	False	False	45	2.202853	10	39	False	3	4	0.000
MtetrA	GAC	False	False	48	1.911867	10	44	False	1	1	0.000
CsaraA	GAC	False	False	4	2.564165	3	3	False	1	1	0.001
	GAT	False	False	58	2.109676	9	56	False	2	2	0.000
CtropB	GAT	False	False	37	2.067827	8	36	False	1	1	0.055
CsuavA	GAC	False	False	5	2.249049	3	4	False	1	1	0.001
	GAT	False	False	56	1.971535	9	54	False	1	1	0.000
CtropA	GAT	False	False	36	2.090994	8	35	False	1	1	0.055
LelonA	GAC	False	False	6	2.420382	2	5	False	1	1	0.000
CalaiA	GAT	False	False	57	2.080242	8	54	False	2	2	0.000
PcanaA	GAT	False	False	60	2.090398	9	57	False	3	3	0.000
WpijpA	GAT	False	False	33	2.073696	5	31	False	2	2	0.000
BcaliA	GAT	False	False	60	2.056407	9	58	False	2	2	0.000
Cphana	GAT	False	False	53	2.092371	9	50	False	1	2	0.000
YlipoB	GAT	False	False	56	2.009817	9	51	False	1	2	0.000
YlipoA	GAT	False	False	56	2.009817	9	51	False	1	2	0.000
Csalma	GAT	False	False	62	2.130472	9	50	False	4	7	0.183
CjadiA	GAT	False	False	58	2.002454	9	55	False	2	2	0.000
CsubhA	GAC	False	False	59	2.004406	9	57	False	1	1	0.000

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
BcustA	GAT	False	False	56	2.025794	9	52	False	1	2	0.000
CsubhB	GAC	False	False	59	2.004406	9	57	False	1	1	0.000
ScryoA	GAC	False	False	14	1.661180	3	10	False	1	1	0.053
	GAT	False	False	27	1.765537	6	20	False	2	2	0.053
CneerA	GAT	False	False	55	2.058235	9	54	False	1	1	0.055
HuvarA	GAT	False	False	38	1.552665	5	28	False	2	2	0.000
CjiufA	GAT	False	False	51	2.050043	9	50	False	1	1	0.000
ScastA	GAT	False	False	42	1.954410	5	39	False	1	1	0.000
CalimA	GAT	False	False	51	2.109583	9	48	False	1	2	0.000
CzempA	GAT	False	False	35	2.340722	5	28	False	4	6	0.242
CmaltA	GAT	False	False	57	2.099411	9	55	False	1	2	0.055
SoctoA	GAC	False	False	8	1.420798	4	5	False	1	1	0.062
	GAT	False	False	32	1.832788	4	25	False	2	2	0.061
DhansA	GAT	False	False	55	2.042089	9	54	False	1	1	0.000