

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
CnorvA	TTA	False	False	345	-0.980447	13	319	False	1	1	0.000
MingeA	TTA	False	False	335	-0.954254	13	283	False	2	2	0.000
CbuenA	TTA	False	False	293	-0.982991	13	274	False	1	1	0.000
MstarA	TTA	False	False	333	-0.960651	13	285	False	2	2	0.000
SjapoA	CTT	False	False	40	-0.728102	5	26	True	2	2	0.000
	TTA	False	False	153	-0.840866	7	99	False	3	4	0.000
NbaciA	CTT	False	False	2	1.380282	2	0	True	1	1	0.022
	TTA	False	False	186	-0.961846	7	173	False	1	1	0.001
WmucoA	CTA	False	False	20	-0.844622	7	16	True	1	1	0.000
AacerA	CTA	True	True	16	2.582133	2	0	True	2	11	0.998
	CTT	True	True	5	2.242938	2	0	True	2	3	0.996
MmagnB	CTG	False	False	62	-0.942281	12	52	False	1	1	0.000
	TTA	False	False	28	-0.759427	7	21	False	1	1	0.000
ScereC	CTA	False	False	11	-0.069324	5	2	True	1	1	0.000
MmagnA	CTG	False	False	63	-0.943103	13	53	False	1	1	0.000
	TTA	False	False	24	-0.720016	7	18	False	1	1	0.000
LstarA	TTA	False	False	229	-0.983439	10	194	False	2	2	0.000
SpastA	CTT	False	False	2	1.394366	2	0	True	1	1	0.022
	TTA	False	False	195	-0.954720	7	175	False	1	1	0.001
CsantA	CTA	False	False	191	-0.913255	13	152	True	2	2	0.000
	CTT	False	False	60	-0.907842	11	50	True	2	2	0.000
MtetrA	CTG	False	False	124	-0.978665	12	104	False	1	1	0.000
	TTA	False	False	11	-0.432356	7	9	False	1	1	0.000
CsaraA	TTA	False	False	288	-0.981960	13	277	False	1	1	0.000
ScereD	CTA	False	False	10	-0.069923	5	2	True	1	1	0.000
ScereE	CTA	False	False	11	-0.062992	6	2	True	1	1	0.001
MfariA	TTA	False	False	312	-0.974805	13	293	False	1	1	0.000
CcastA	CTA	False	False	12	0.089444	5	0	True	1	1	0.001
PcanaA	CTA	False	False	6	-0.417460	3	5	True	1	1	0.001
WpijpA	CTA	False	False	29	-0.830207	6	28	True	1	1	0.000
BcaliA	TTA	False	False	346	-0.980015	13	319	False	1	1	0.000
CsalmA	TTA	False	False	294	-0.949769	12	228	False	4	4	0.000
AgossA	CTA	True	True	24	1.987008	5	2	True	4	13	0.999
	CTT	True	True	15	2.165922	4	0	True	4	10	0.998
ScereA	CTA	False	False	11	-0.069324	5	2	True	1	1	0.000
AgossB	CTA	True	True	24	1.987008	5	2	True	4	13	0.999
	CTT	True	True	15	2.165922	4	0	True	4	10	0.998
CglabA	CTT	False	False	1	2.961165	1	0	True	1	1	0.120

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
	TTA	False	False	192	-0.958602	7	178	False	1	1	0.000
SoctoA	TTA	False	False	159	-0.939502	7	112	False	2	2	0.000
OanguA	TTA	False	False	306	-0.985083	13	264	False	1	1	0.051
NdelpA	CTT	False	False	2	1.131707	2	0	True	1	1	0.029
	TTA	False	False	187	-0.955745	7	171	False	1	1	0.000
ScryoA	CTT	False	False	22	-0.887582	6	16	True	1	1	0.000
	TTA	False	False	136	-0.924592	7	97	False	1	1	0.000
SarboA	CTA	False	False	11	-0.069324	5	2	True	1	1	0.000
	CTA	False	False	23	-0.124791	6	1	True	1	1	0.000
HuvarA	TTA	False	False	166	-0.928291	6	122	False	1	1	0.000
	CTT	False	False	2	1.436893	1	0	True	1	1	0.042
ScastA	TTA	False	False	193	-0.978092	7	169	False	1	1	0.001
CzempA	TTA	False	False	193	-0.966416	7	141	False	1	1	0.000
SpombA	TTA	False	False	157	-0.937090	7	108	False	2	2	0.000