

CoReTracker report

		Clad. Val	Reassigned	Tot. count	Telford	Tot. Genes	Use. count	Trans. Val	Rea. Genes	Rea. count	Prob
BarxA	AAT	False	False	18	0.058586	1	14	False	1	1	0.042
MmagnB	AAC	False	False	74	-0.081683	12	62	False	1	1	0.000
MmagnA	AAC	False	False	74	-0.058172	12	63	False	1	1	0.000
MingeA	AAT	False	False	80	-0.150061	12	64	False	1	1	0.000
MstarA	AAT	False	False	79	-0.123336	12	66	False	1	1	0.000
CsalmA	AAT	False	False	75	-0.166288	10	51	False	1	1	0.000
BcustA	AAT	False	False	68	-0.116427	11	57	False	1	1	0.000
CsantA	AAC	False	False	29	-0.232283	9	19	False	1	1	0.000
Mtetra	AAC	False	False	81	-0.103672	12	68	False	1	1	0.000
SjapoA	AAT	False	False	70	-0.387587	7	36	False	1	1	0.000
SpombA	AAC	False	False	8	-0.231051	4	1	False	1	1	0.000
	AAT	False	False	58	-0.087193	6	40	False	1	2	0.000
CzempA	AAT	False	False	64	-0.185108	7	45	False	1	1	0.000
SoctoA	AAT	False	False	53	-0.019859	6	33	False	3	3	0.000
PkluyA	AAC	False	False	25	0.107475	9	19	False	1	1	0.000
OanguA	AAT	False	False	68	0.002891	12	53	False	1	1	0.000
NdelpA	AAT	False	False	51	-0.046551	7	45	False	1	1	0.000
ScryoA	AAC	False	False	25	-0.077422	7	14	False	2	2	0.000
	AAT	False	False	41	-0.120290	6	27	False	1	1	0.000