

AFF1

Ex-L5or6-CC

Posterior
inclusion
probability

1.00
0.75
0.50
0.25
0.00

eQTL
p-value

1.0e+00
3.2e-03
1.0e-05
3.2e-08
1.0e-10

87,000

87,500

88,000

88,500

89,000

genomic location (kb) on chr4

SMC

87,000

87,500

88,000

88,500

89,000