

Accuracy 10000 iterations LGG 1.0 -0.9 ngenes 91 101 114 0.8 -130 accuracy 152 182 227 0.7 -303 455 910 1819 0.6 -0.5 -DNA\_methylation\_clusters -

variable

Accuracy 10000 iterations GBM+LGG 1.0 -0.9 ngenes 91 101 114 0.8 -130 accuracy 152 182 0.7 -227 303 455 910 0.6 -1819 0.5 -IDH\_TOT-DNA\_methylation\_clusters variable