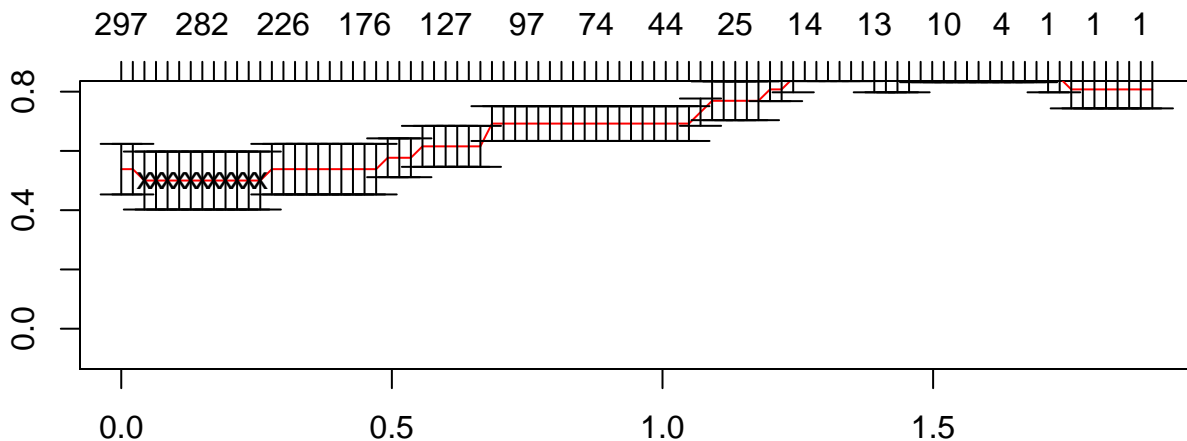
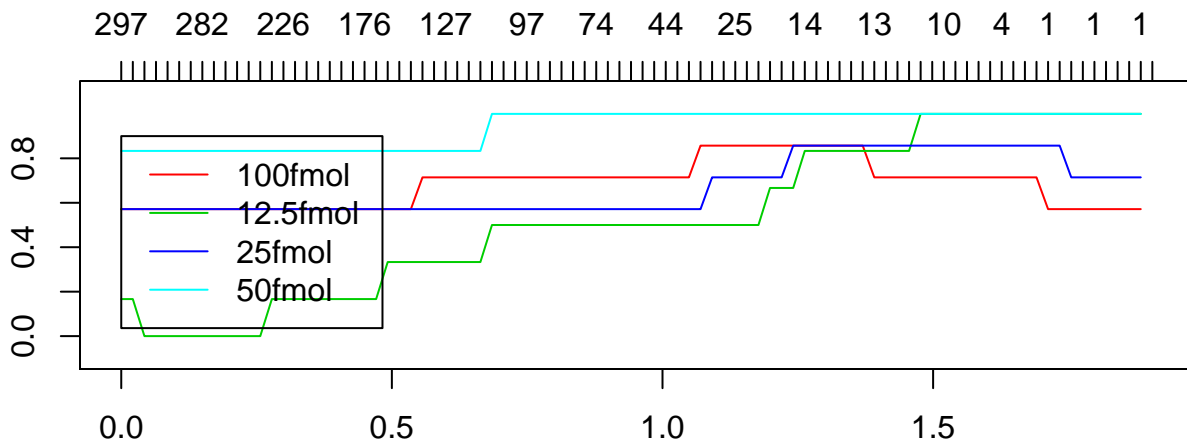


Number of genes

Misclassification Error



Misclassification Error



Value of threshold