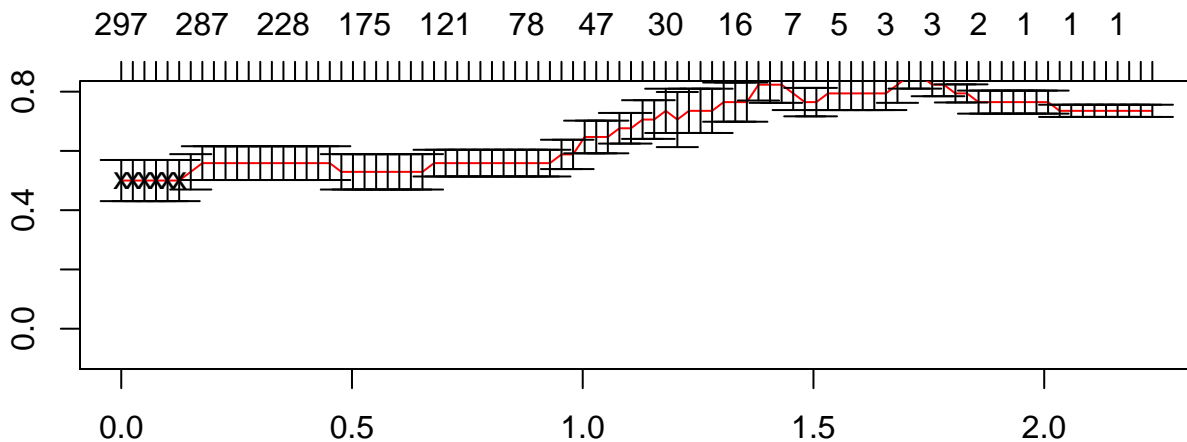


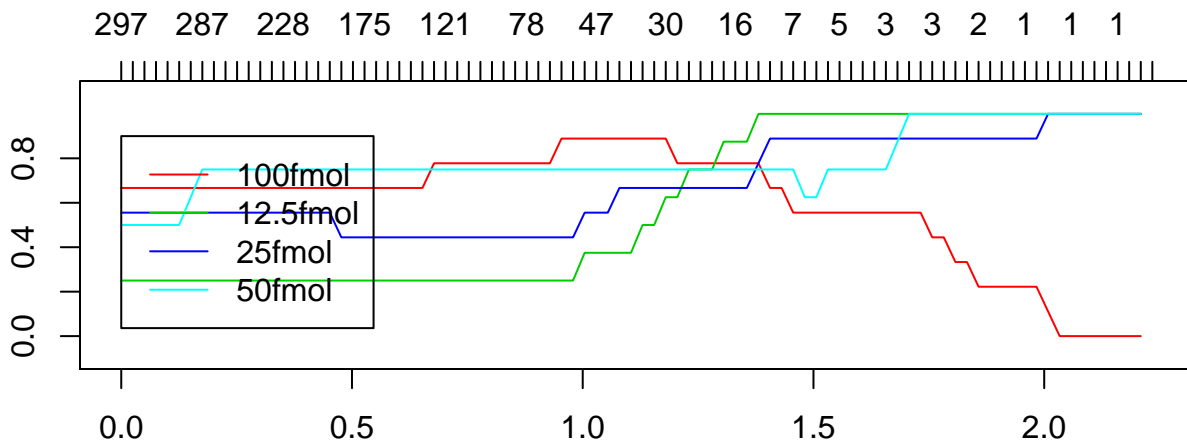
Number of genes

Misclassification Error



Value of threshold

Misclassification Error



Value of threshold