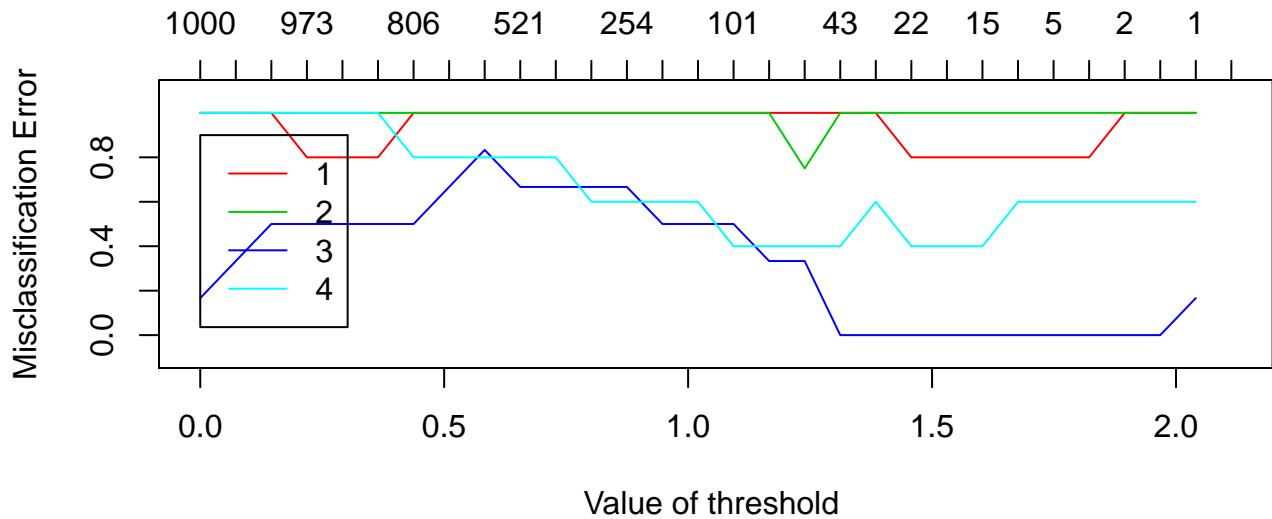
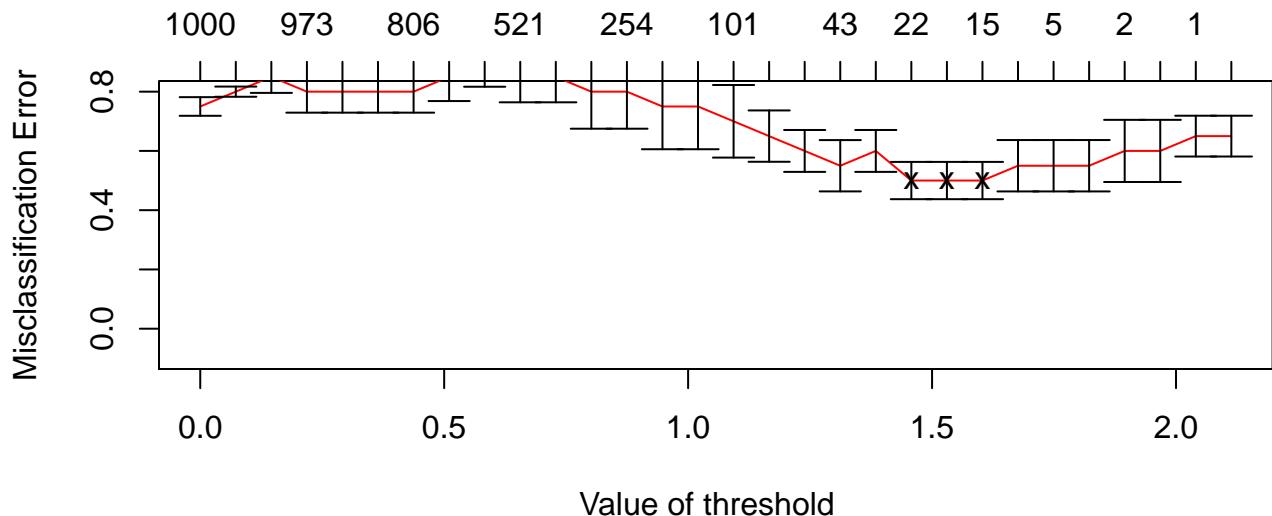


Number of genes



cross-validated probabilities

1.0
0.8
0.6
0.4
0.2
0.0

1

2

3

4

5 10 15 20

sample

