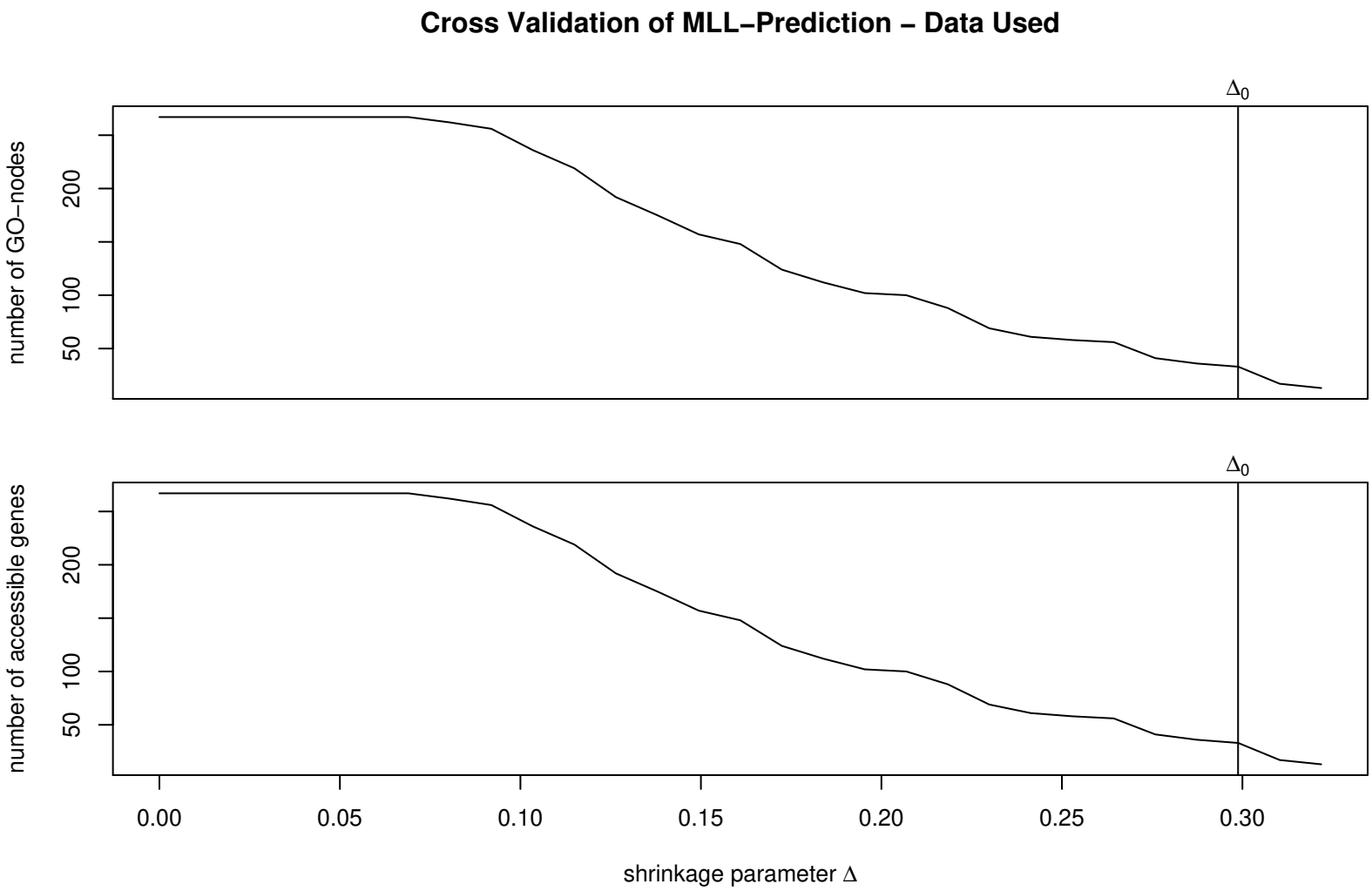
Cross Validation of MLL−Prediction − Data Used
Δ_0
number of GO−nodes
50
100
200
Δ_0
number of accessible genes
50
100
200
0.00
0.05
0.10
0.15
0.20
0.25
0.30
shrinkage parameter Δ