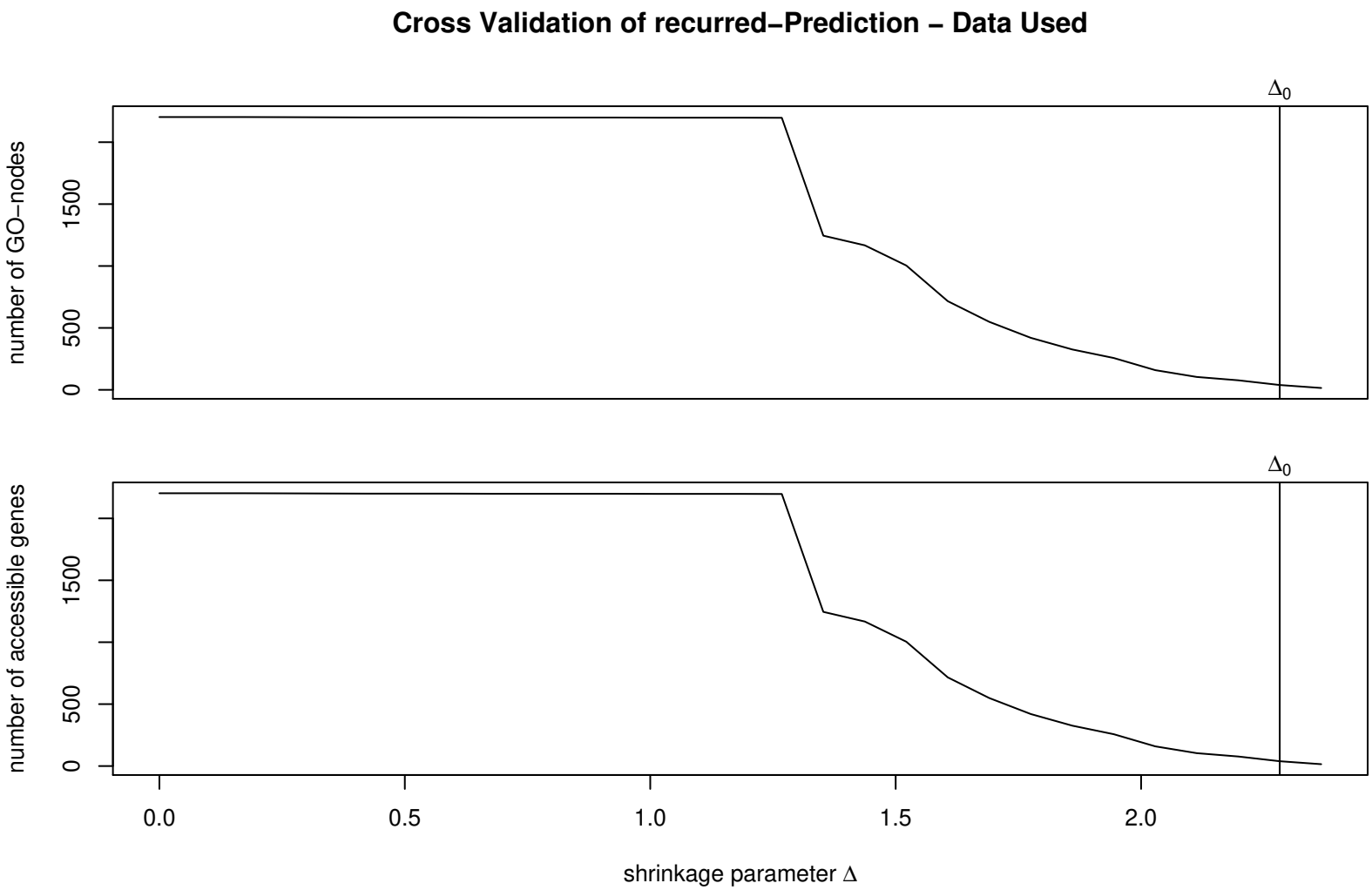
Cross Validation of recurred−Prediction − Data Used
number of GO−nodes
0
500
1500
Δ_0
number of accessible genes
0
500
1500
Δ_0
0.0
0.5
1.0
1.5
2.0
shrinkage parameter Δ