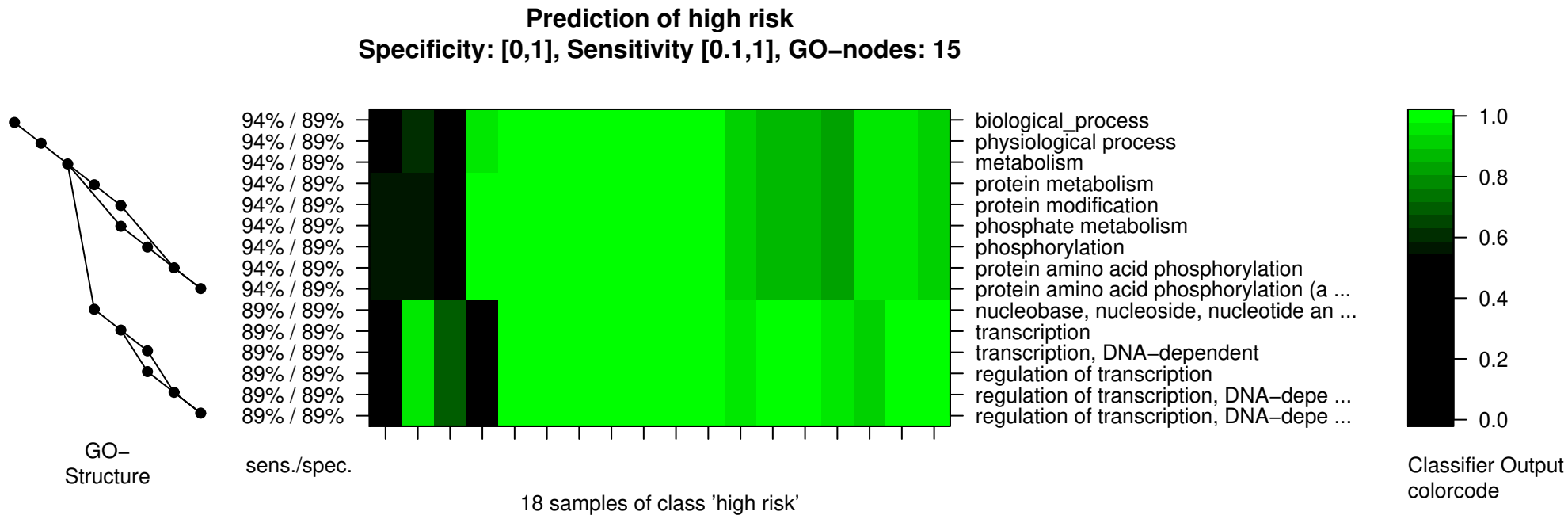
Prediction of high risk
Specificity: [0,1], Sensitivity [0.1,1], GO−nodes: 15
94% / 89%
94% / 89%
94% / 89%
94% / 89%
94% / 89%
94% / 89%
94% / 89%
94% / 89%
94% / 89%
89% / 89%
89% / 89%
89% / 89%
89% / 89%
89% / 89%
89% / 89%
biological_process
physiological process
metabolism
protein metabolism
protein modification
phosphate metabolism
phosphorylation
protein amino acid phosphorylation
protein amino acid phosphorylation (a ...
nucleobase, nucleoside, nucleotide an ...
transcription
transcription, DNA−dependent
regulation of transcription
regulation of transcription, DNA−depe ...
regulation of transcription, DNA−depe ...
1.0
0.8
0.6
0.4
0.2
0.0
GO−
Structure
sens./spec.
18 samples of class 'high risk'
Classifier Output
colorcode