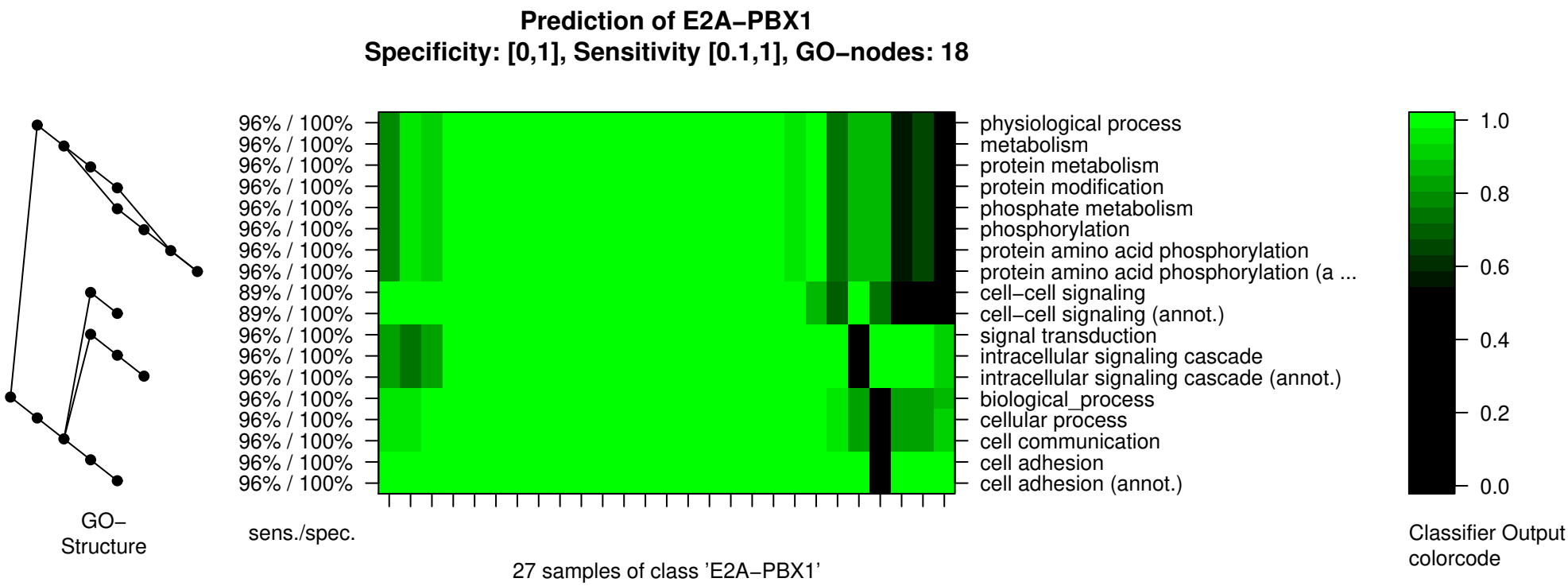
Prediction of E2A−PBX1
Specificity: [0,1], Sensitivity [0.1,1], GO−nodes: 18
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
89% / 100%
89% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
96% / 100%
physiological process
metabolism
protein metabolism
protein modification
phosphate metabolism
phosphorylation
protein amino acid phosphorylation
protein amino acid phosphorylation (a ...
cell−cell signaling
cell−cell signaling (annot.)
signal transduction
intracellular signaling cascade
intracellular signaling cascade (annot.)
biological_process
cellular process
cell communication
cell adhesion
cell adhesion (annot.)
GO−
Structure
sens./spec.
27 samples of class 'E2A−PBX1'
1.0
0.8
0.6
0.4
0.2
0.0
Classifier Output
colorcode