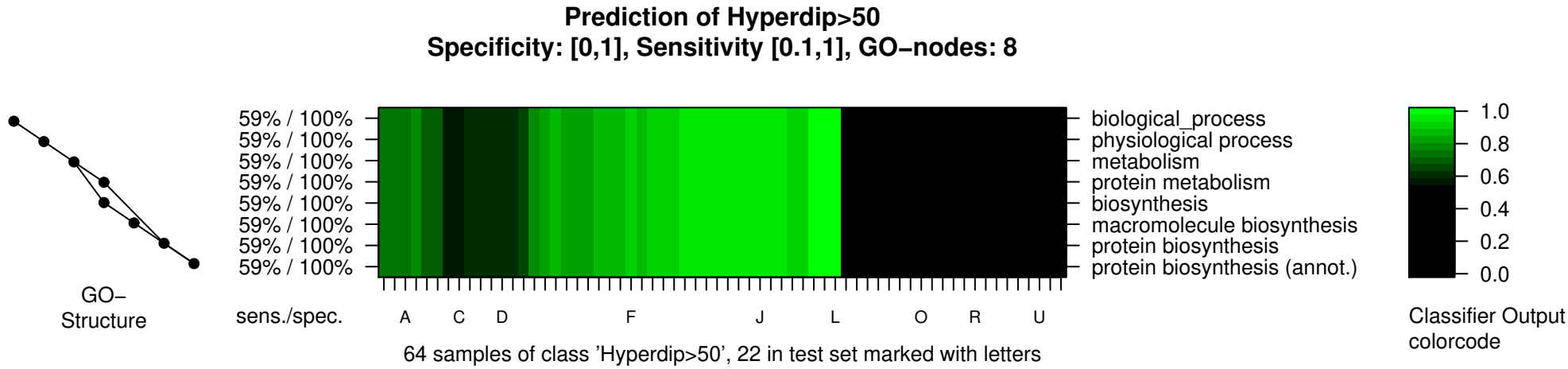
Prediction of Hyperdip>50
Specificity: [0,1], Sensitivity [0.1,1], GO−nodes: 8
GO−
Structure
59% / 100%
59% / 100%
59% / 100%
59% / 100%
59% / 100%
59% / 100%
59% / 100%
59% / 100%
sens./spec.
A
C
D
F
J
L
O
R
U
biological_process
physiological process
metabolism
protein metabolism
biosynthesis
macromolecule biosynthesis
protein biosynthesis
protein biosynthesis (annot.)
1.0
0.8
0.6
0.4
0.2
0.0
Classifier Output
colorcode
64 samples of class 'Hyperdip>50', 22 in test set marked with letters