ROC 10–fold cross–validation transcriptome classification (Accuracy = 70.59%) 1.00 -0.75 -Methylation class Classical (N = 14, AUC = 0.86) Mesenchymal (N = 13, AUC = 0.94) 0.50 -Neural (N = 7, AUC = 0.60) Proneural (N = 17, AUC = 0.91) 0.25 -0.00 -

0.75

1.00

0.50

False positive rate

True positive rate

0.25

0.00