ROC 10-fold cross-validation transcriptome classification (Accuracy = 72.55%) 1.00 -0.75 -Methylation class Classical (N = 14, AUC = 0.86) Mesenchymal (N = 13, AUC = 0.94) 0.50 -Neural (N = 7, AUC = 0.67) Proneural (N = 17, AUC = 0.92) 0.25 -0.00 -

0.75

1.00

True positive rate

0.25

0.00

0.50

False positive rate