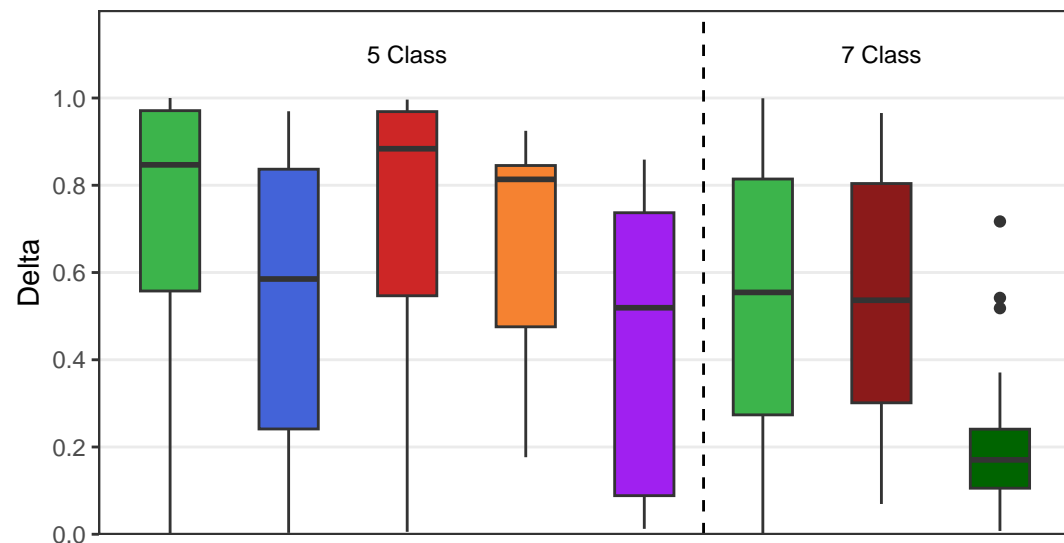
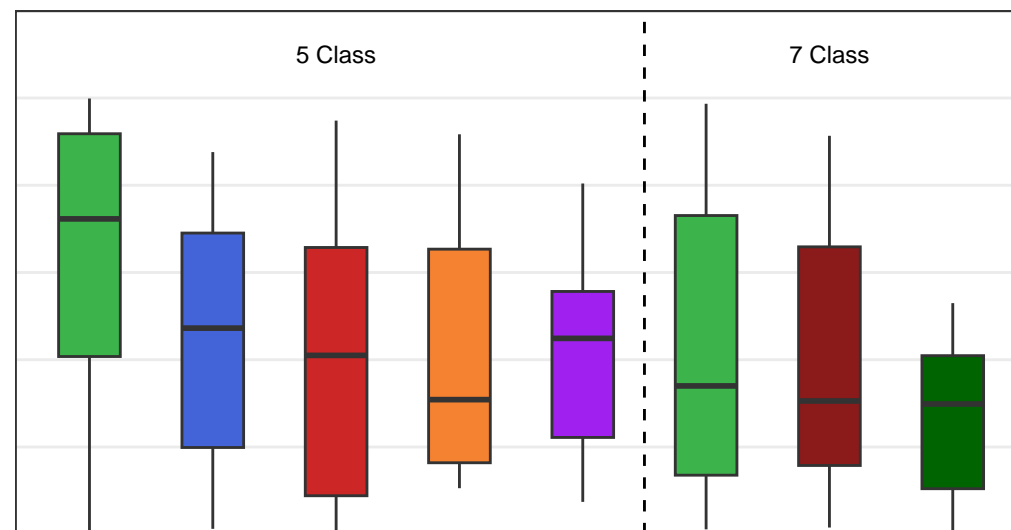


**A**

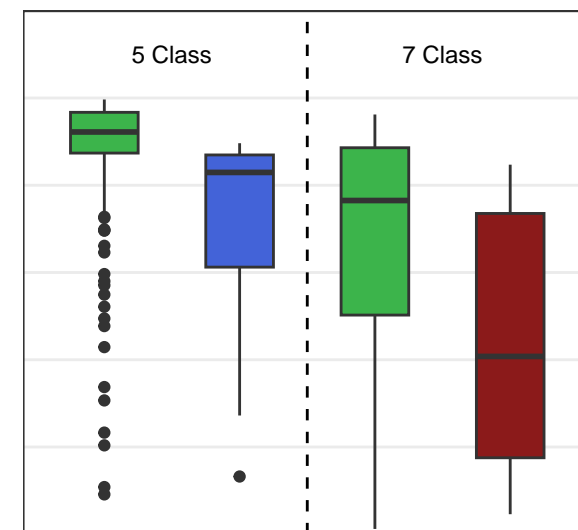
TCGA

**B**

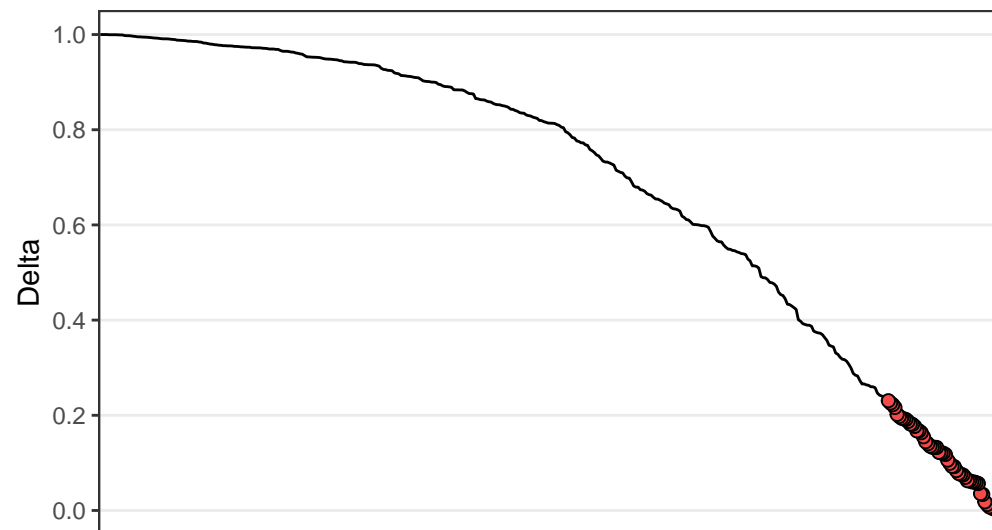
UC Genome

**C**

Leeds

**D**

Prediction-calls Delta Distribution

**E**

Library Size With Ambiguous Prediction Calls (TCGA)

