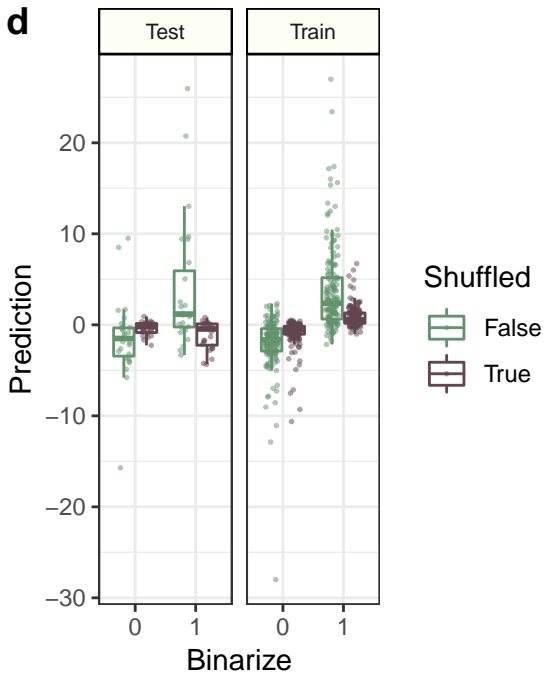
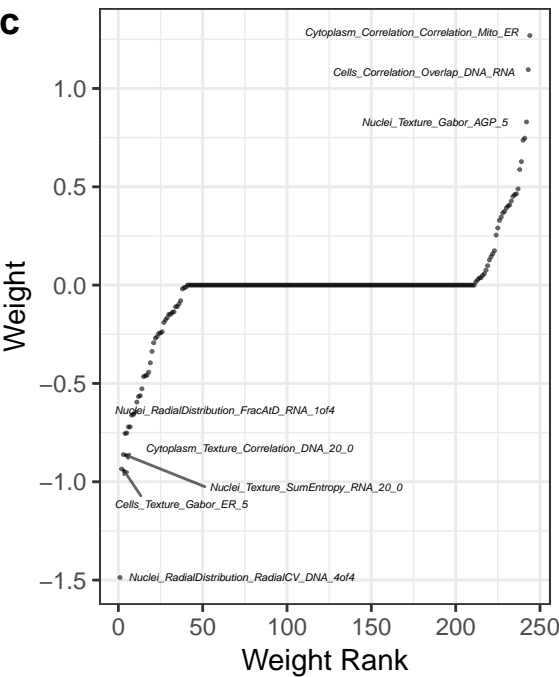
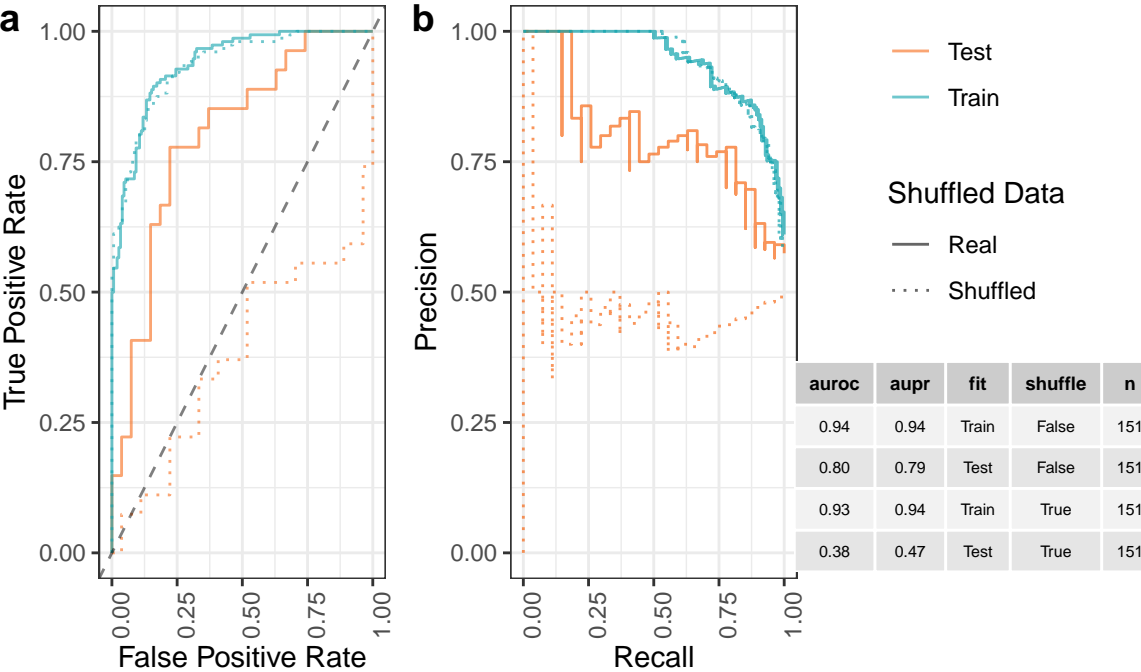
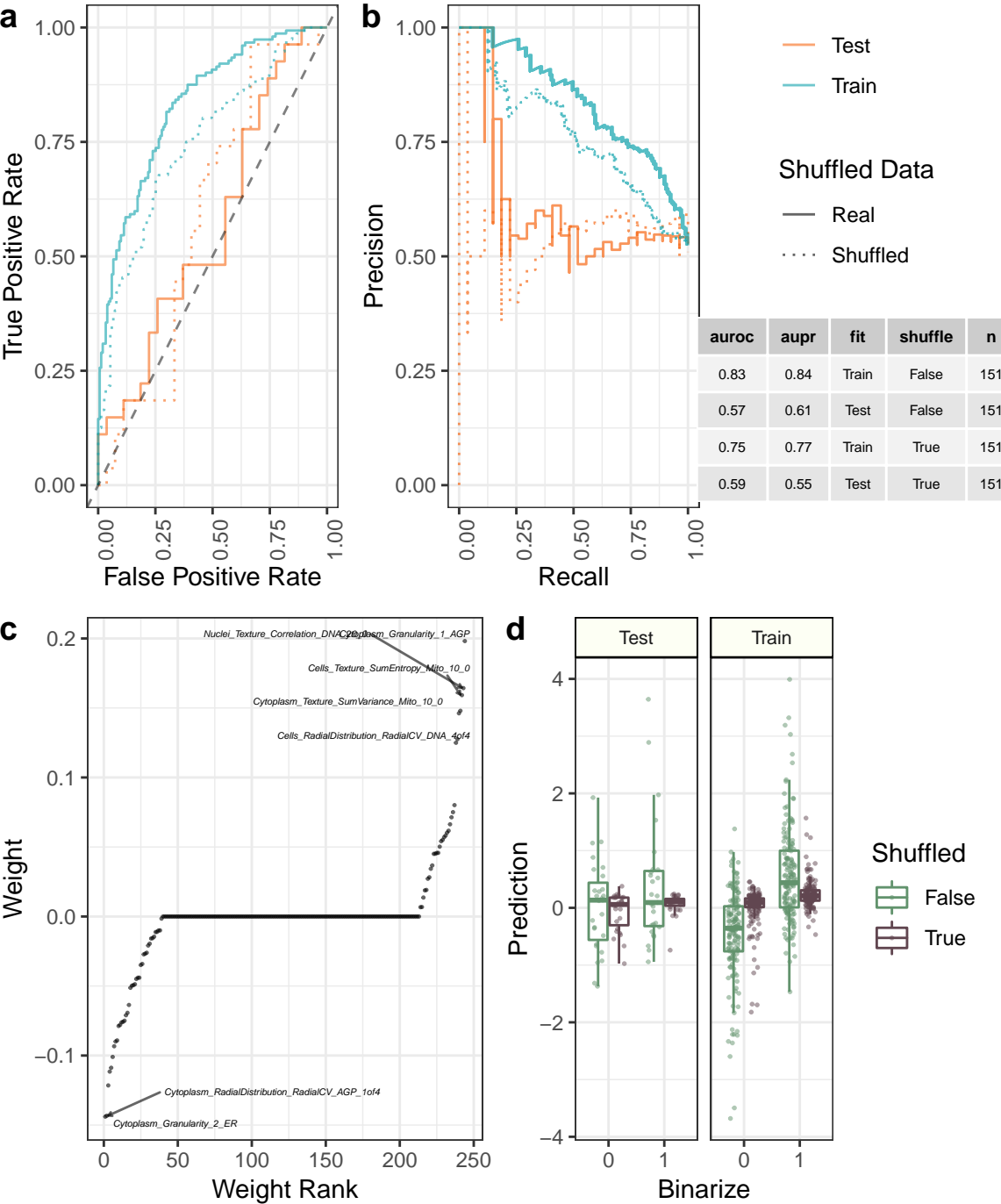


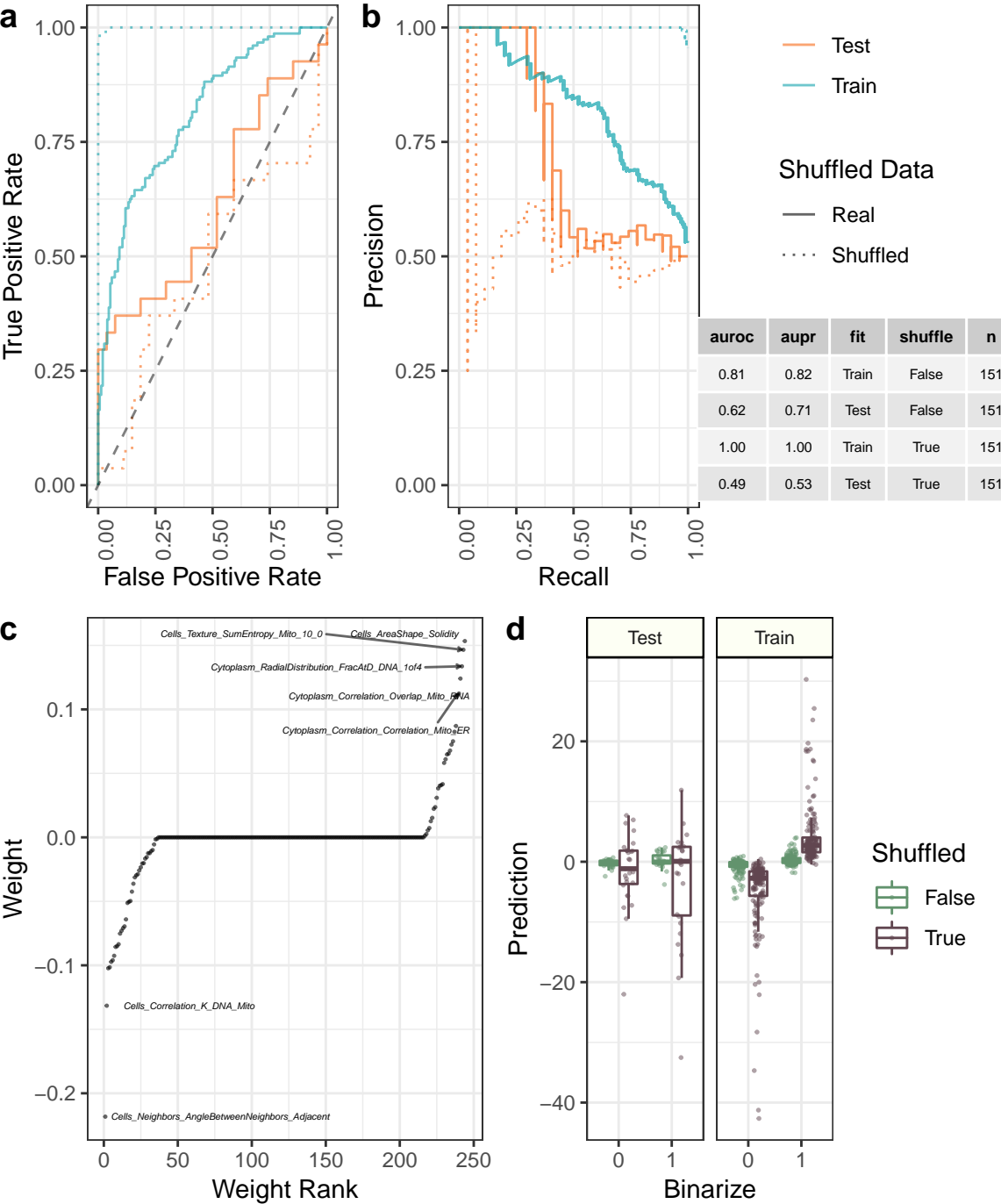
Performance: cc\_all\_high\_n\_spots\_h2ax\_mean



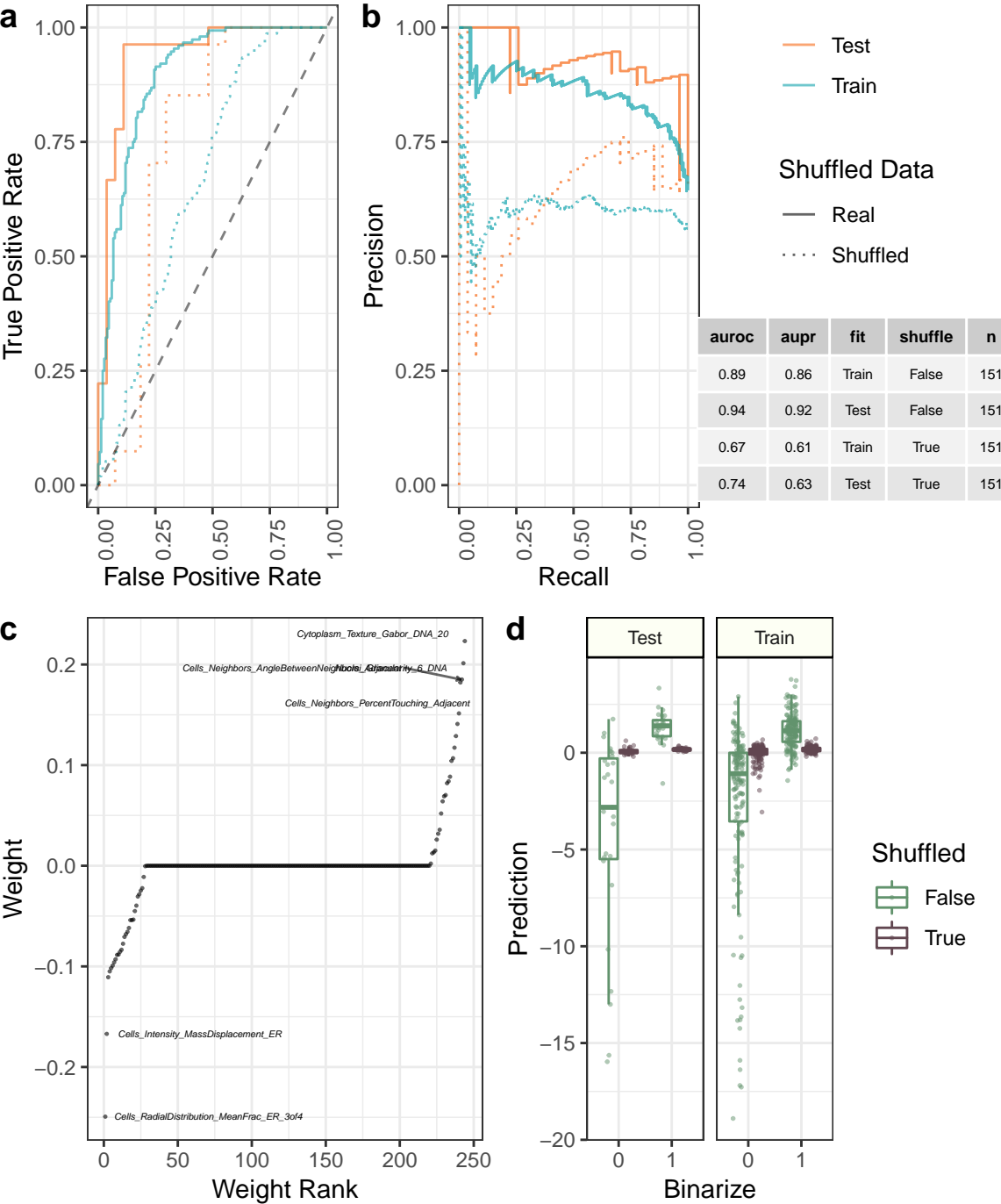
Performance: cc\_all\_large\_notround\_polynuclear\_mean



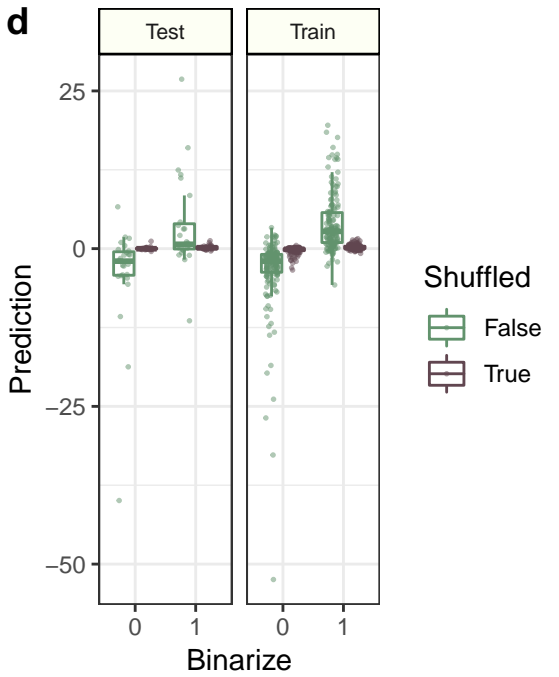
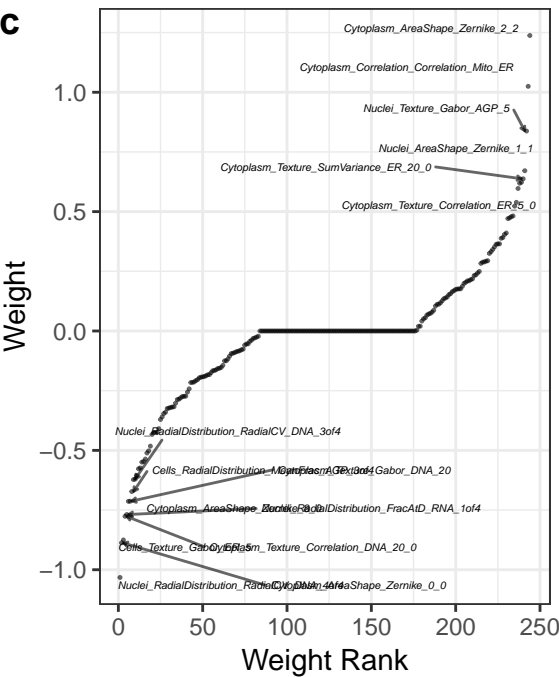
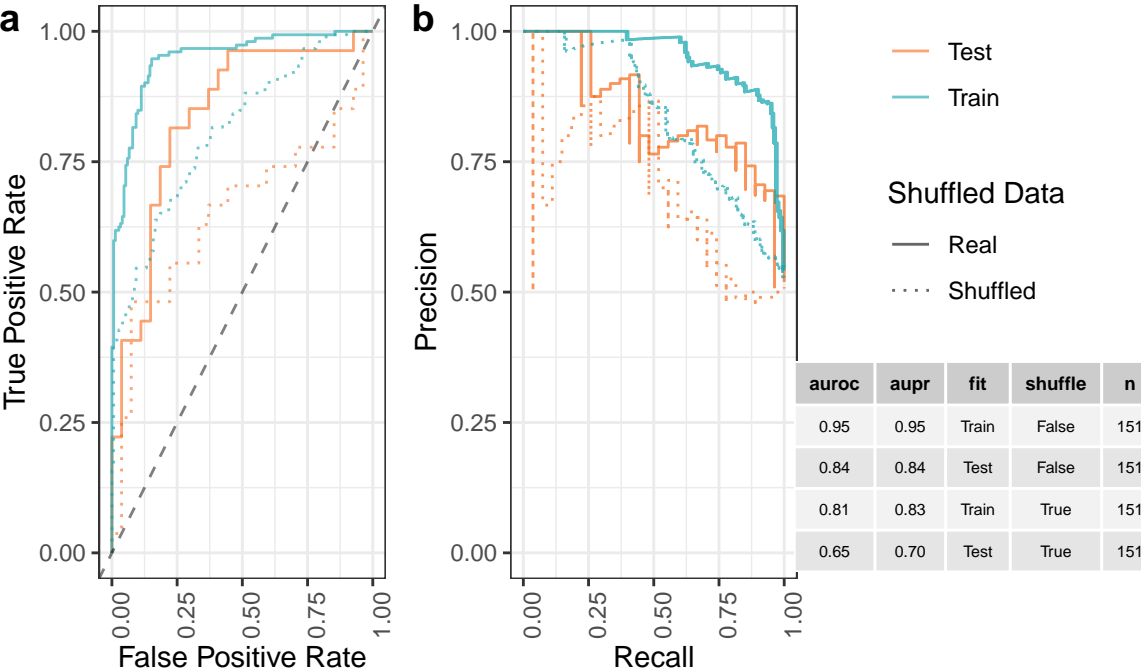
Performance: cc\_all\_large\_round\_polyloid\_mean



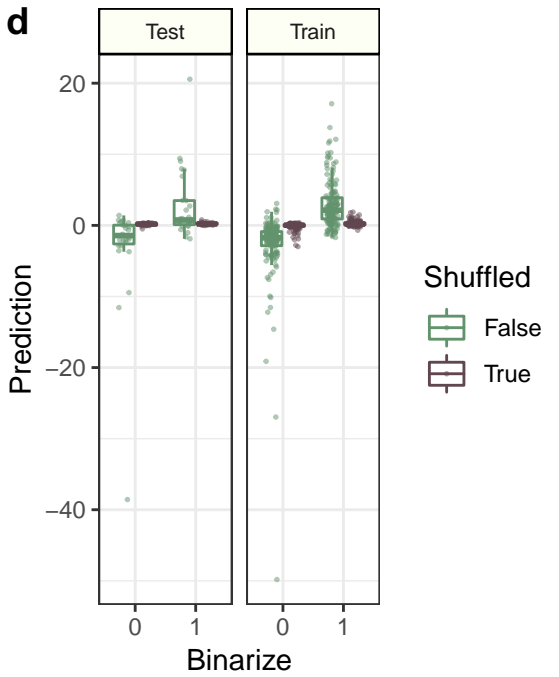
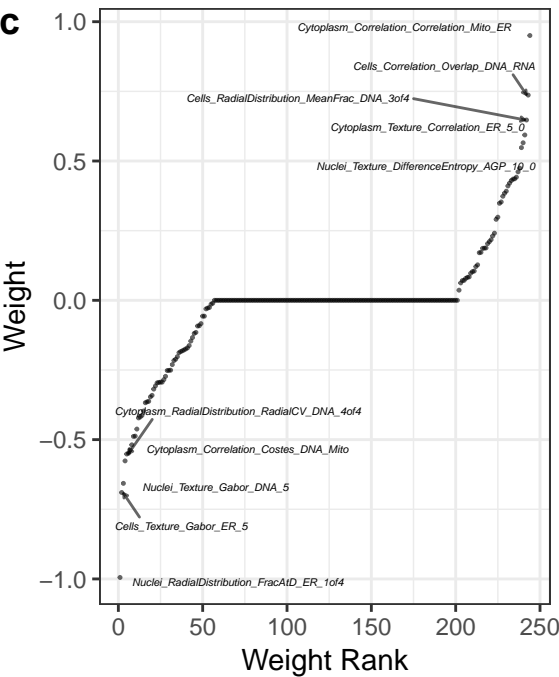
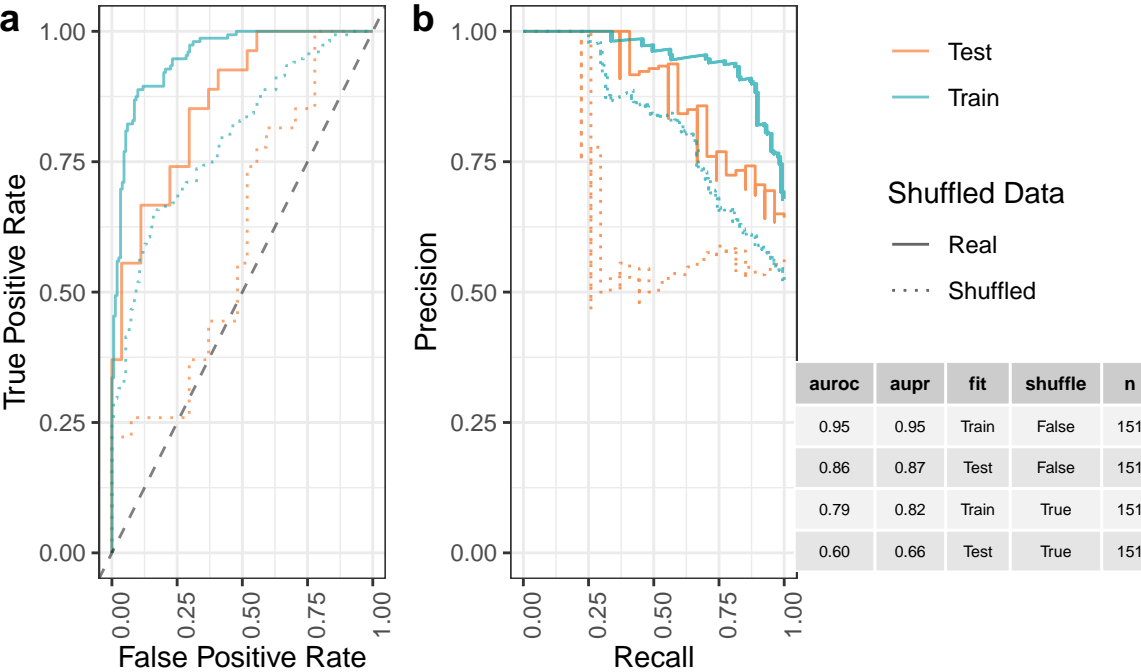
# Performance: cc\_all\_n\_objects



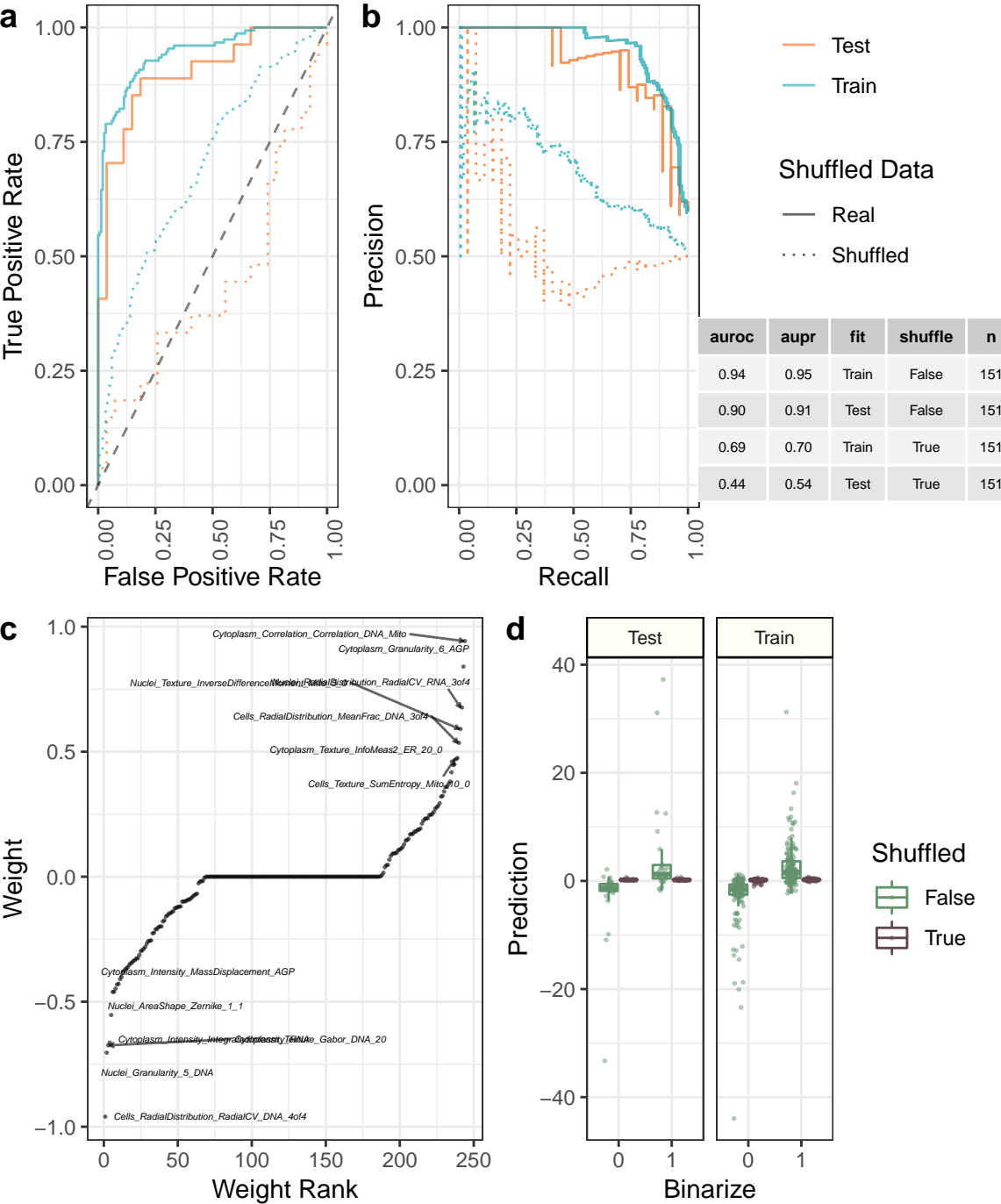
Performance: cc\_all\_n\_spots\_mean



Performance: cc\_all\_n\_spots\_per\_nucleus\_area\_mean



Performance: cc\_all\_nucleus\_area\_mean



**c**

Weight

Weight Rank

**d**

Prediction

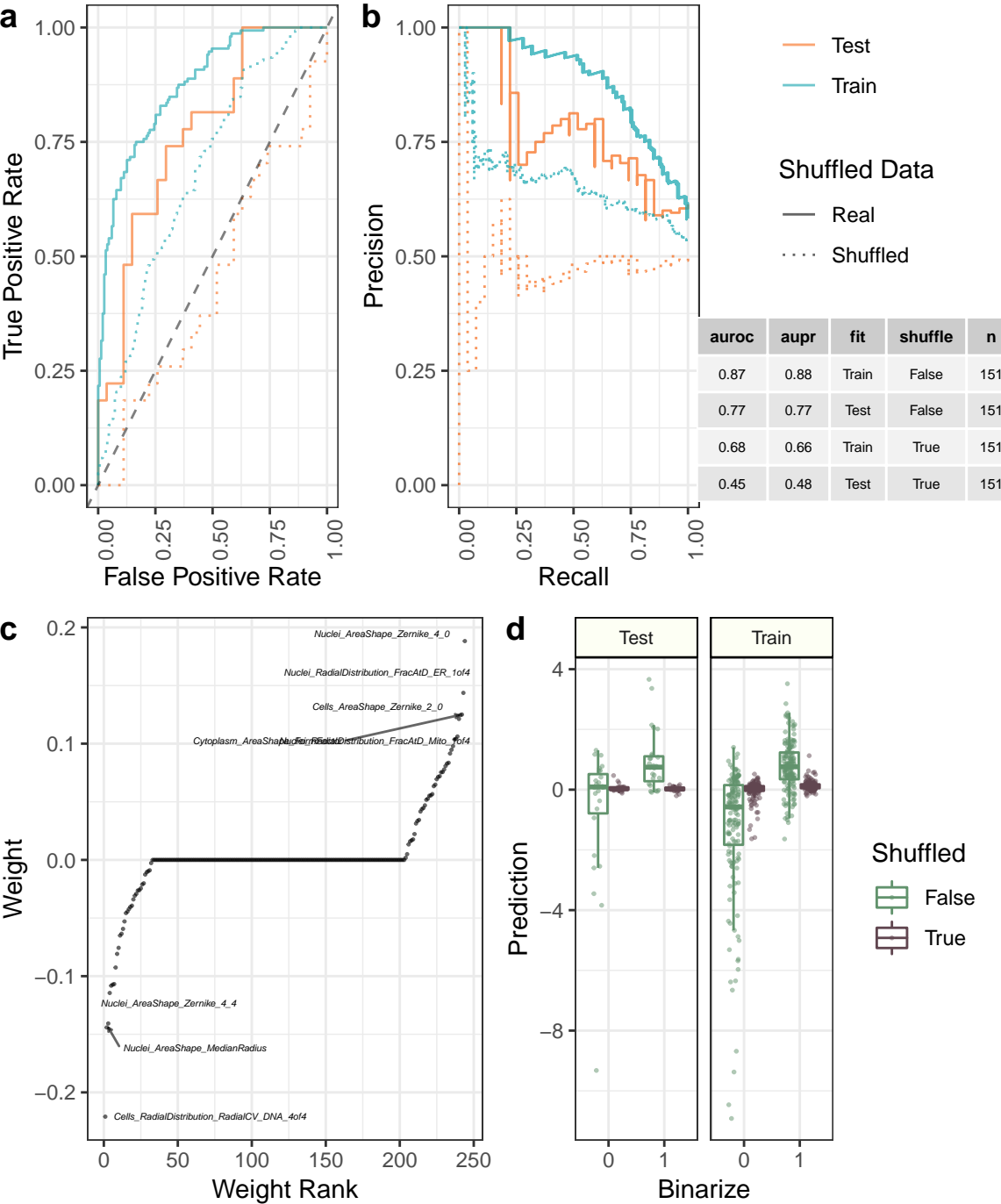
Binarize

Shuffled

False

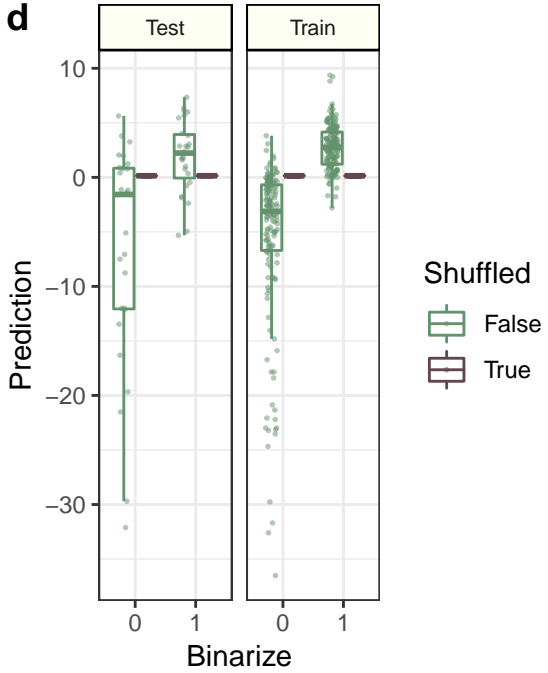
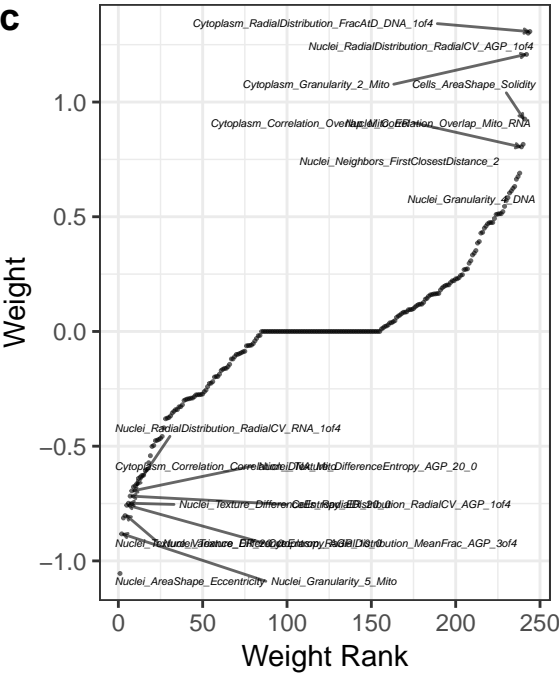
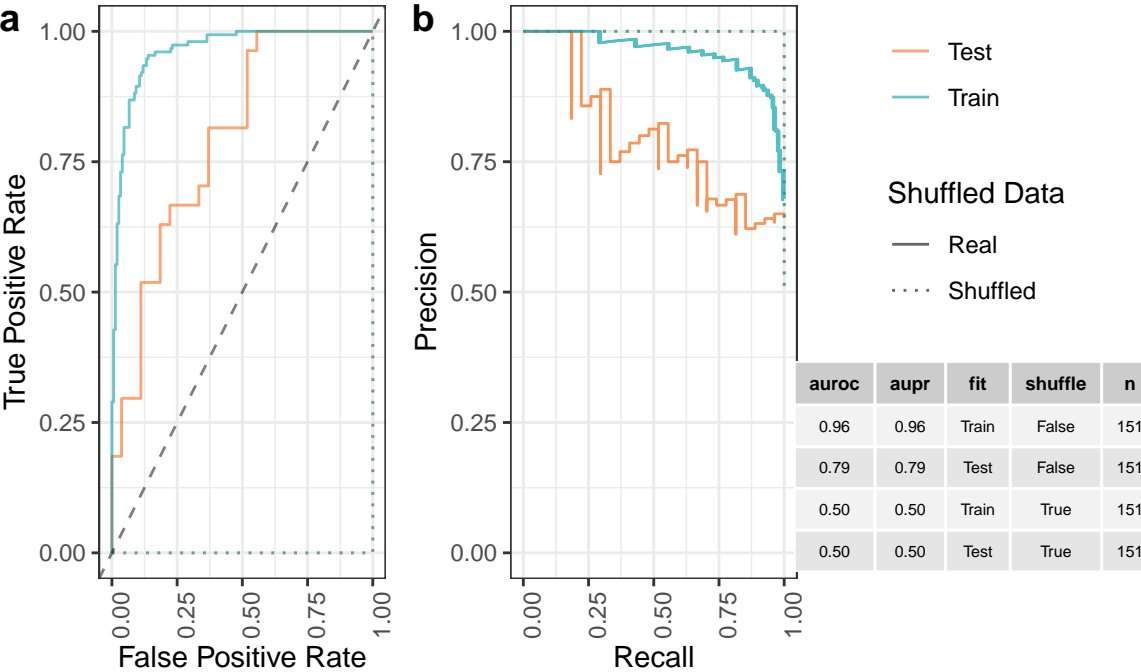
True

Performance: cc\_all\_nucleus\_roundness\_mean

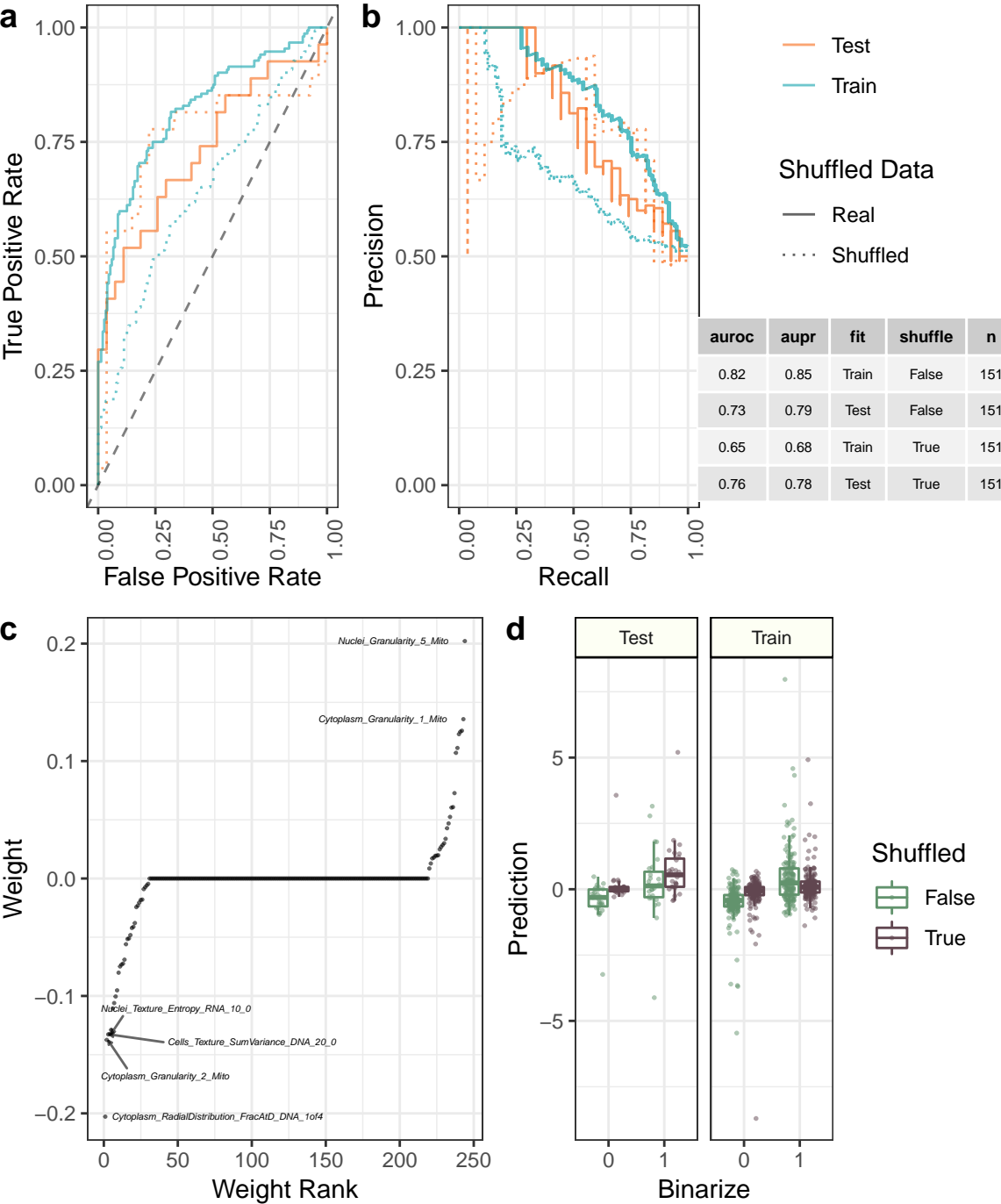




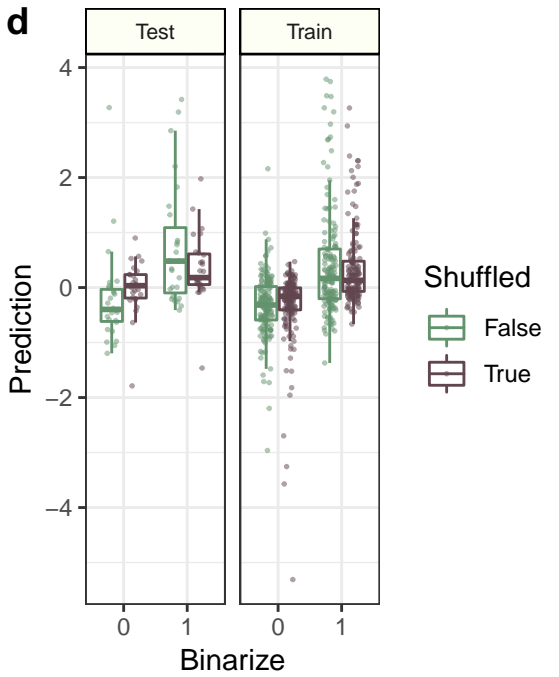
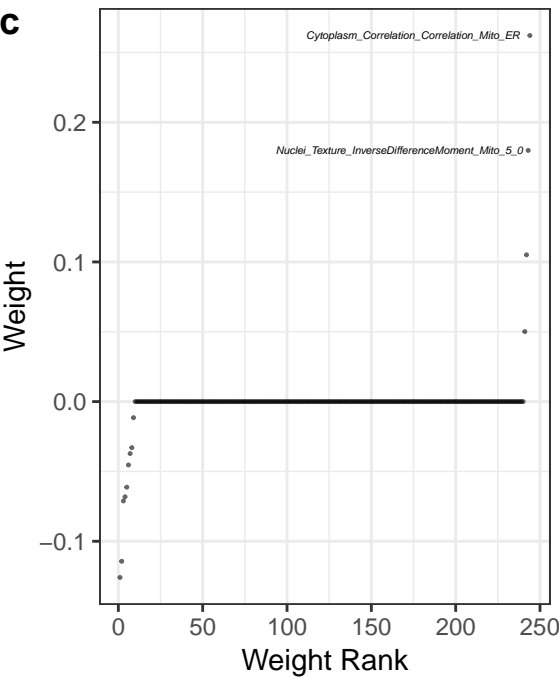
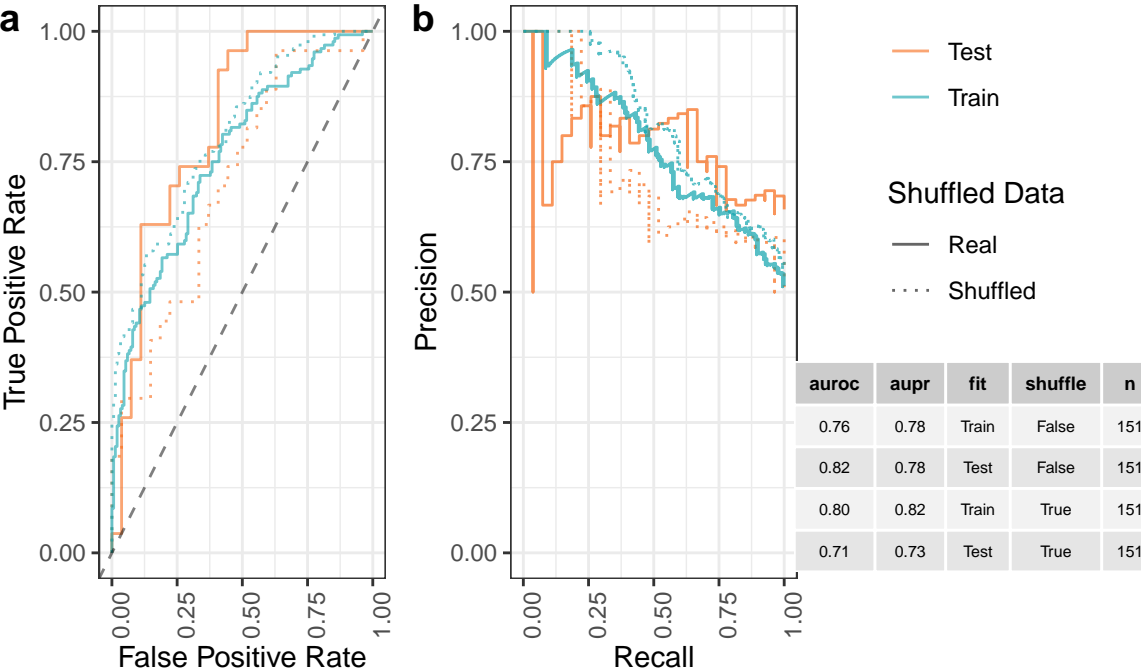
Performance: cc\_cc\_edu\_pos\_mean



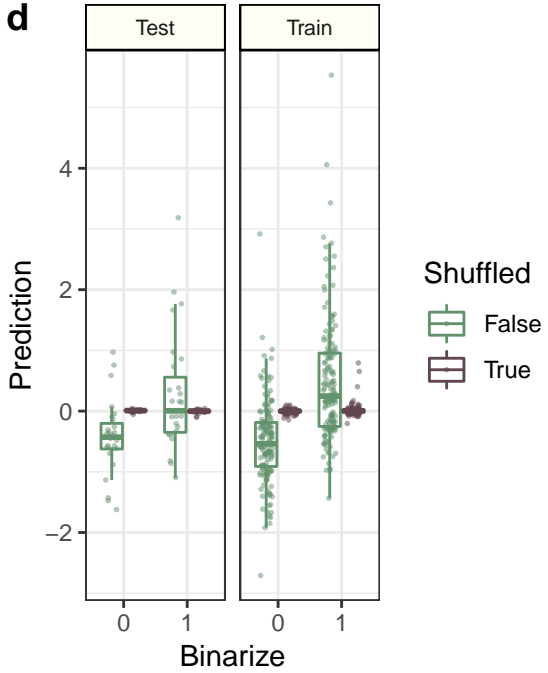
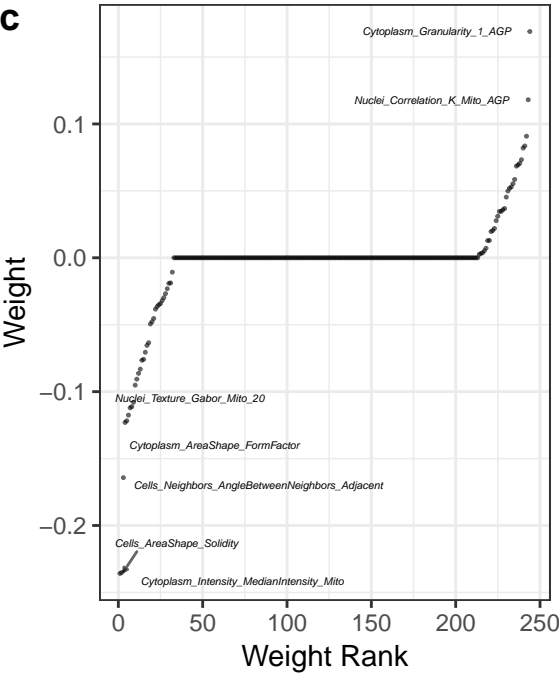
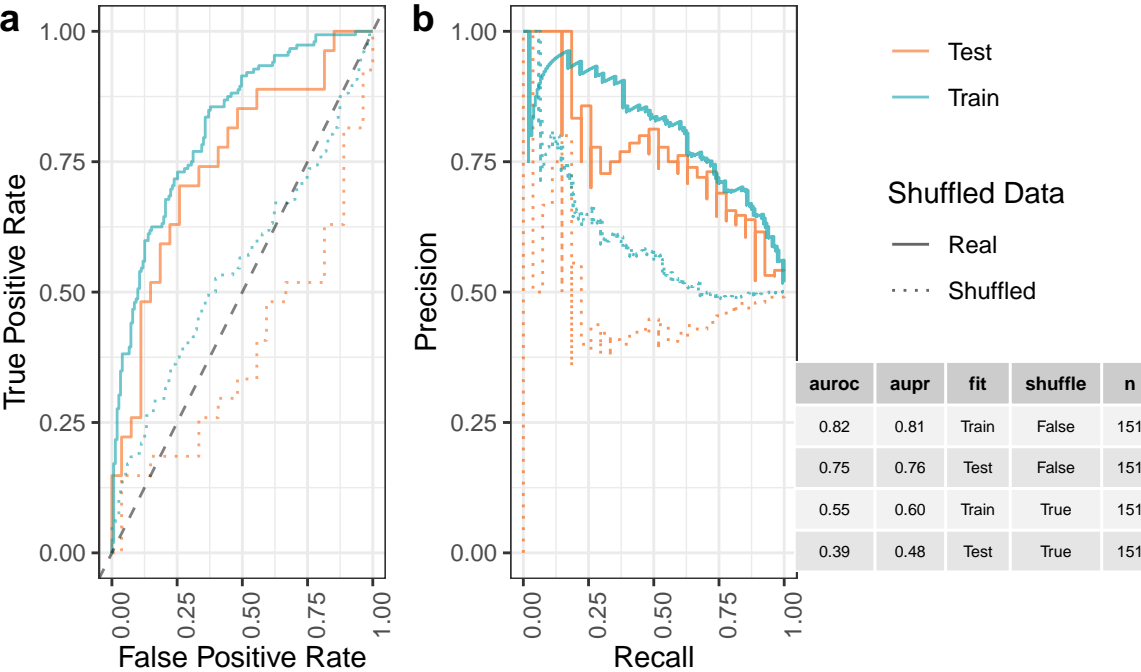
# Performance: cc\_cc\_g1\_mean



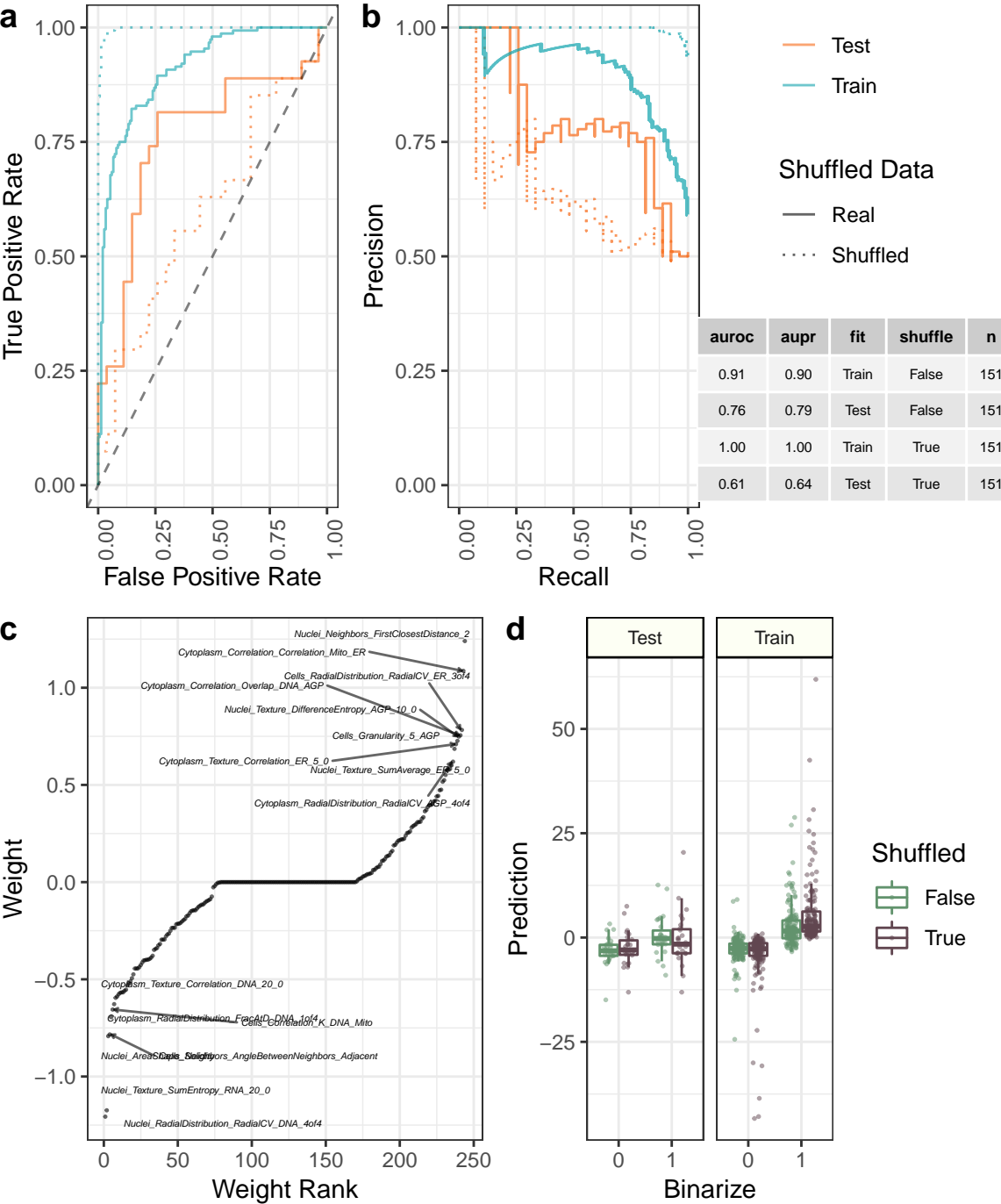
Performance: cc\_cc\_g2\_ph3\_neg\_mean



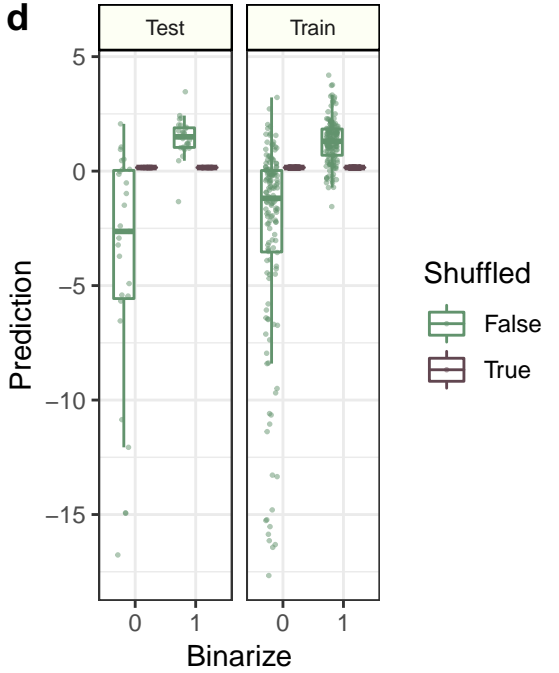
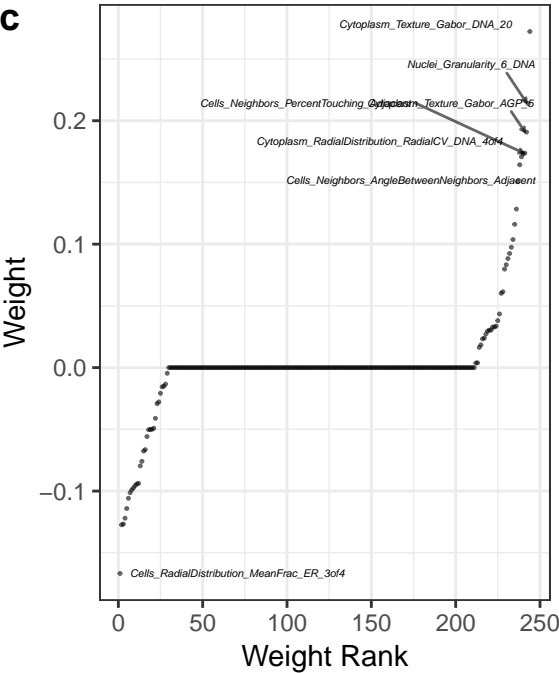
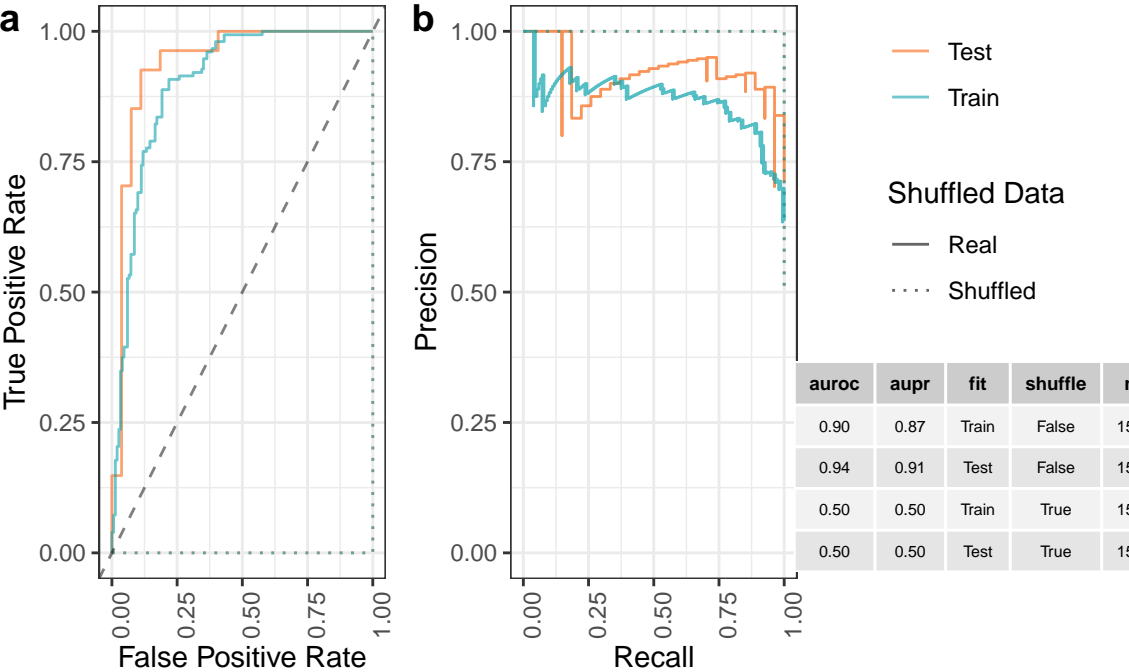
Performance: cc\_cc\_g2\_ph3\_pos\_early\_mitosis\_mean



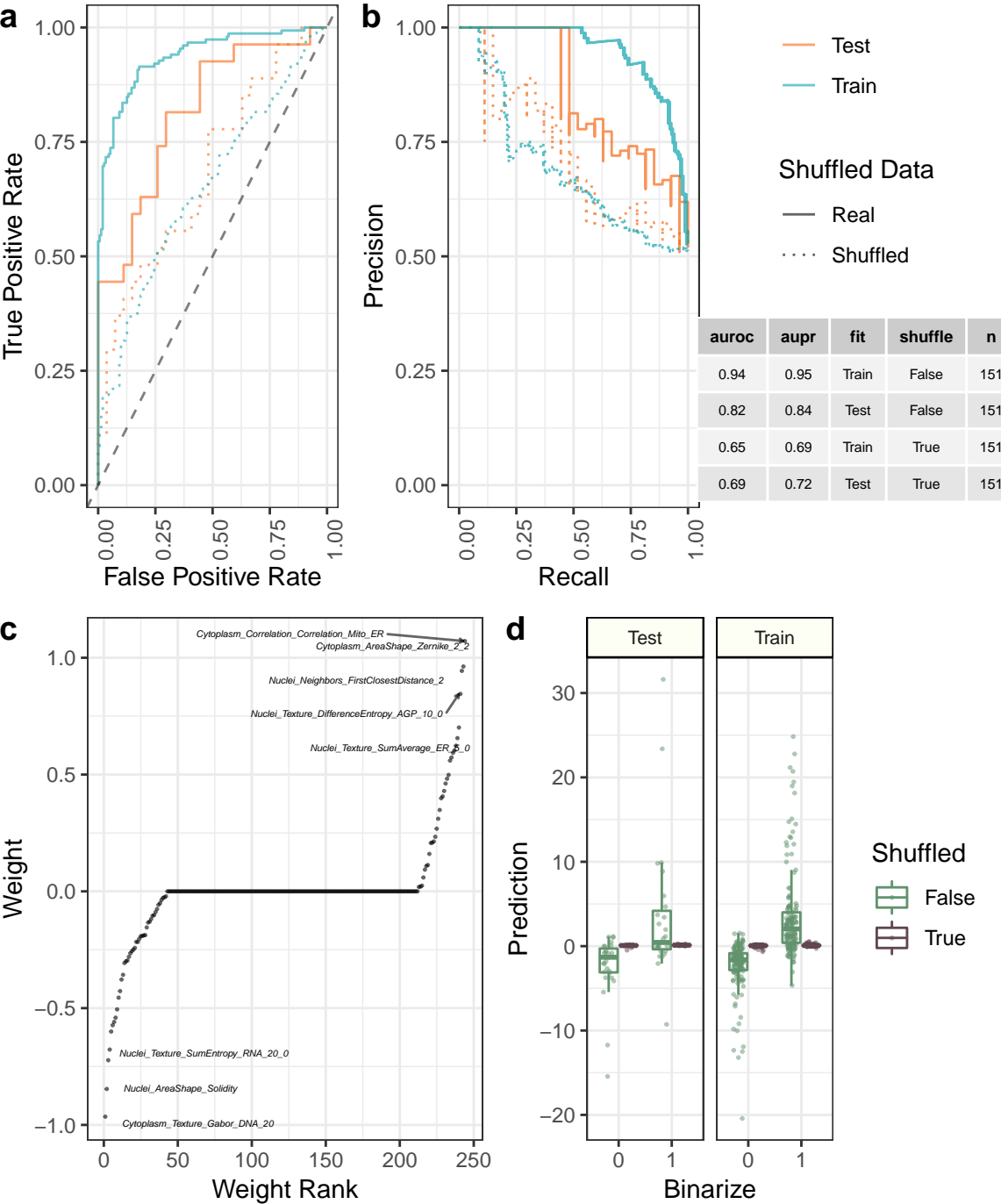
Performance: cc\_cc\_high\_n\_spots\_h2ax\_mean



# Performance: cc\_cc\_n\_objects



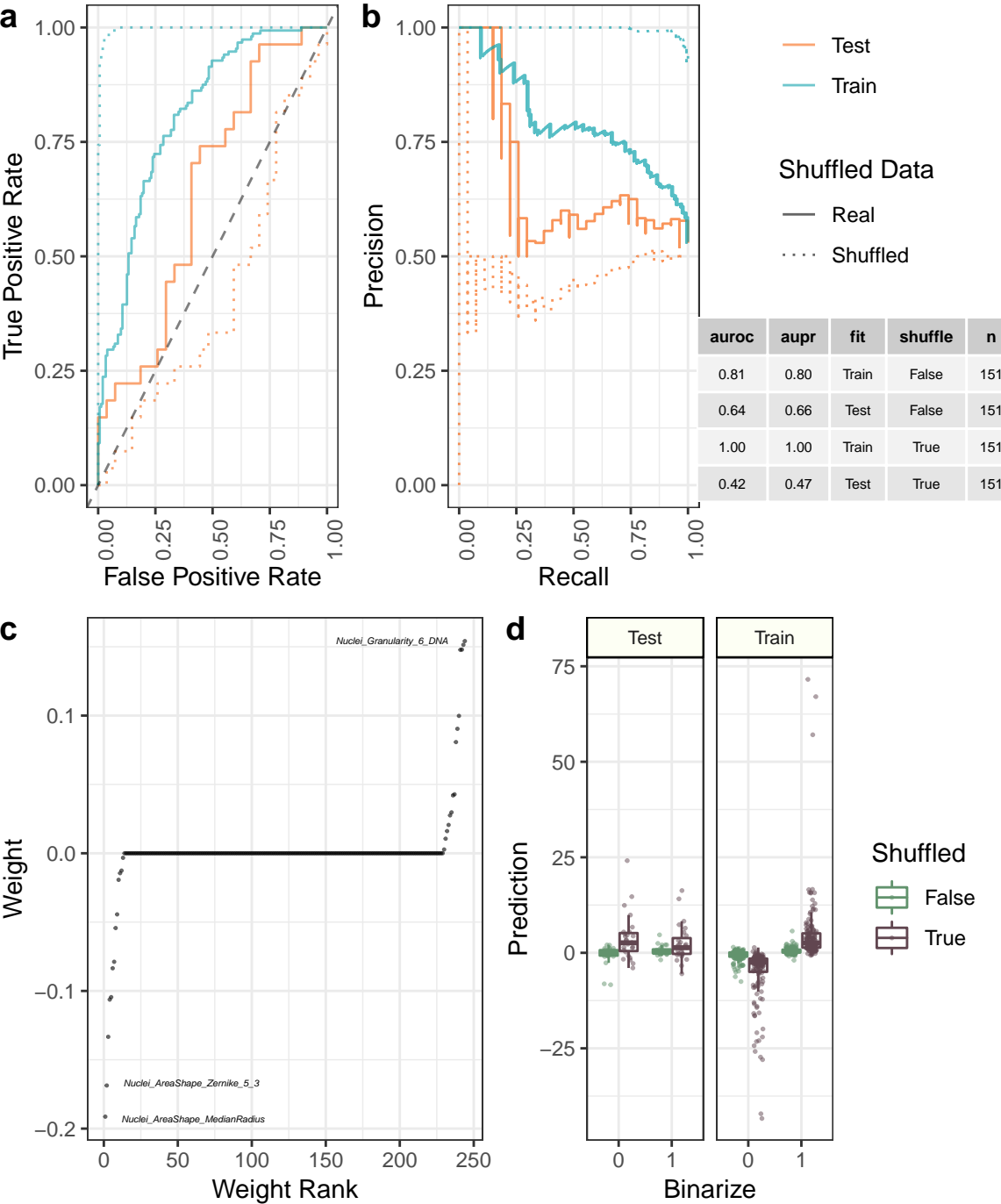
# Performance: cc\_cc\_n\_spots\_mean



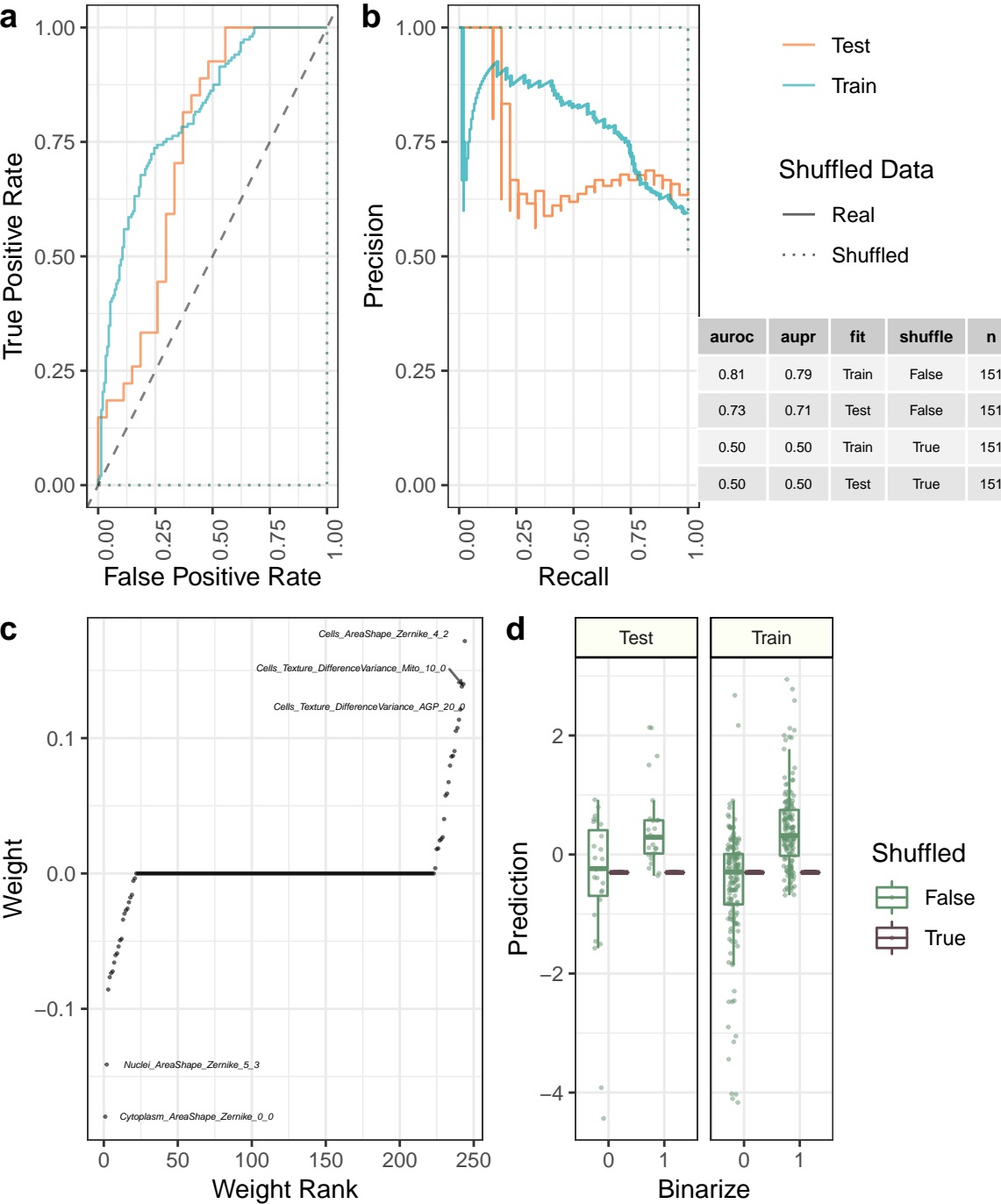




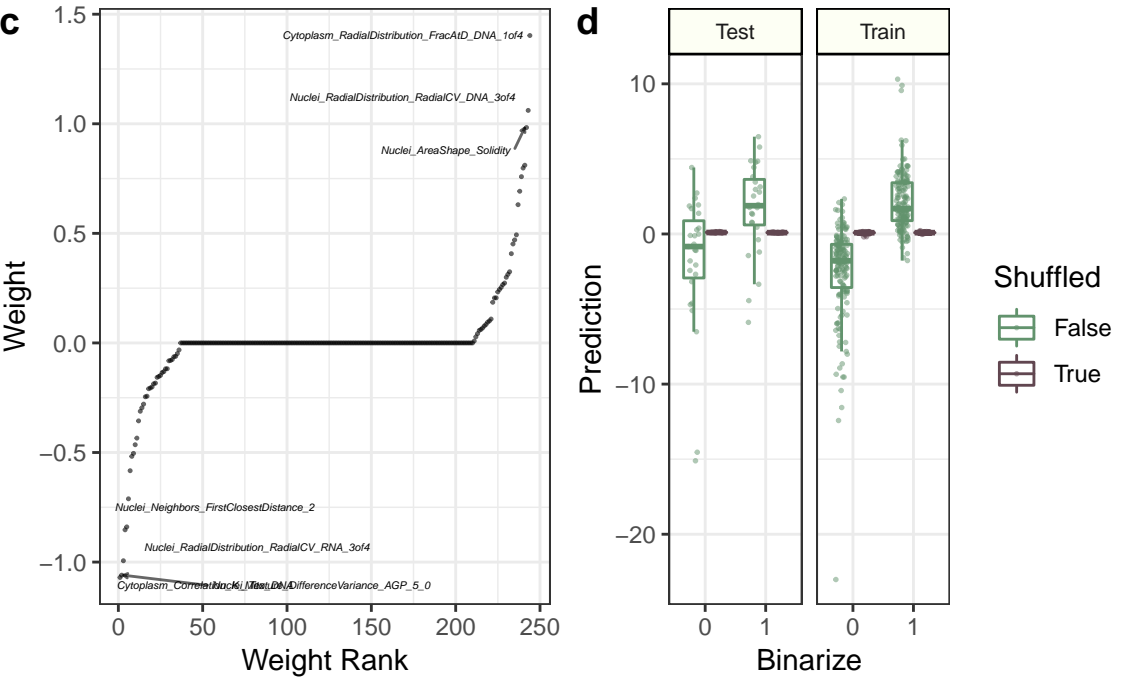
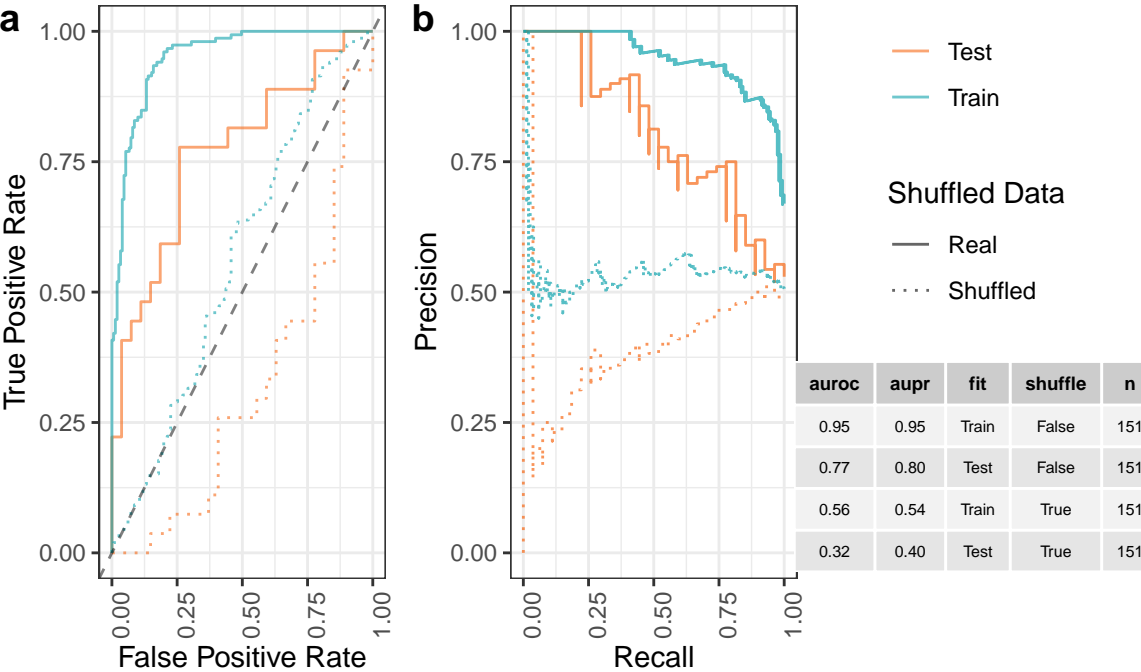
# Performance: cc\_cc\_ph3\_neg\_hoechst\_late\_mitosis\_mean



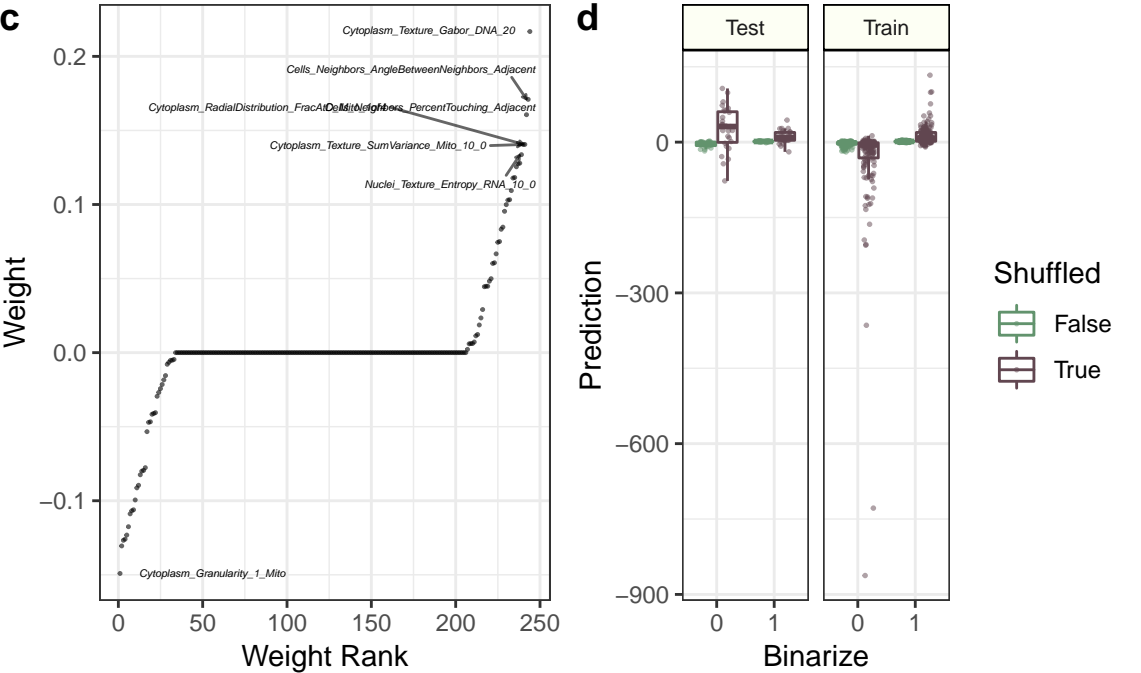
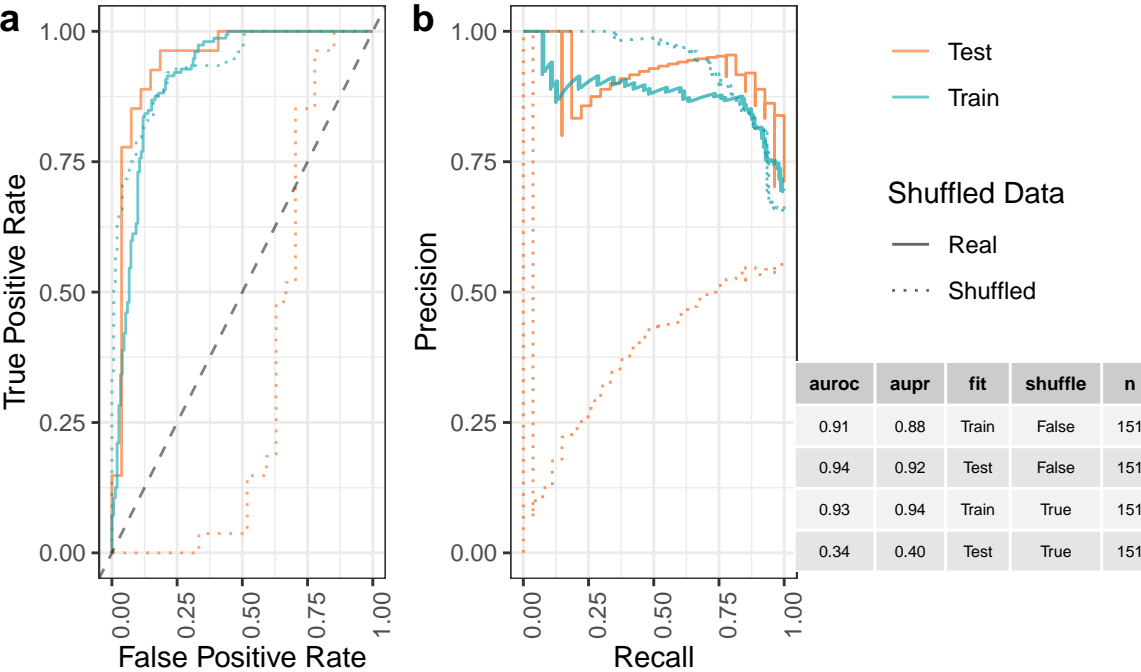
Performance: cc\_cc\_ph3\_pos\_hoechst\_mitosis\_mean



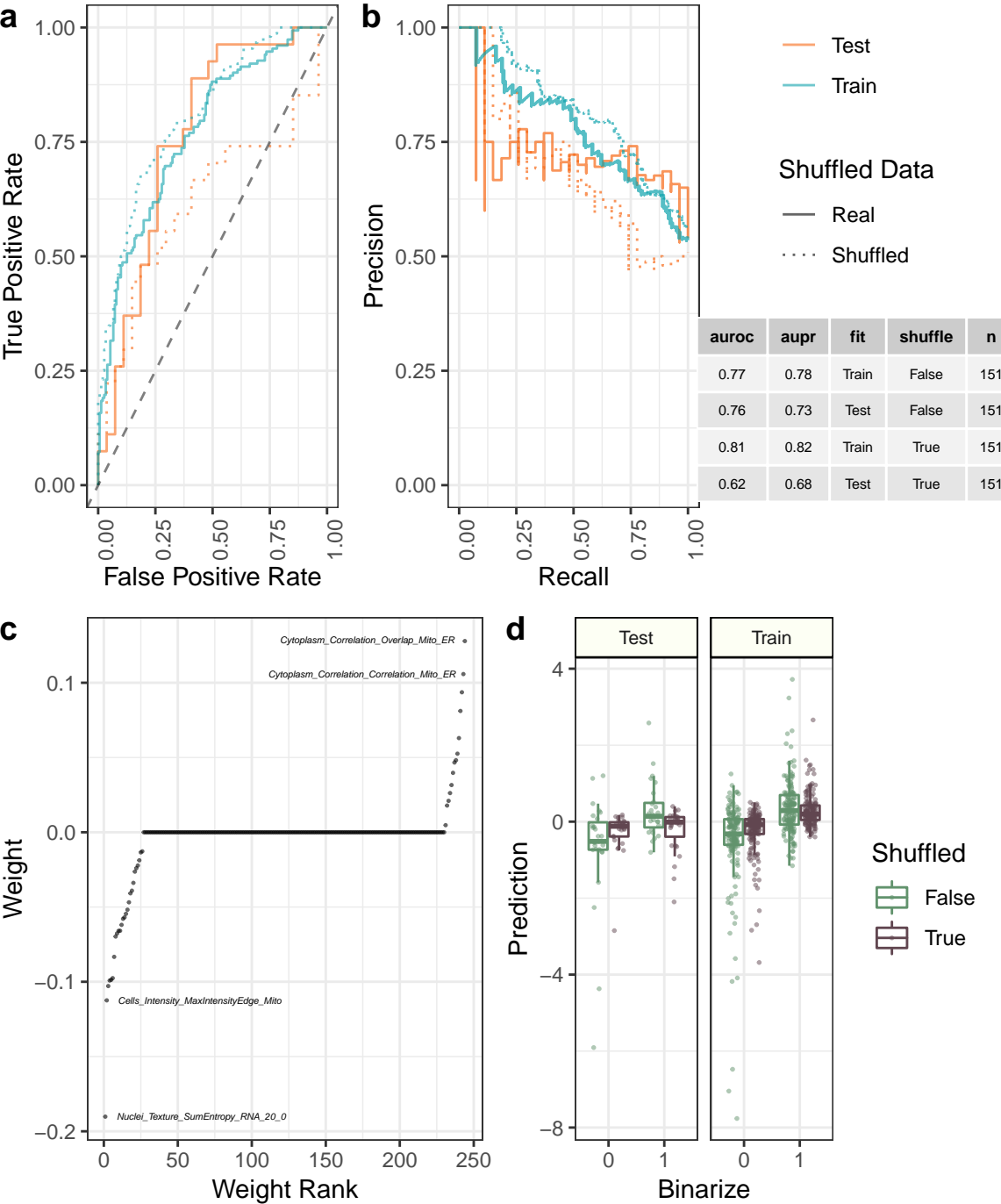
Performance: cc\_edu\_pos\_alexa647\_intensity\_nucleus\_area\_m



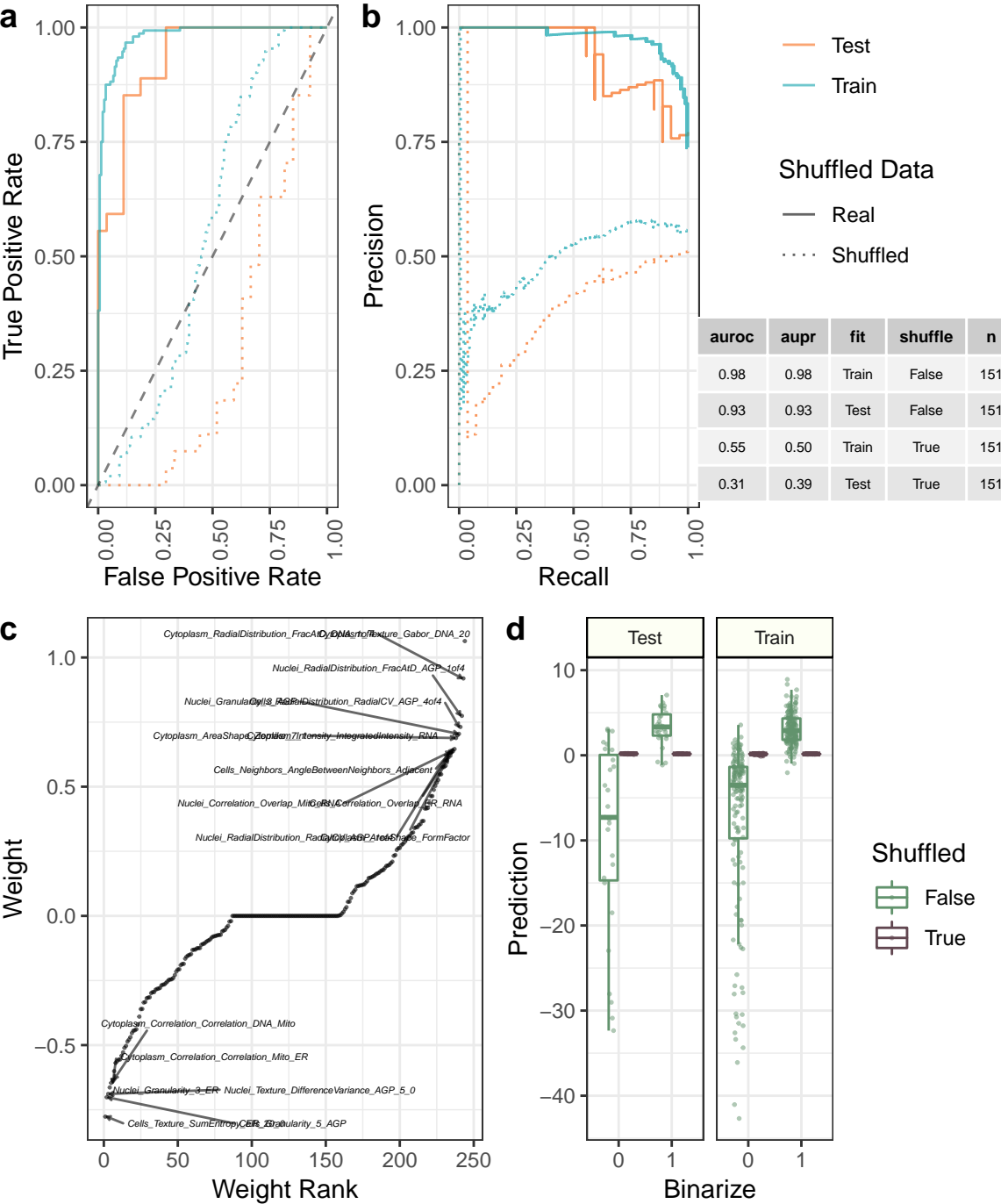
Performance: cc\_edu\_pos\_alexa647\_intensity\_nucleus\_area\_su



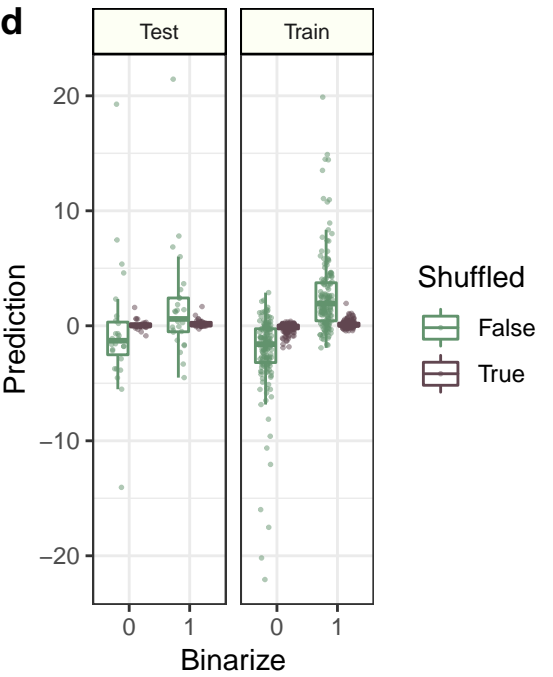
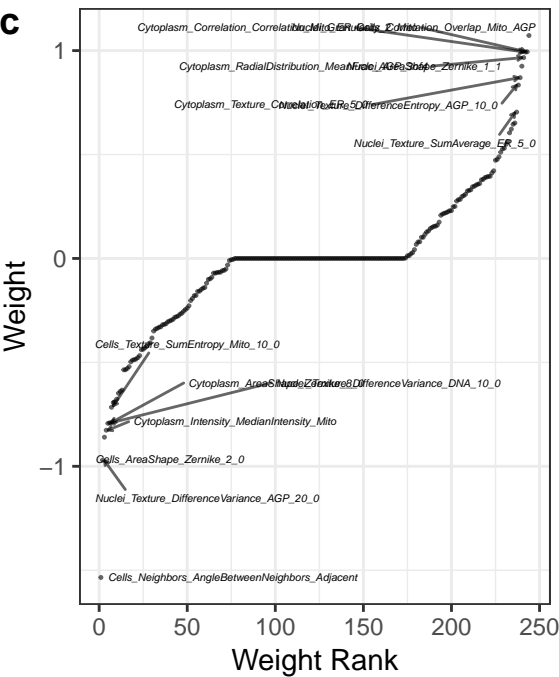
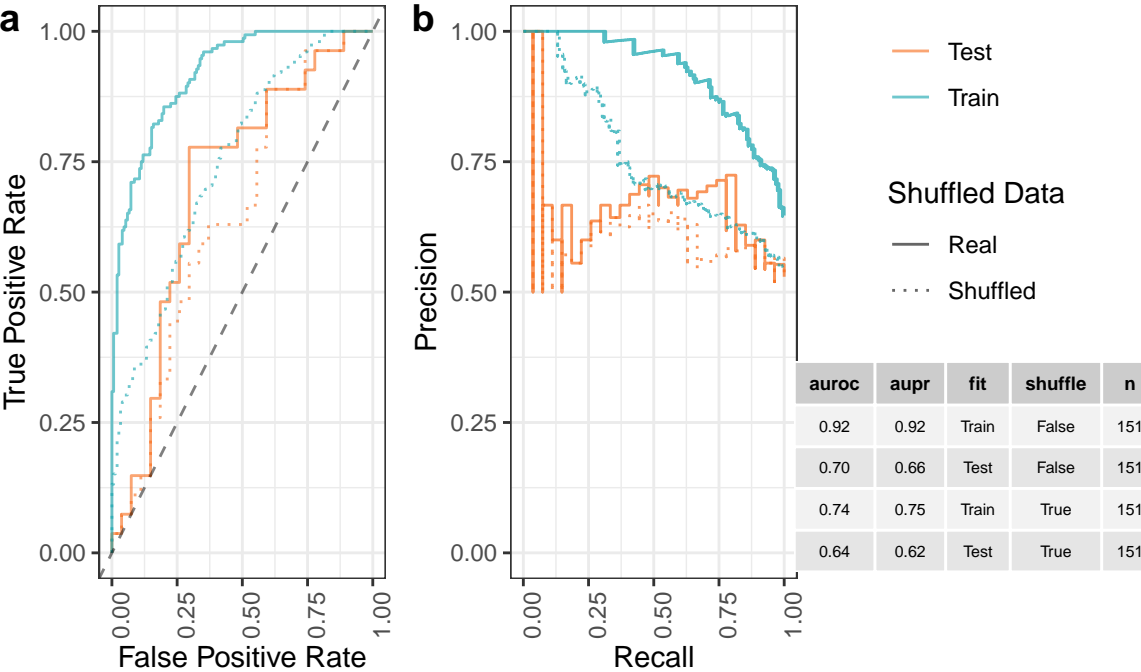
Performance: cc\_edu\_pos\_high\_n\_spots\_h2ax\_mean



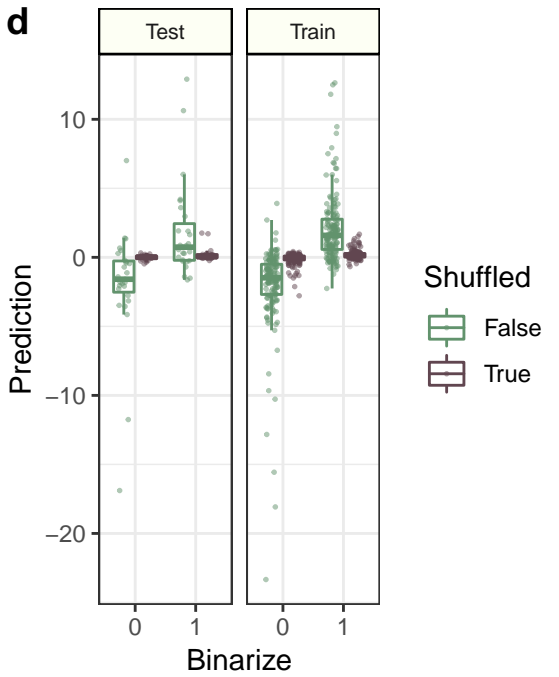
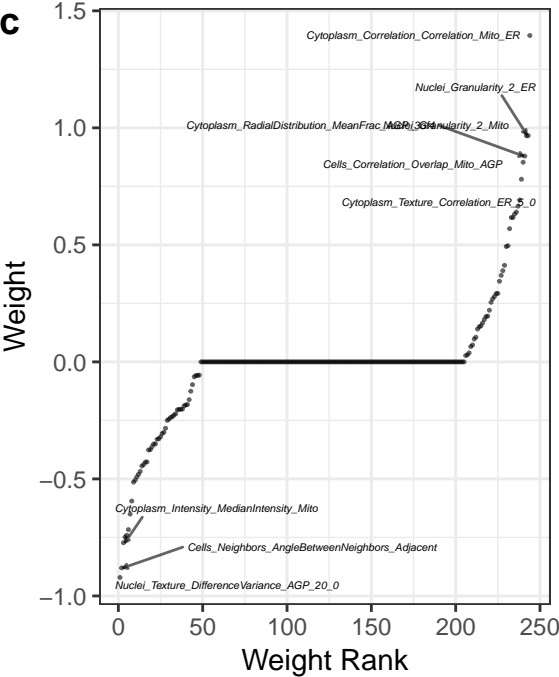
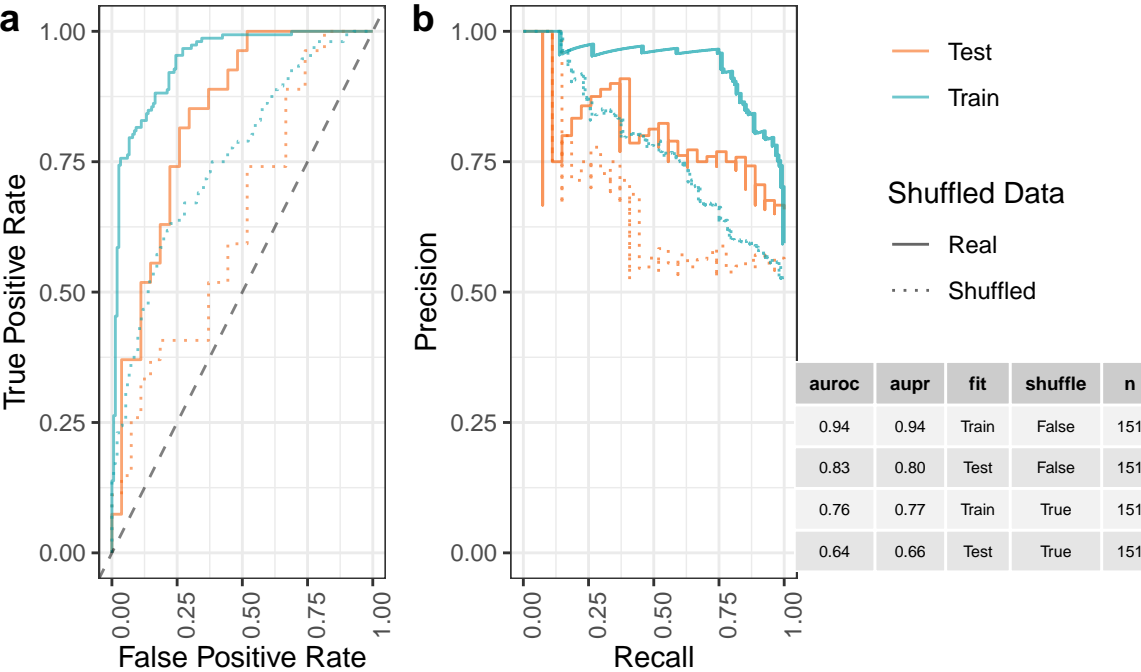
Performance: cc\_edu\_pos\_n\_objects



Performance: cc\_edu\_pos\_n\_spots\_mean

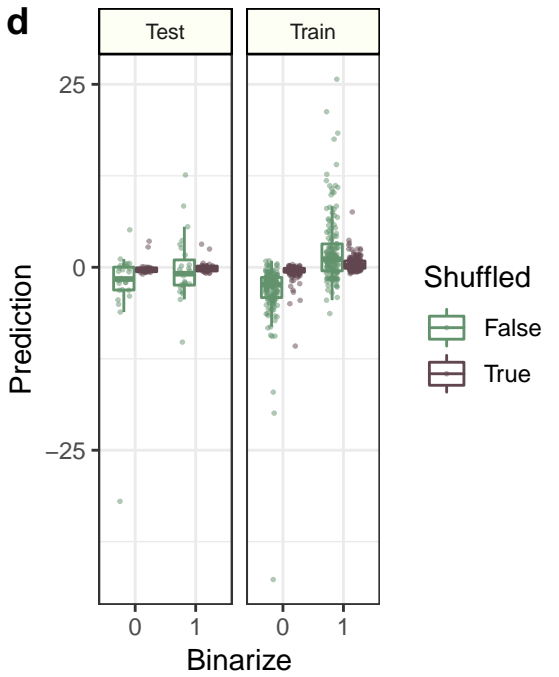
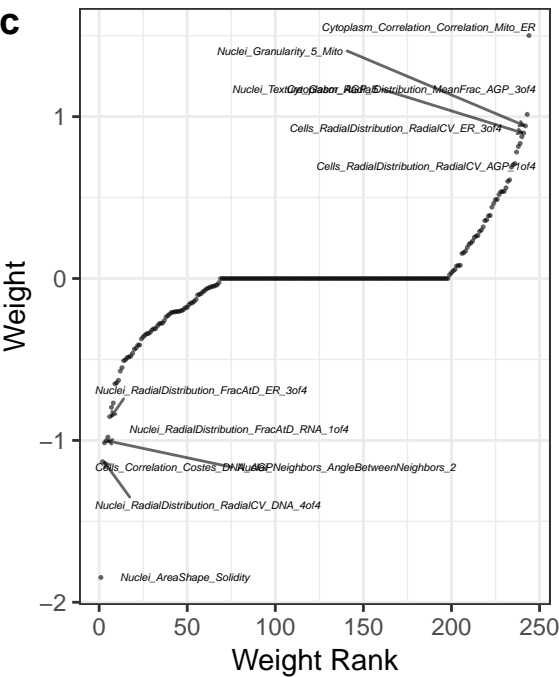
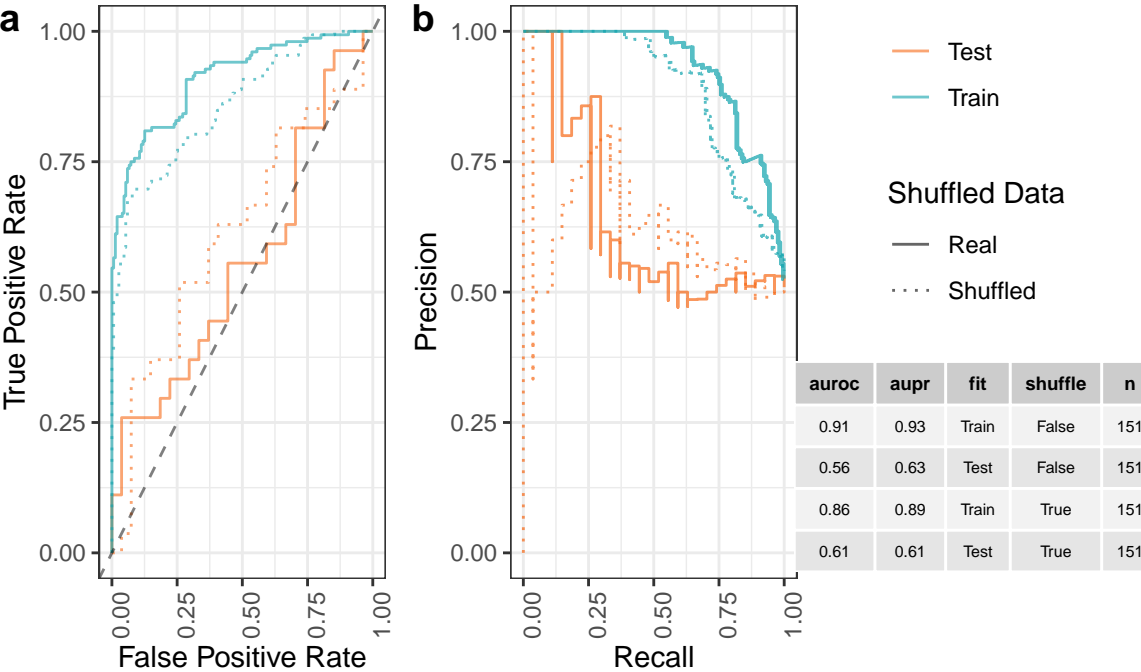


Performance: cc\_edu\_pos\_n\_spots\_per\_nucleus\_area\_mean

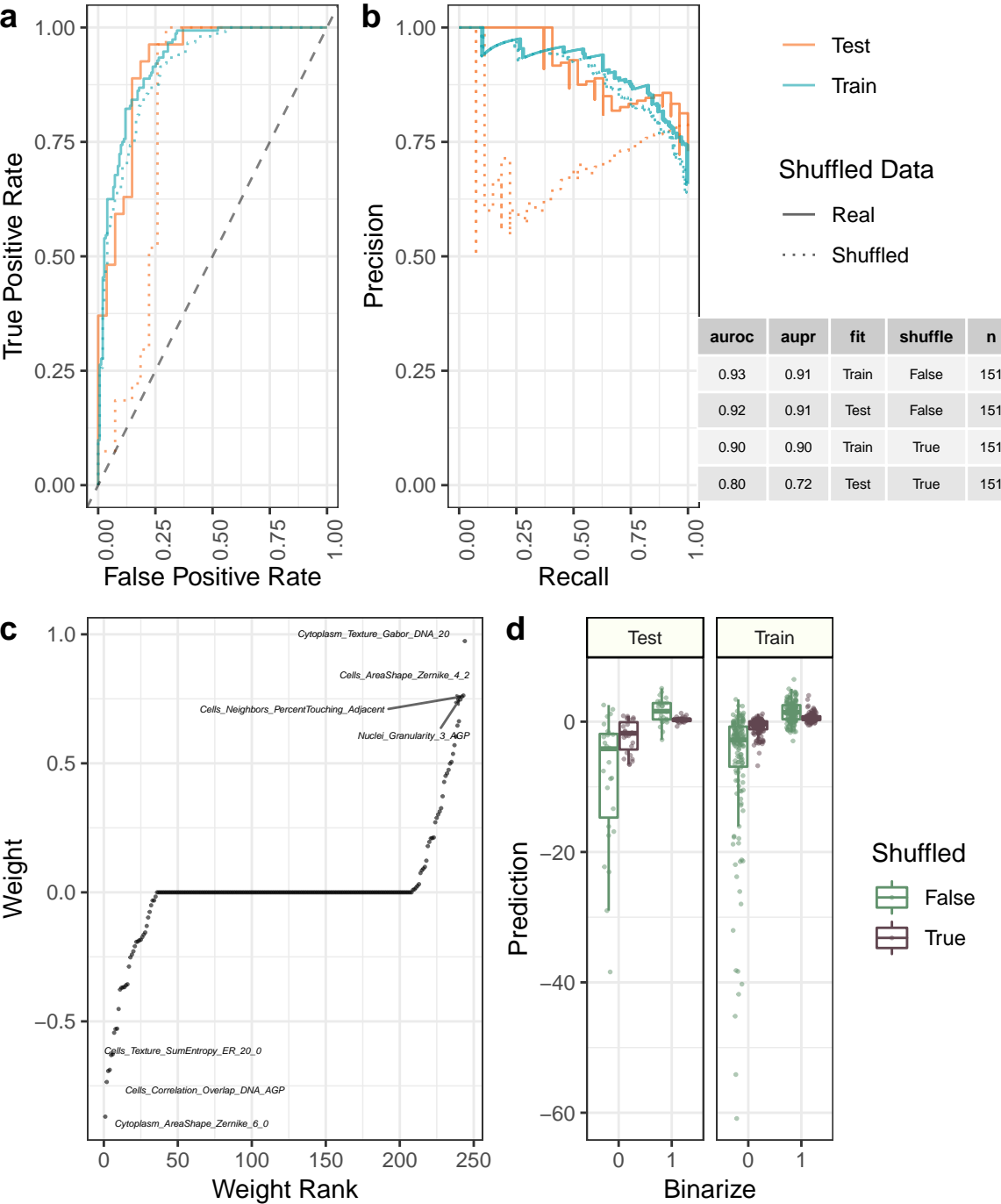




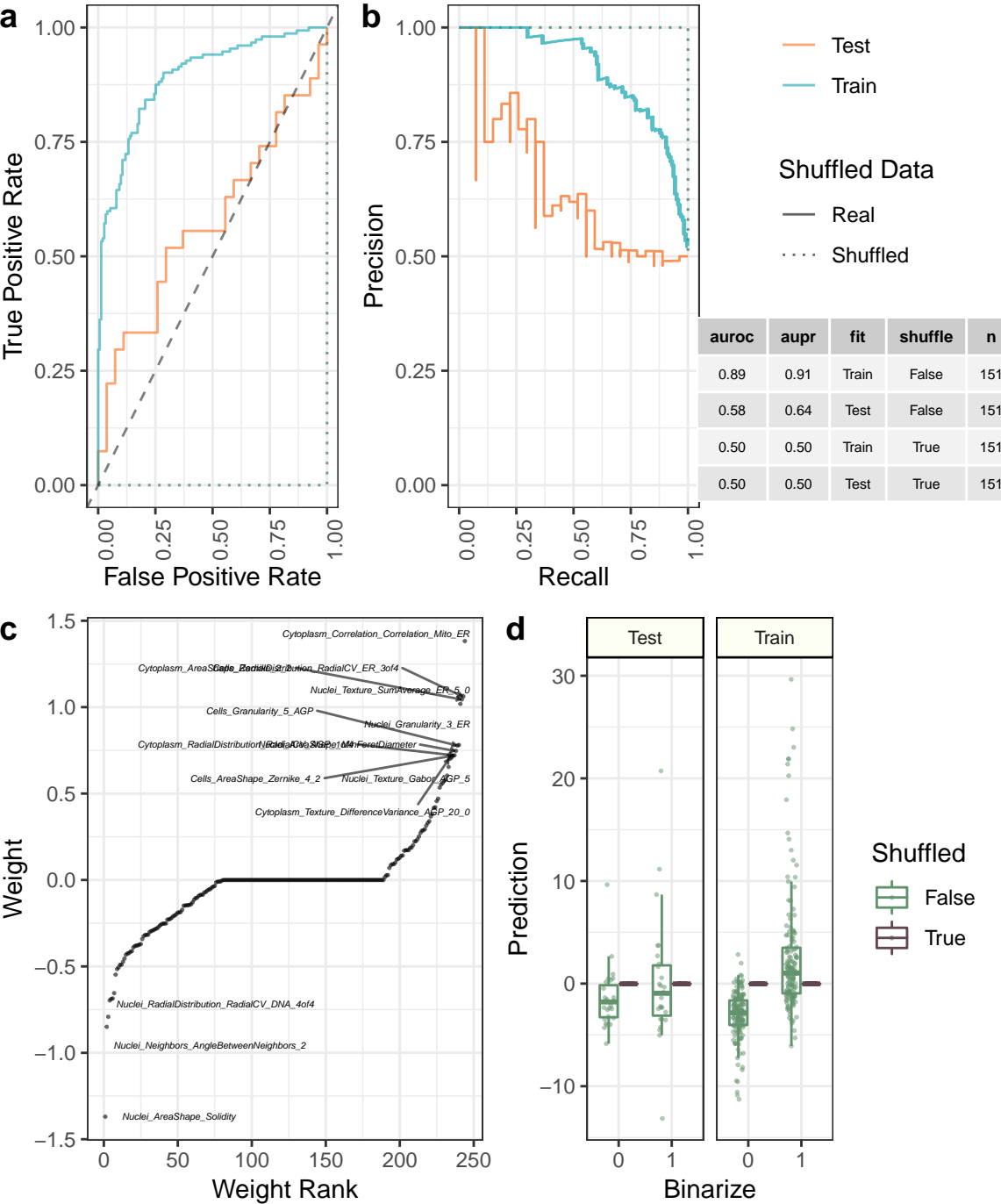
Performance: cc\_g1\_high\_n\_spots\_h2ax\_mean



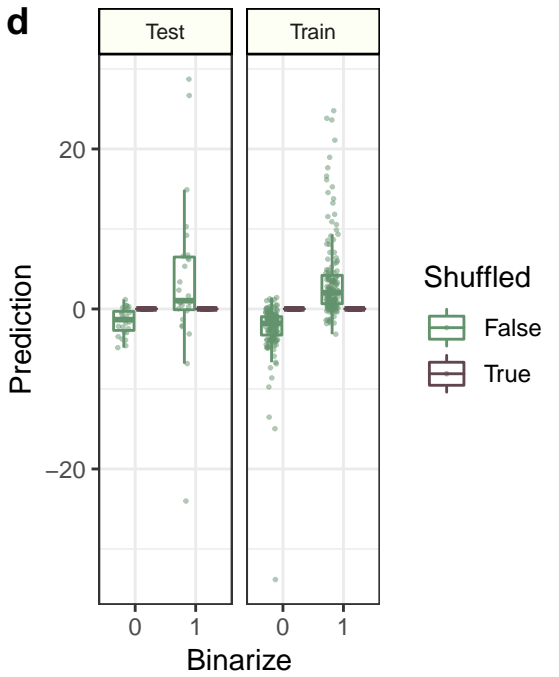
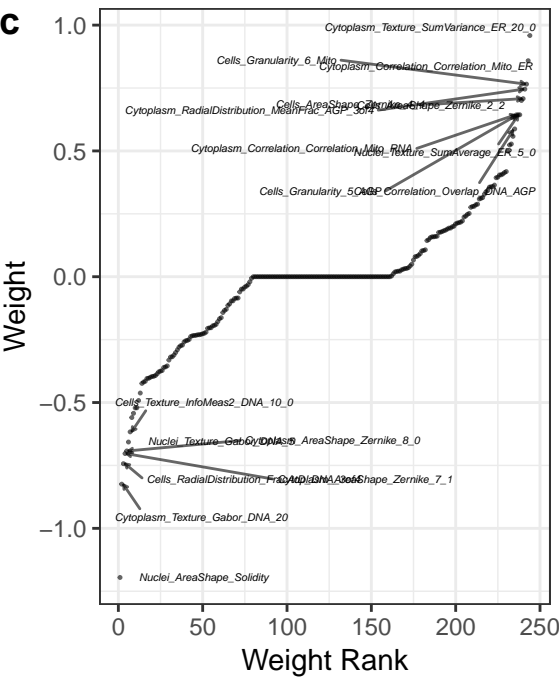
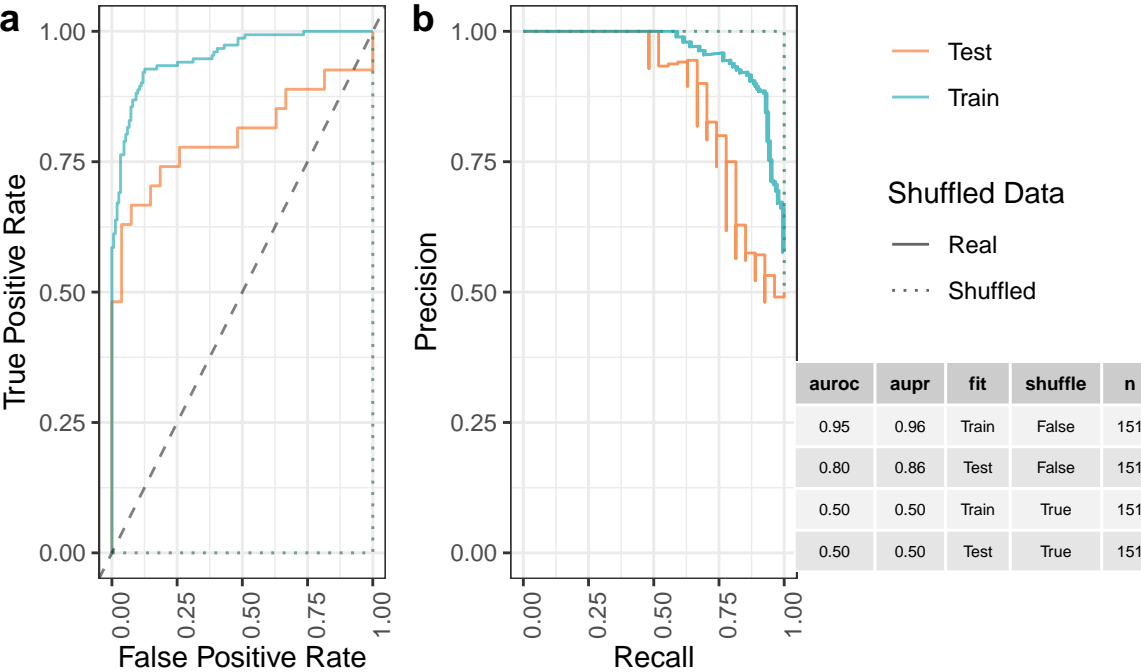
# Performance: cc\_g1\_n\_objects



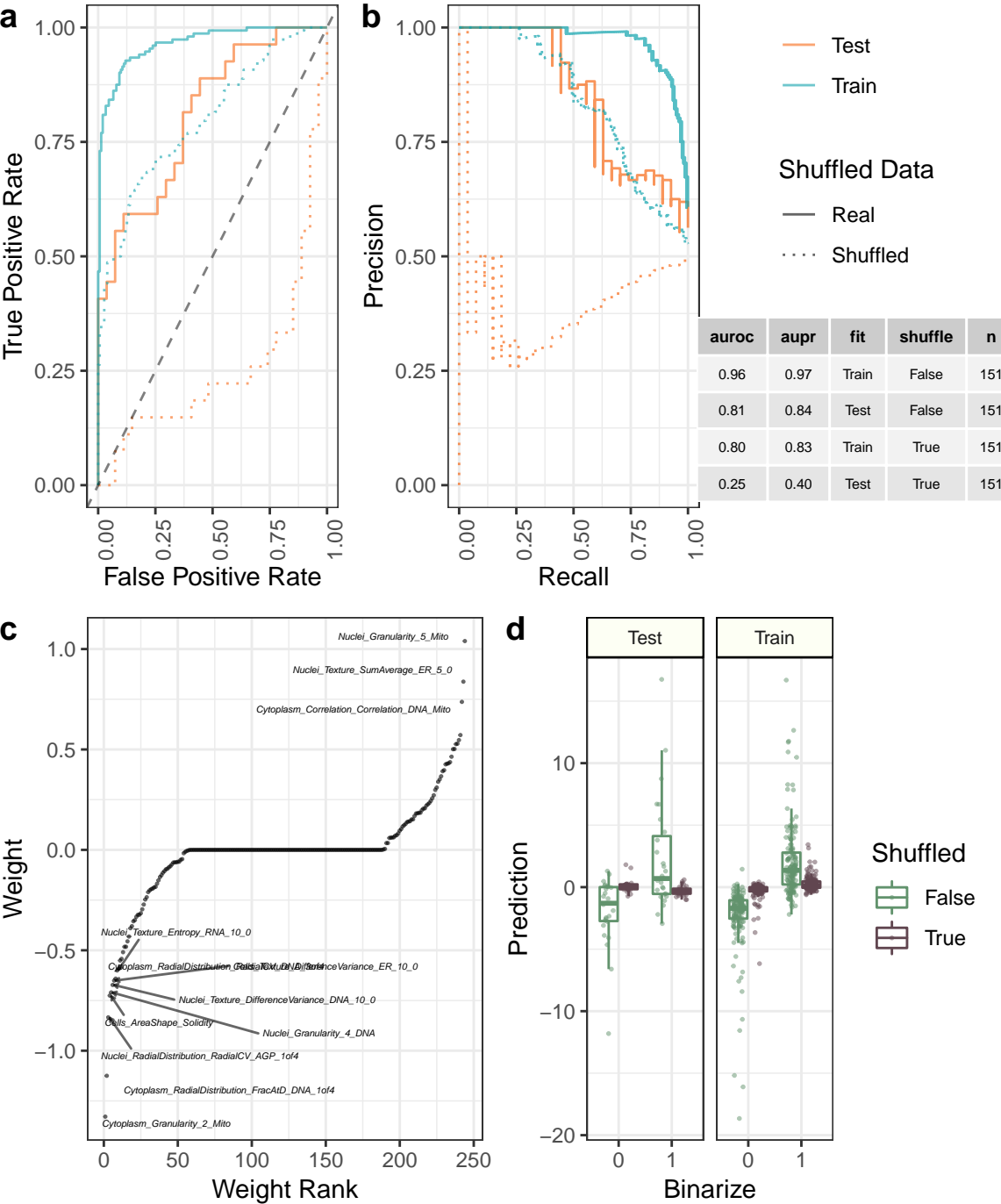
Performance: cc\_g1\_n\_spots\_mean



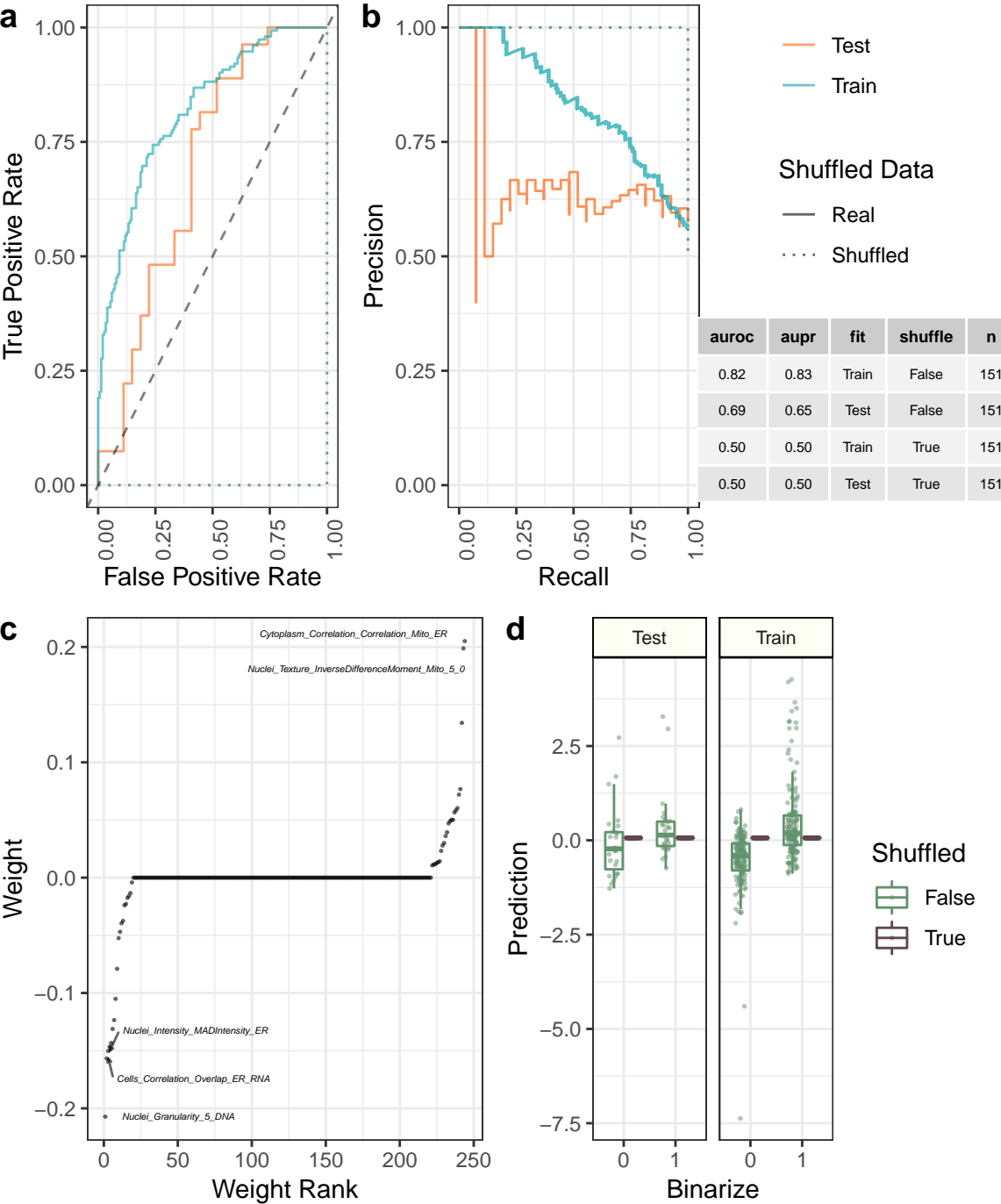
Performance: cc\_g1\_n\_spots\_per\_nucleus\_area\_mean



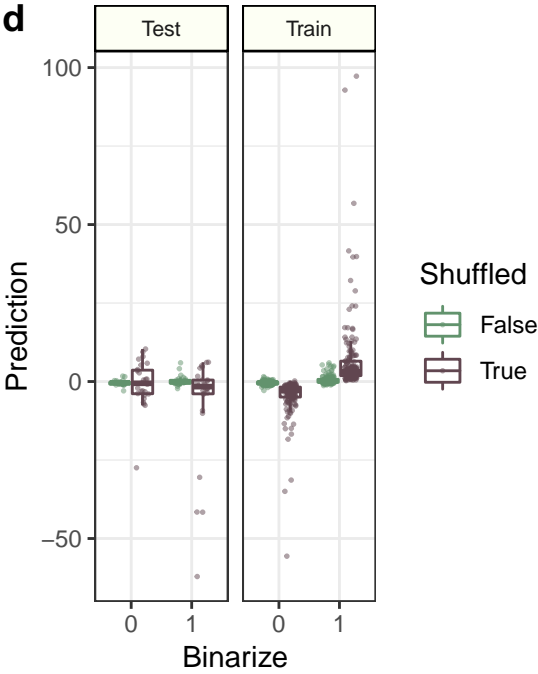
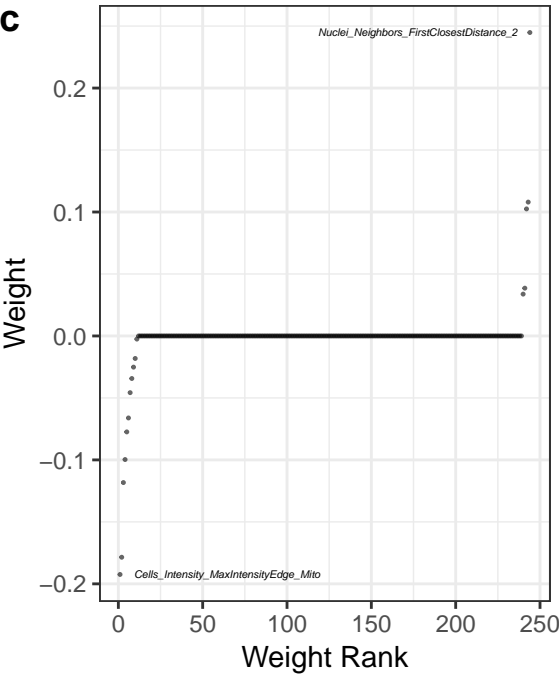
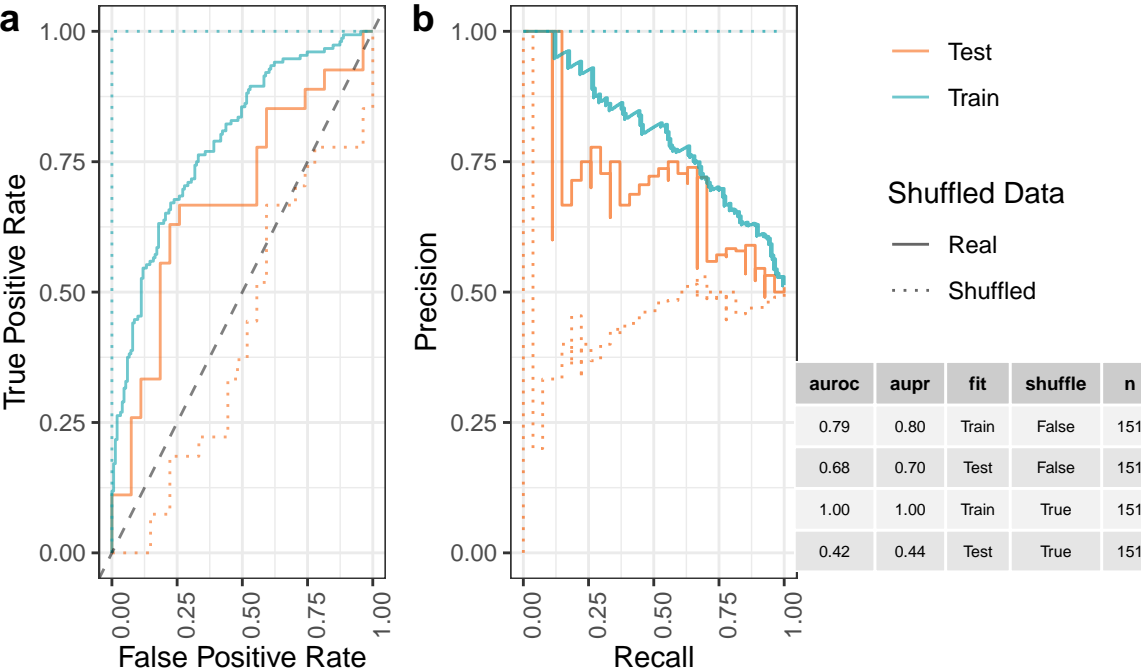
# Performance: cc\_g1\_plus\_g2



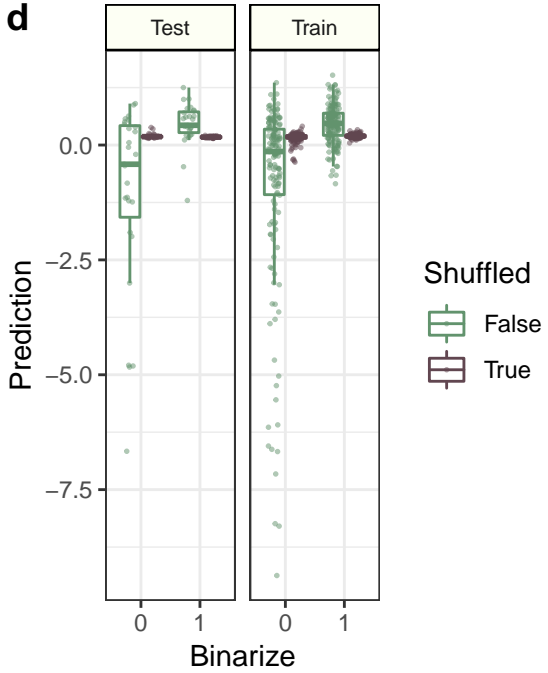
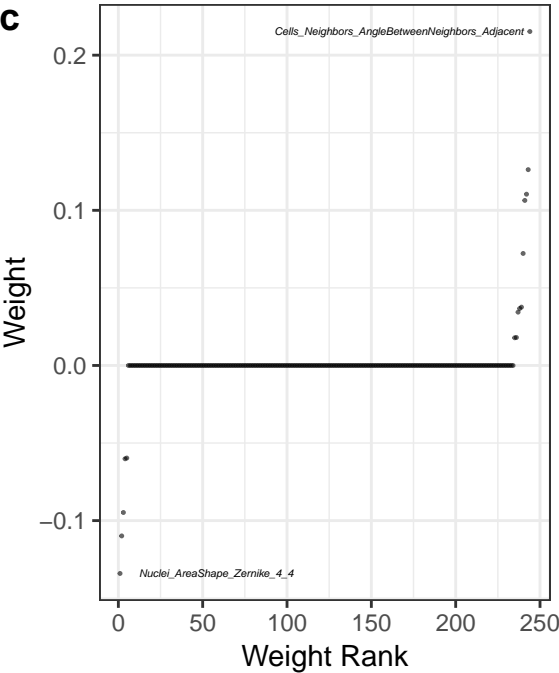
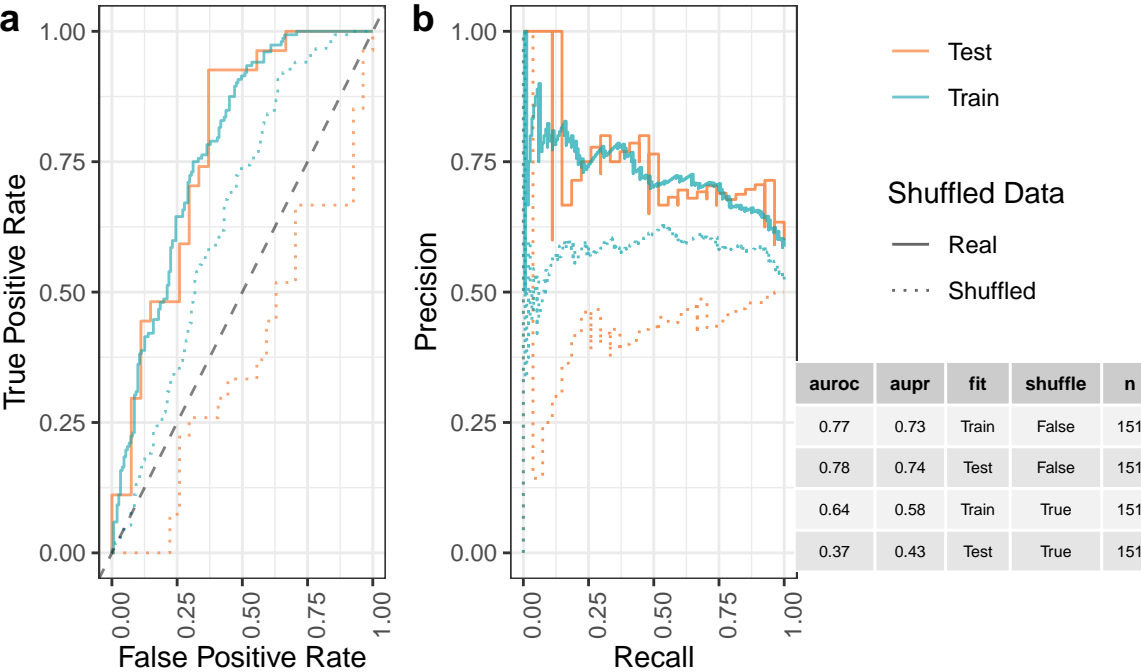
# Performance: cc\_g2\_g1



# Performance: cc\_g2\_ph3\_neg\_high\_n\_spots\_h2ax\_mean

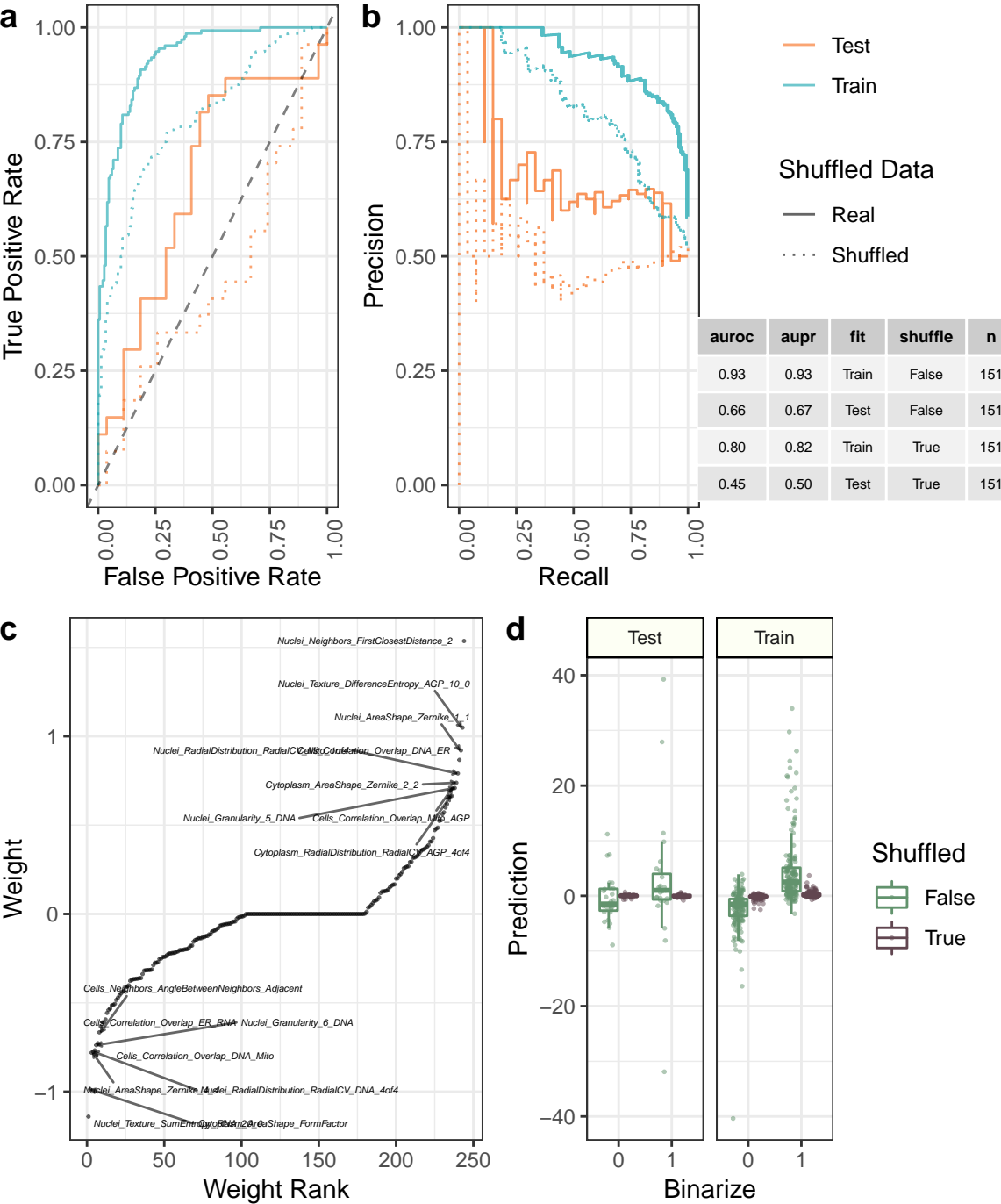


# Performance: cc\_g2\_ph3\_neg\_n\_objects

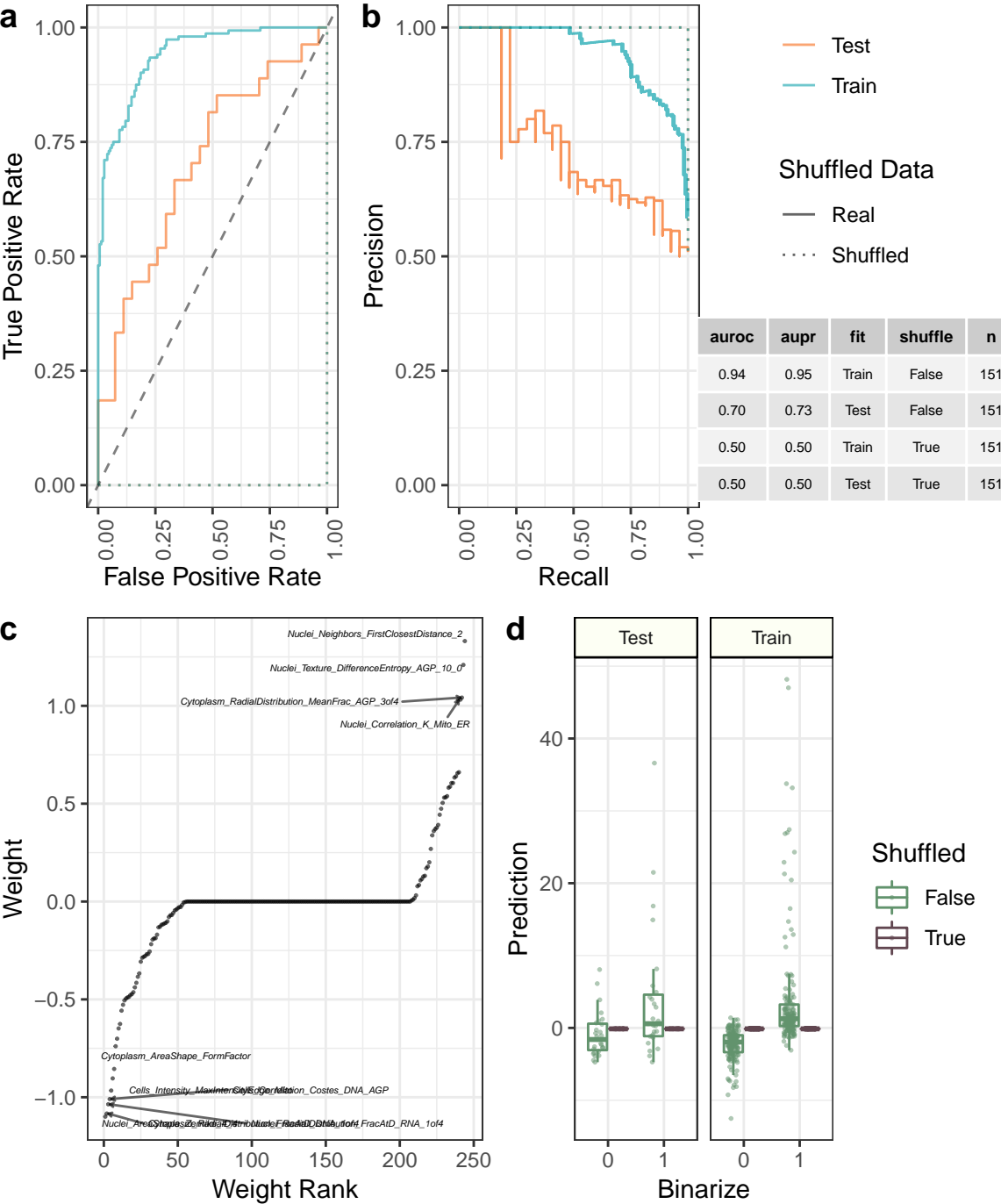




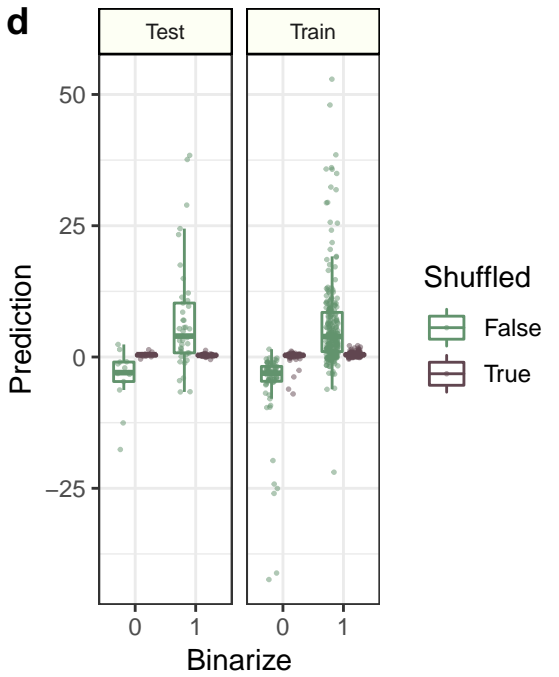
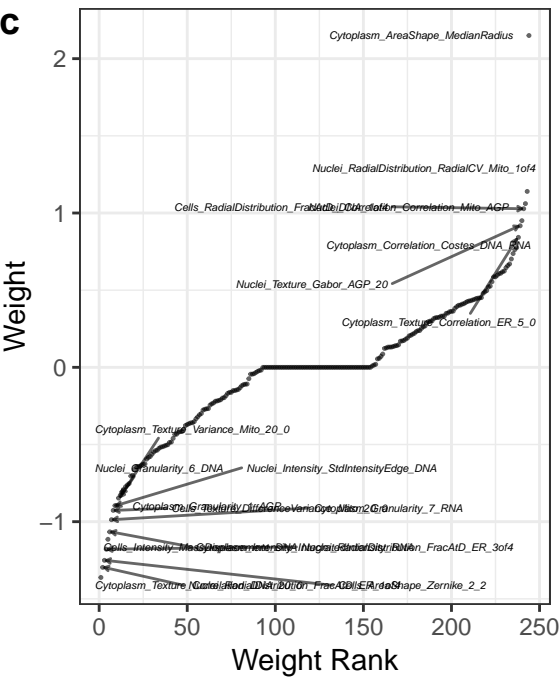
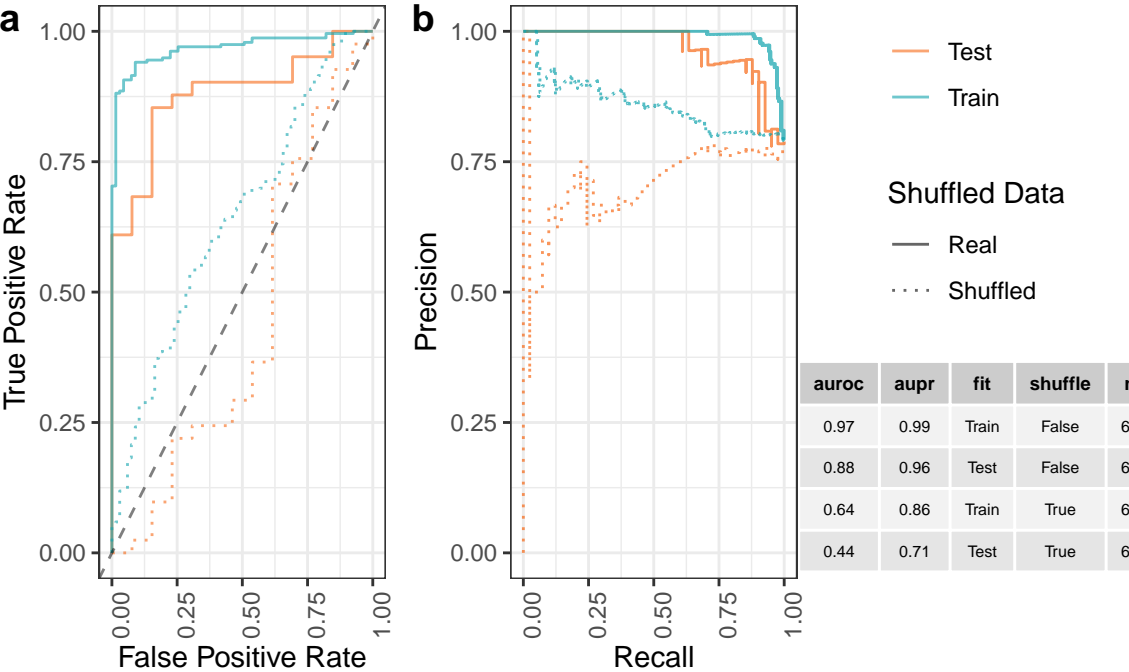
Performance: cc\_g2\_ph3\_neg\_n\_spots\_mean



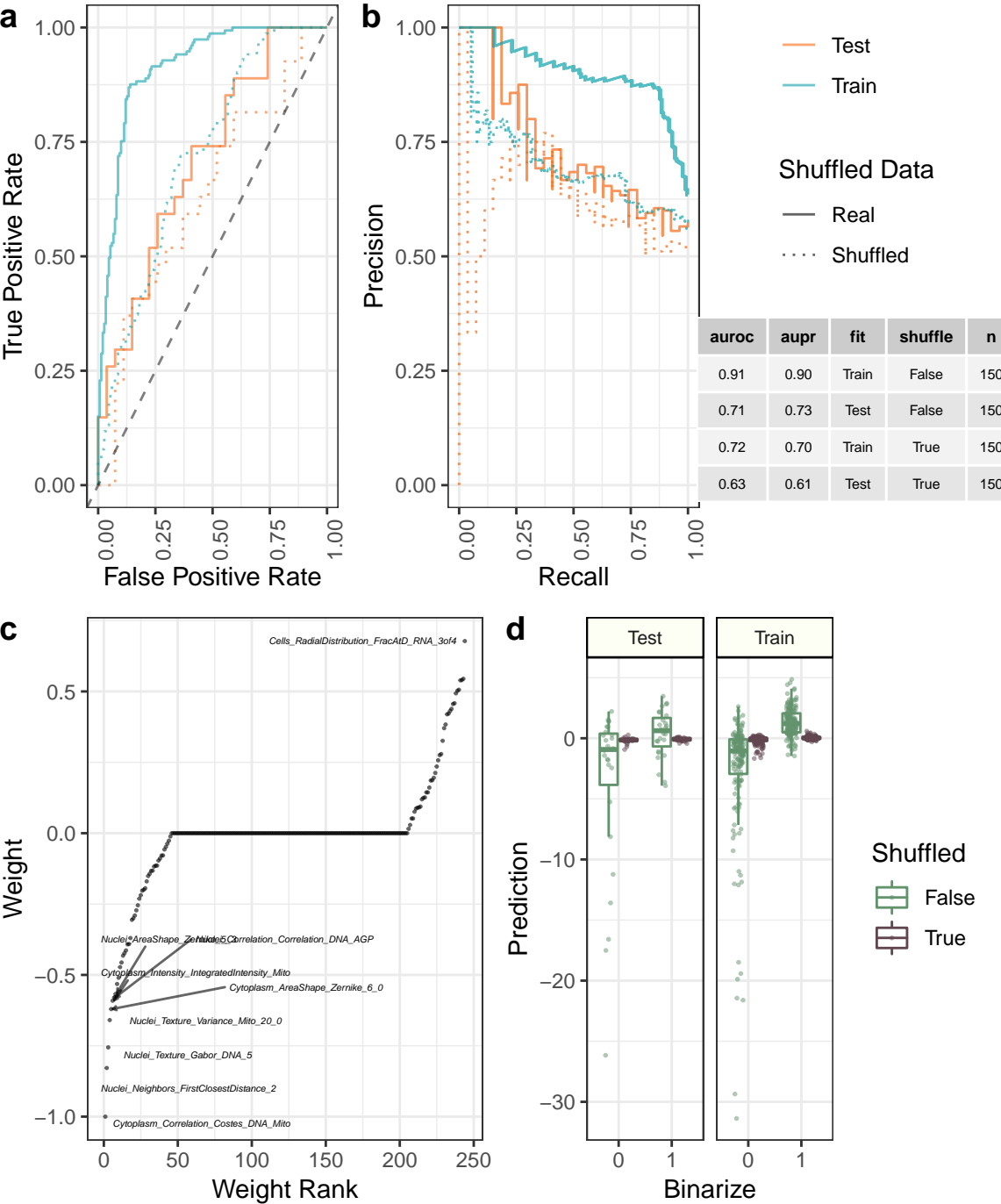
Performance: cc\_g2\_ph3\_neg\_n\_spots\_per\_nucleus\_area\_mean



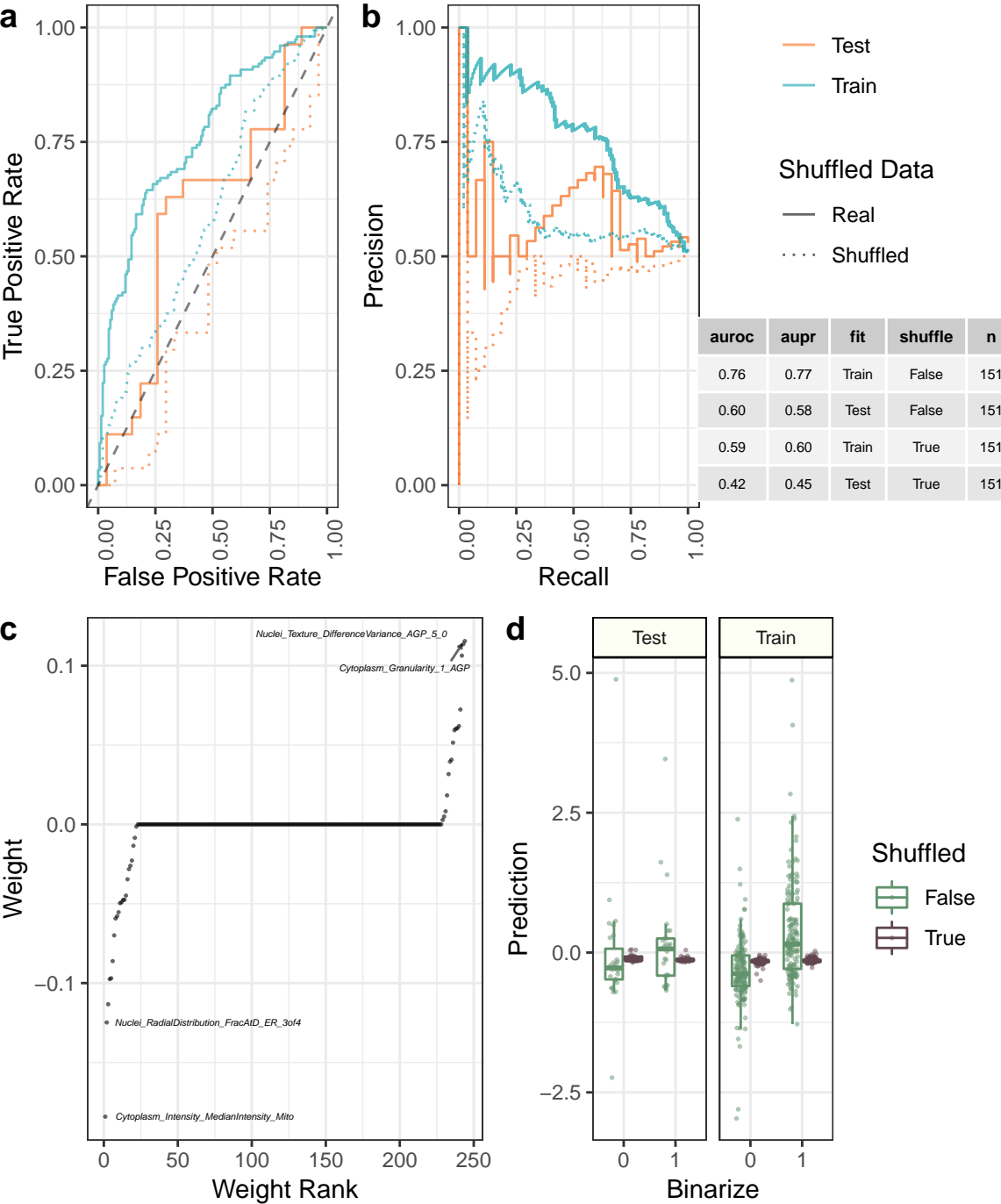
Performance: cc\_g2\_ph3\_pos\_high\_n\_spots\_h2ax\_mean



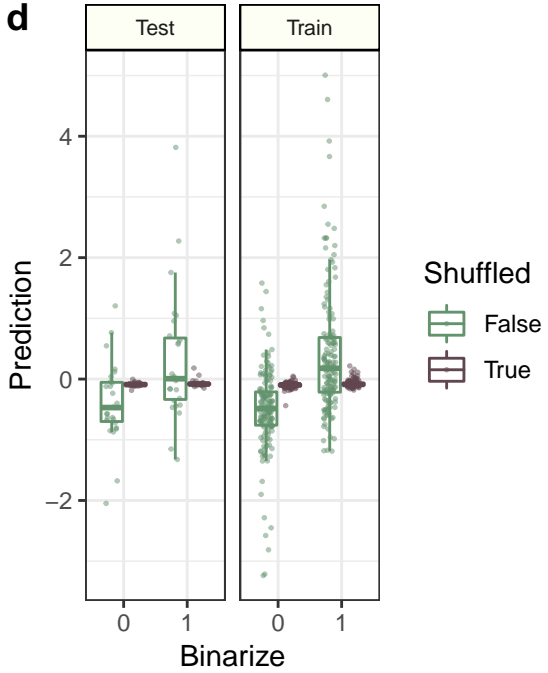
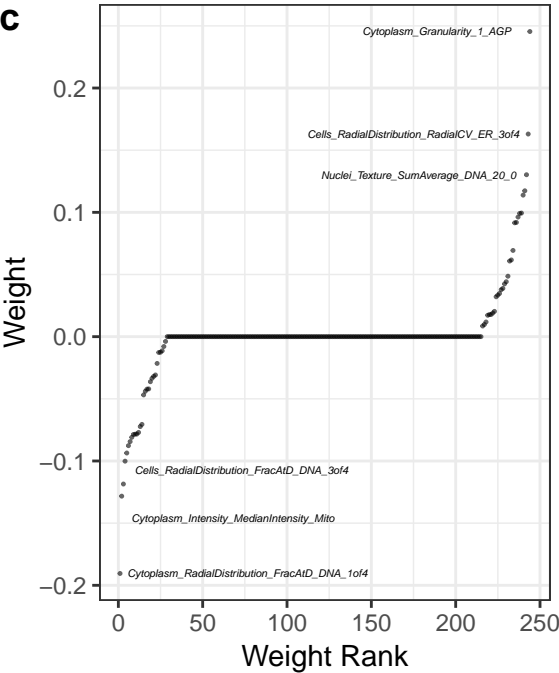
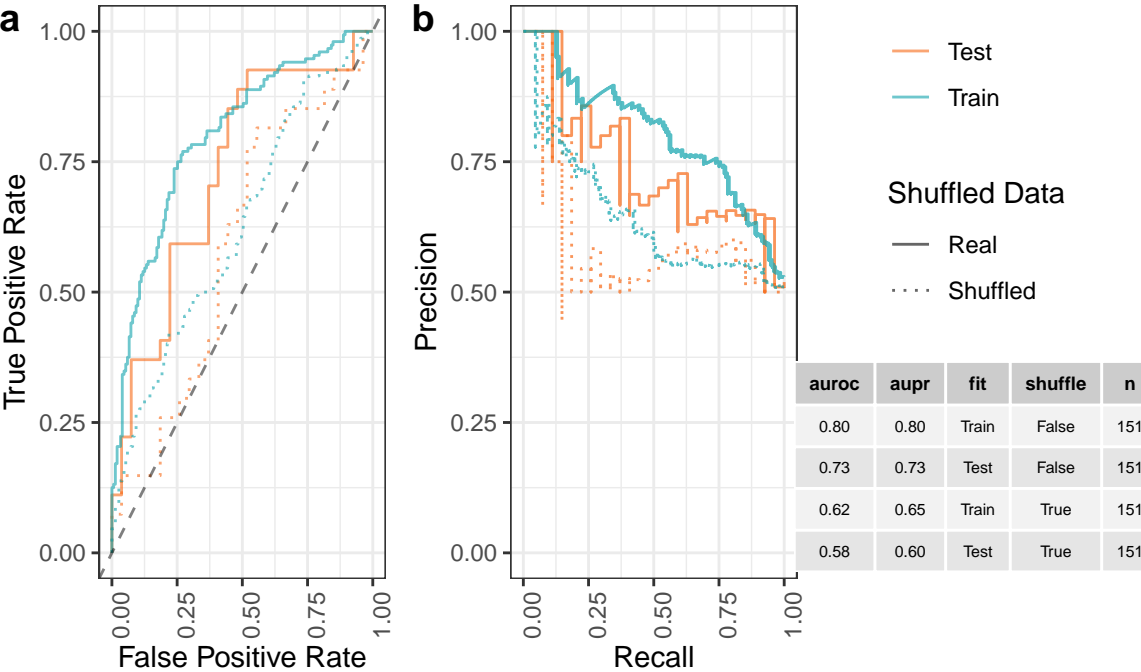
Performance: cc\_g2\_ph3\_pos\_n\_objects



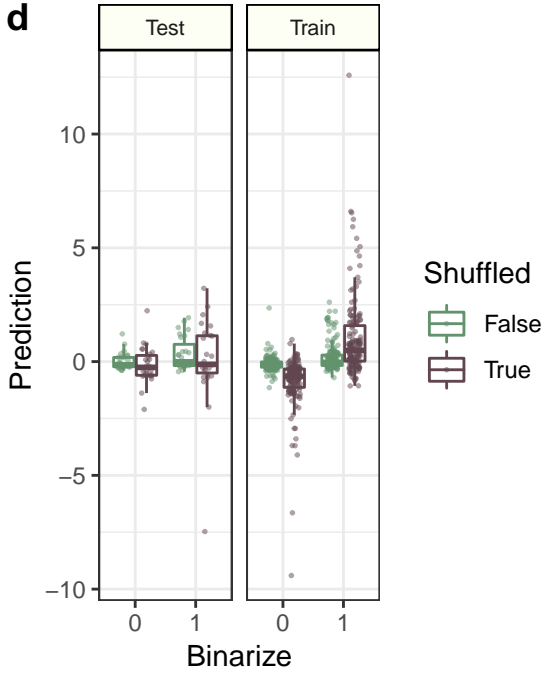
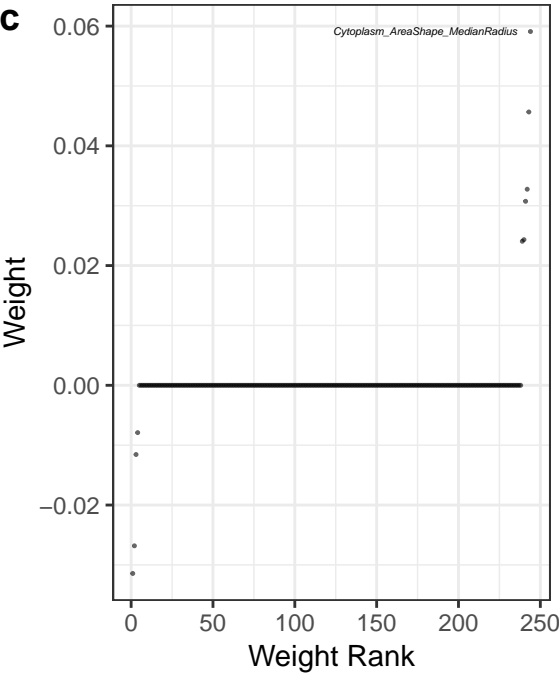
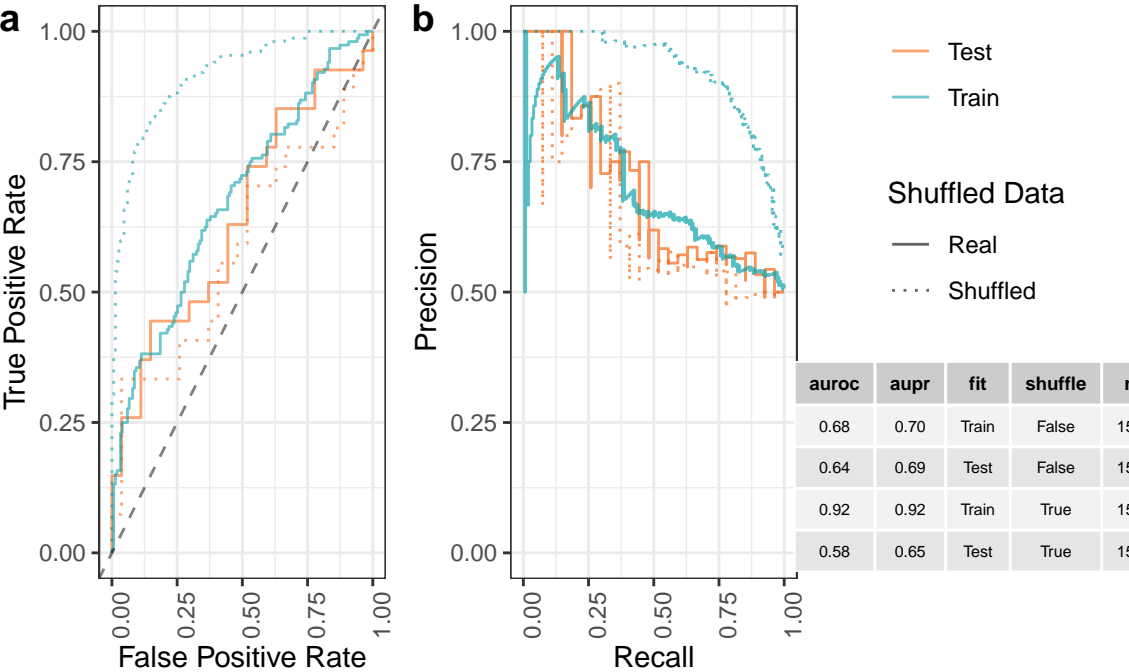
Performance: cc\_g2\_ph3\_pos\_n\_spots\_mean



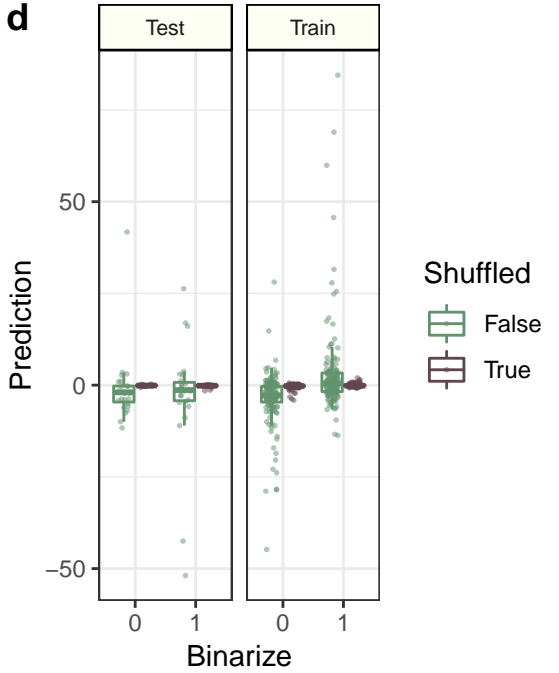
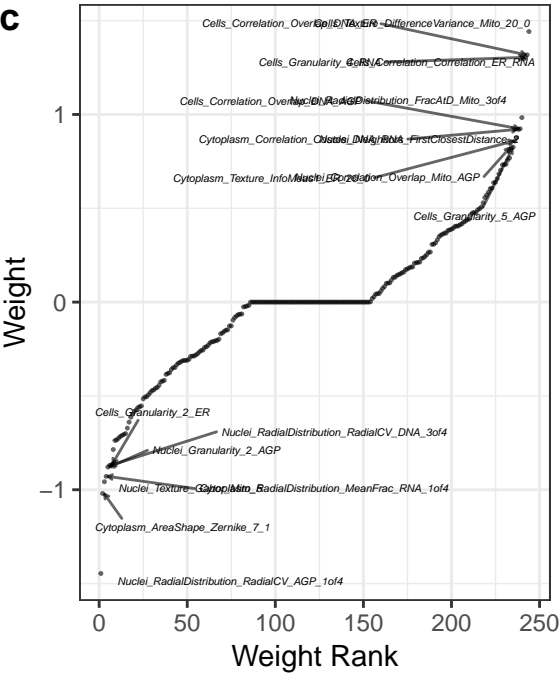
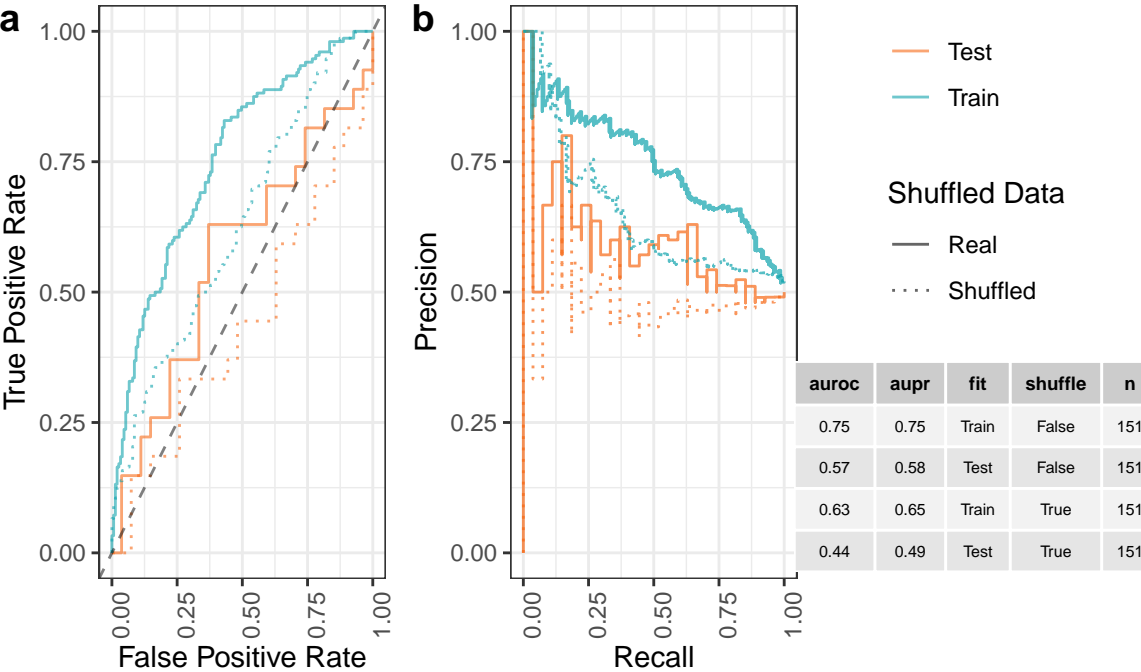
Performance: cc\_g2\_ph3\_pos\_n\_spots\_per\_nucleus\_area\_mean



# Performance: cc\_g2\_plus\_all\_m

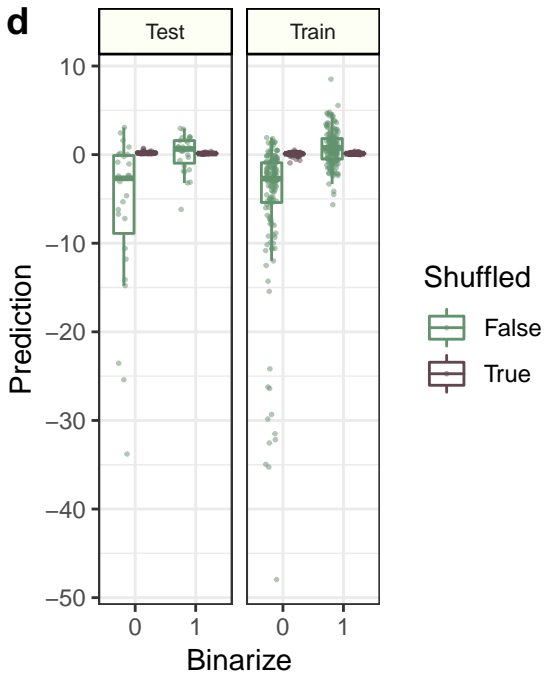
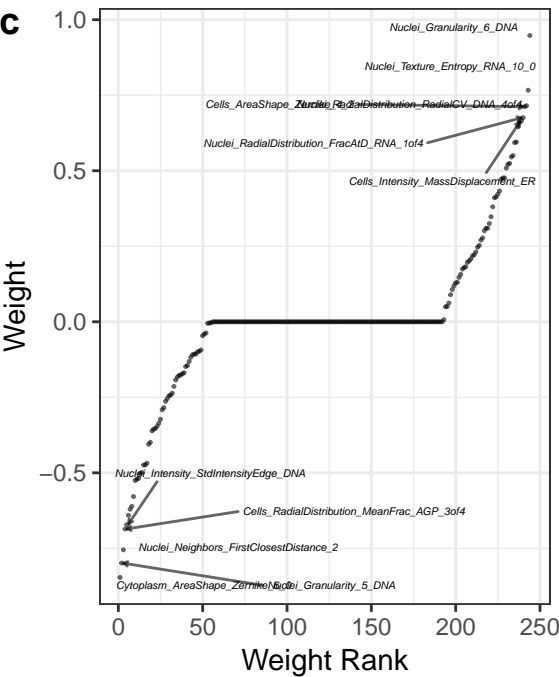
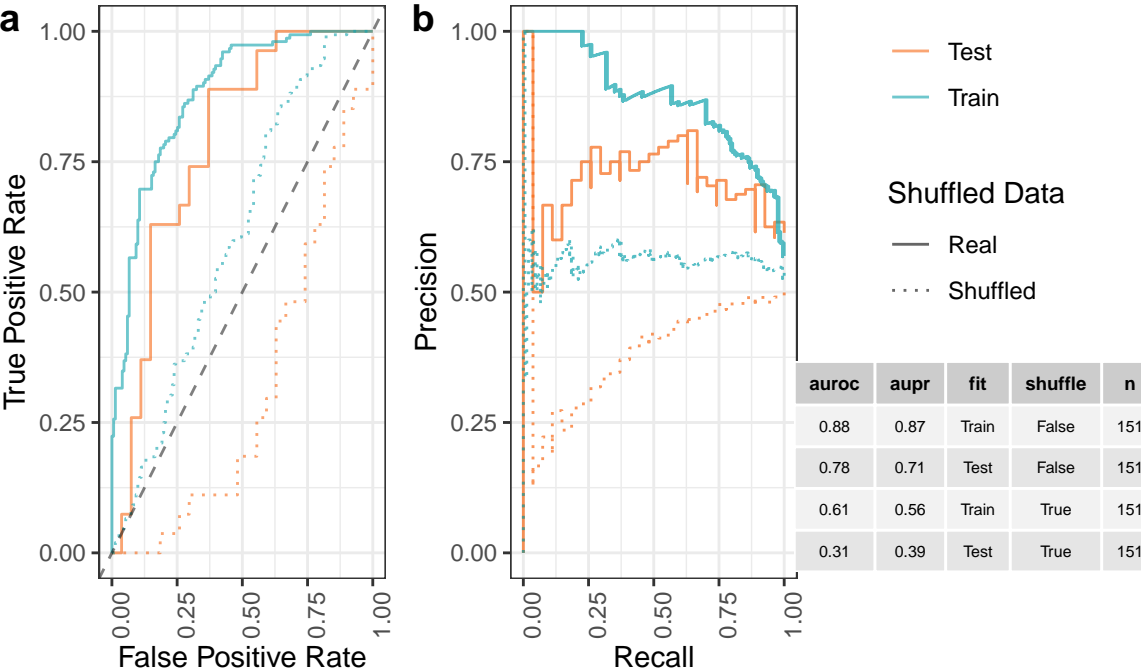


# Performance: cc\_infection\_percentage

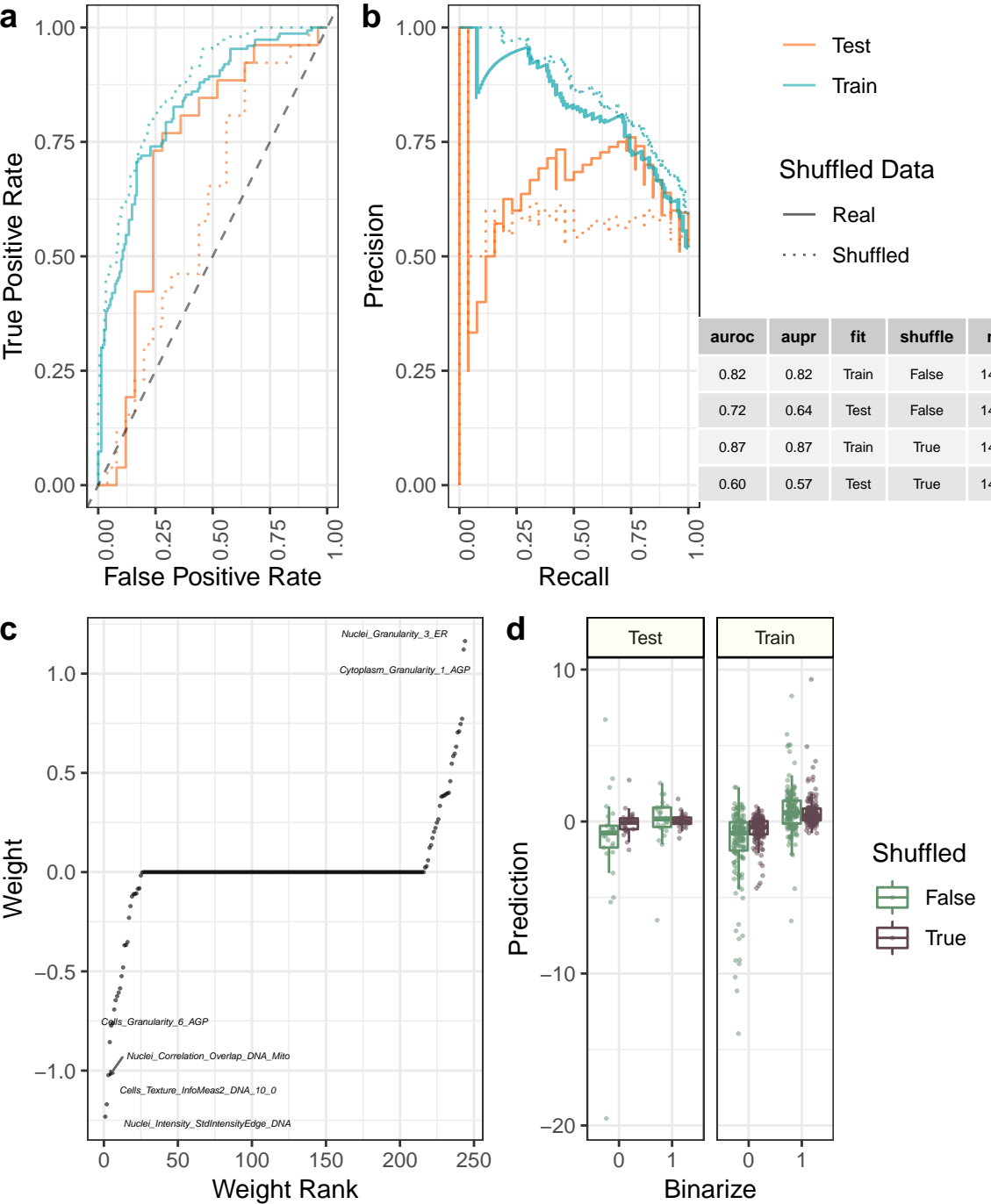




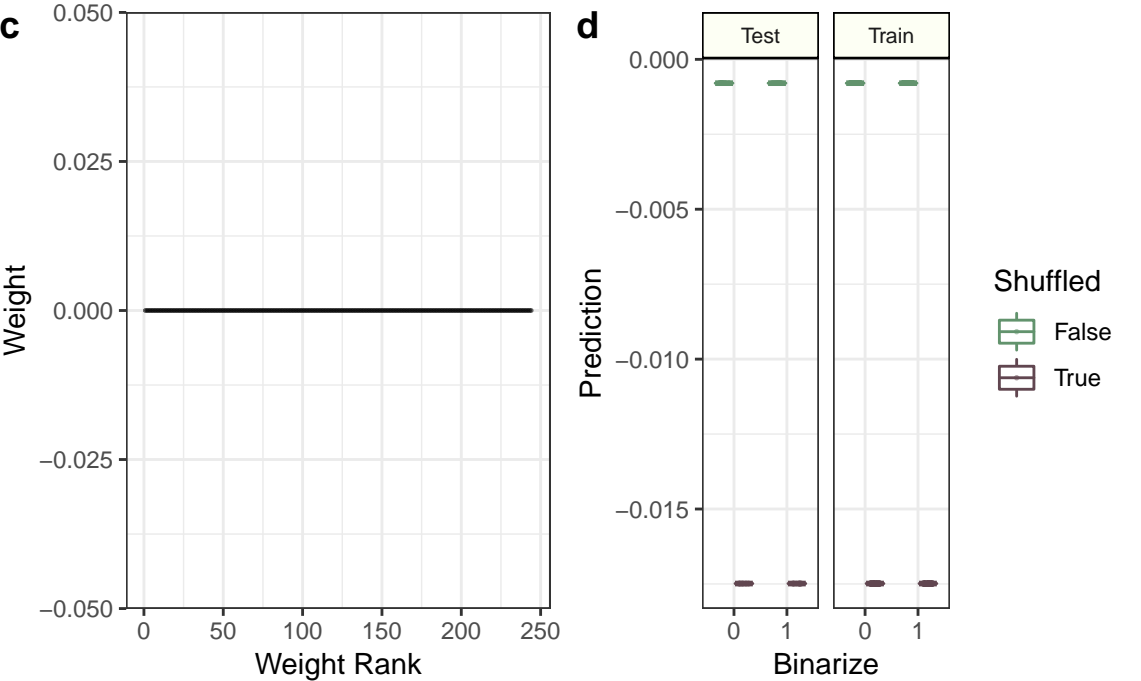
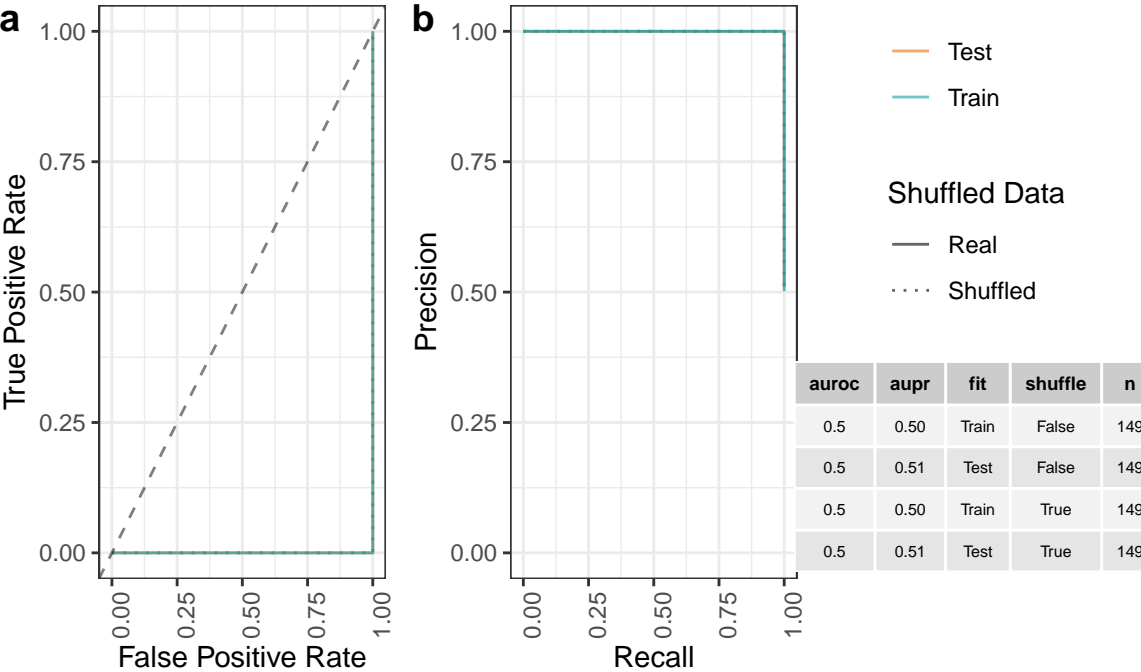
Performance: cc\_mitosis\_ph3\_neg\_n\_objects



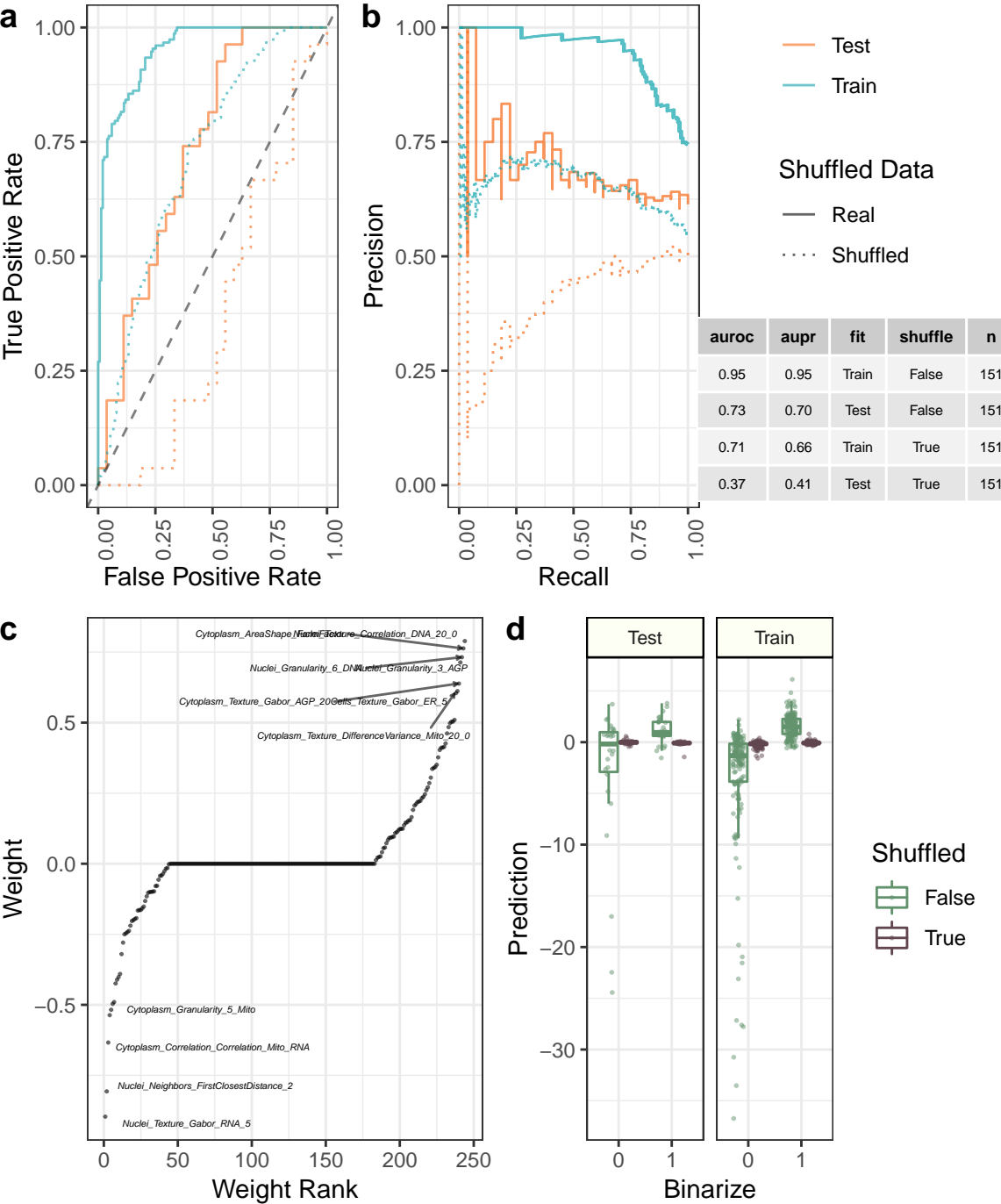
# Performance: cc\_mitosis\_ph3\_neg\_n\_spots\_mean



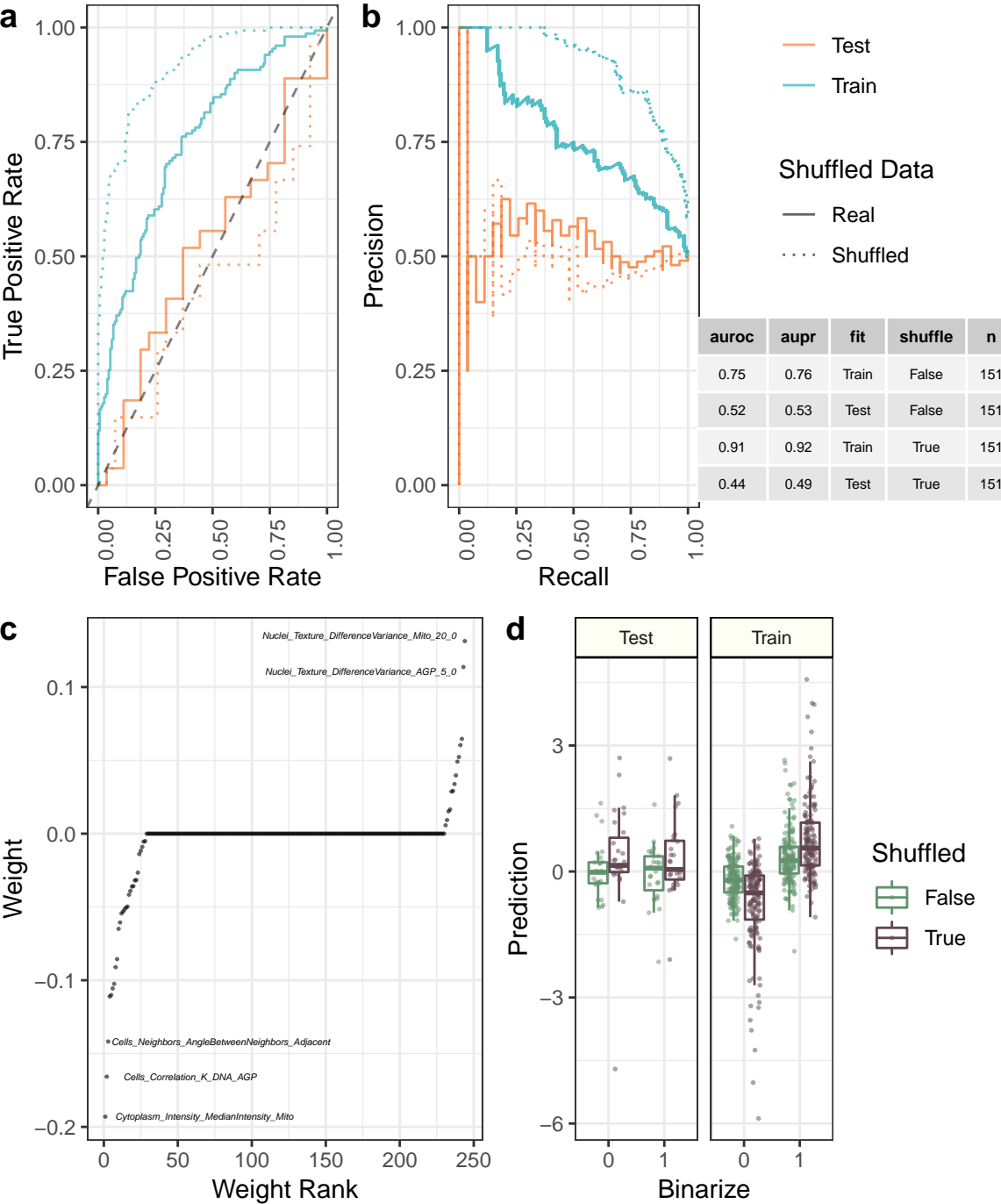
Performance: cc\_mitosis\_ph3\_neg\_n\_spots\_per\_nucleus\_area



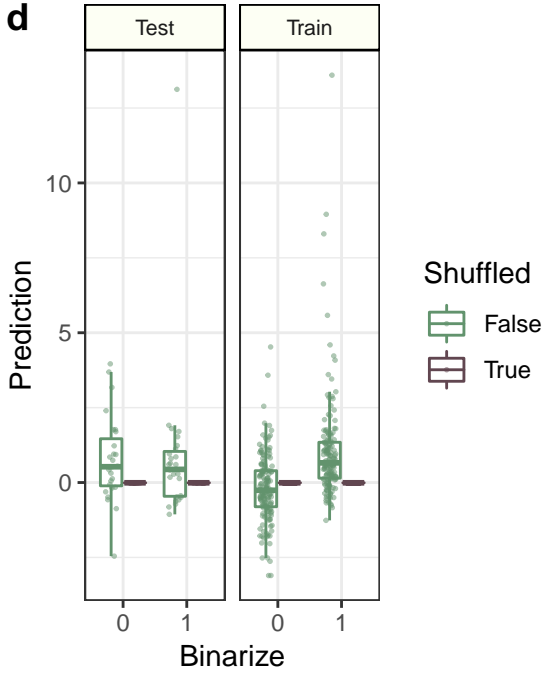
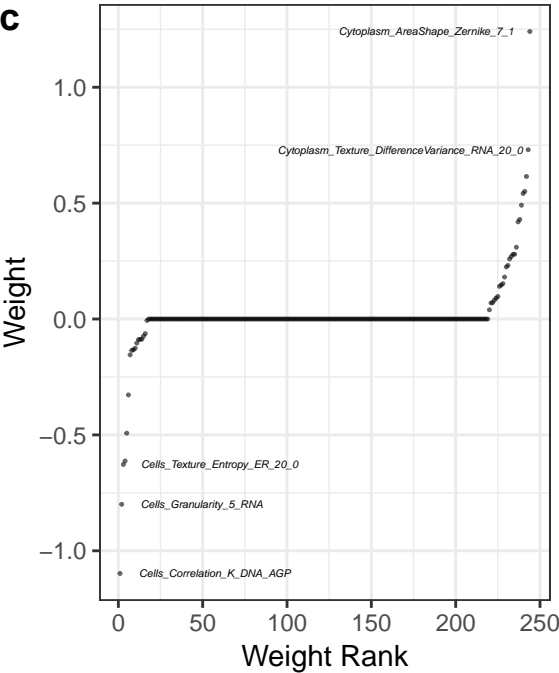
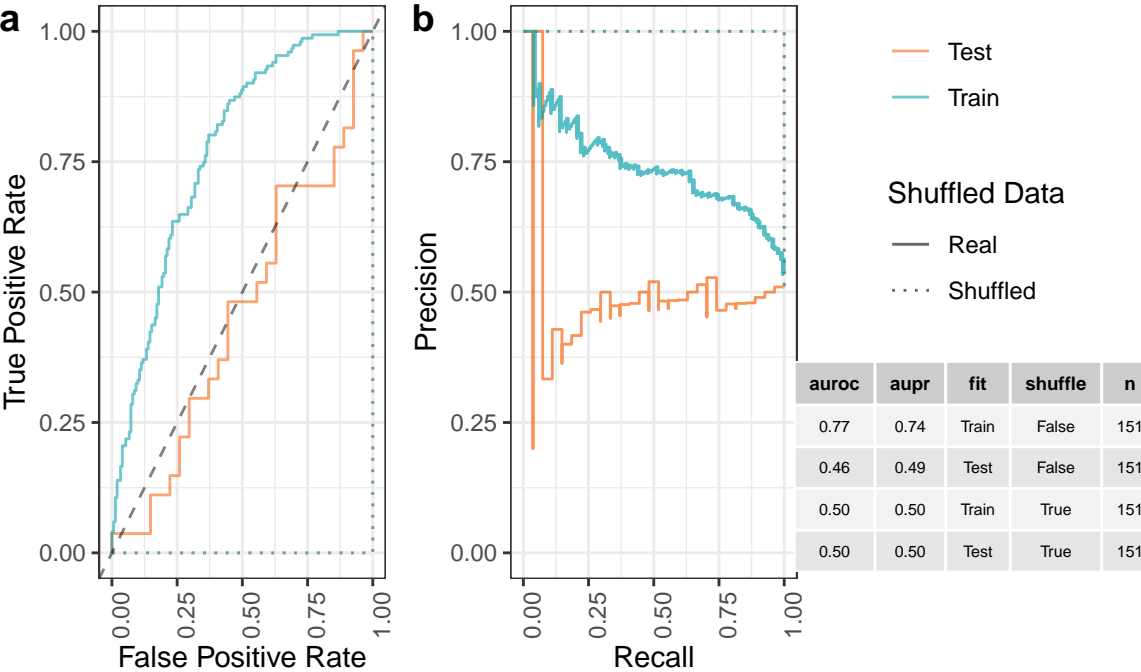
Performance: cc\_mitosis\_ph3\_pos\_n\_objects



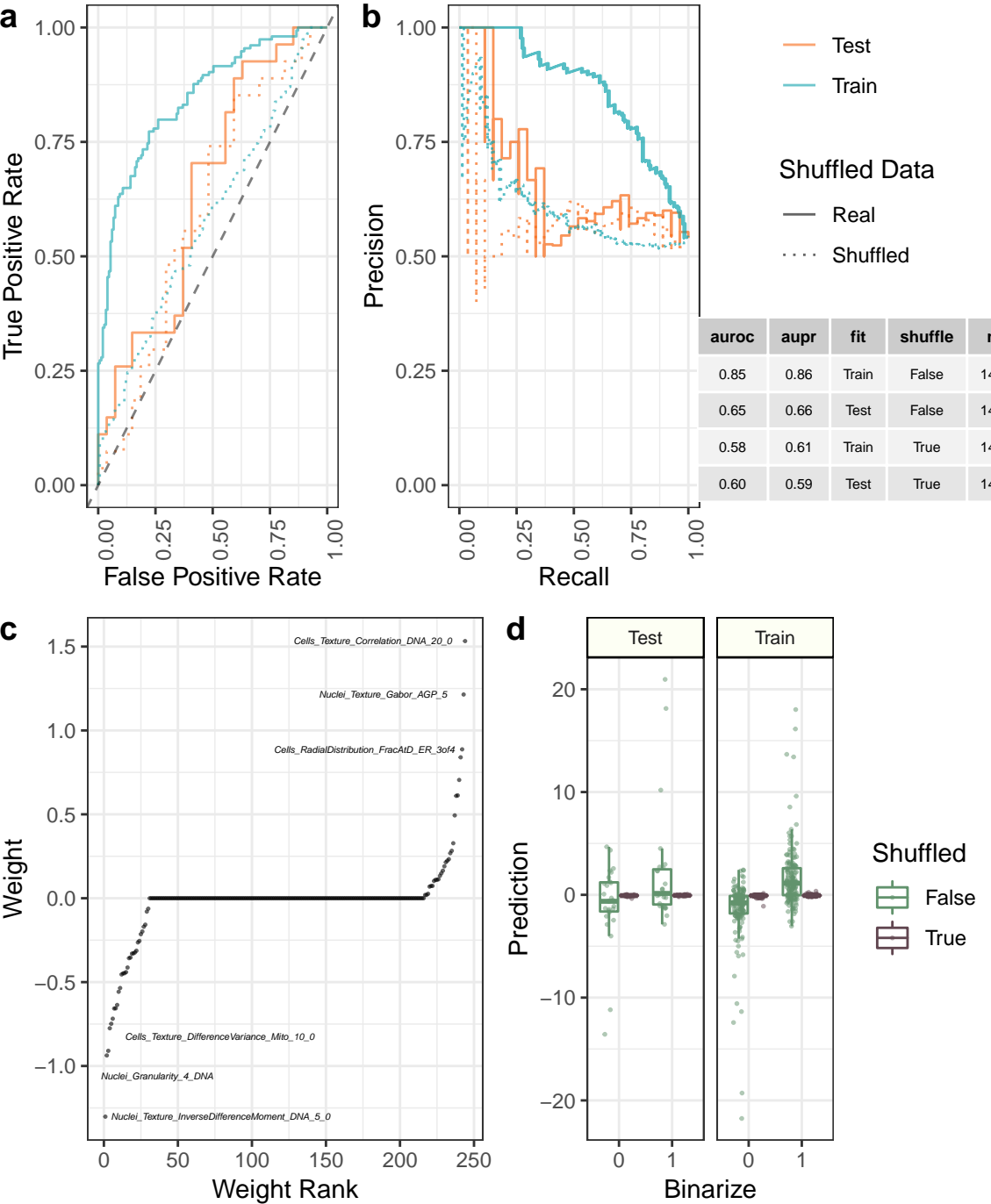
Performance: cc\_mitosis\_ph3\_pos\_n\_spots\_mean



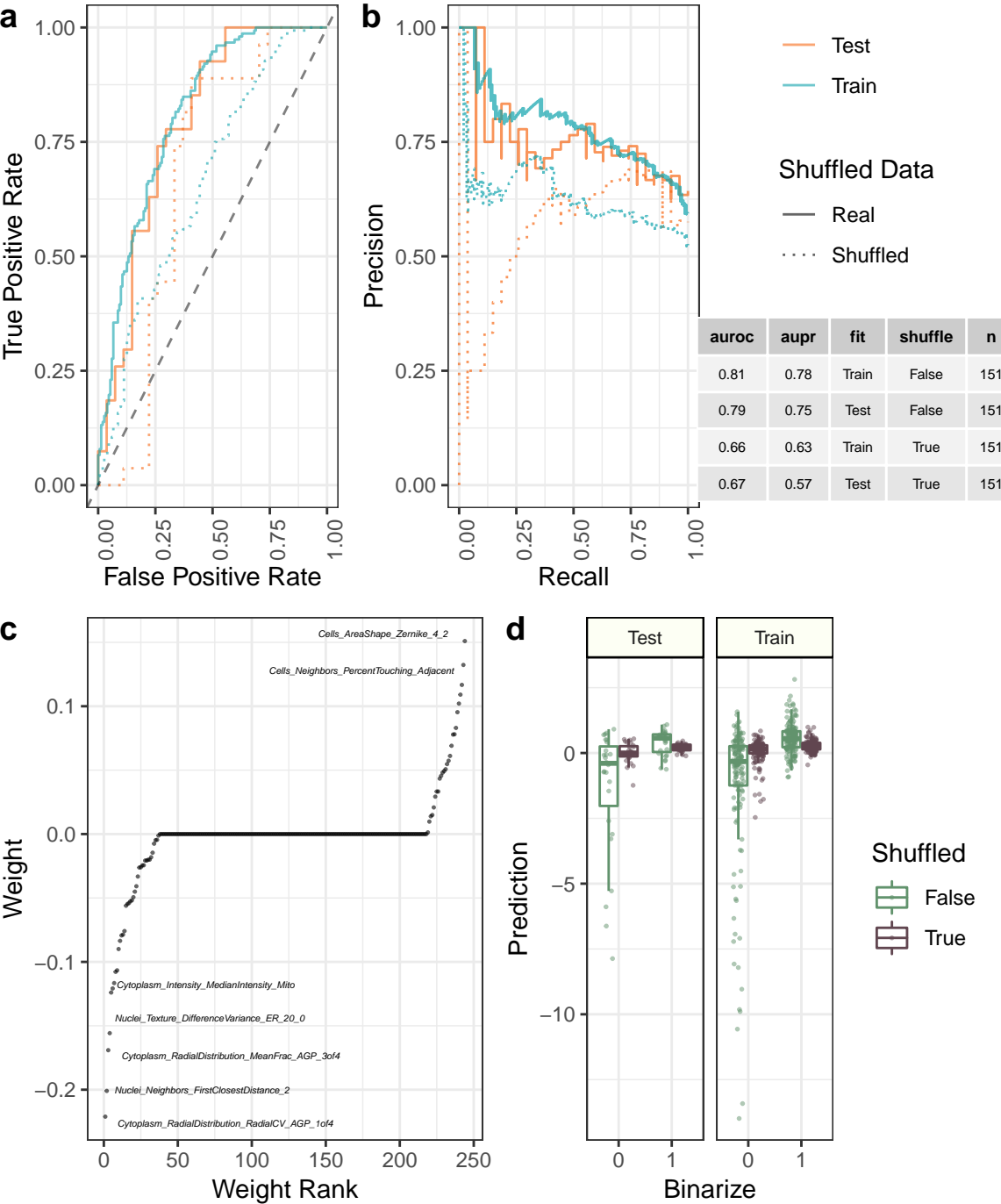
Performance: cc\_mitosis\_ph3\_pos\_n\_spots\_per\_nucleus\_area



Performance: cc\_polynuclear\_high\_n\_spots\_h2ax\_mean

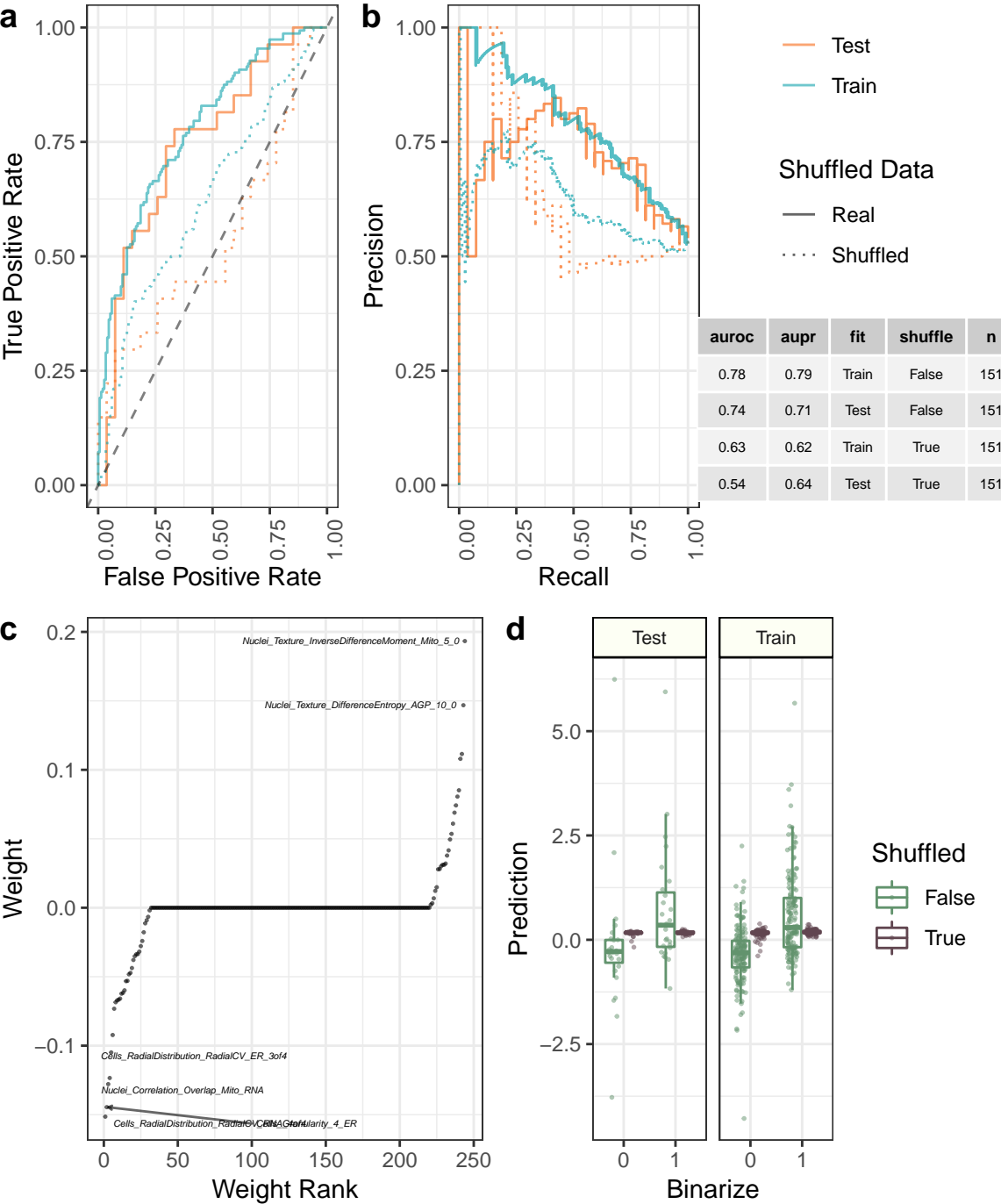


# Performance: cc\_polynuclear\_n\_objects

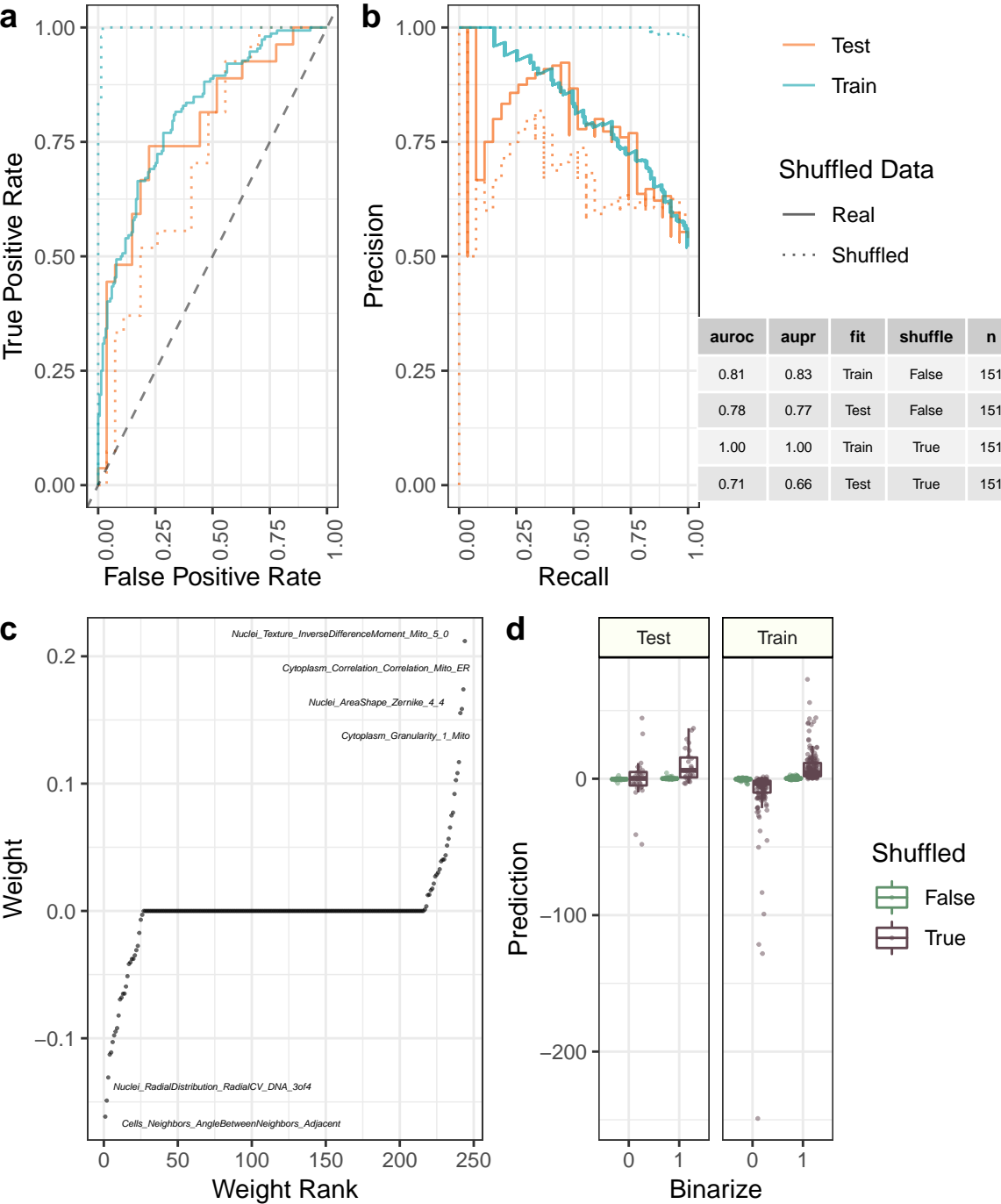




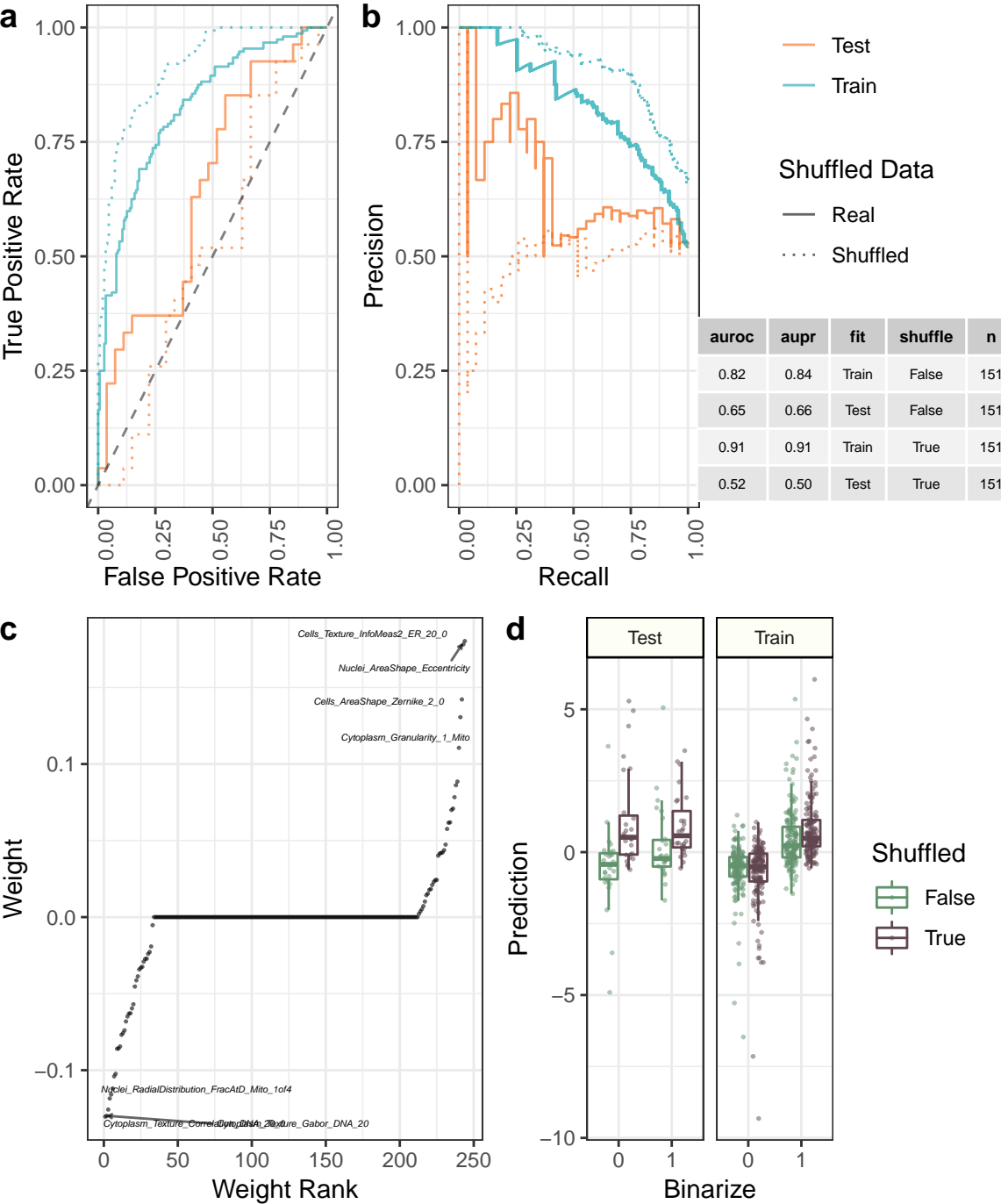
Performance: cc\_polynuclear\_n\_spots\_mean



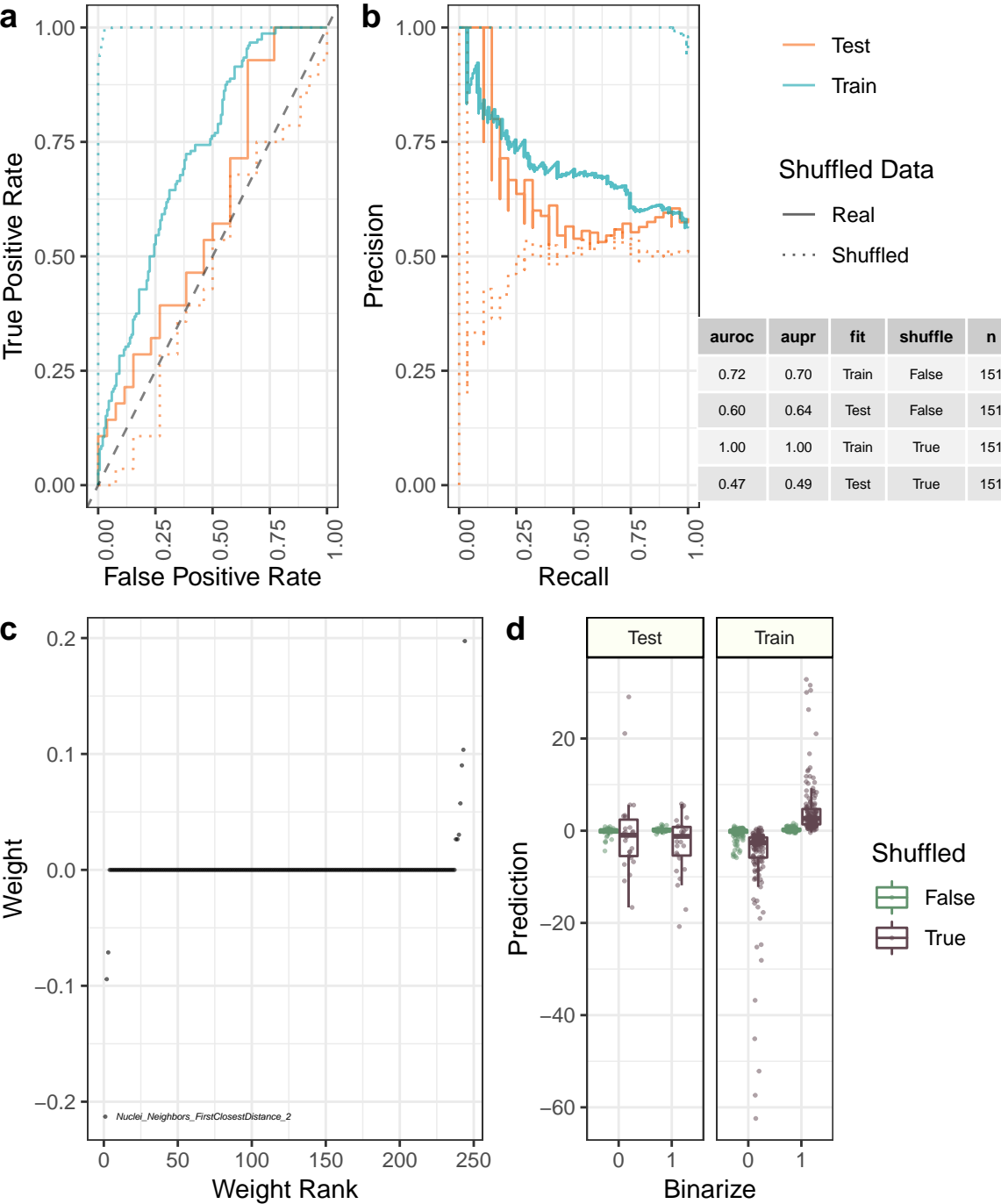
Performance: cc\_polynuclear\_n\_spots\_per\_nucleus\_area\_mean



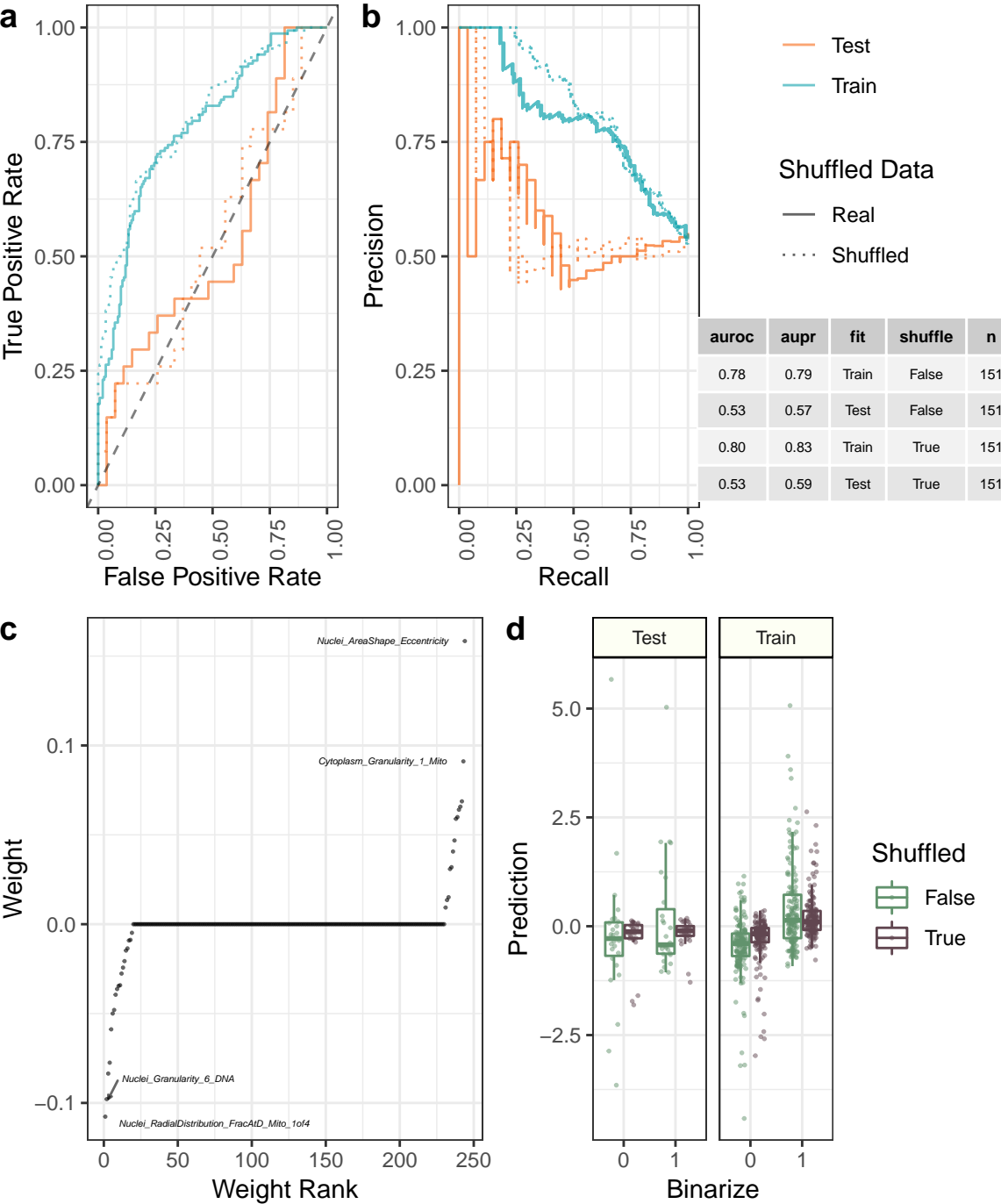
Performance: cc\_polyloid\_high\_n\_spots\_h2ax\_mean



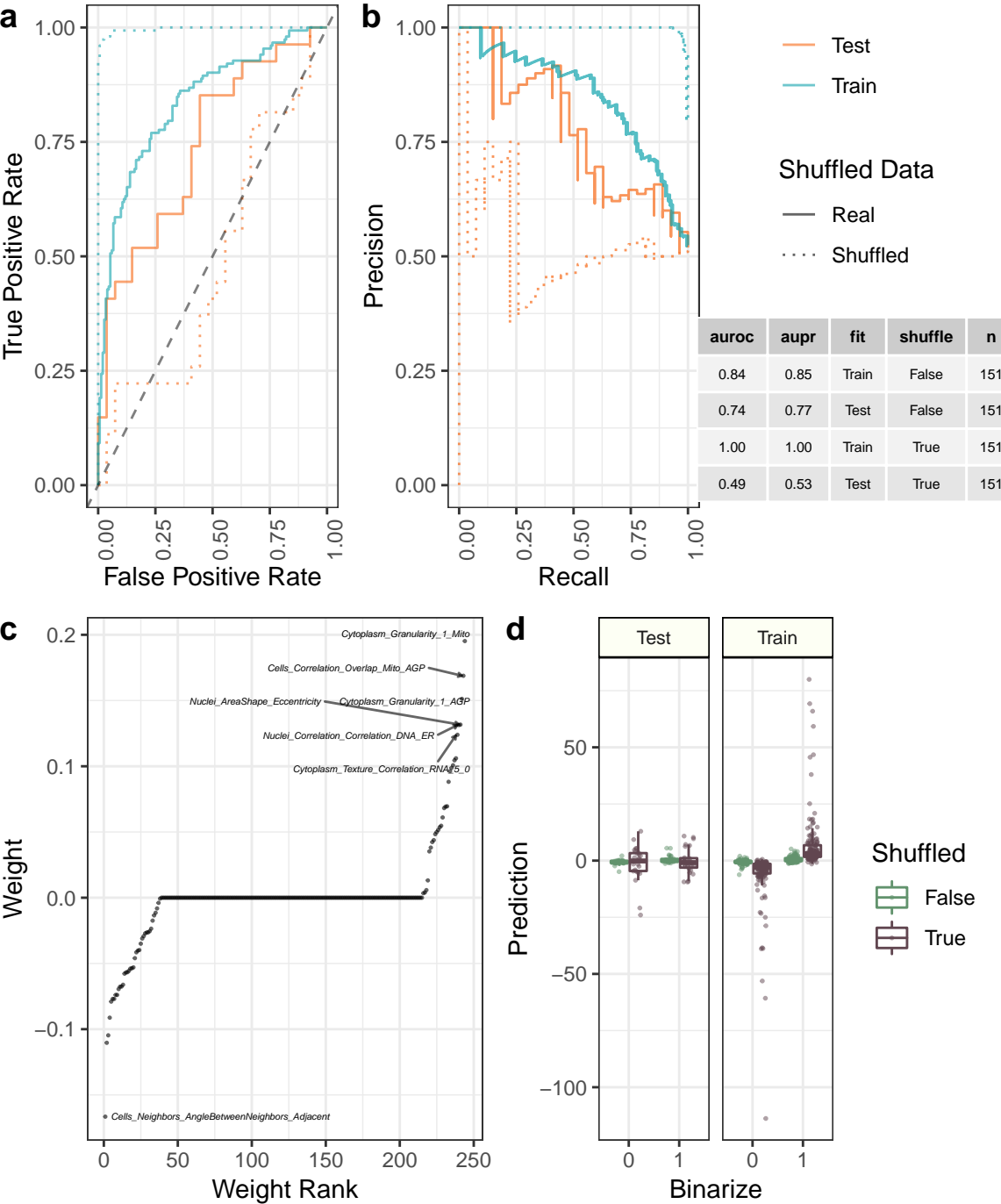
# Performance: cc\_polyloid\_n\_objects



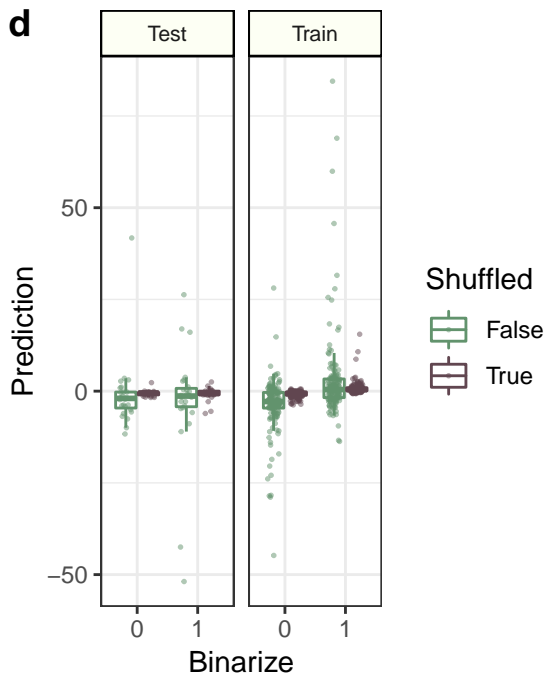
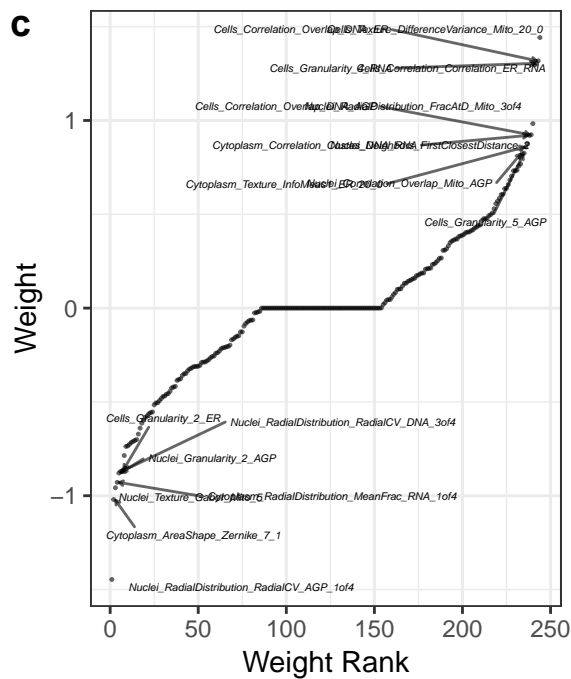
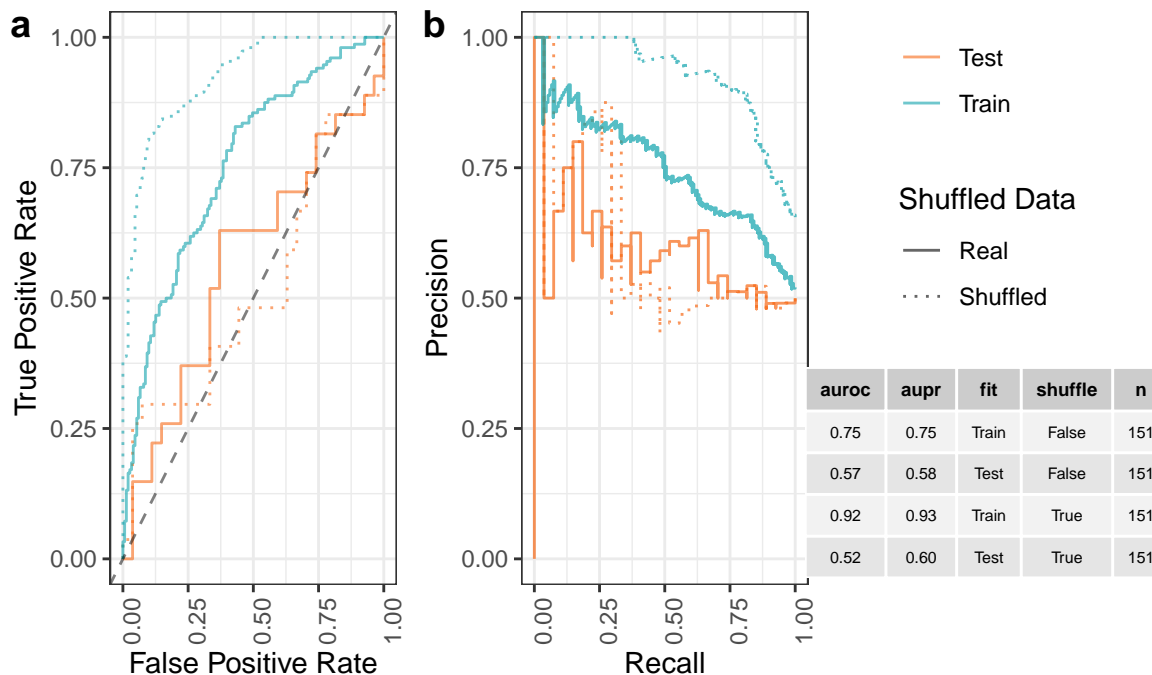
# Performance: cc\_polyloid\_n\_spots\_mean



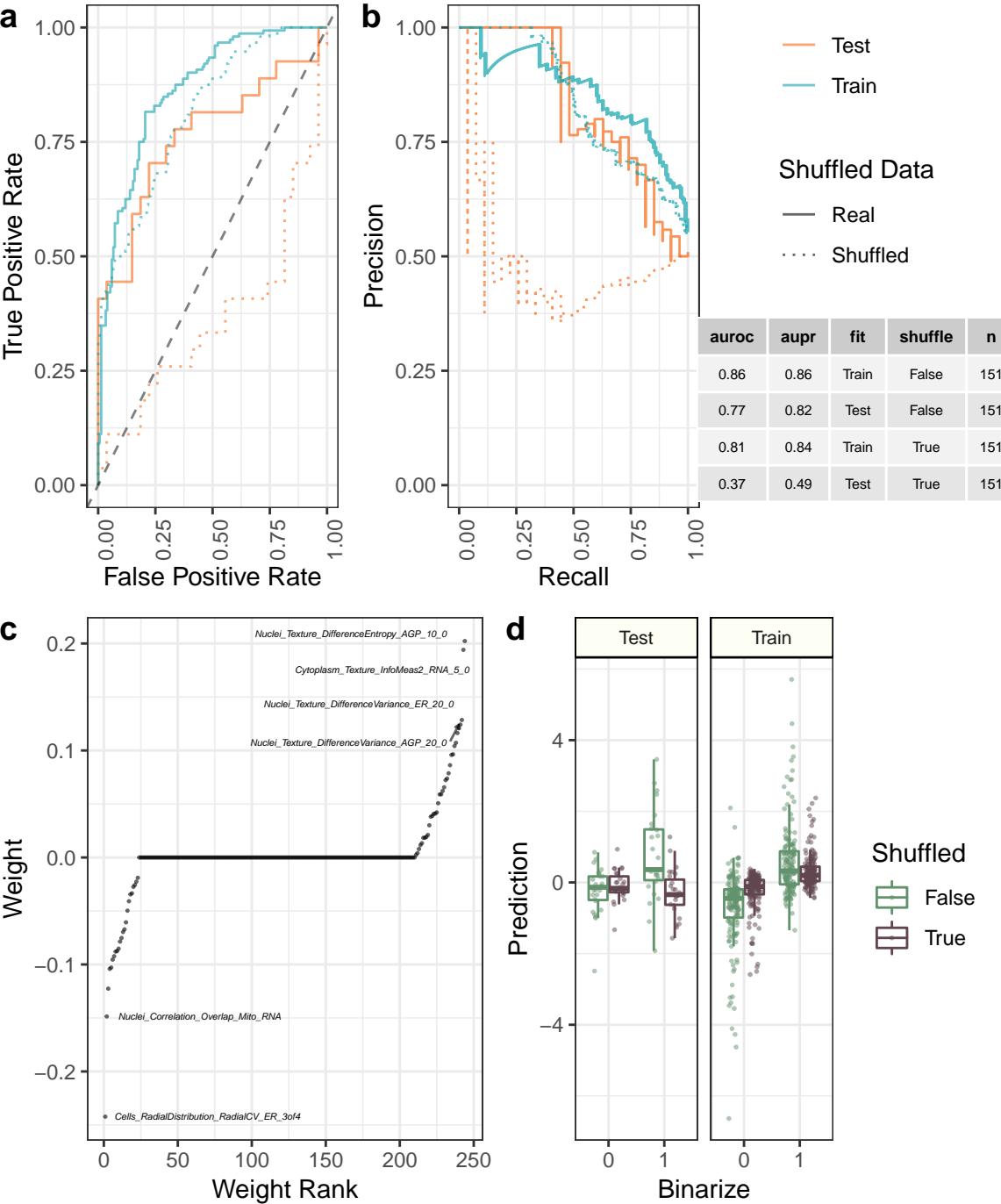
Performance: cc\_polyploid\_n\_spots\_per\_nucleus\_area\_mean



### Performance: vb\_infection\_percentage



# Performance: vb\_live\_cell\_area



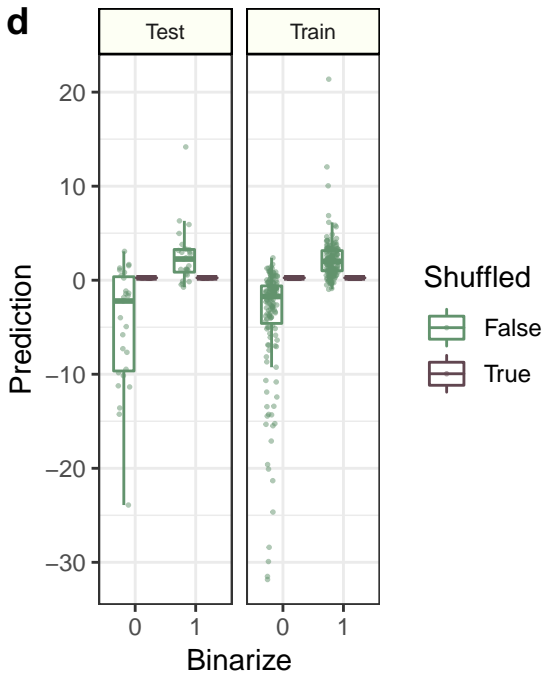
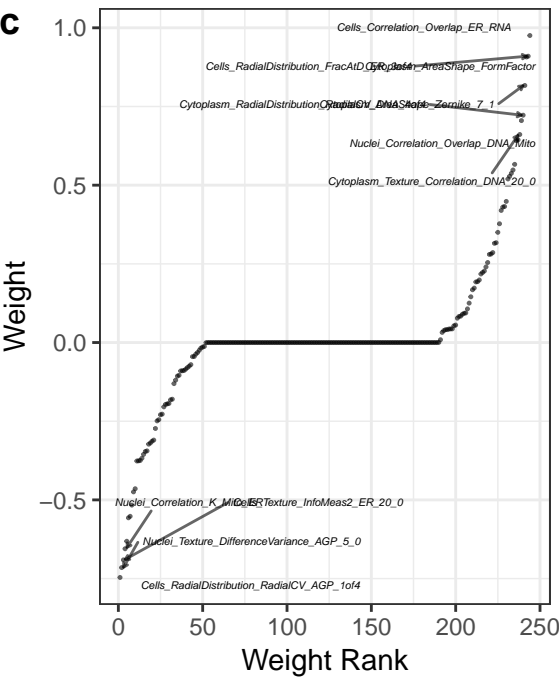
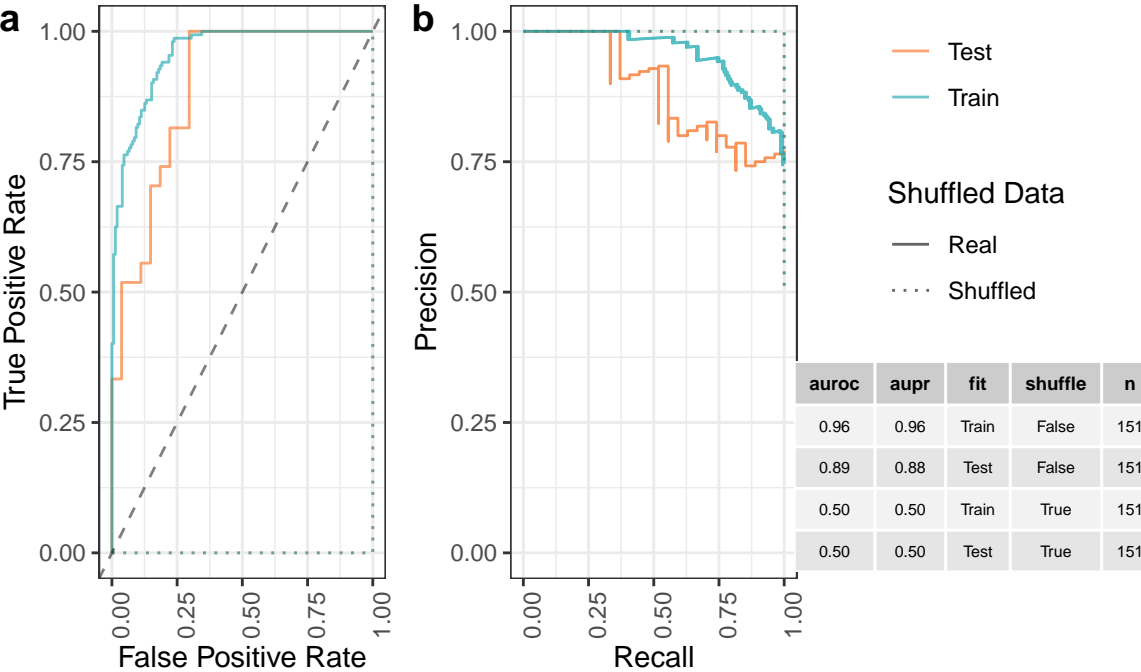
Shuffled

False

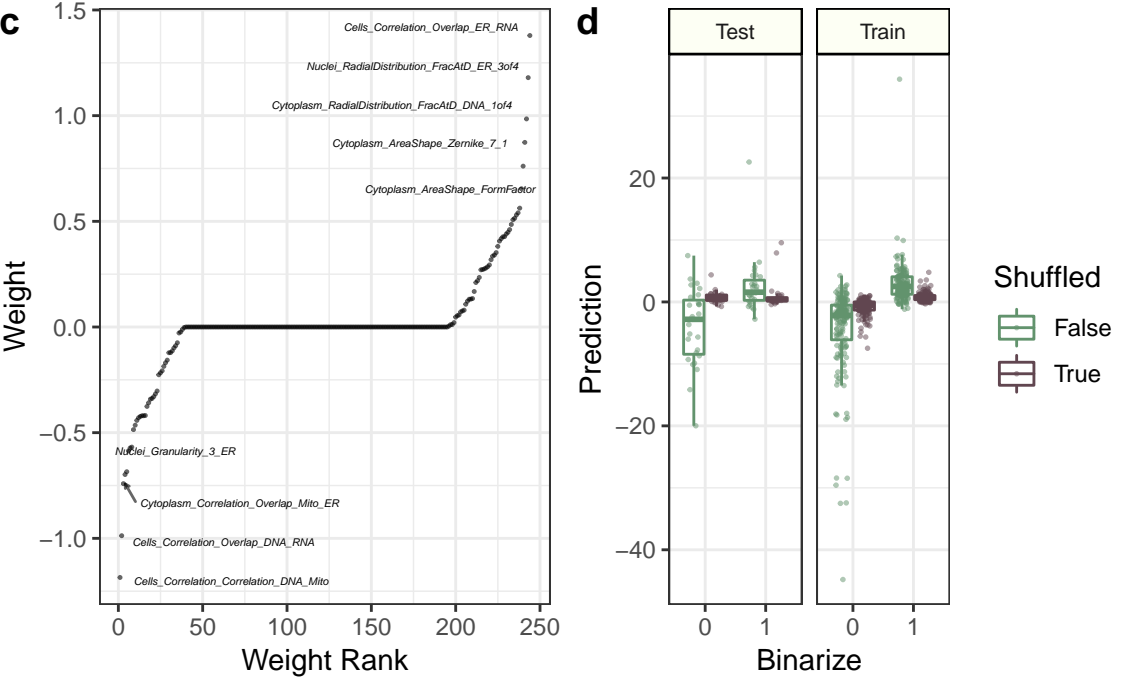
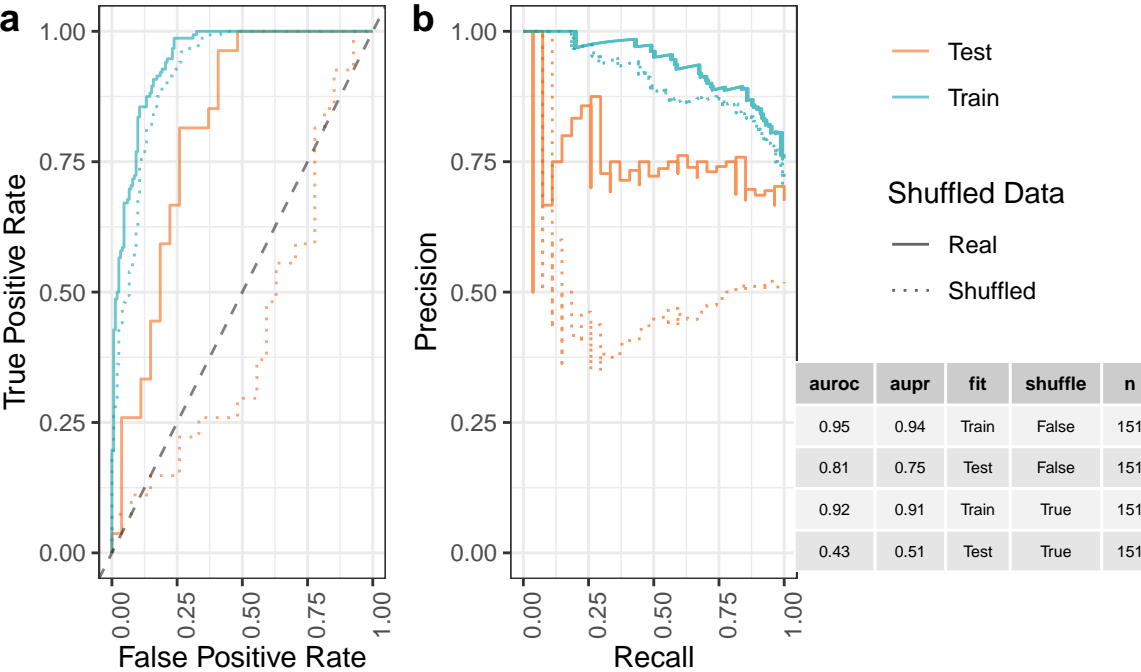
True



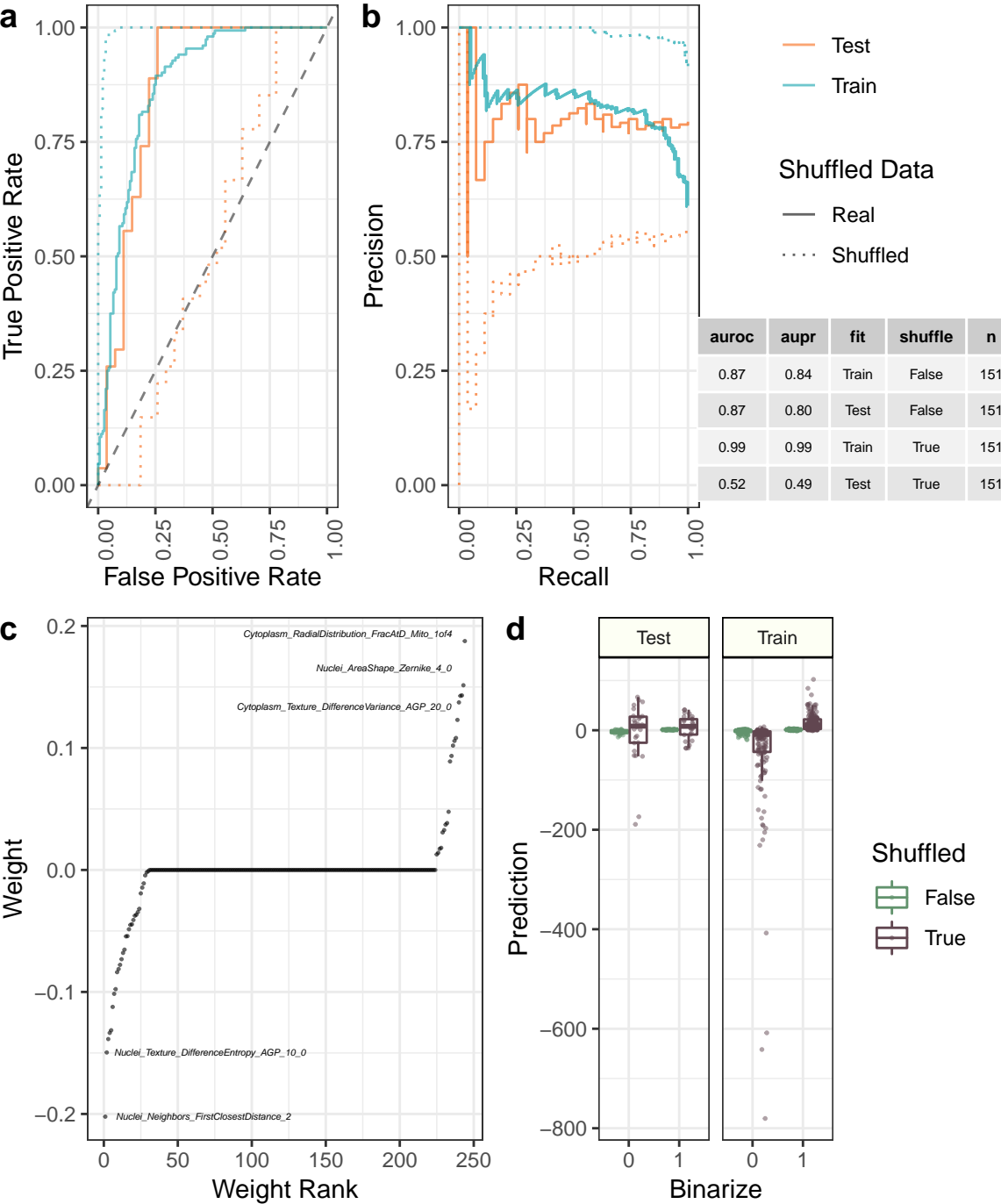
Performance: vb\_live\_cell\_roundness



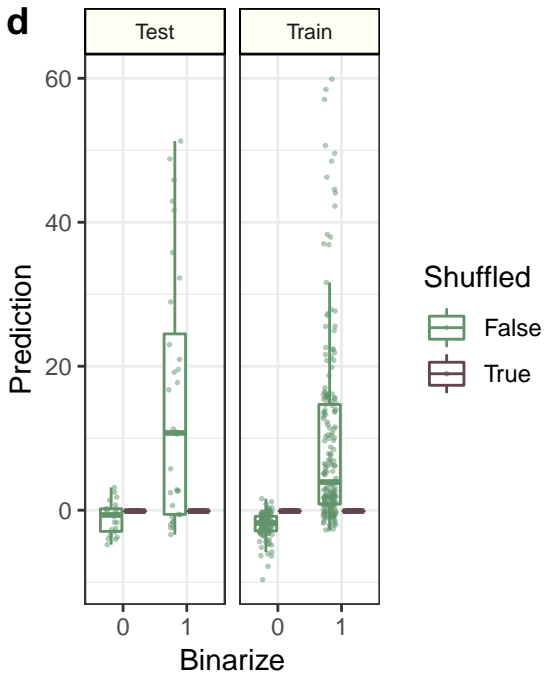
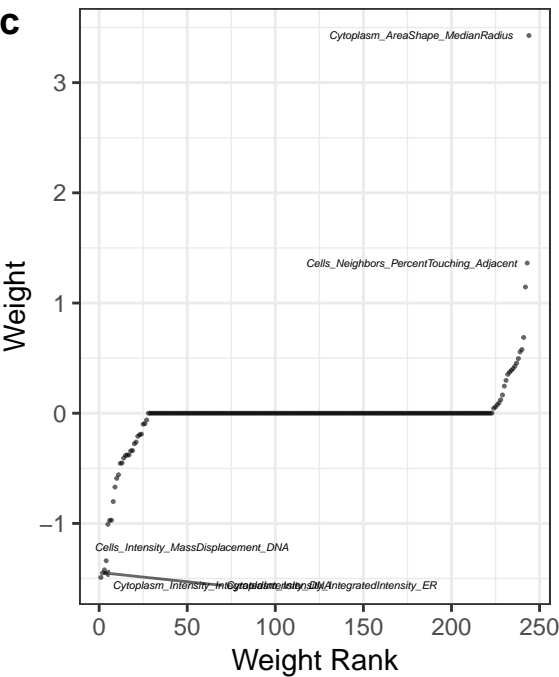
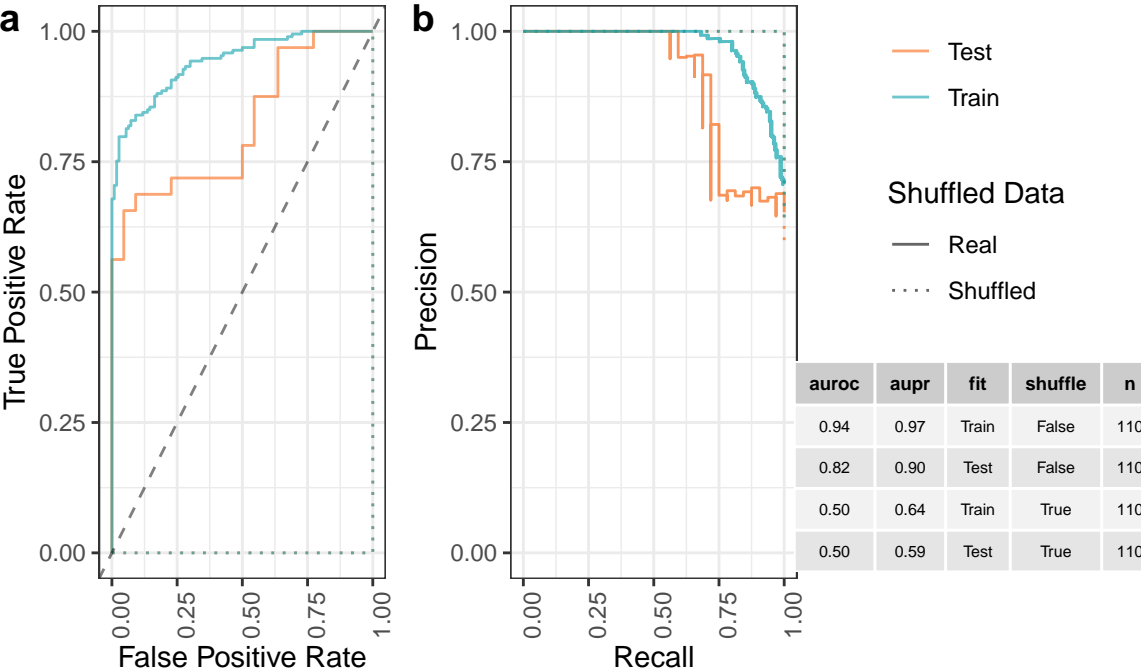
# Performance: vb\_live\_cell\_width\_length



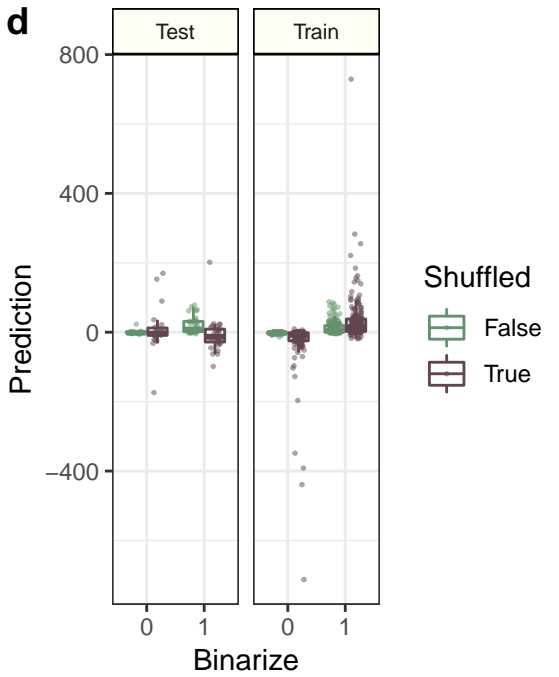
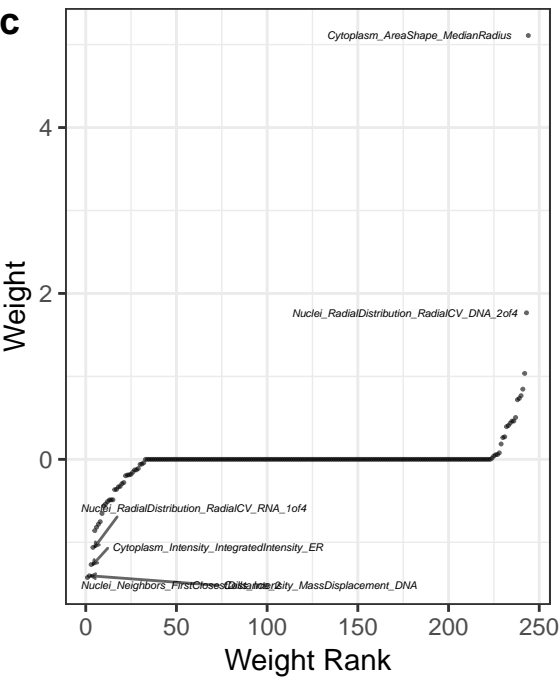
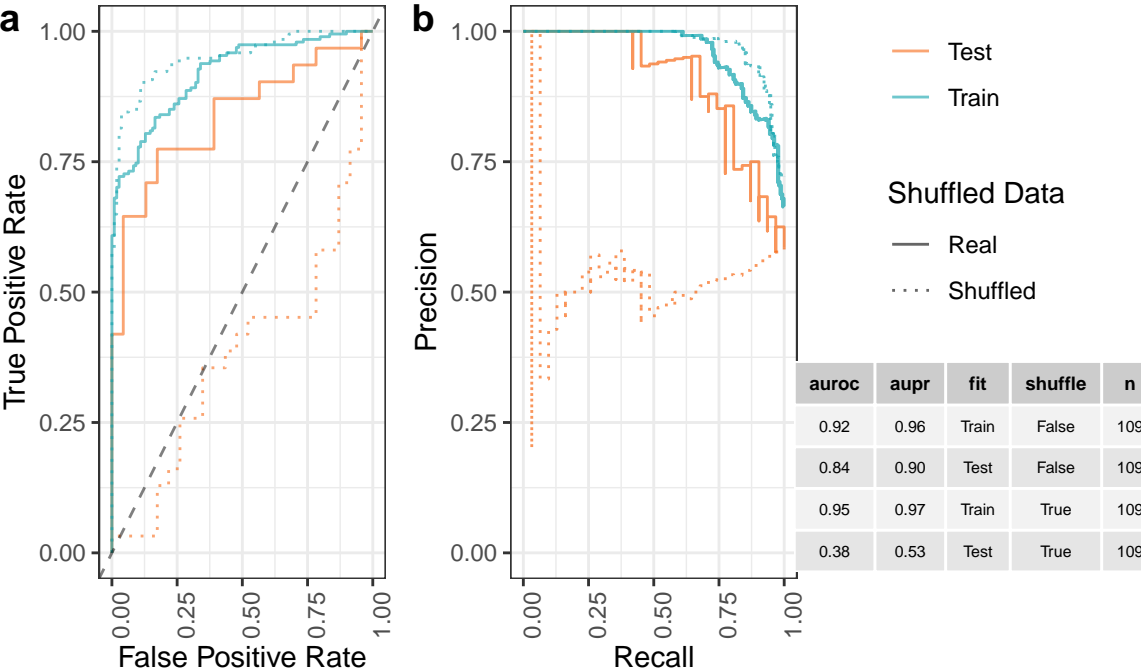
# Performance: vb\_num\_live\_cells



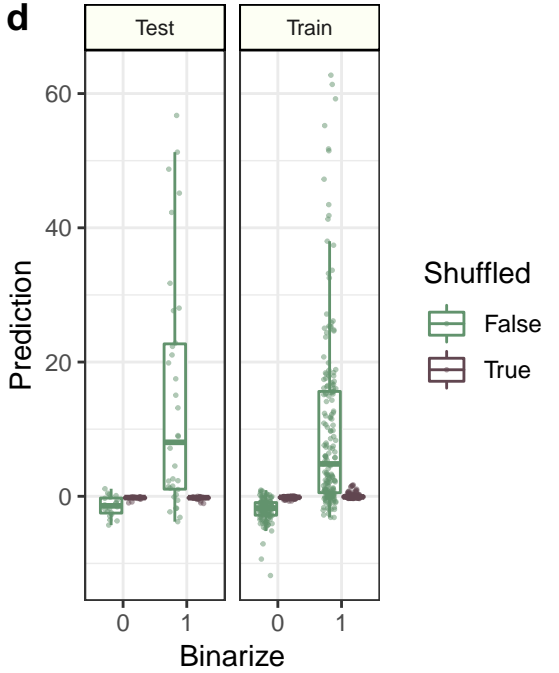
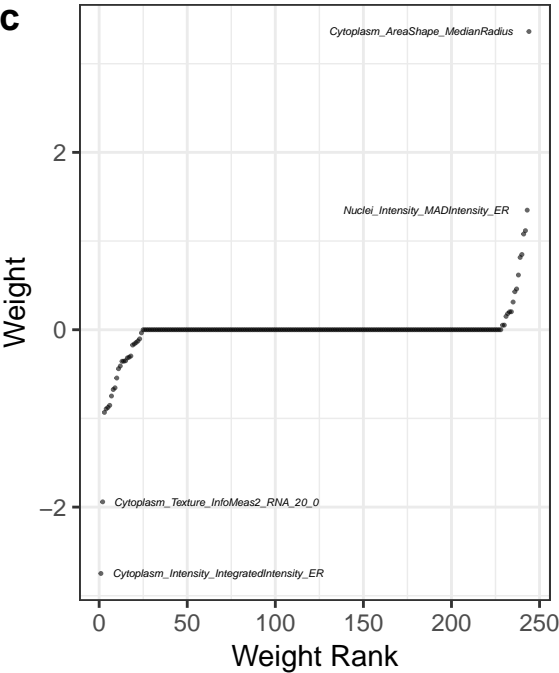
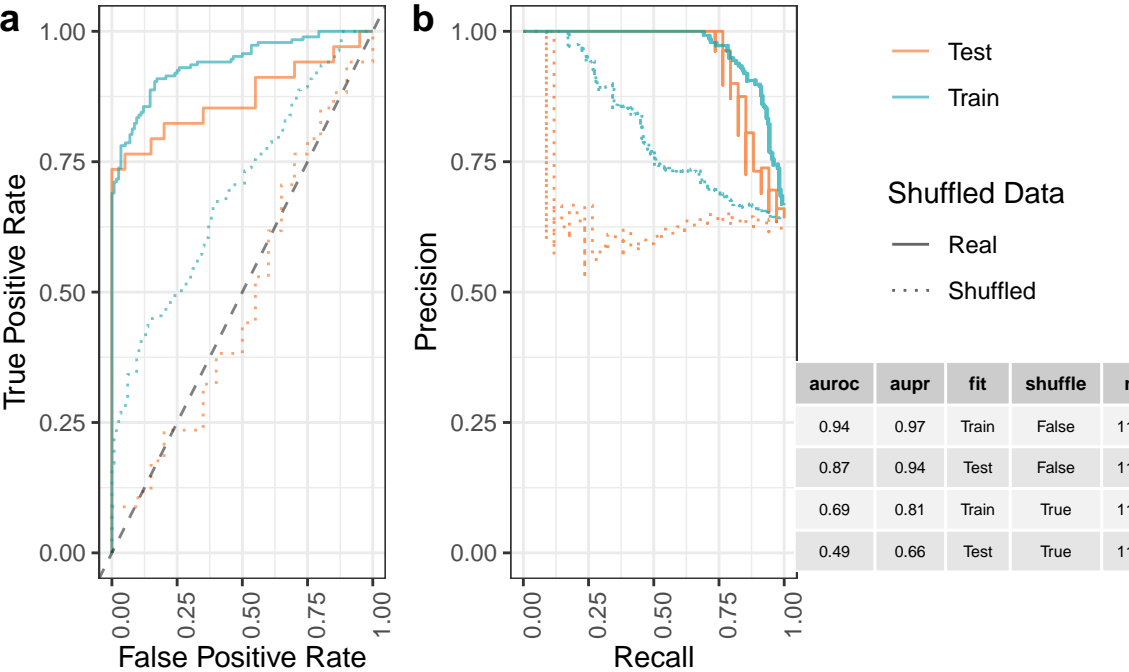
# Performance: vb\_percent\_all\_apoptosis



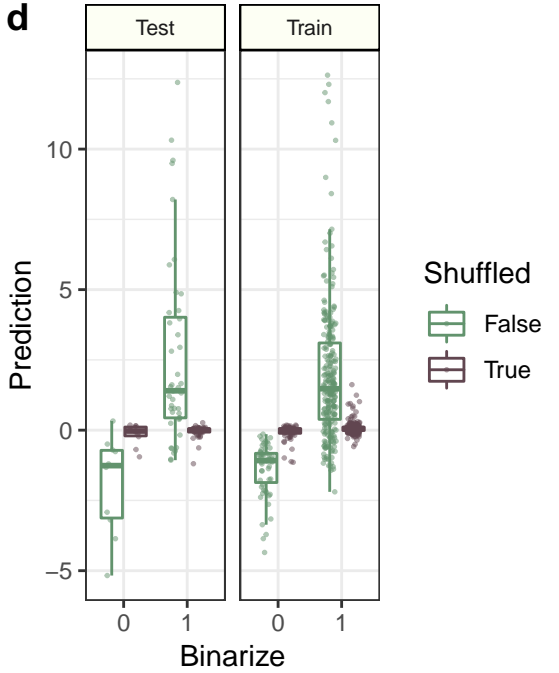
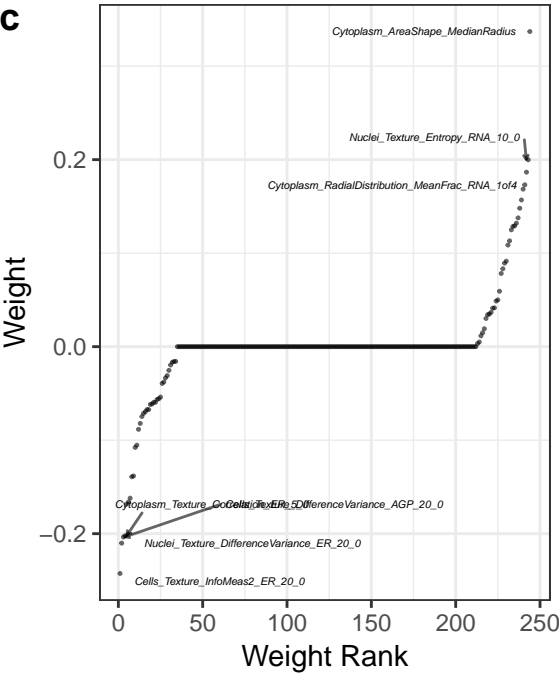
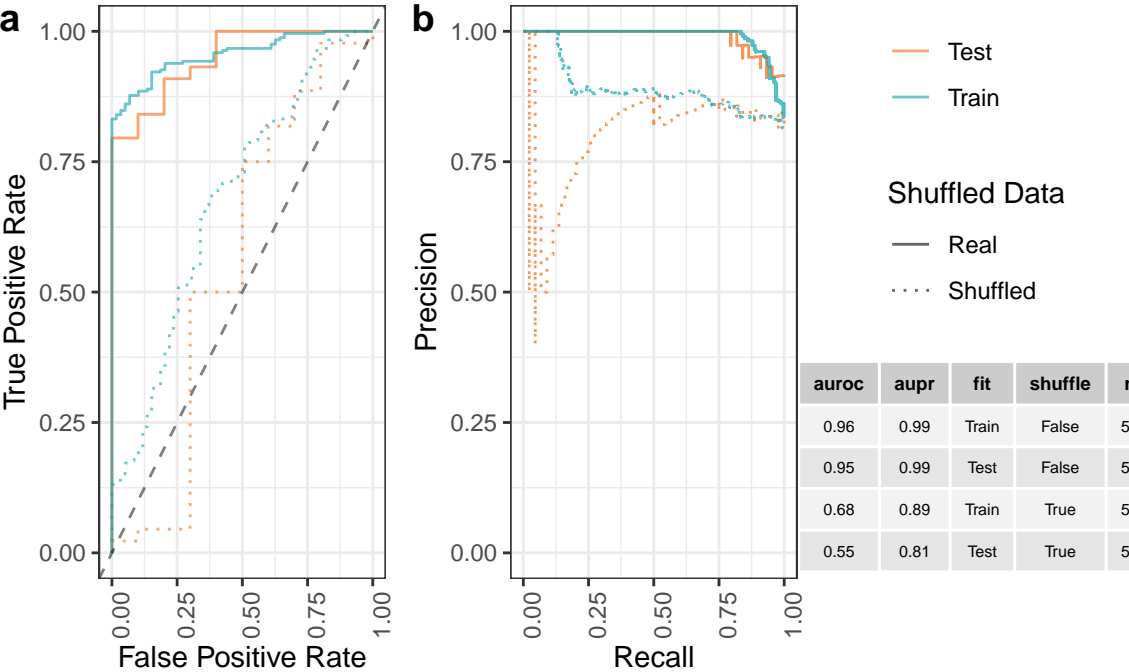
# Performance: vb\_percent\_all\_early\_apoptosis



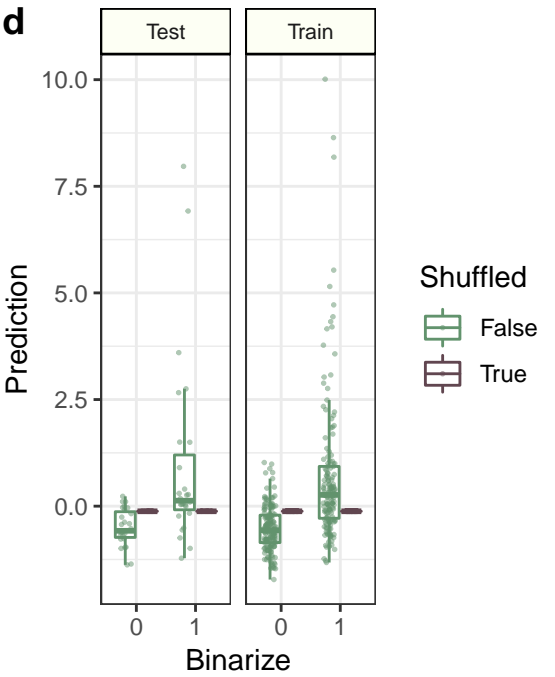
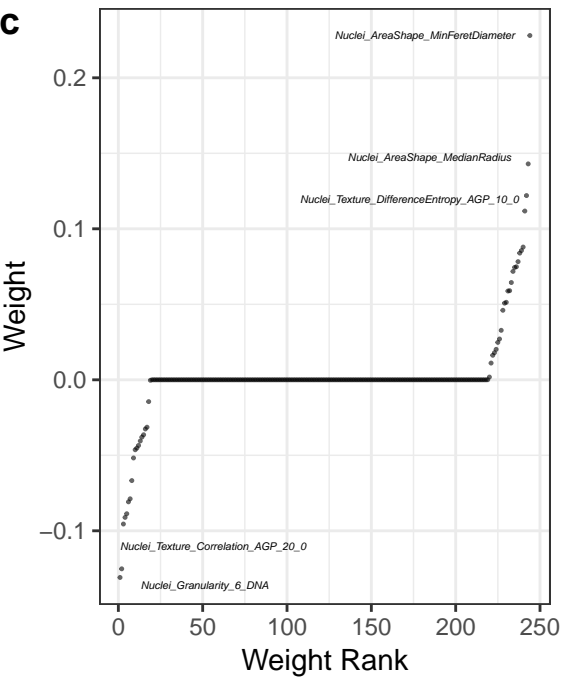
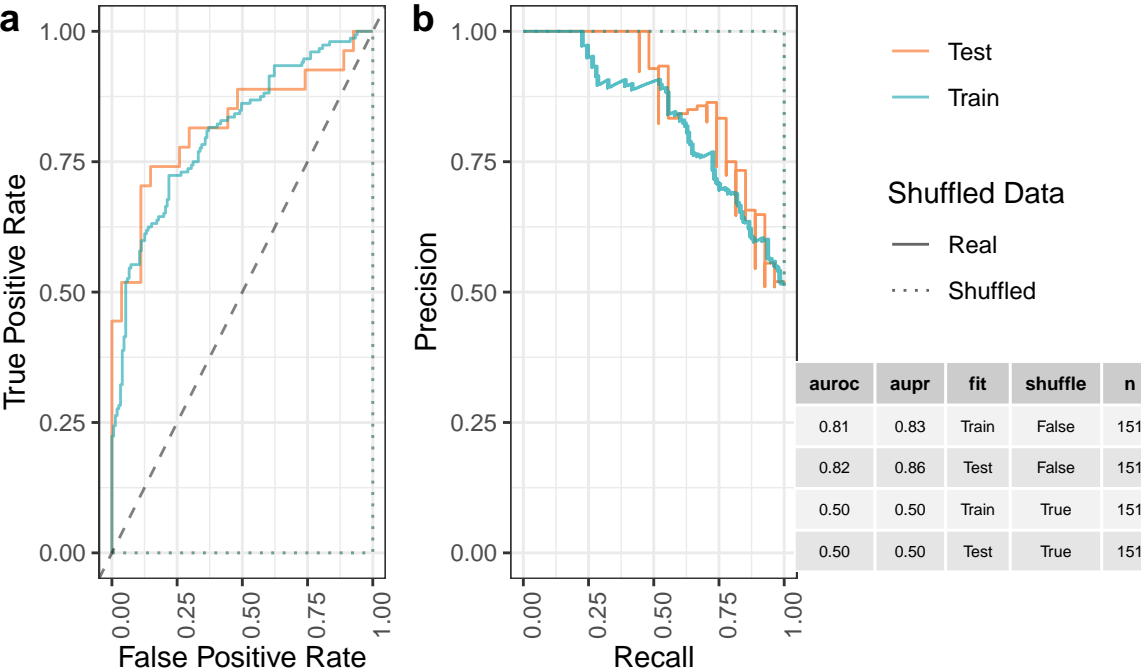
# Performance: vb\_percent\_all\_late\_apoptosis



Performance: vb\_percent\_caspase\_dead\_only

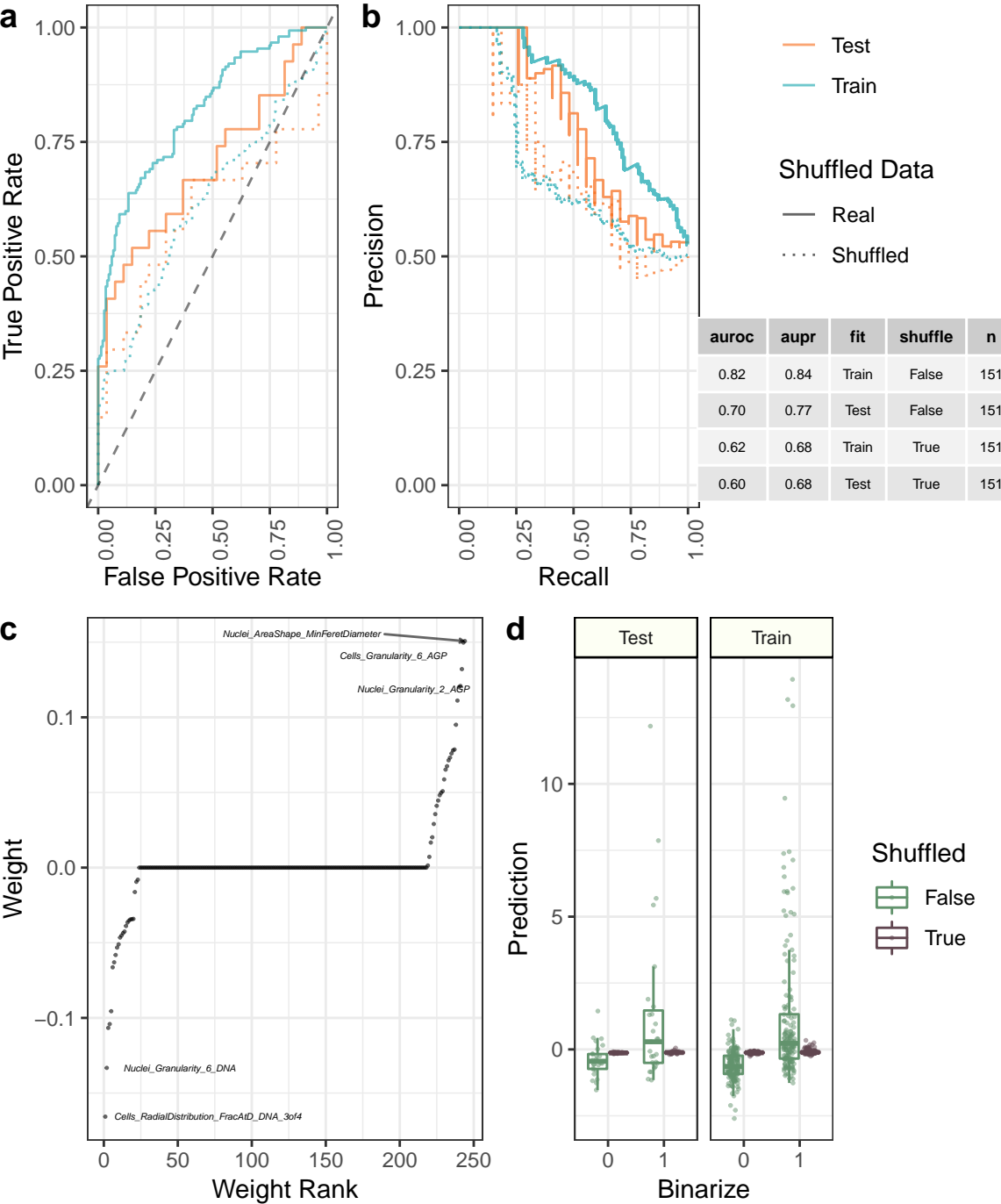


# Performance: vb\_percent\_dead

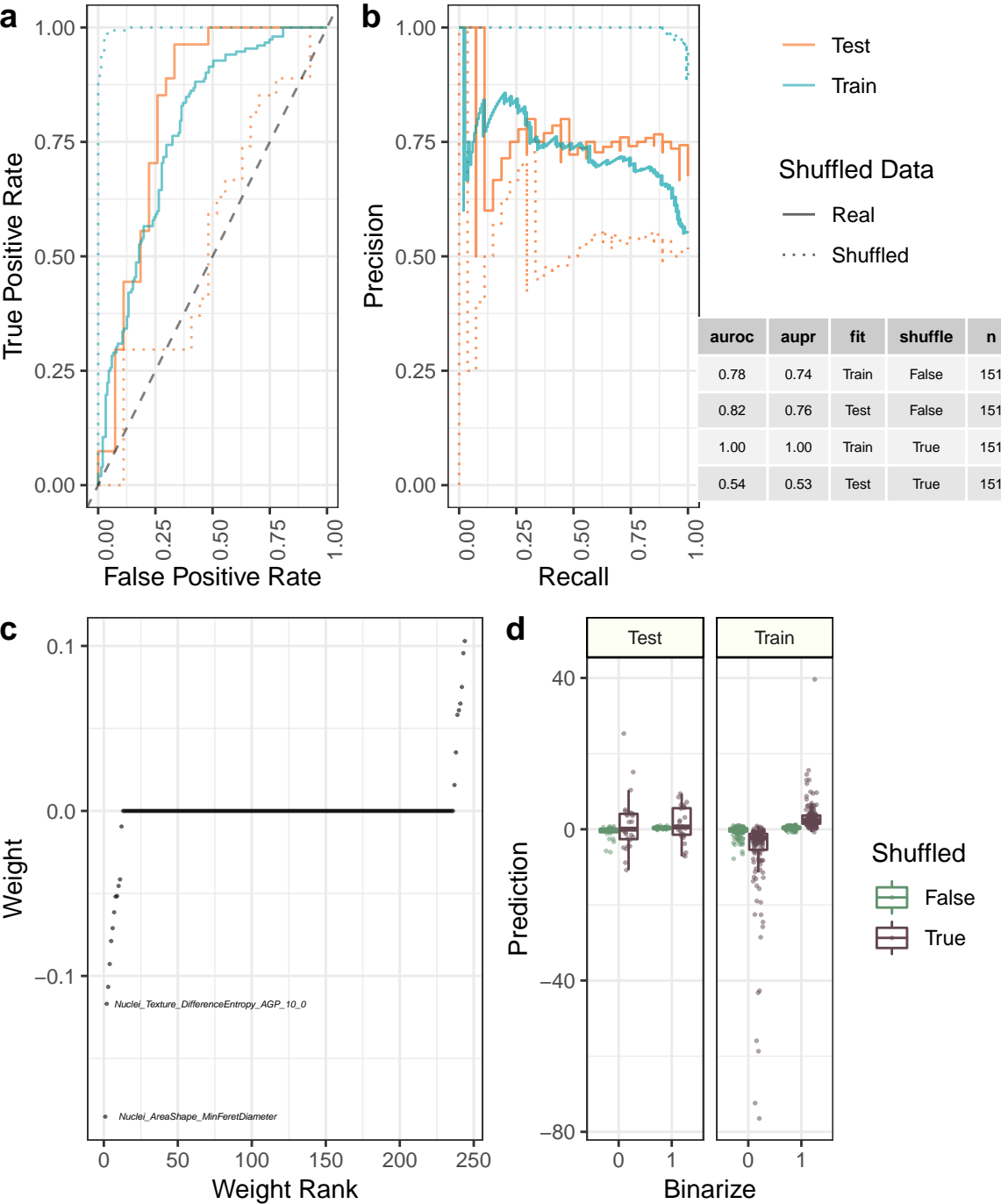




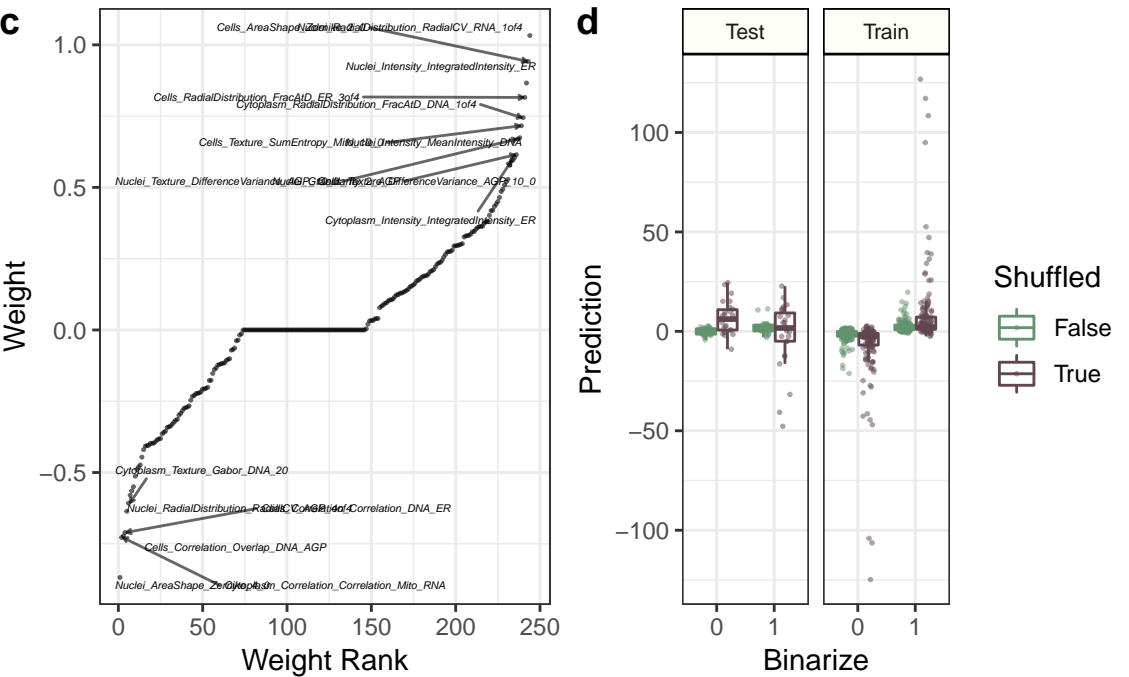
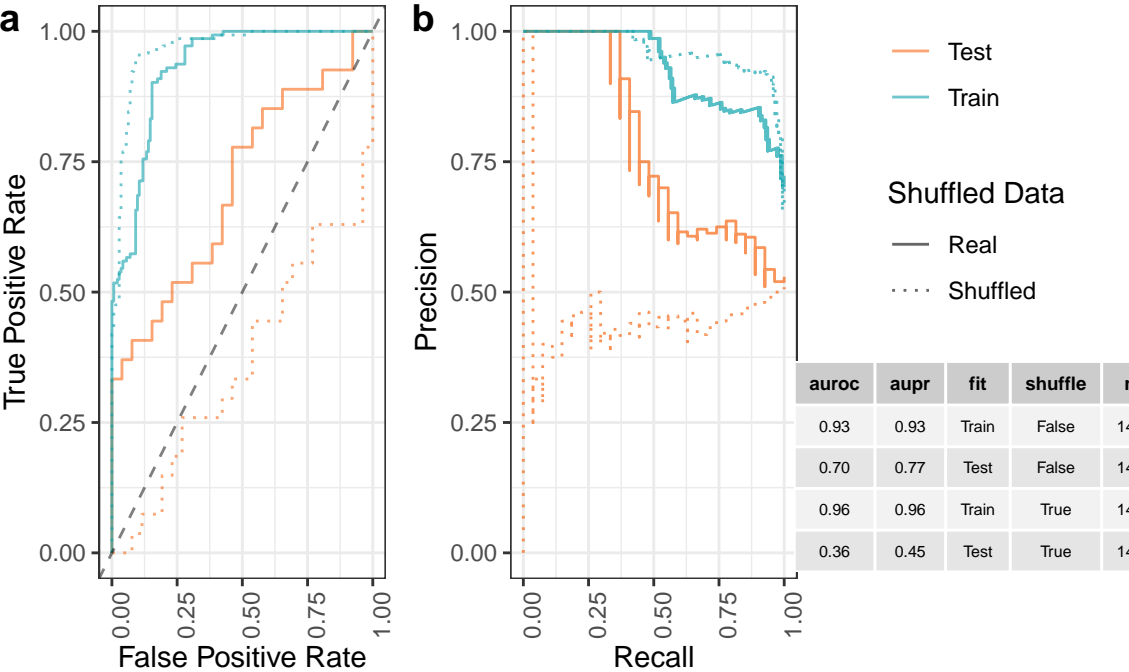
# Performance: vb\_percent\_dead\_only



# Performance: vb\_percent\_live



# Performance: vb\_ros\_back\_mean



# Performance: vb\_ros\_mean

