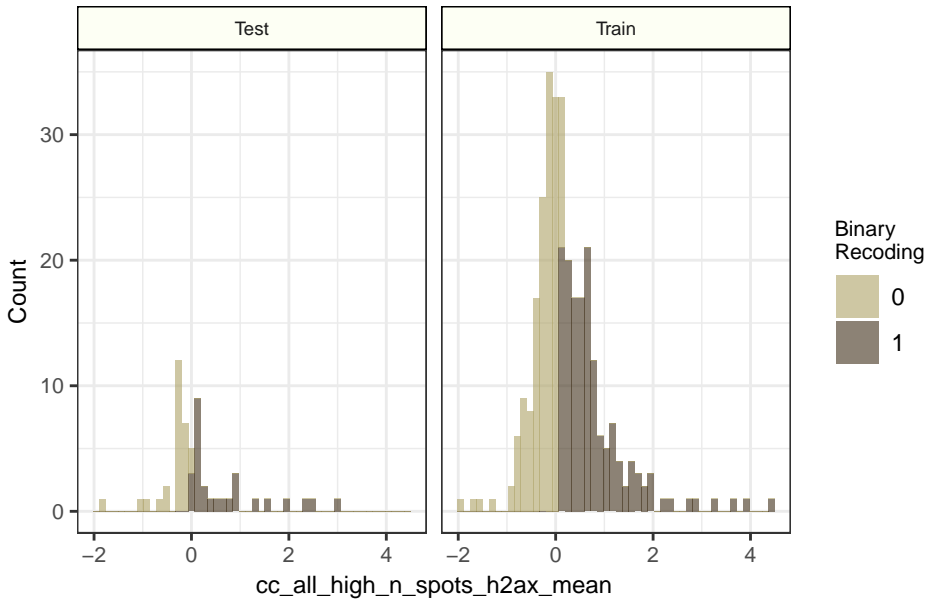
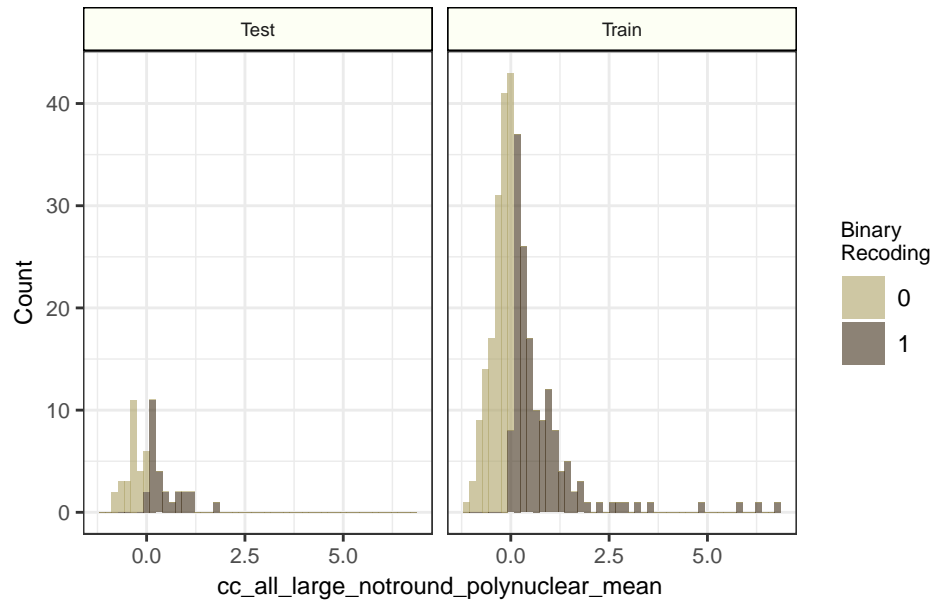


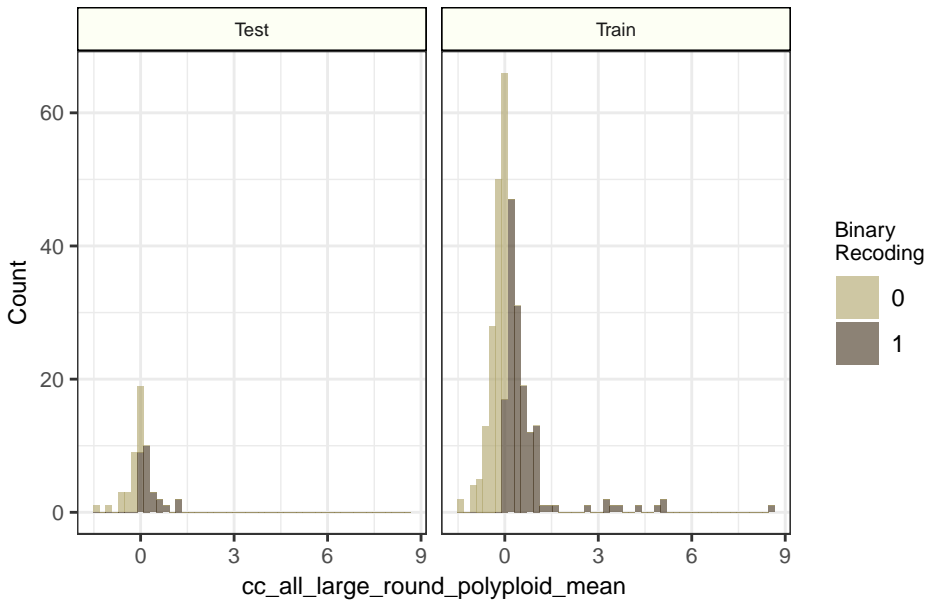
cc\_all\_high\_n\_spots\_h2ax\_mean



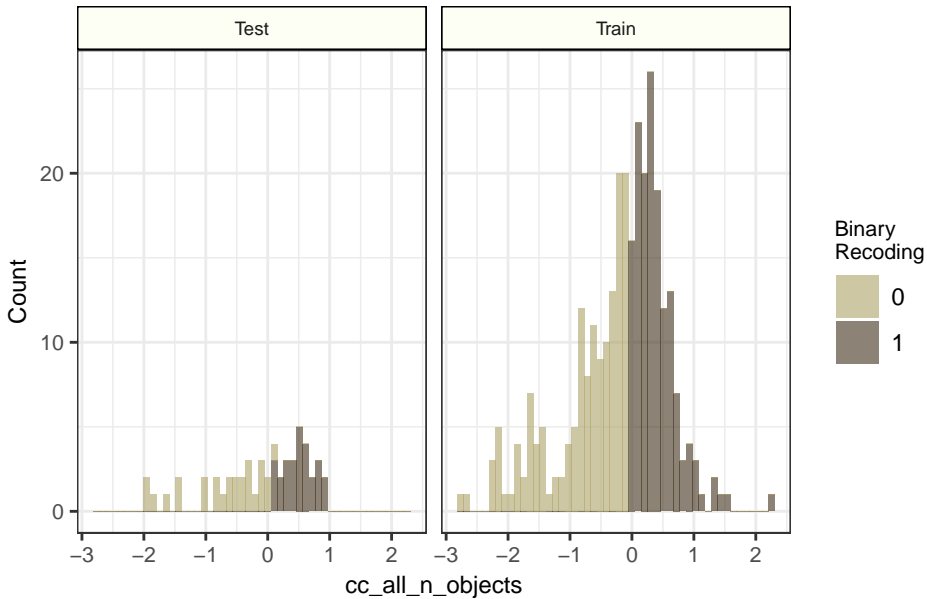
# cc\_all\_large\_notround\_polynuclear\_mean



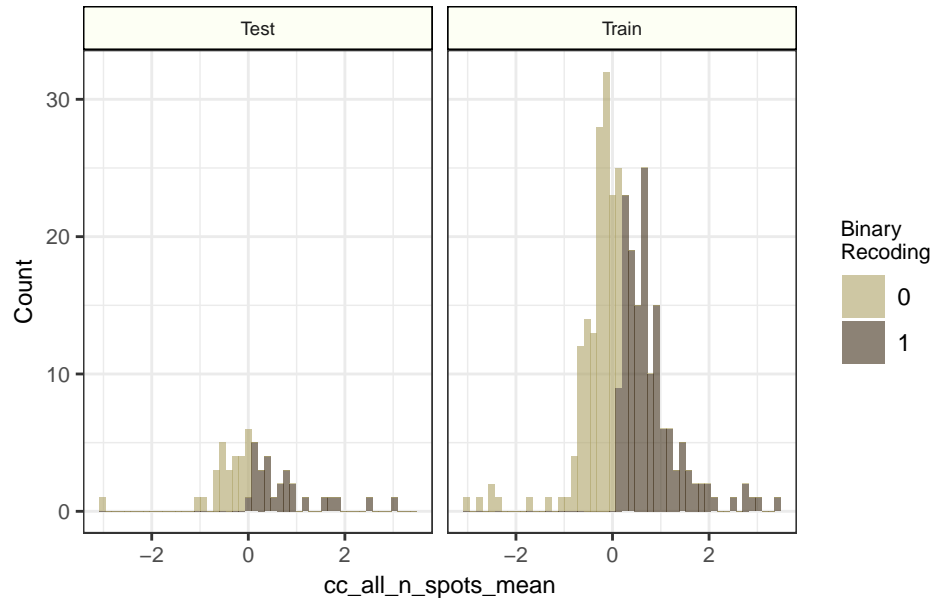
# cc\_all\_large\_round\_polyploid\_mean



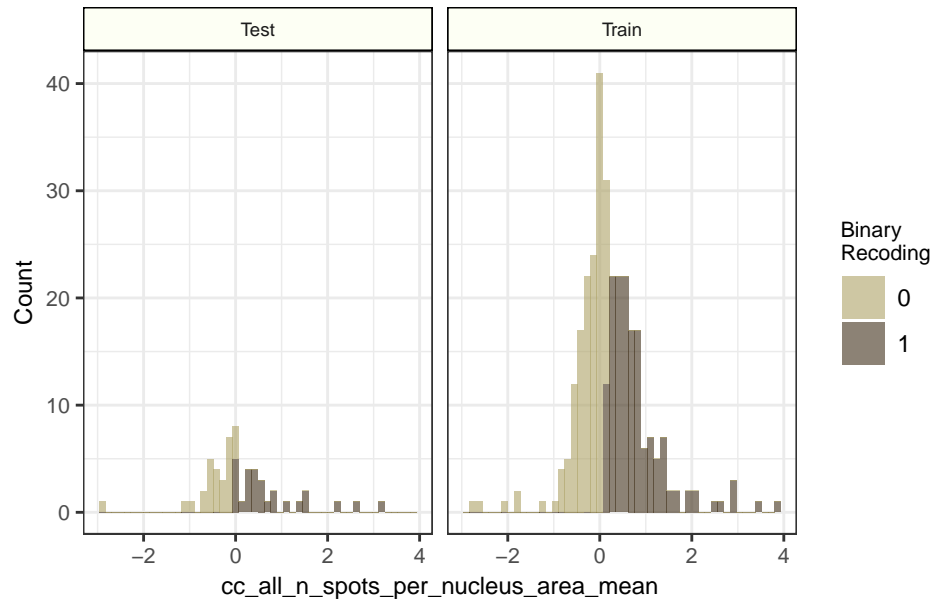
# cc\_all\_n\_objects



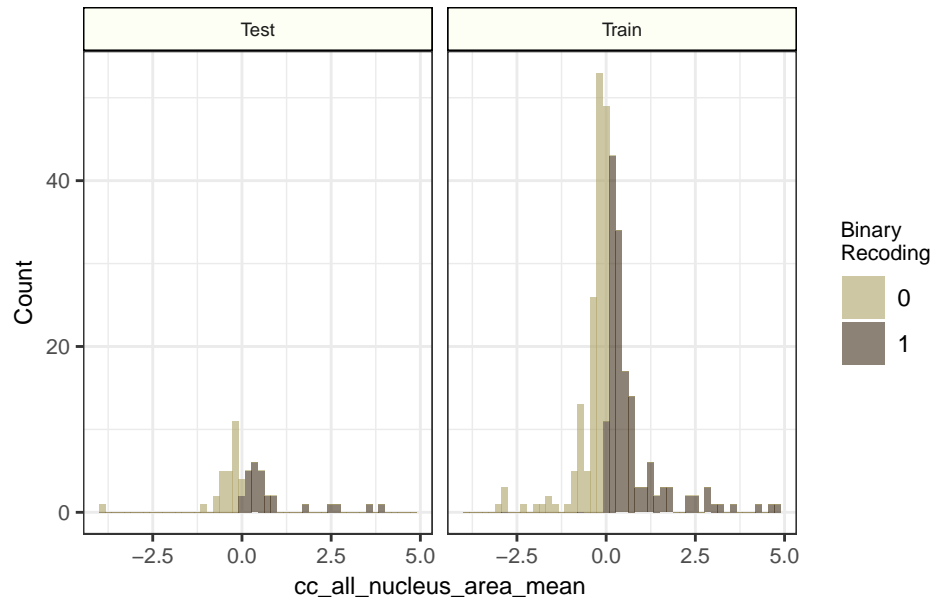
# cc\_all\_n\_spots\_mean



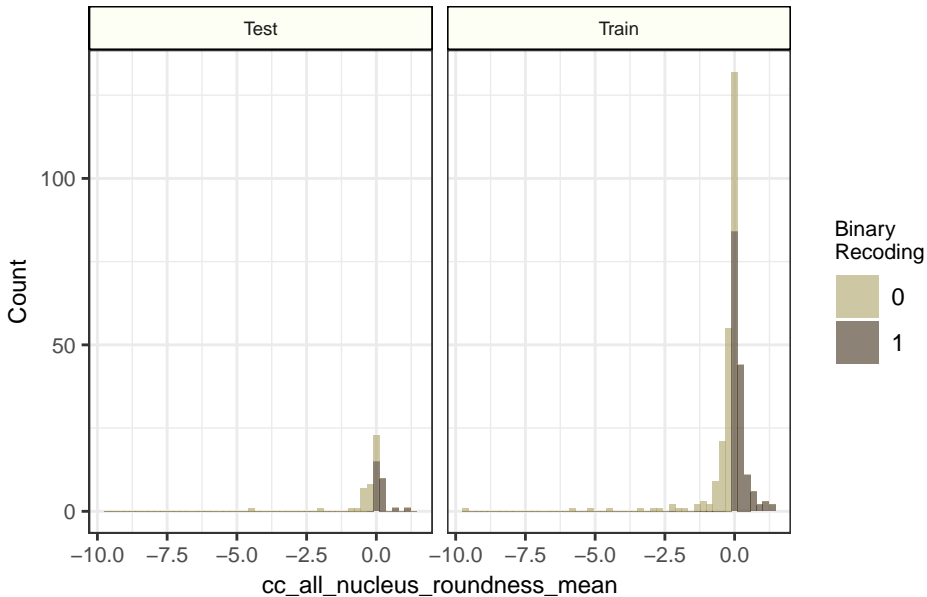
# cc\_all\_n\_spots\_per\_nucleus\_area\_mean



# cc\_all\_nucleus\_area\_mean

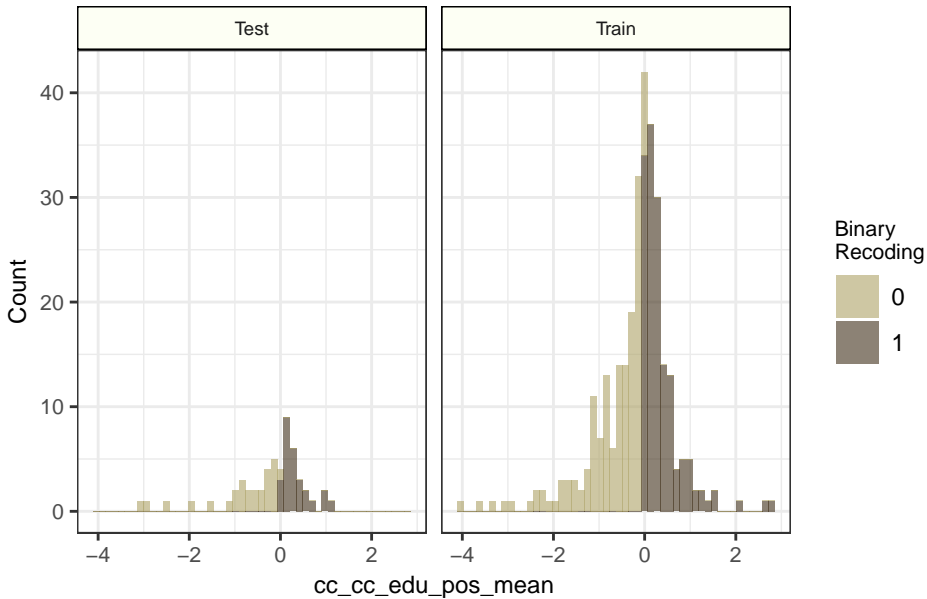


# cc\_all\_nucleus\_roundness\_mean

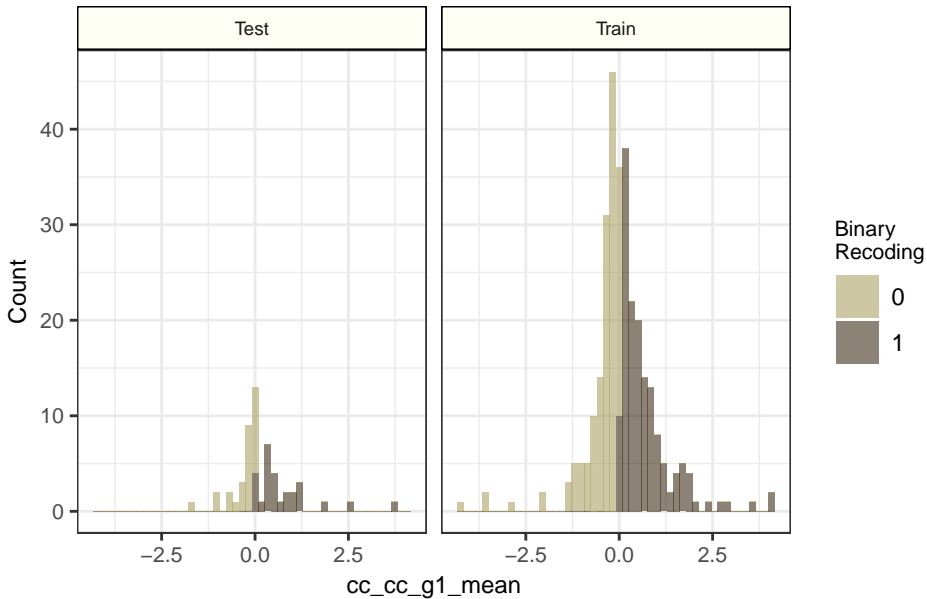




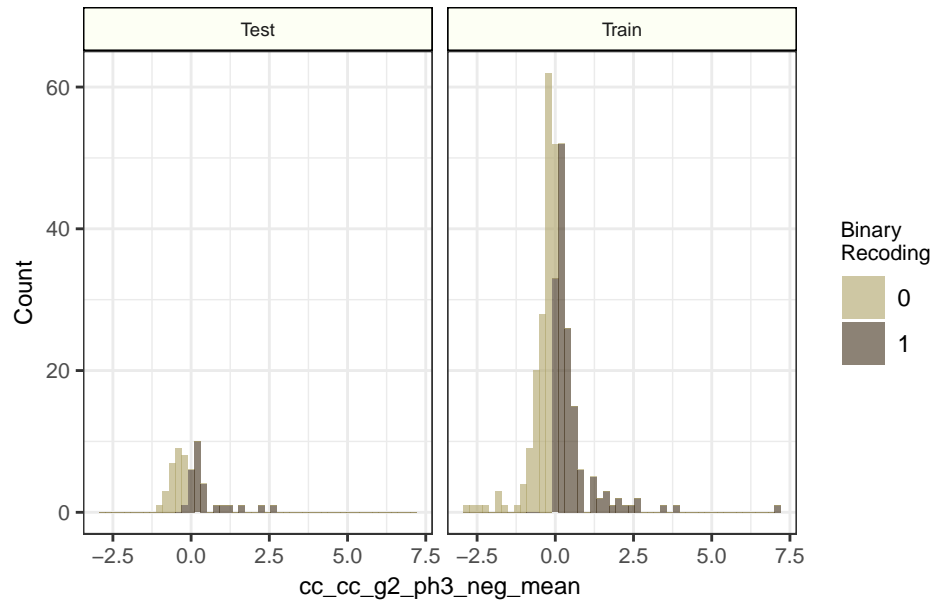
cc\_cc\_edu\_pos\_mean



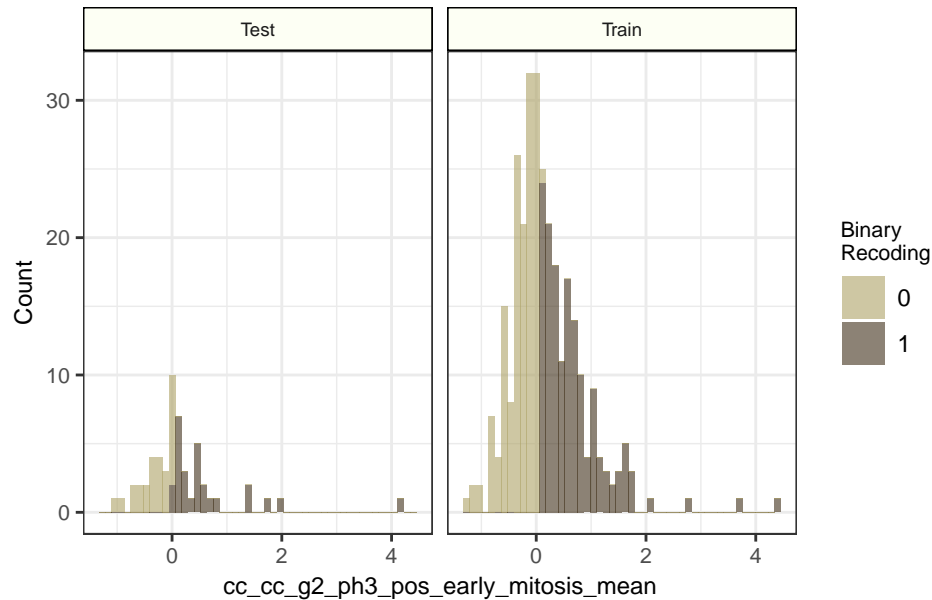
cc\_cc\_g1\_mean



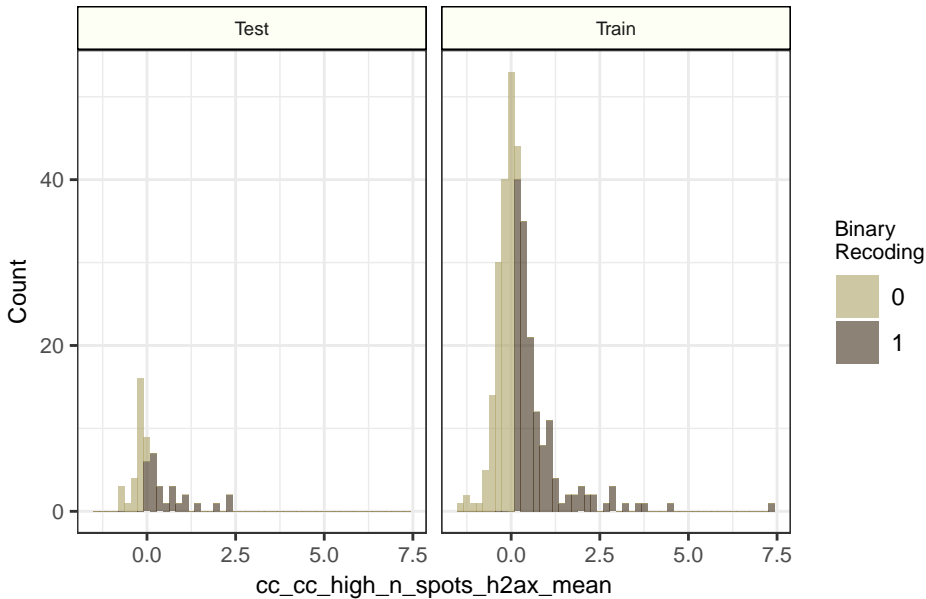
# cc\_cc\_g2\_ph3\_neg\_mean



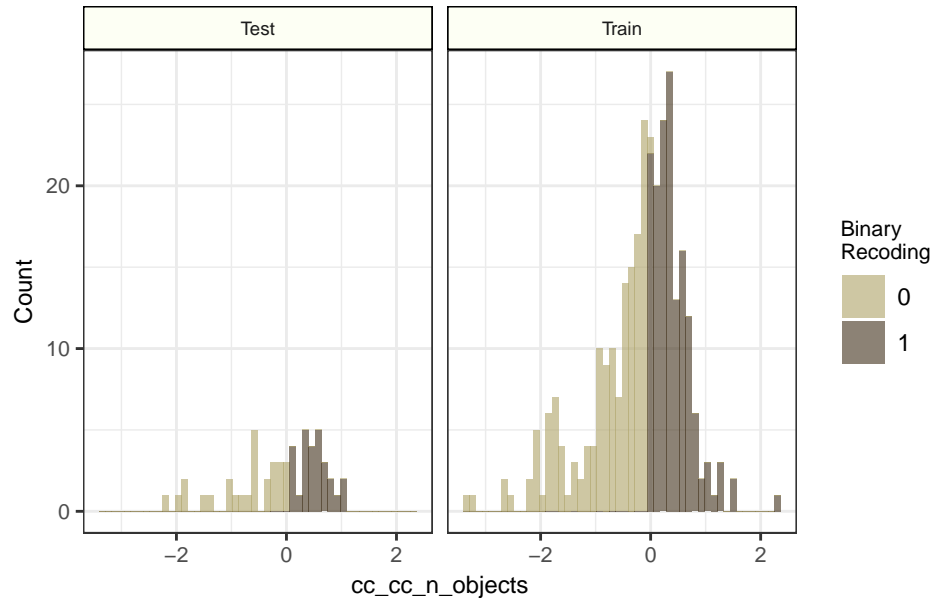
cc\_cc\_g2\_ph3\_pos\_early\_mitosis\_mean



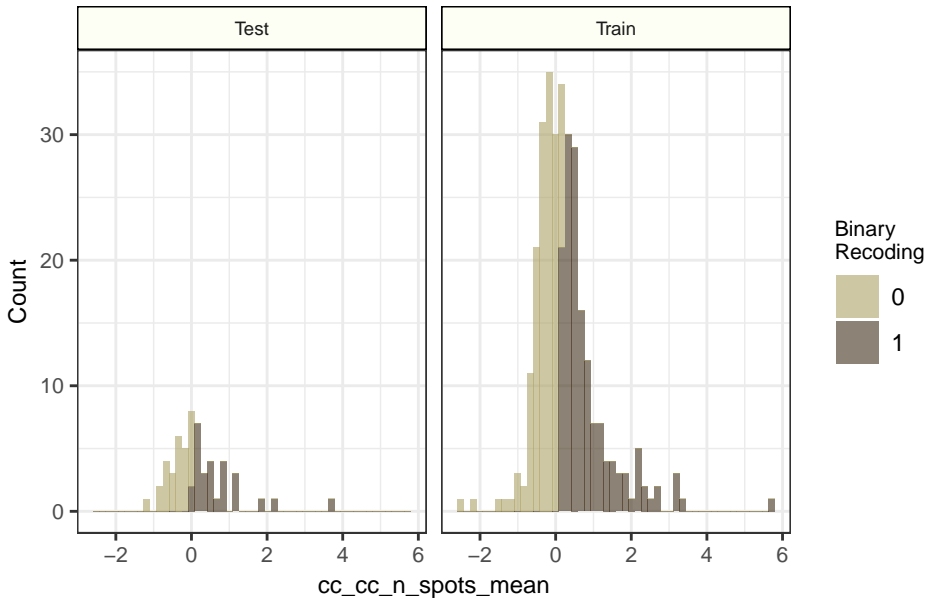
cc\_cc\_high\_n\_spots\_h2ax\_mean



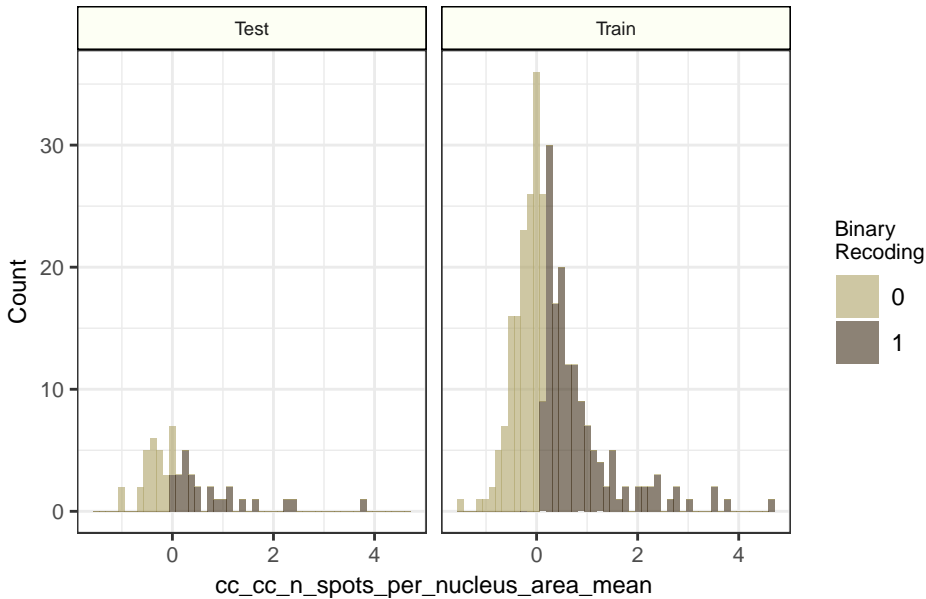
# cc\_cc\_n\_objects



cc\_cc\_n\_spots\_mean

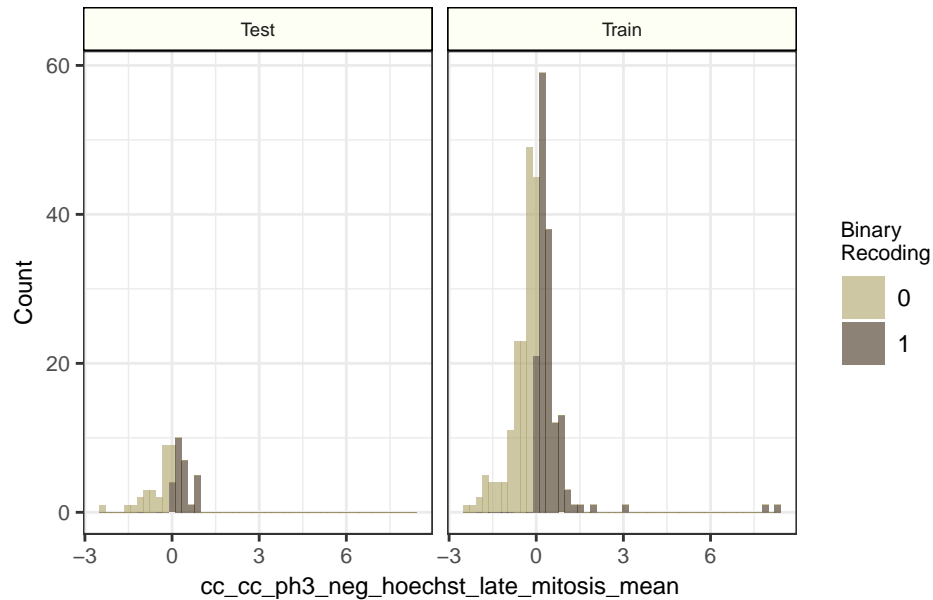


# cc\_cc\_n\_spots\_per\_nucleus\_area\_mean

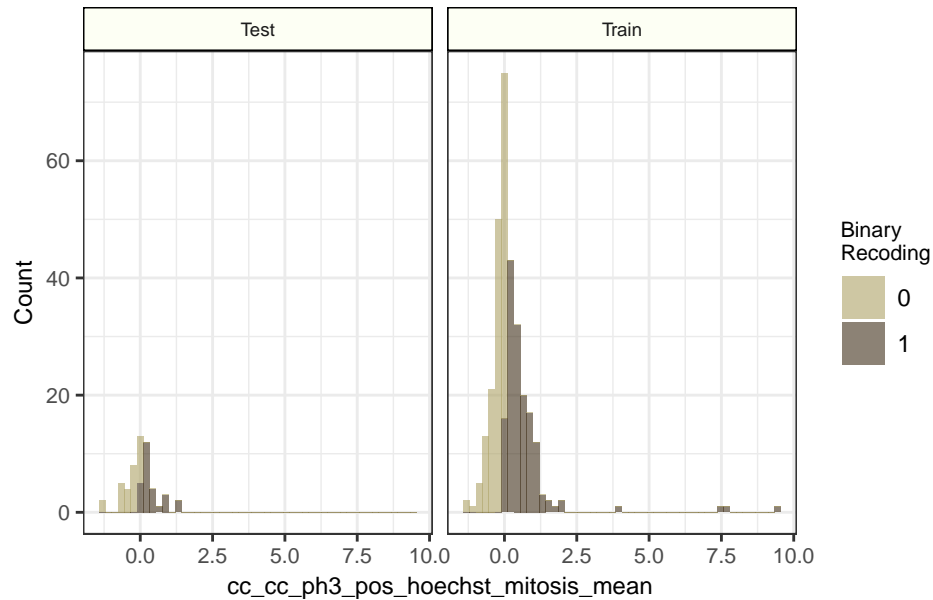




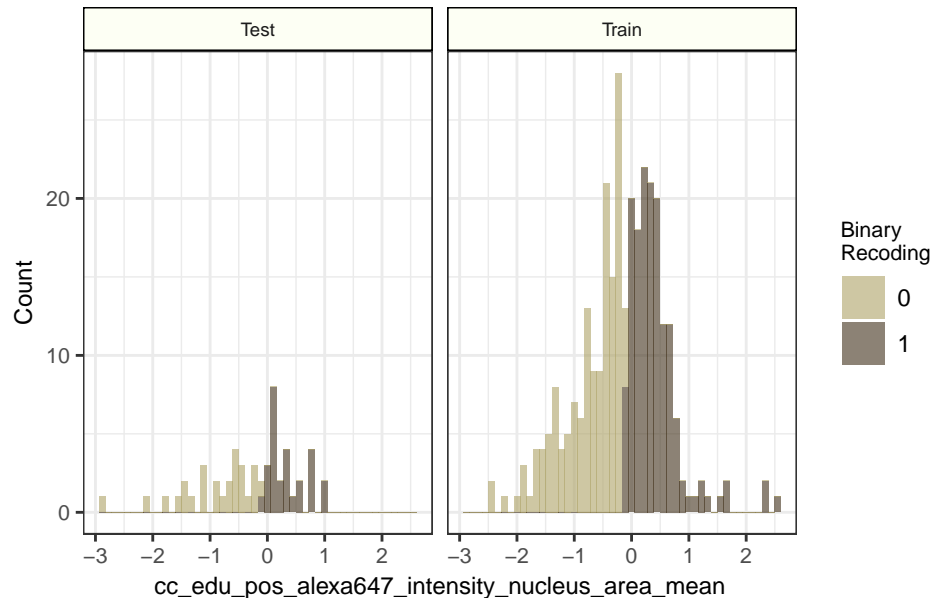
# cc\_cc\_ph3\_neg\_hoechst\_late\_mitosis\_mean



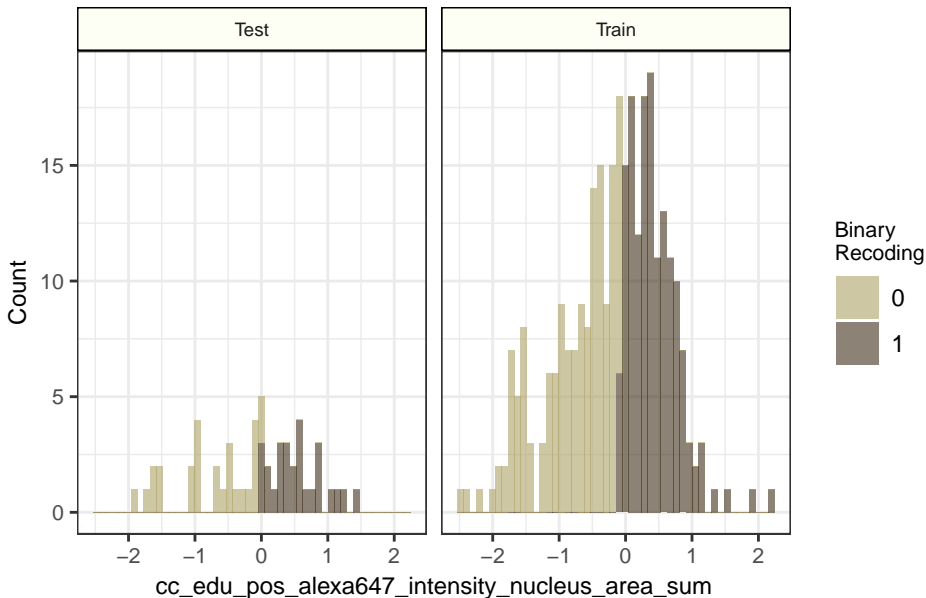
# cc\_cc\_ph3\_pos\_hoechst\_mitosis\_mean



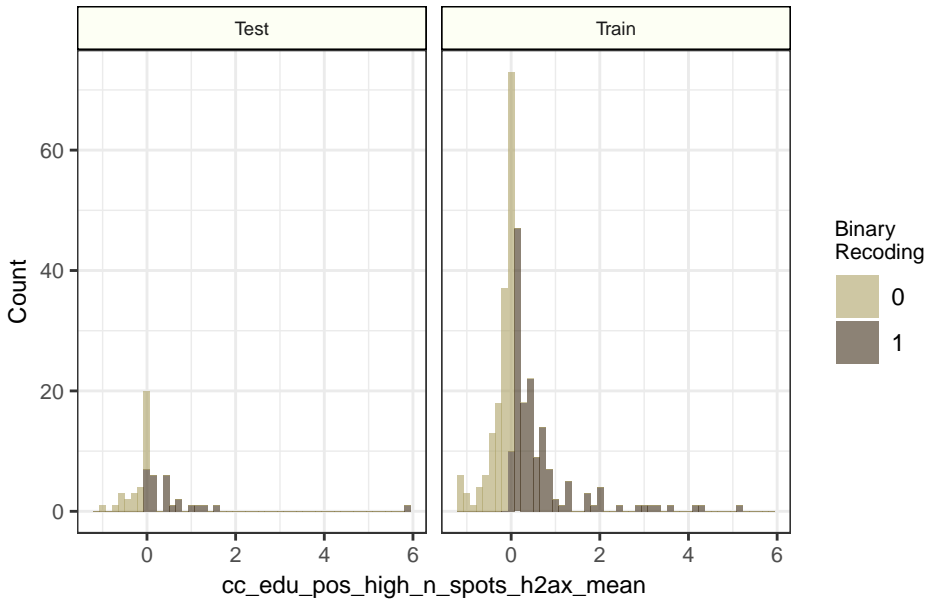
cc\_edu\_pos\_alex647\_intensity\_nucleus\_area\_mean



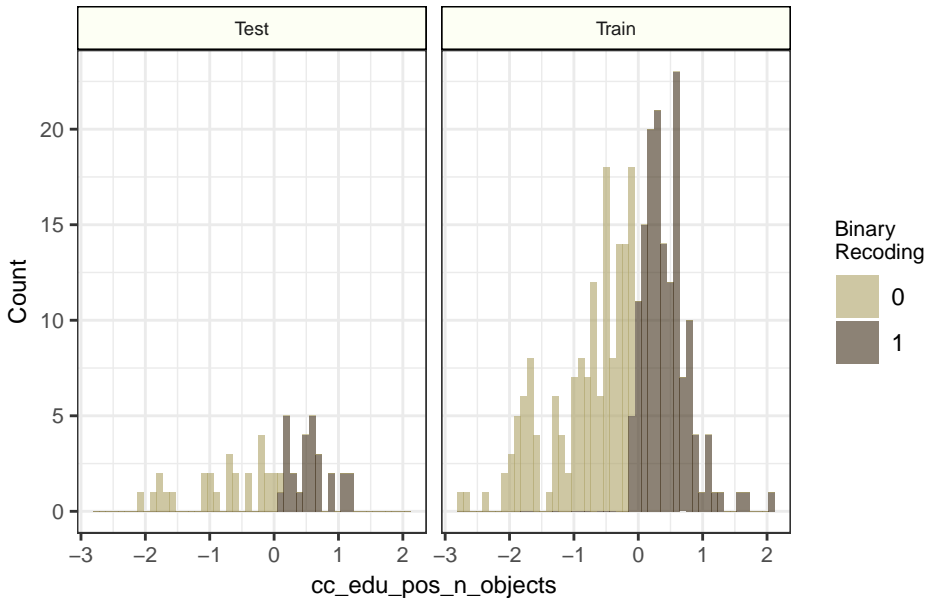
# cc\_edu\_pos\_alex647\_intensity\_nucleus\_area\_sum



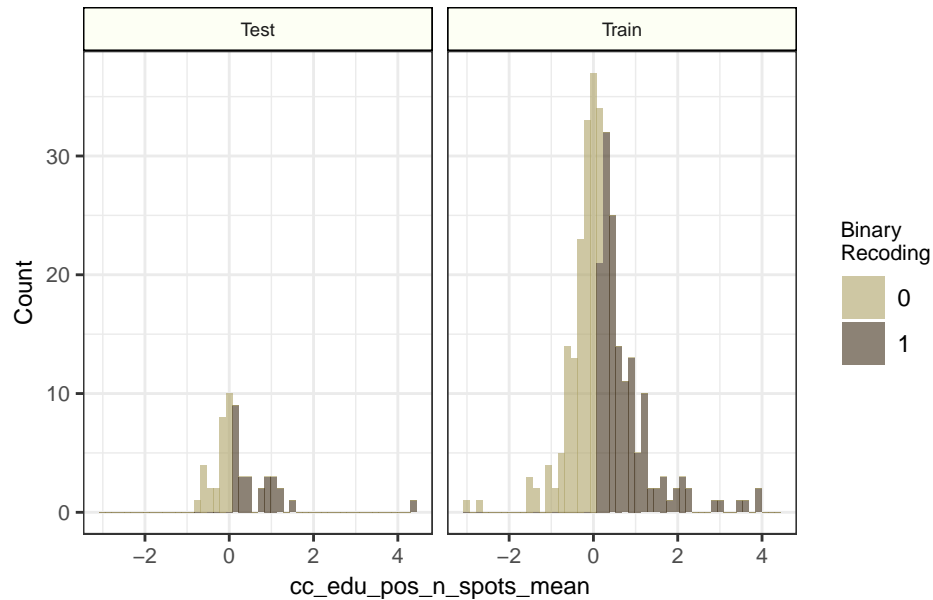
cc\_edu\_pos\_high\_n\_spots\_h2ax\_mean



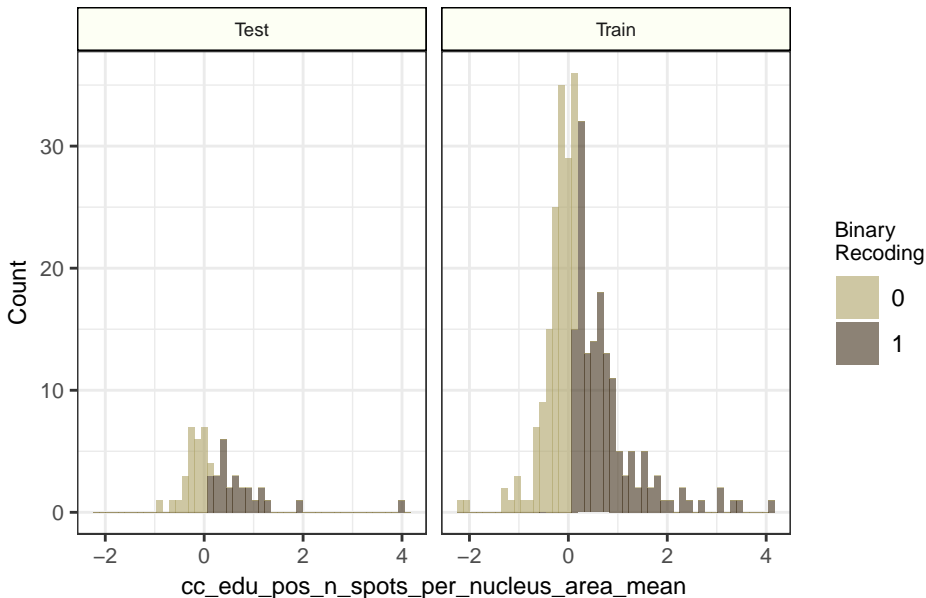
# cc\_edu\_pos\_n\_objects



cc\_edu\_pos\_n\_spots\_mean

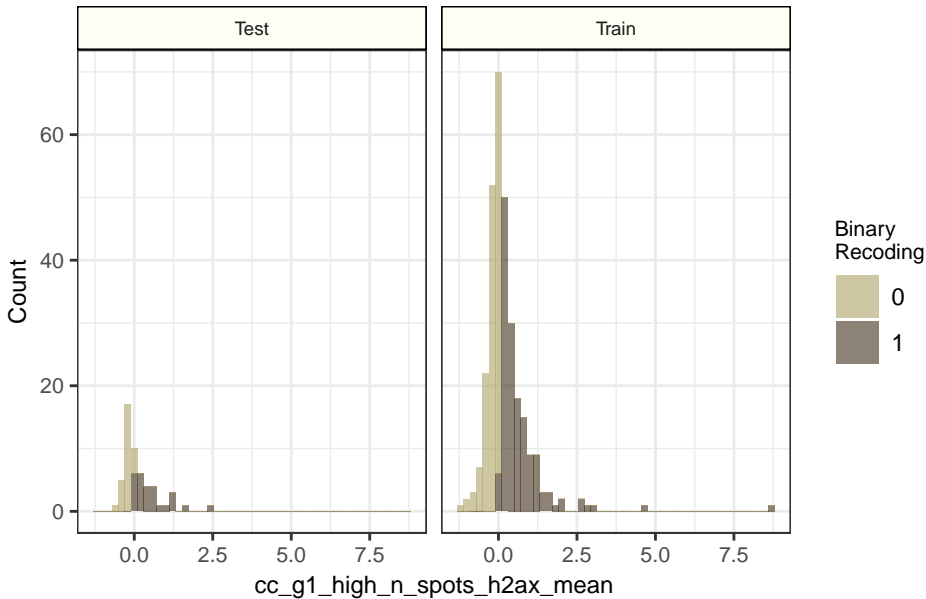


cc\_edu\_pos\_n\_spots\_per\_nucleus\_area\_mean

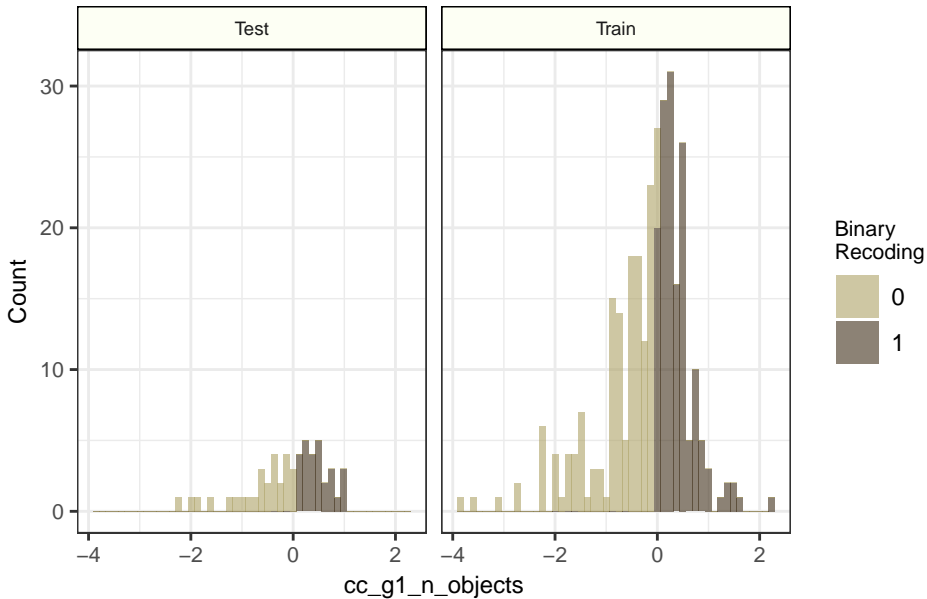




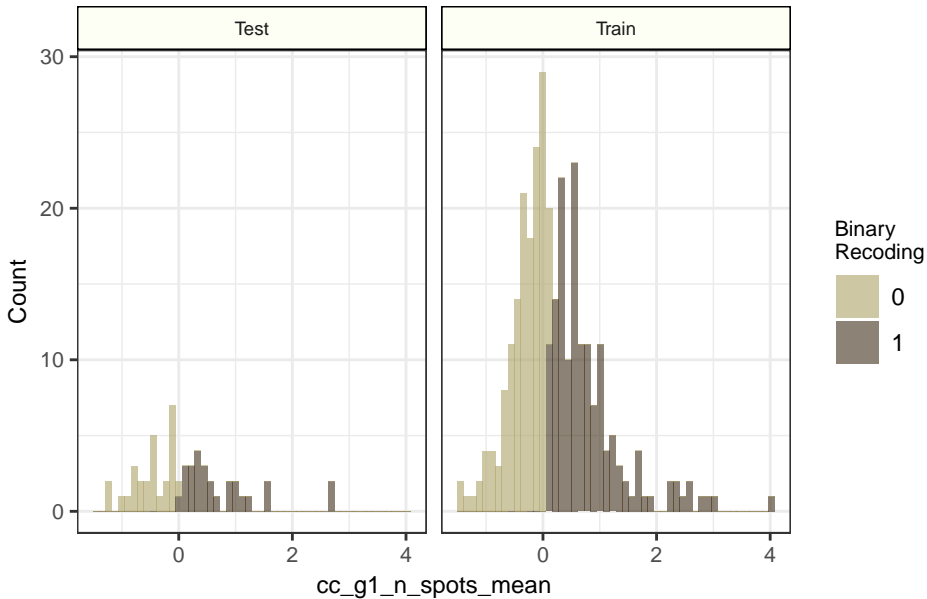
# cc\_g1\_high\_n\_spots\_h2ax\_mean



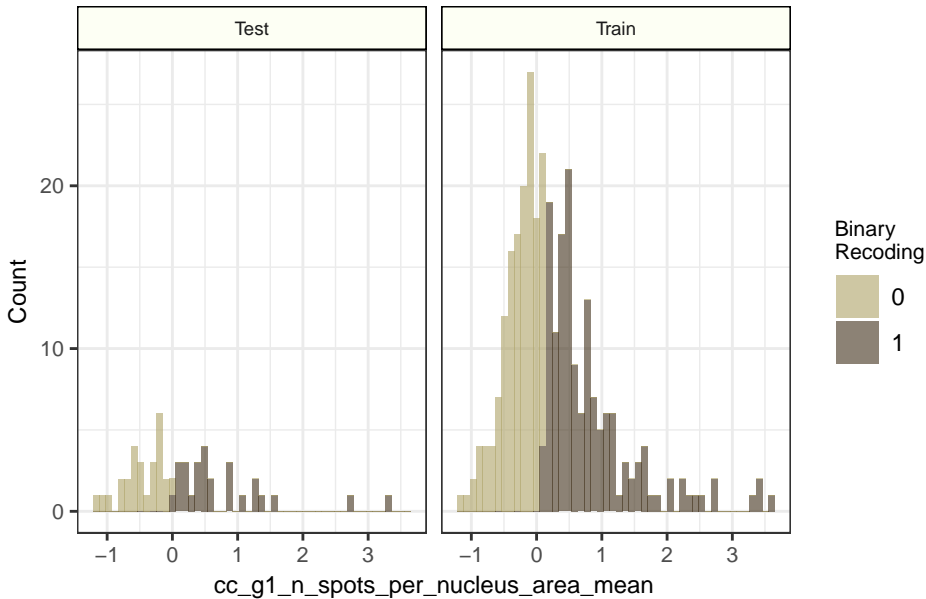
# cc\_g1\_n\_objects



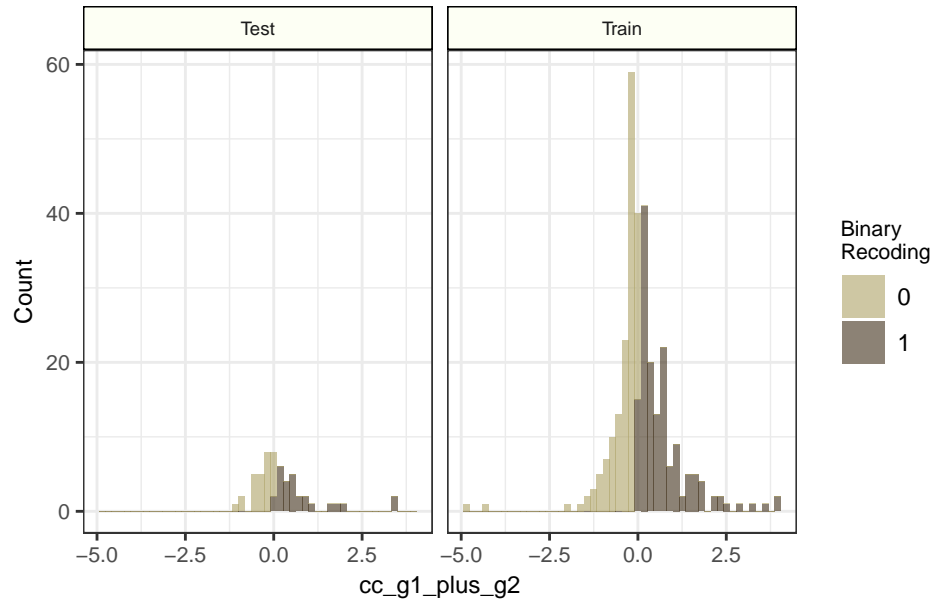
# cc\_g1\_n\_spots\_mean



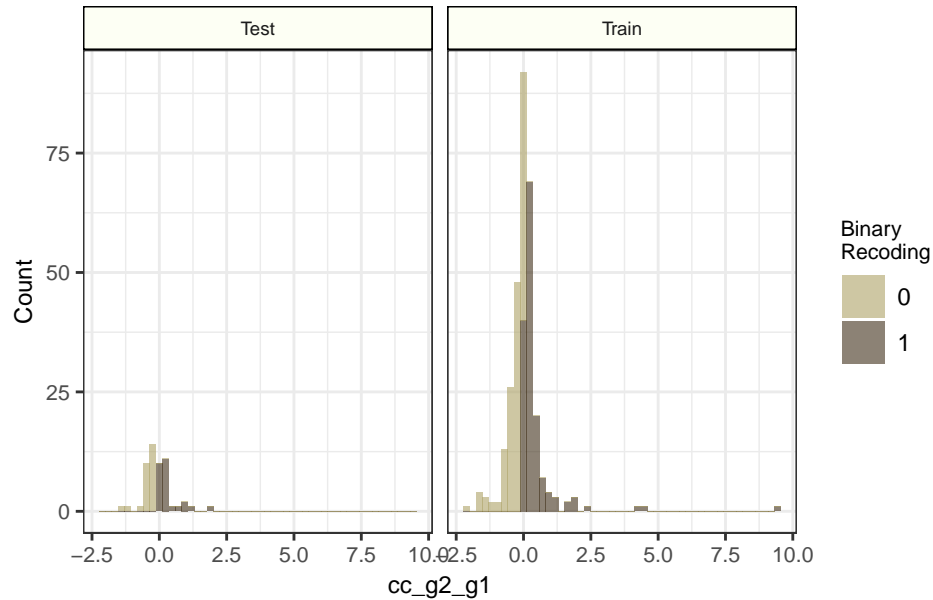
# cc\_g1\_n\_spots\_per\_nucleus\_area\_mean



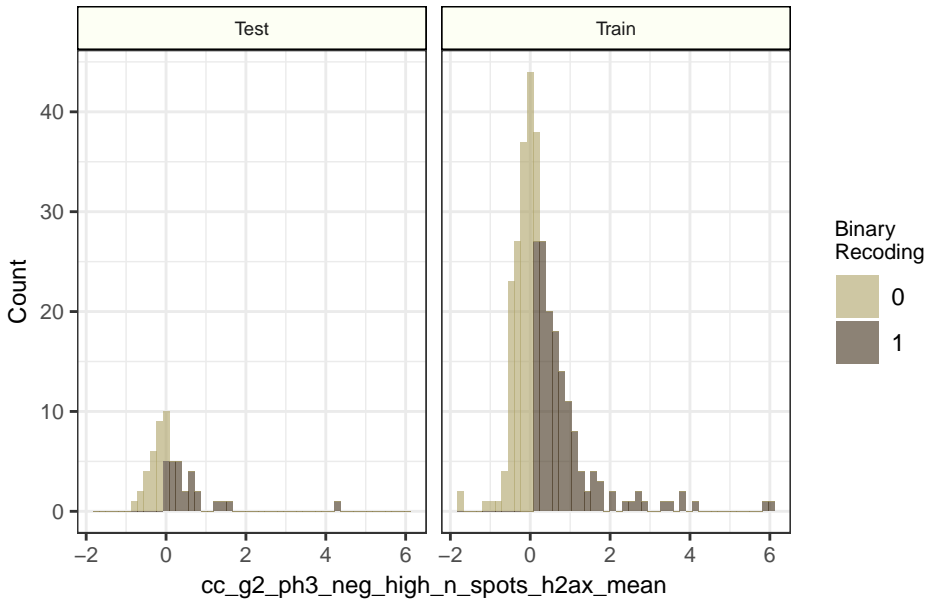
# cc\_g1\_plus\_g2



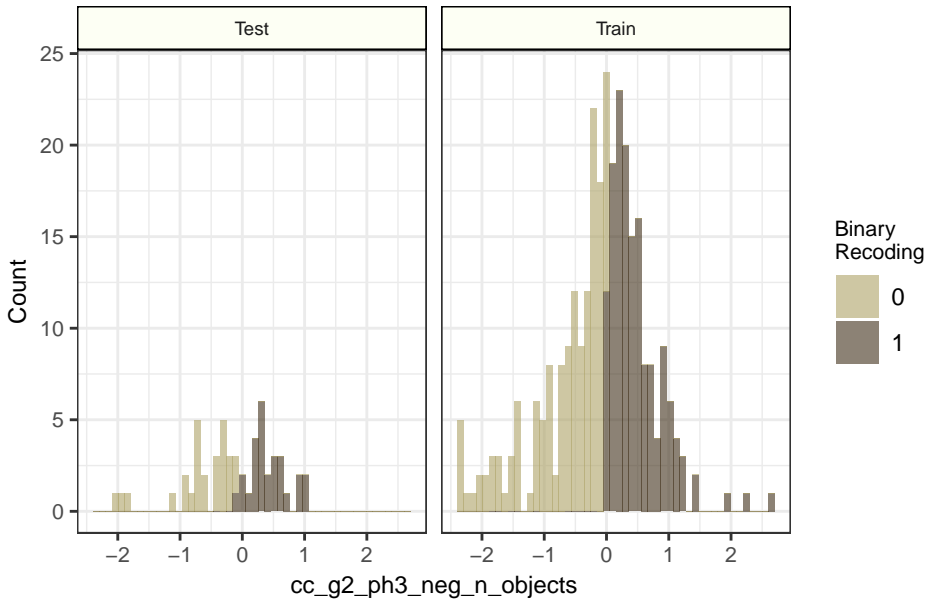
cc\_g2\_g1



cc\_g2\_ph3\_neg\_high\_n\_spots\_h2ax\_mean

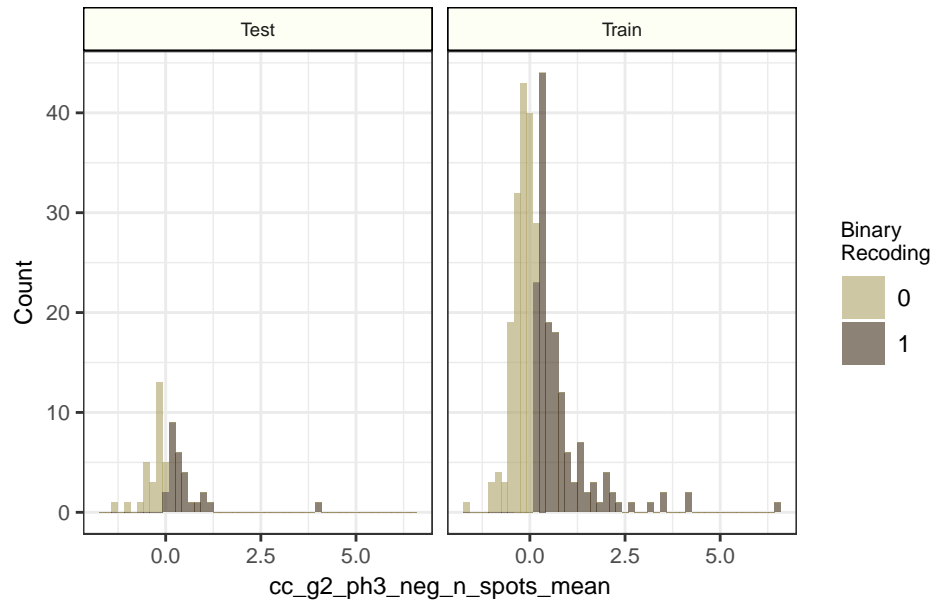


# cc\_g2\_ph3\_neg\_n\_objects

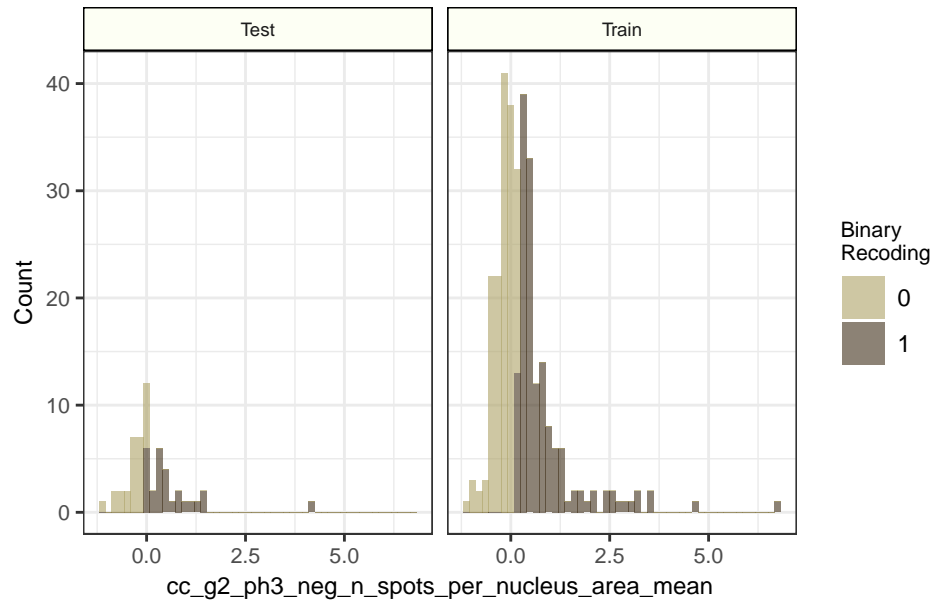




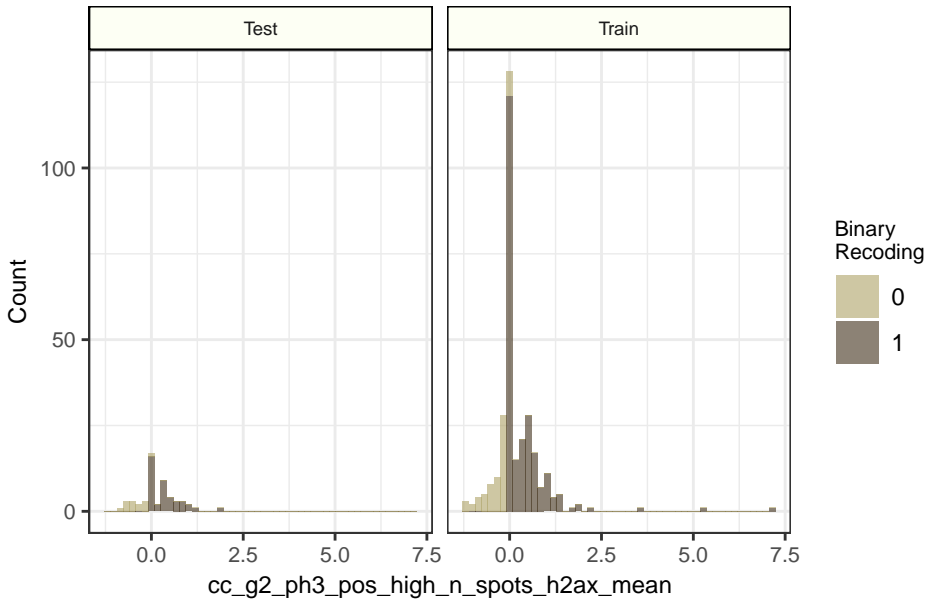
# cc\_g2\_ph3\_neg\_n\_spots\_mean



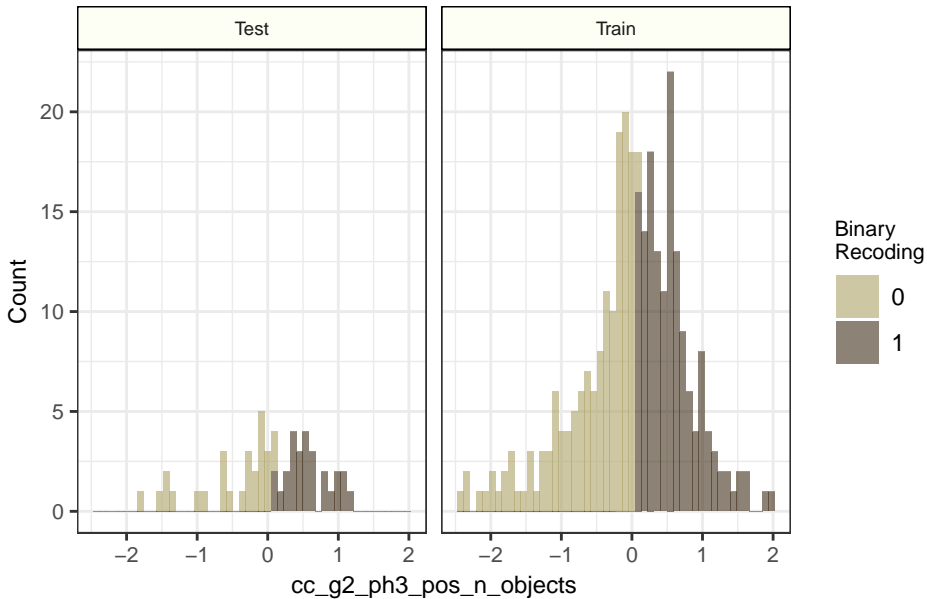
# cc\_g2\_ph3\_neg\_n\_spots\_per\_nucleus\_area\_mean



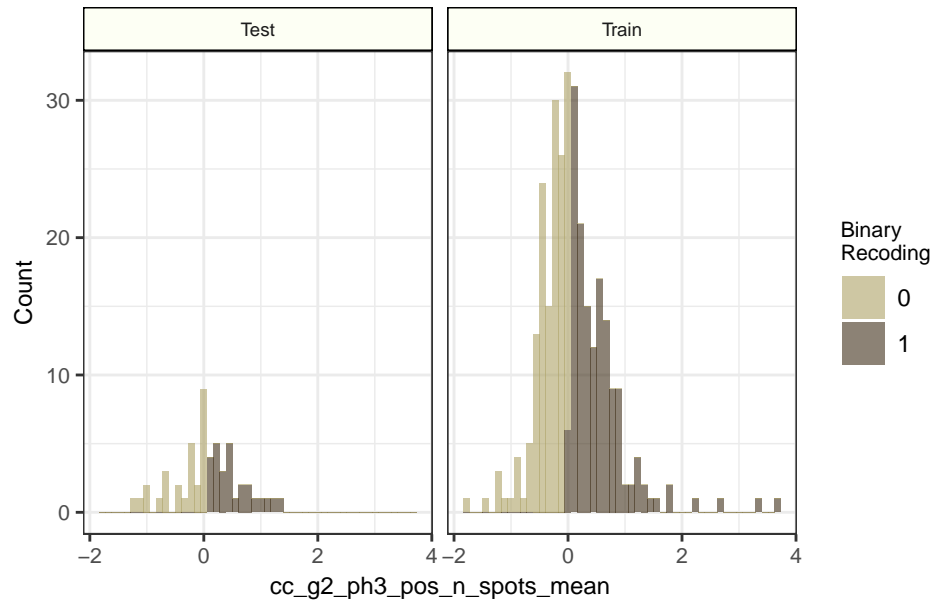
# cc\_g2\_ph3\_pos\_high\_n\_spots\_h2ax\_mean



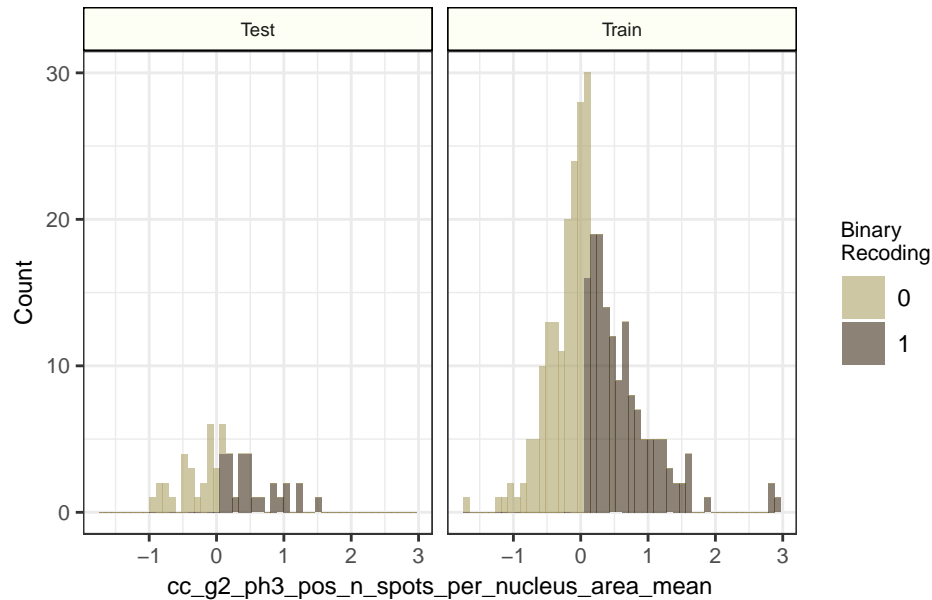
# cc\_g2\_ph3\_pos\_n\_objects



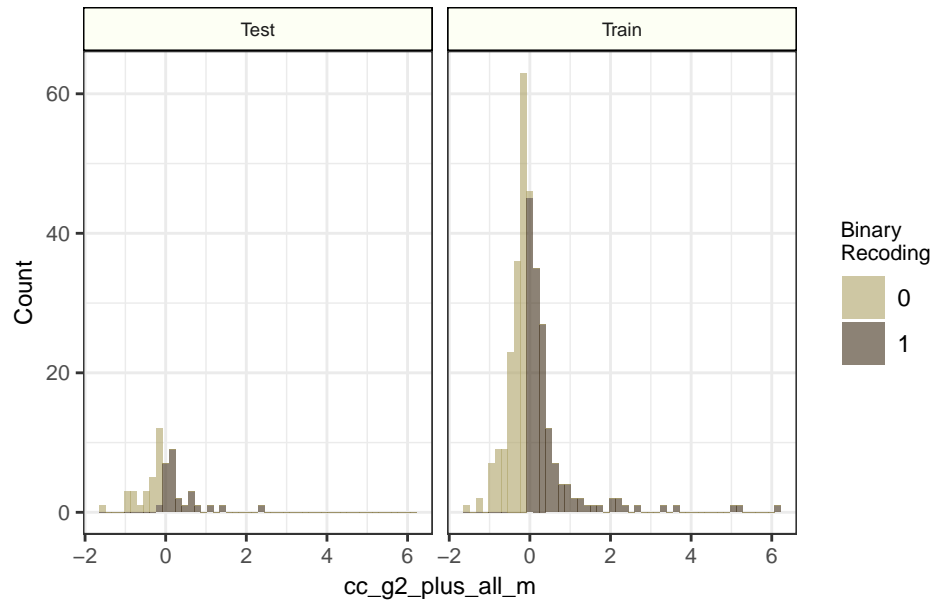
# cc\_g2\_ph3\_pos\_n\_spots\_mean



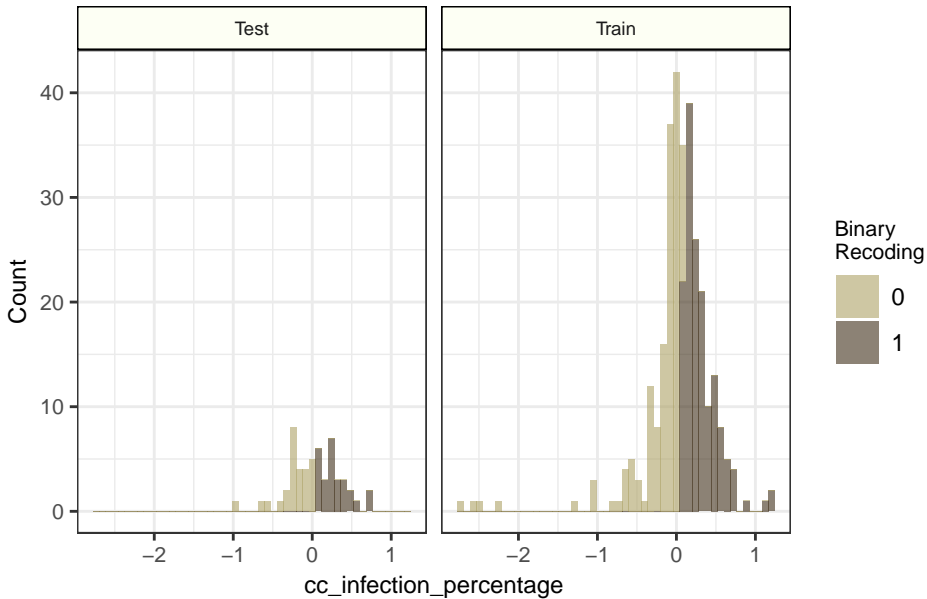
# cc\_g2\_ph3\_pos\_n\_spots\_per\_nucleus\_area\_mean



# cc\_g2\_plus\_all\_m

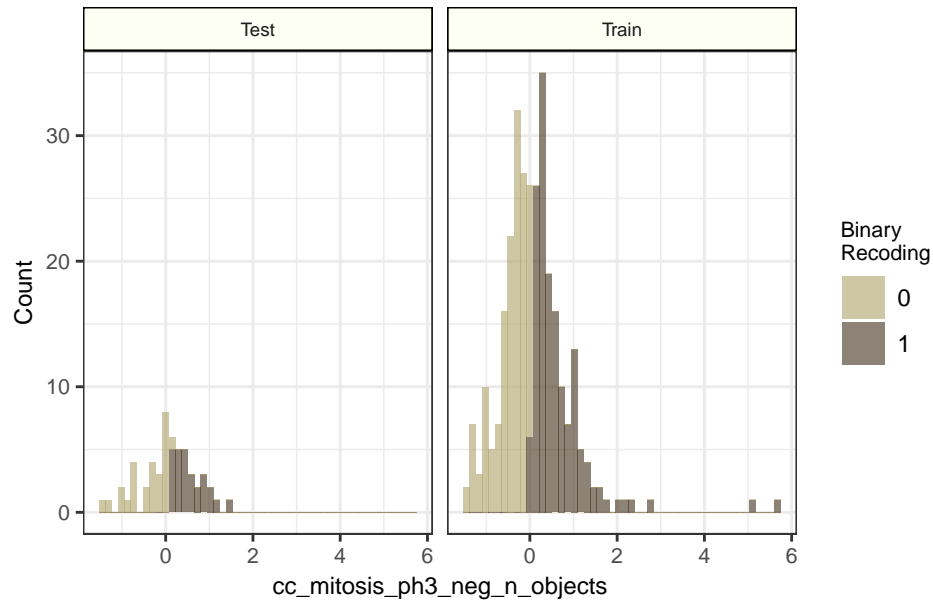


# cc\_infection\_percentage

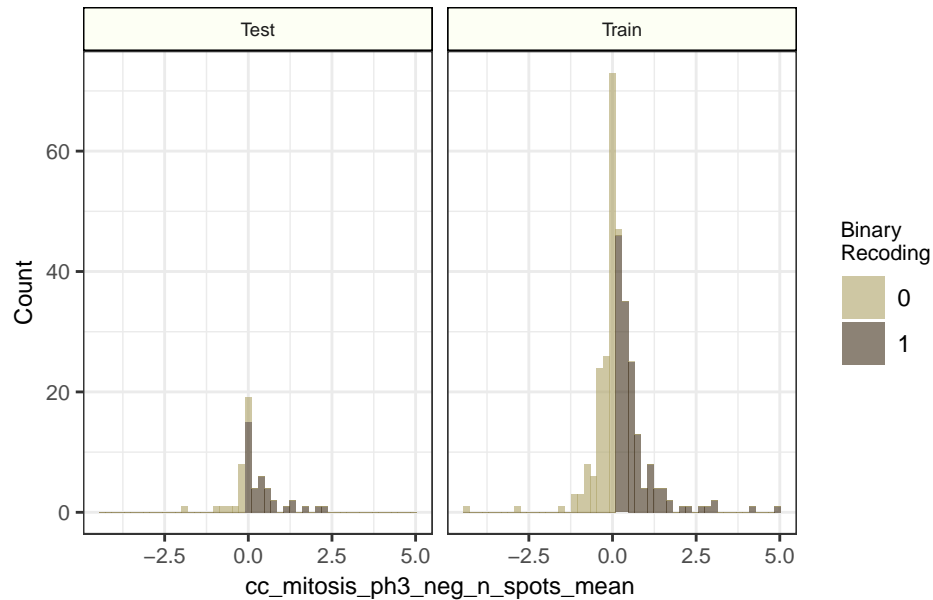




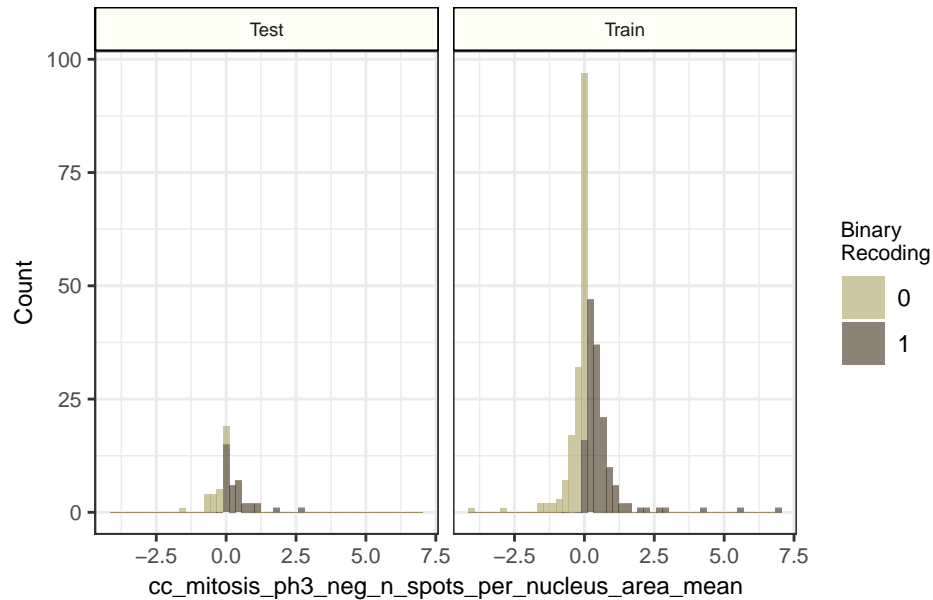
# cc\_mitosis\_ph3\_neg\_n\_objects



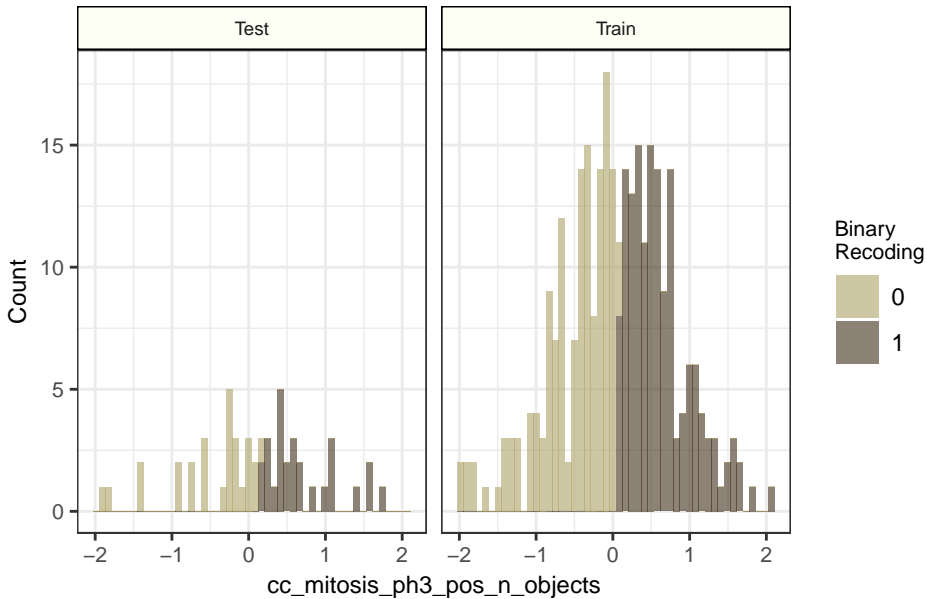
# cc\_mitosis\_ph3\_neg\_n\_spots\_mean



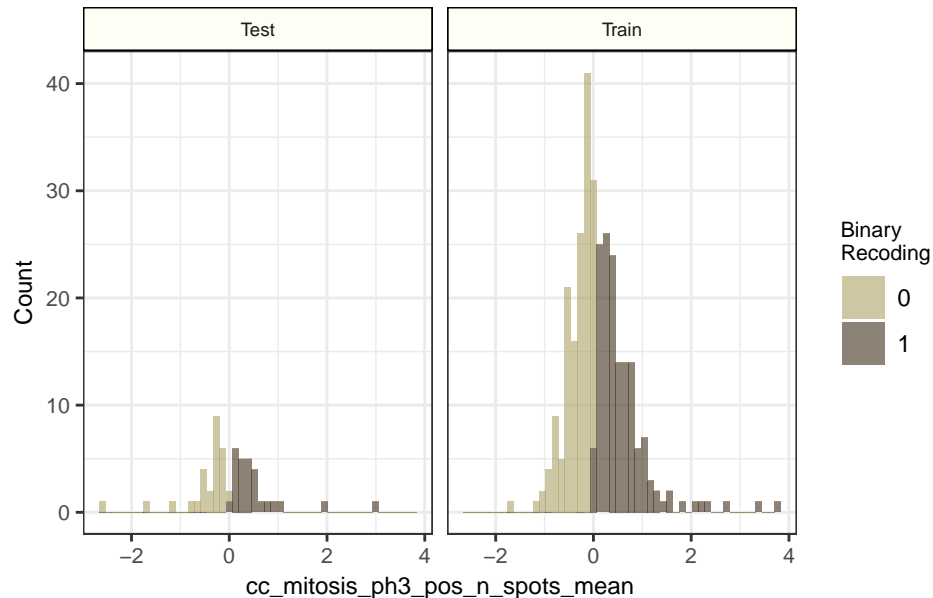
cc\_mitosis\_ph3\_neg\_n\_spots\_per\_nucleus\_area\_m



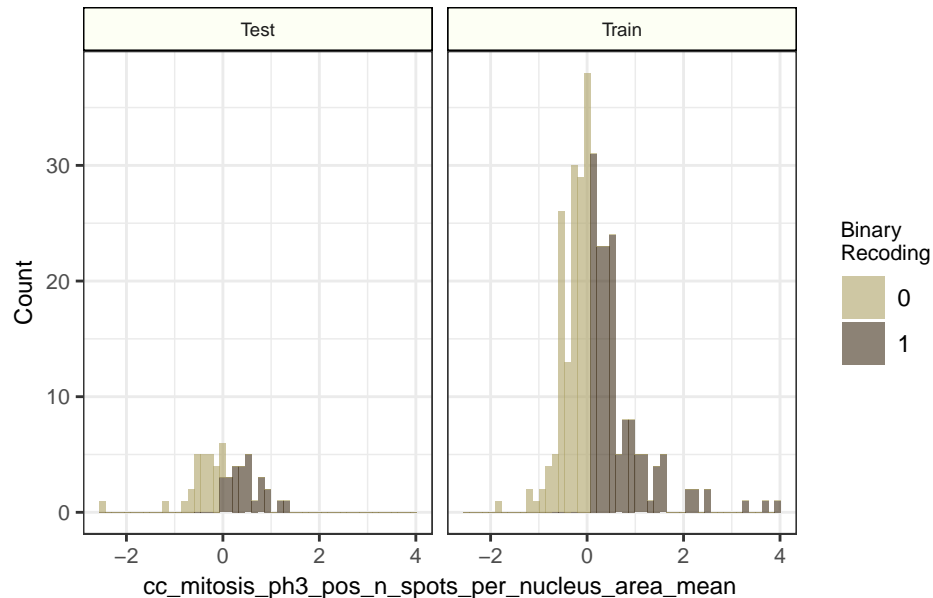
# cc\_mitosis\_ph3\_pos\_n\_objects



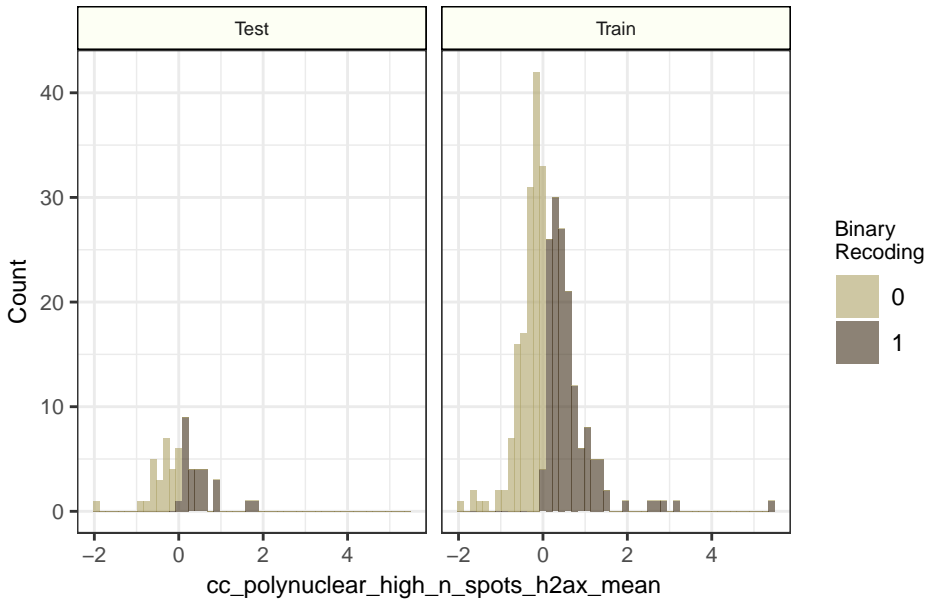
# cc\_mitosis\_ph3\_pos\_n\_spots\_mean



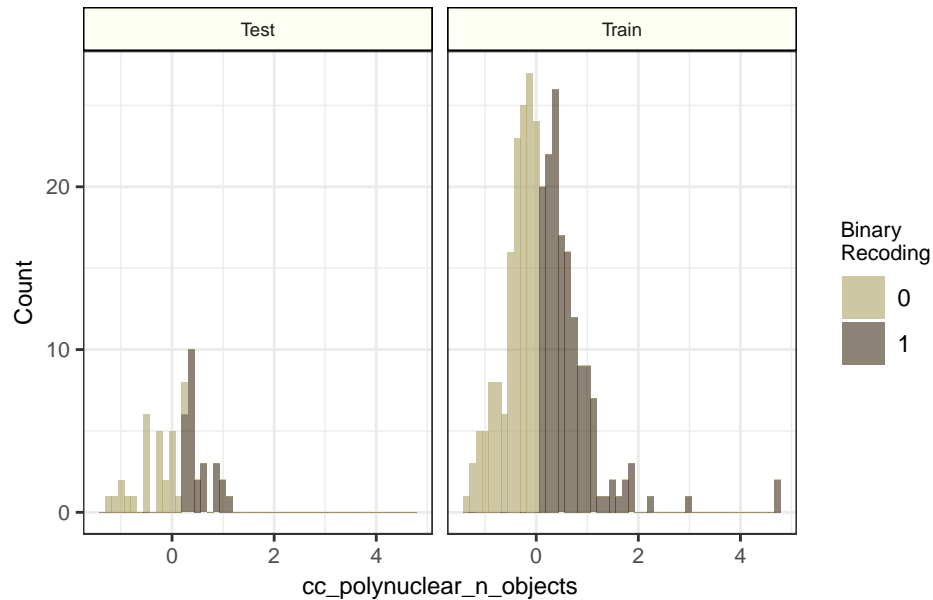
cc\_mitosis\_ph3\_pos\_n\_spots\_per\_nucleus\_area\_me



# cc\_polynuclear\_high\_n\_spots\_h2ax\_mean

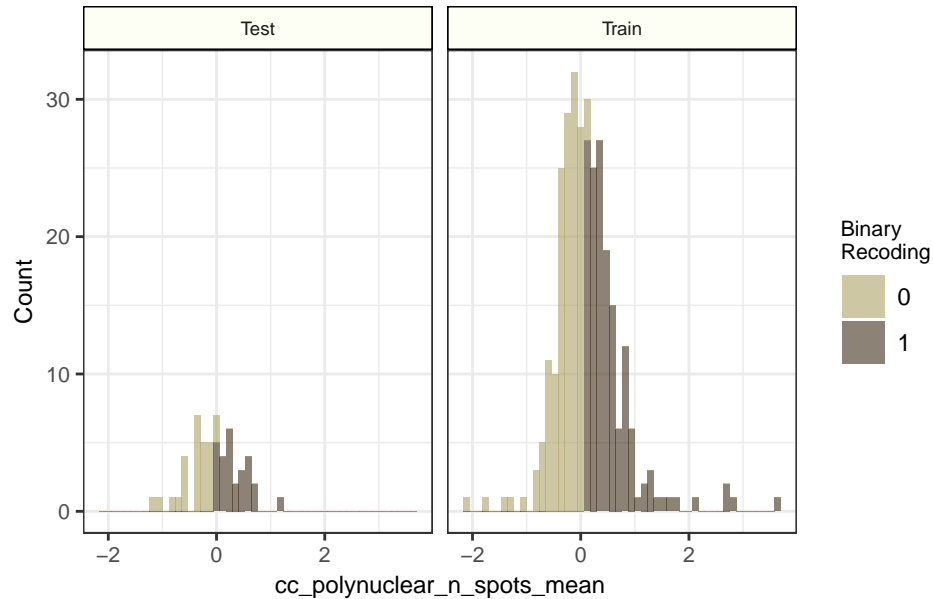


# cc\_polynuclear\_n\_objects

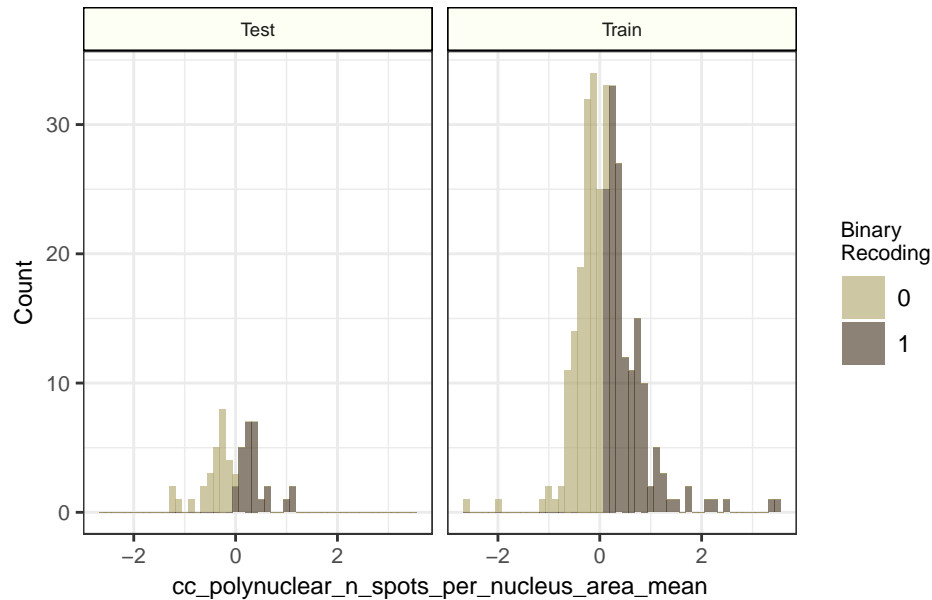




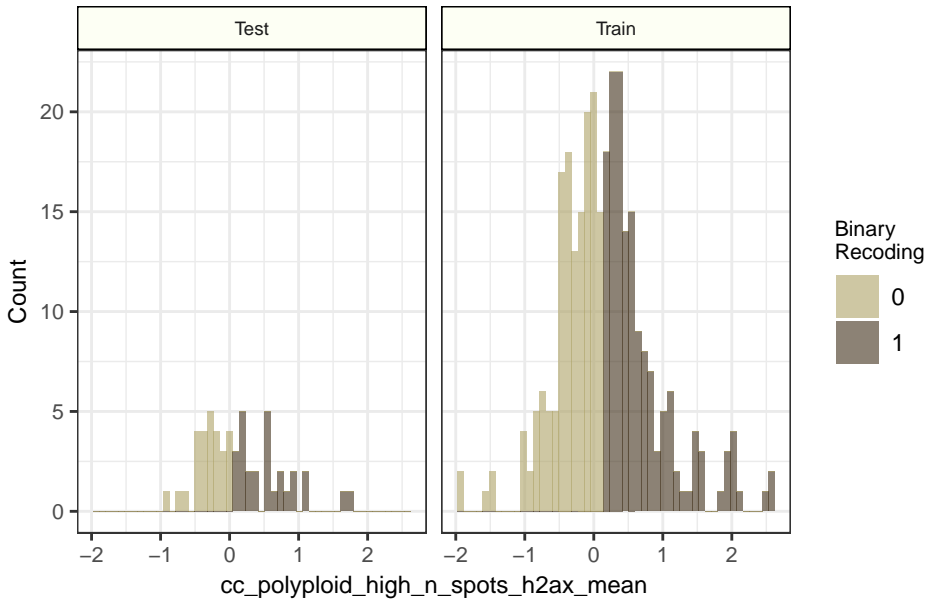
# cc\_polynuclear\_n\_spots\_mean



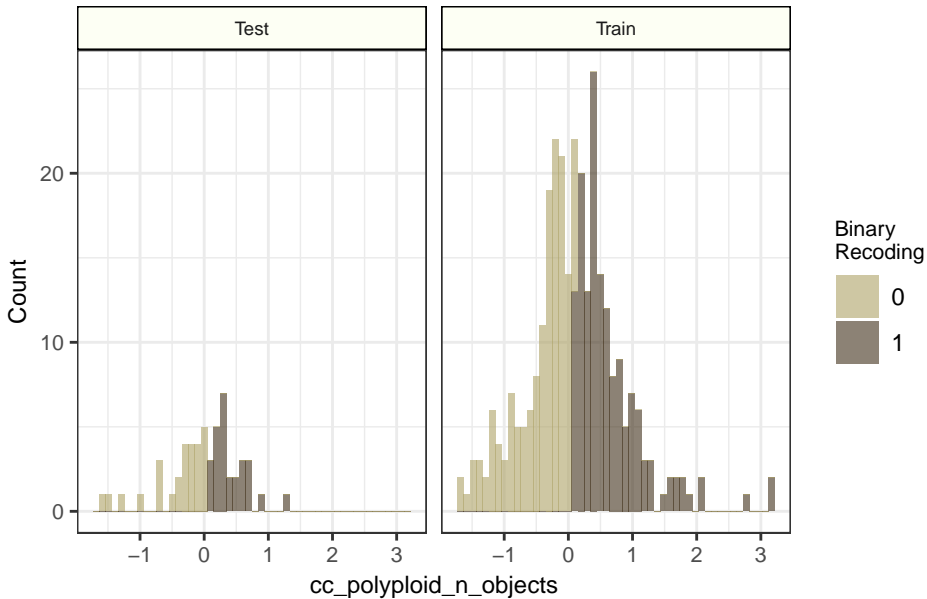
# cc\_polynuclear\_n\_spots\_per\_nucleus\_area\_mean



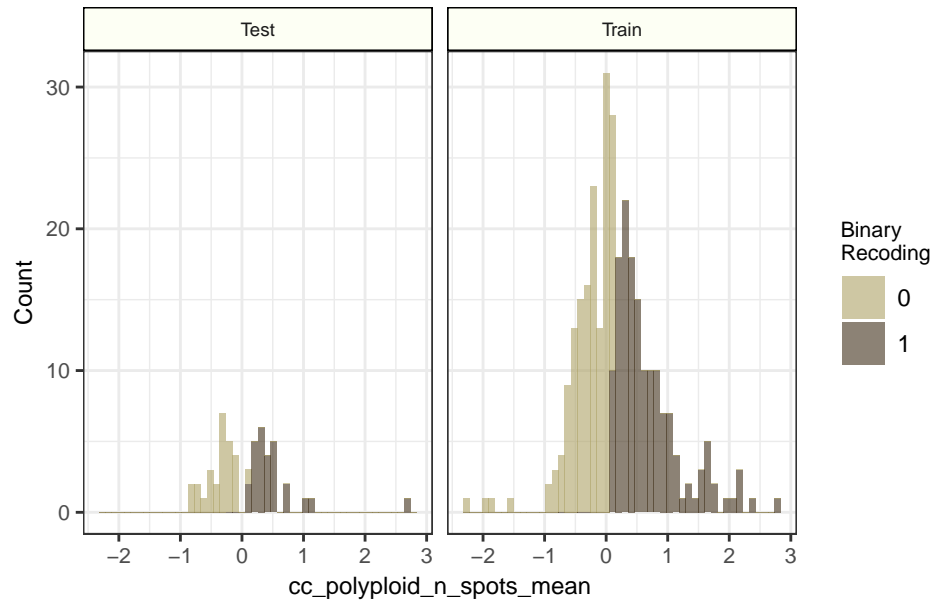
# cc\_polyploid\_high\_n\_spots\_h2ax\_mean



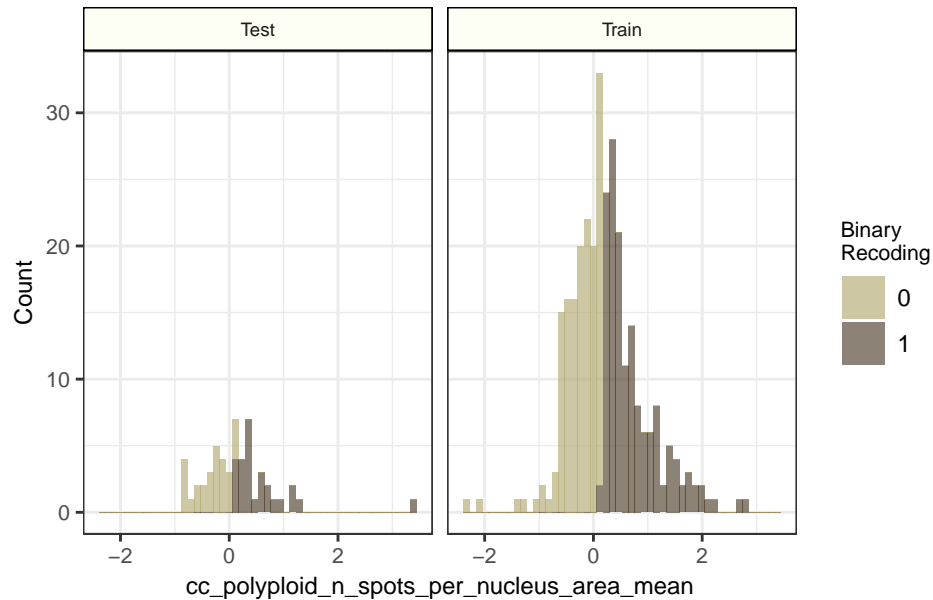
# cc\_polyploid\_n\_objects



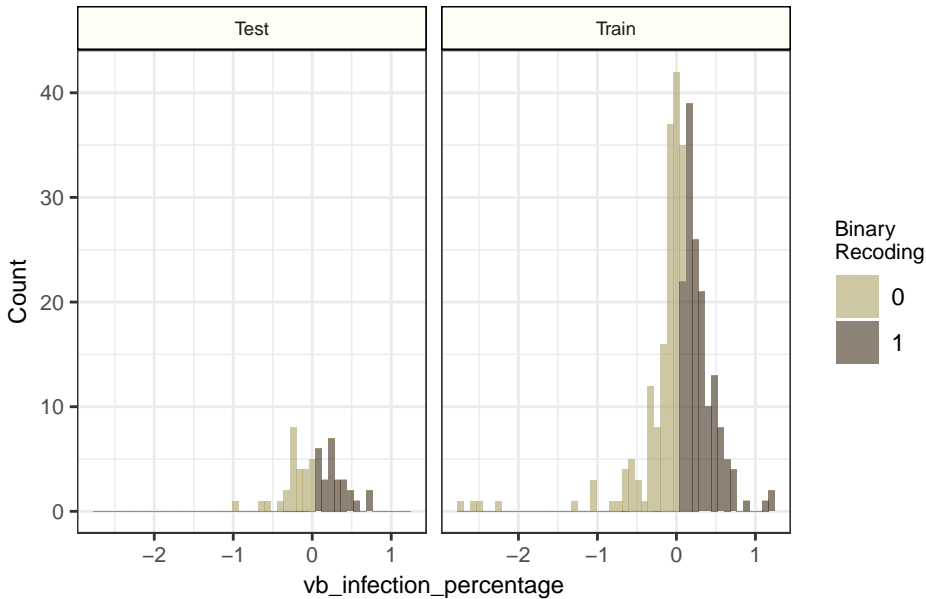
# cc\_polyploid\_n\_spots\_mean



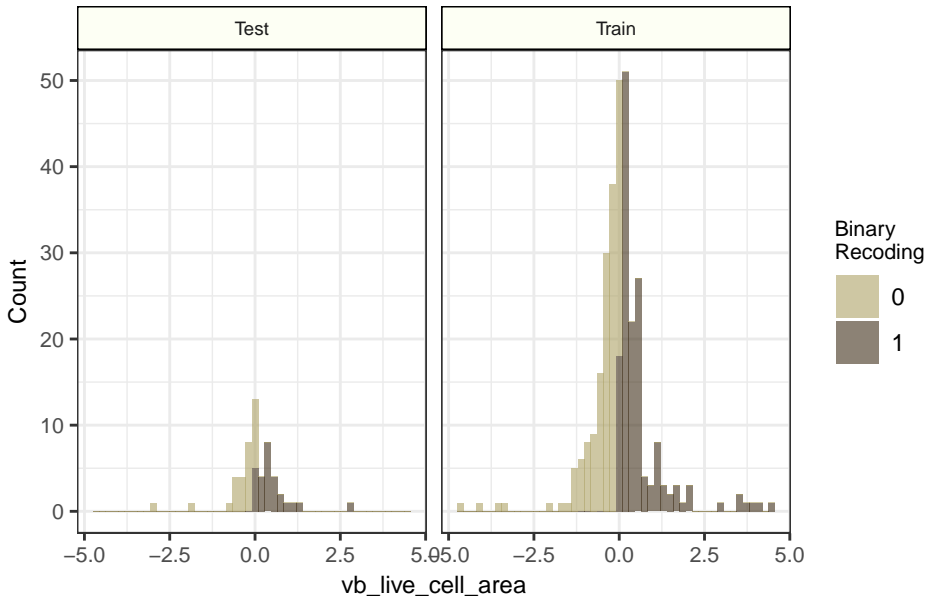
# cc\_polyploid\_n\_spots\_per\_nucleus\_area\_mean



# vb\_infection\_percentage

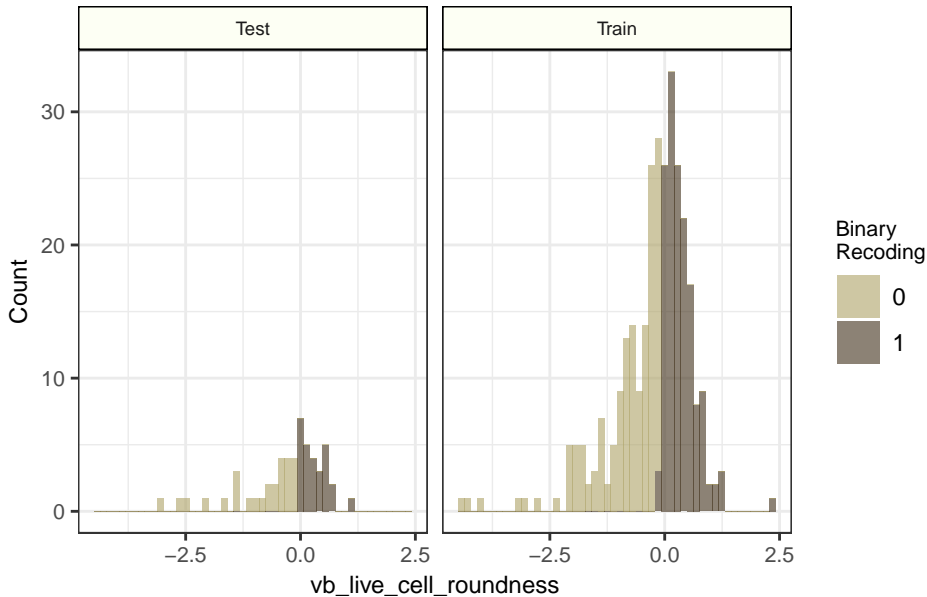


# vb\_live\_cell\_area

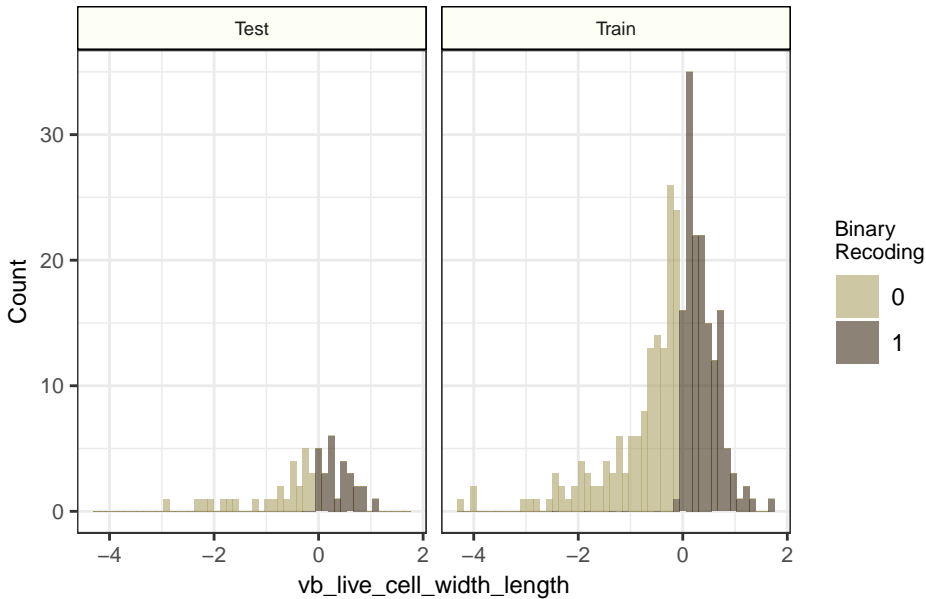




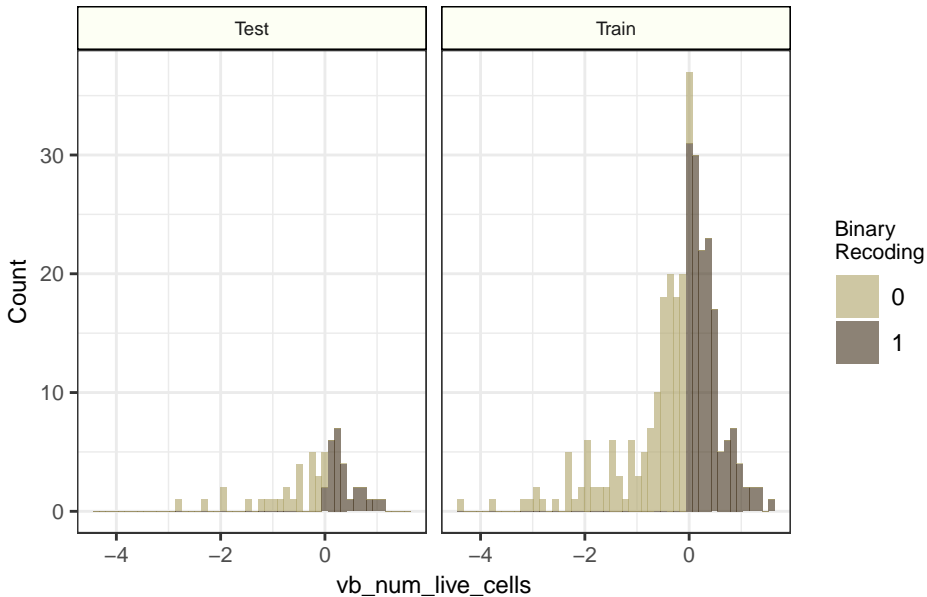
# vb\_live\_cell\_roundness



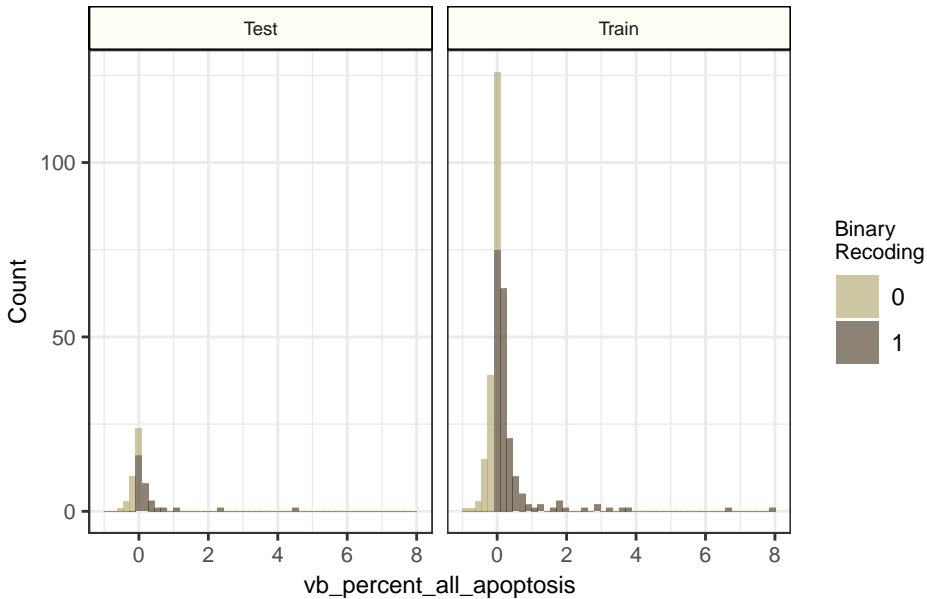
# vb\_live\_cell\_width\_length



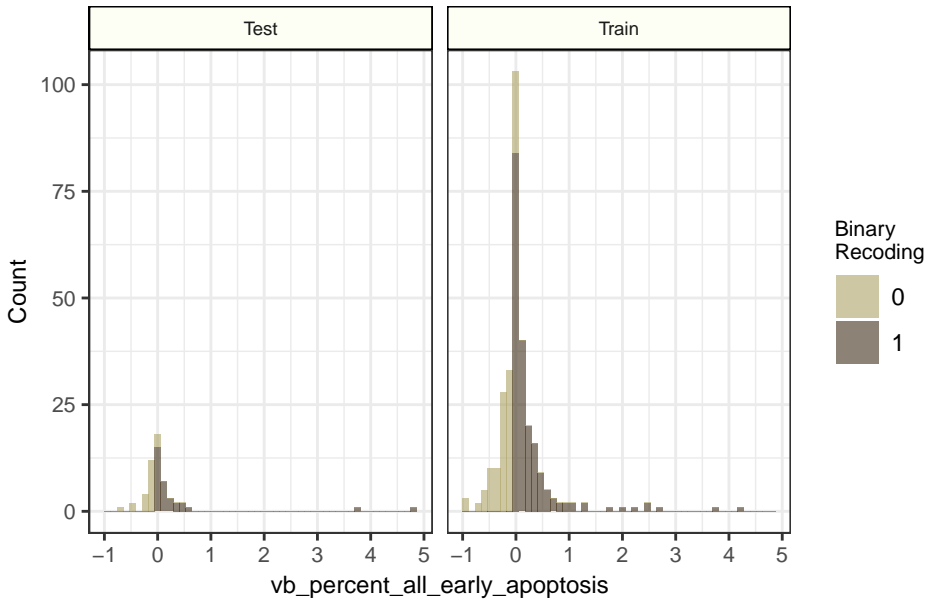
# vb\_num\_live\_cells



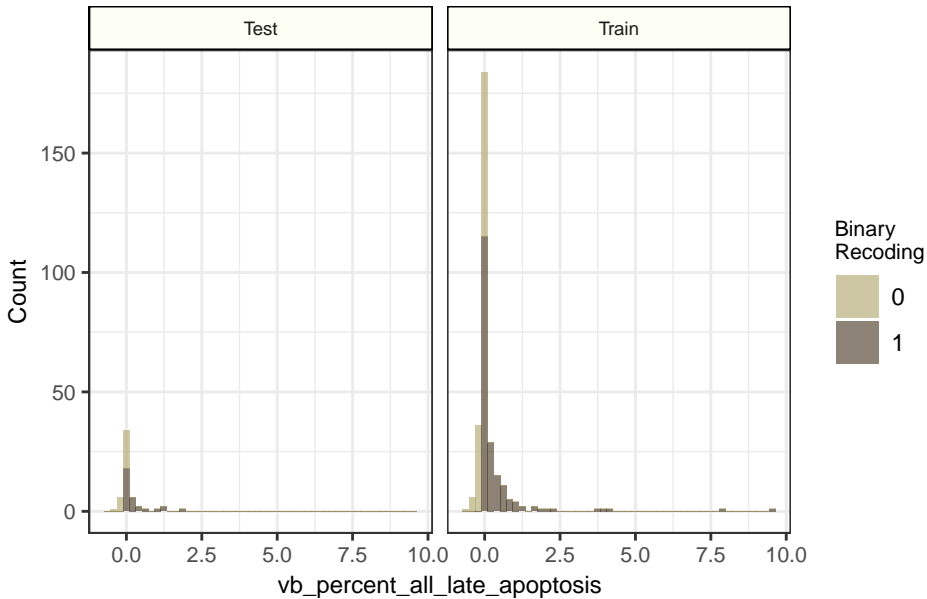
# vb\_percent\_all\_apoptosis



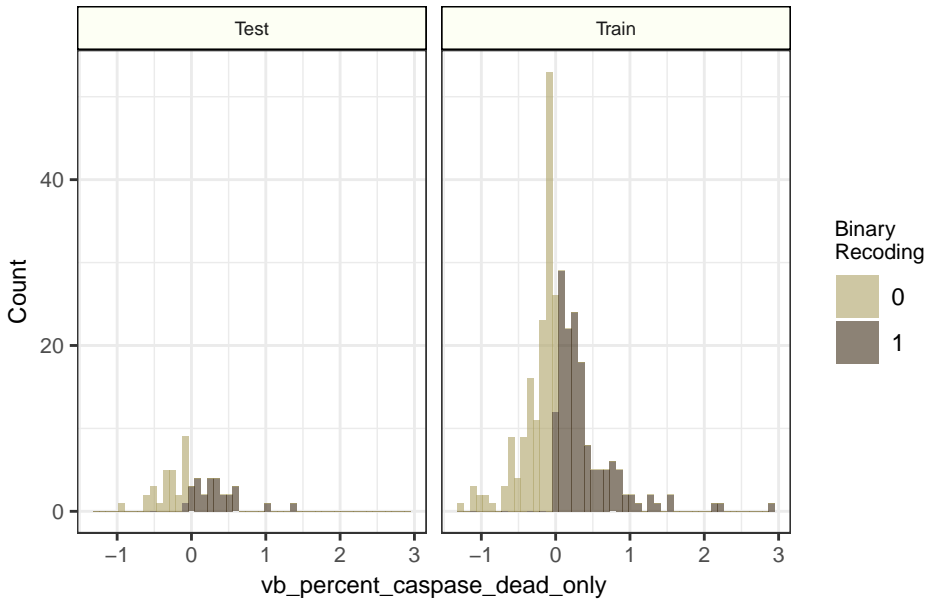
# vb\_percent\_all\_early\_apoptosis



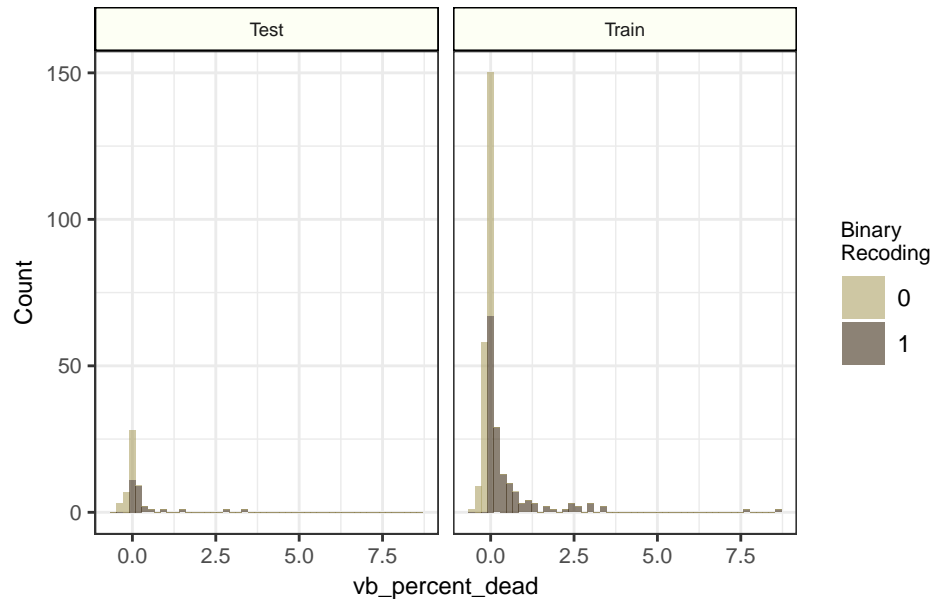
# vb\_percent\_all\_late\_apoptosis



# vb\_percent\_caspase\_dead\_only

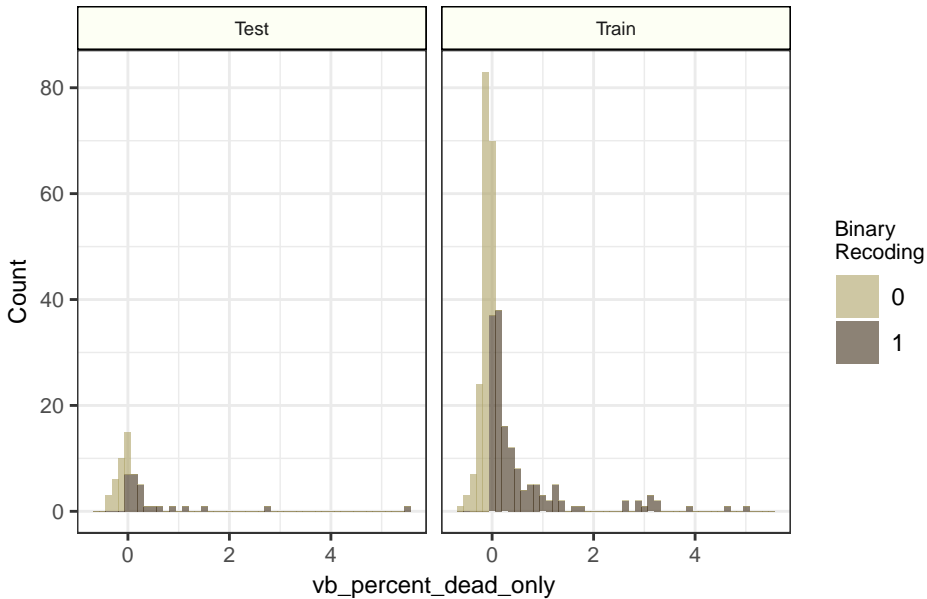


# vb\_percent\_dead

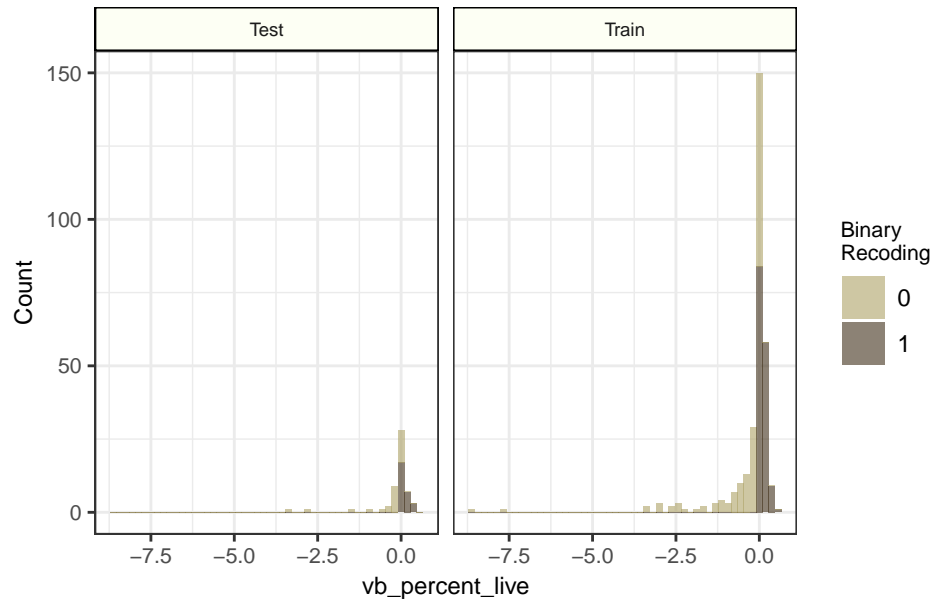




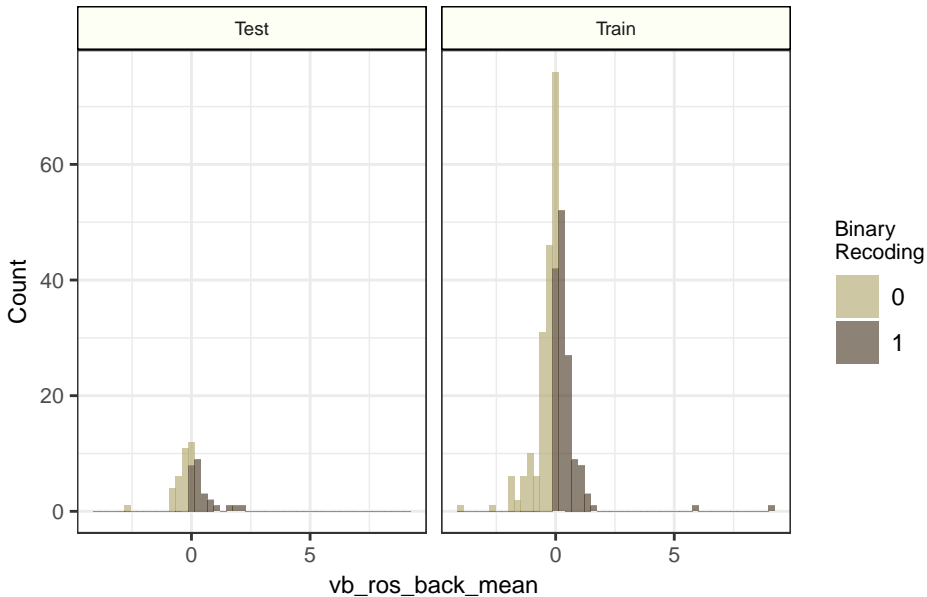
# vb\_percent\_dead\_only



# vb\_percent\_live



# vb\_ros\_back\_mean



# vb\_ros\_mean

