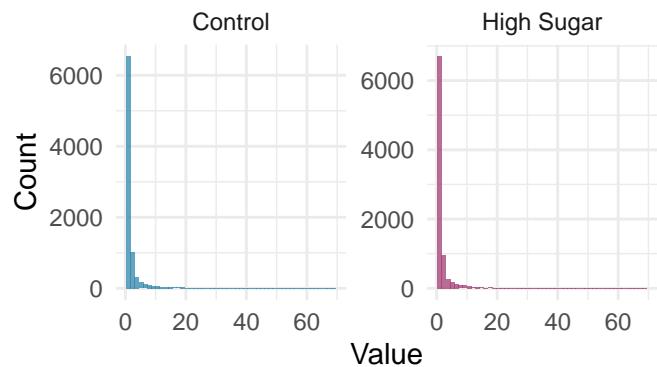
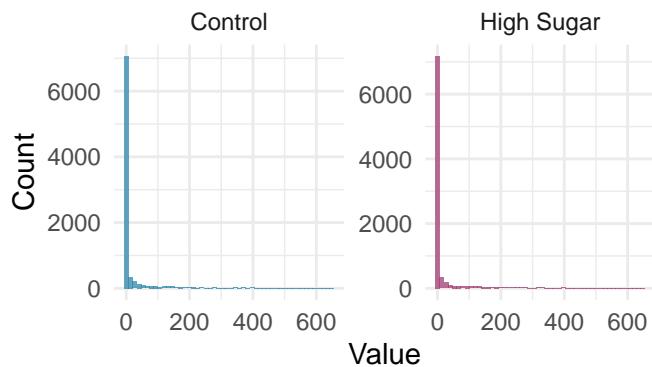


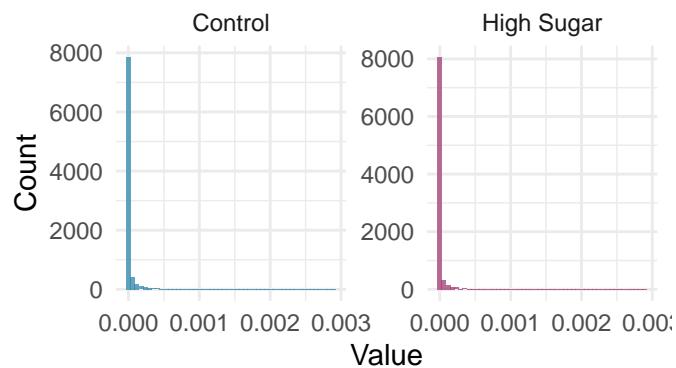
## weighted connectivity



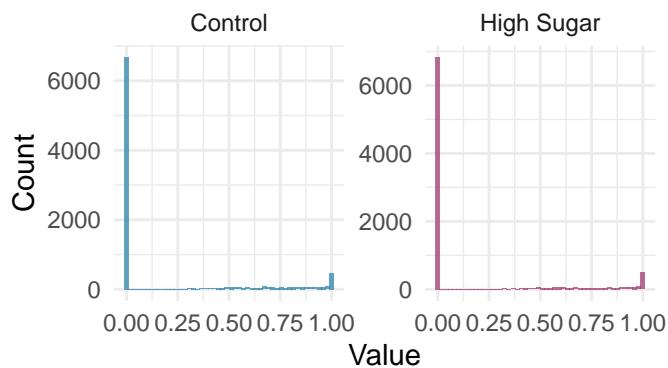
## degree



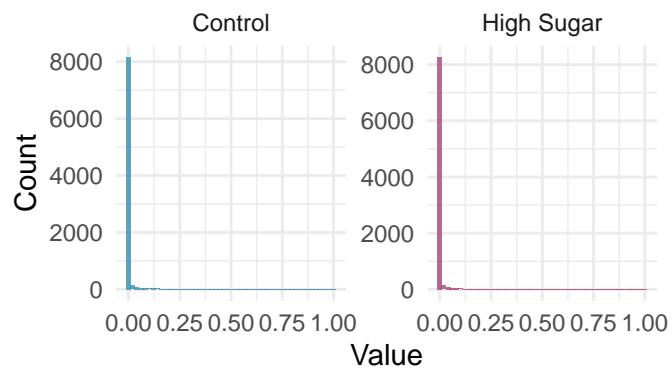
## betweenness centrality



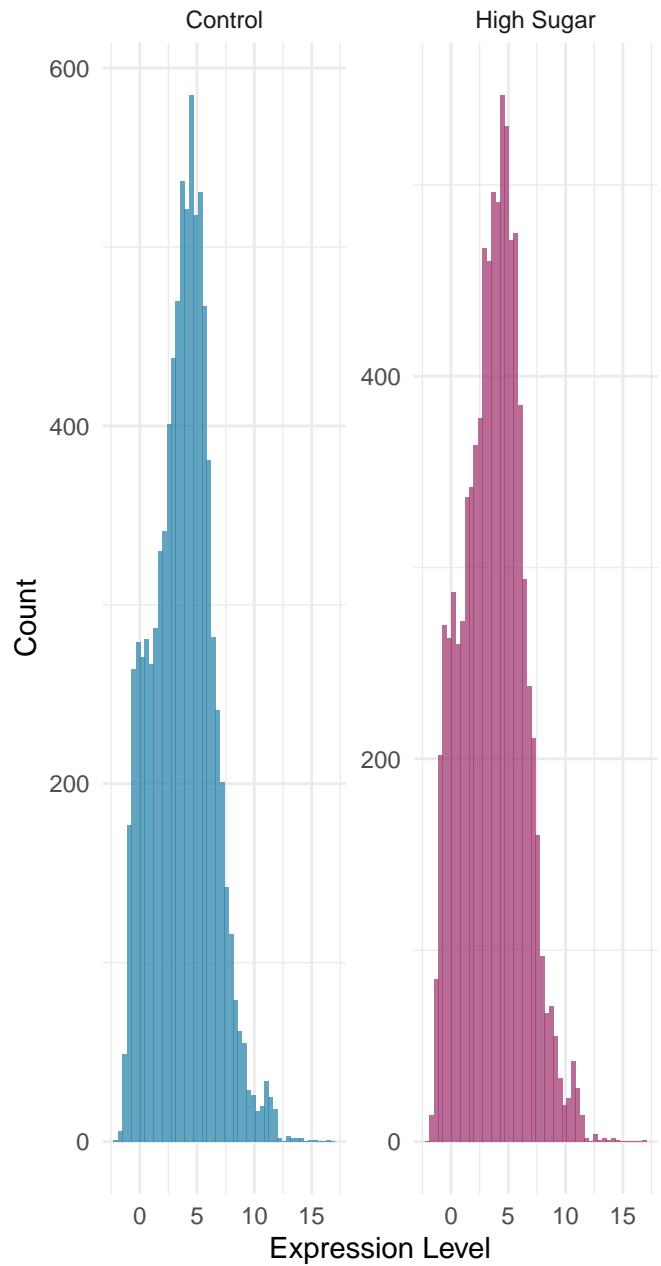
## clustering coefficient



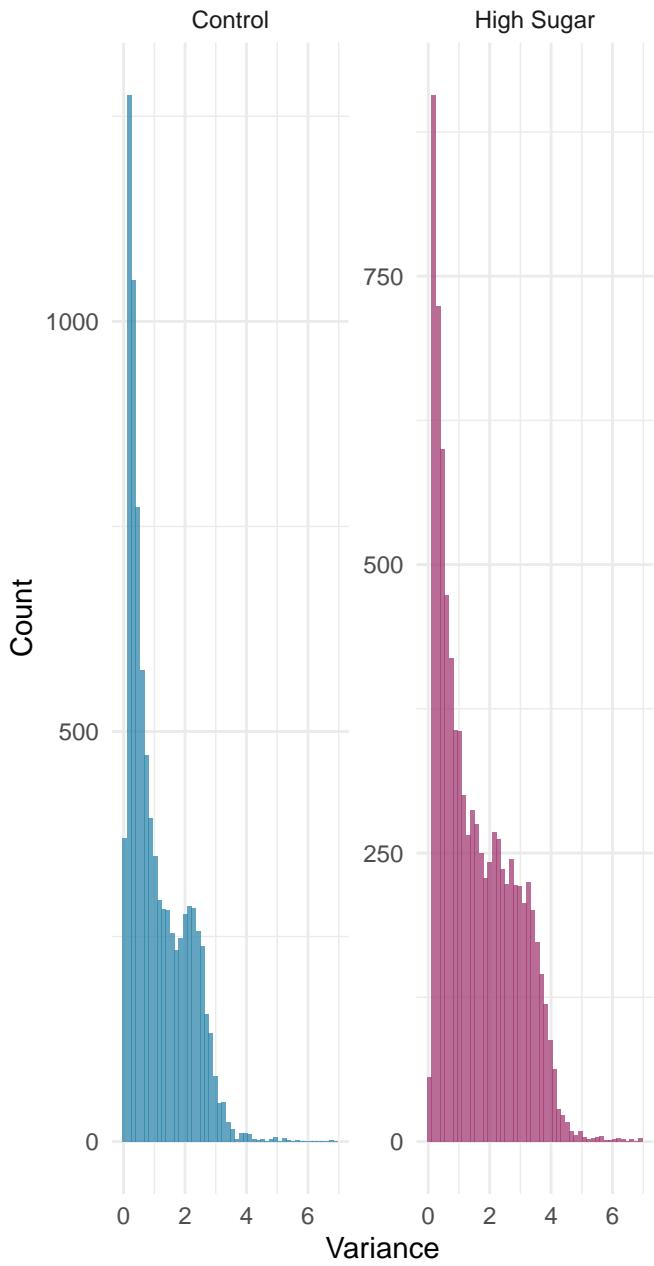
## eigenvector centrality



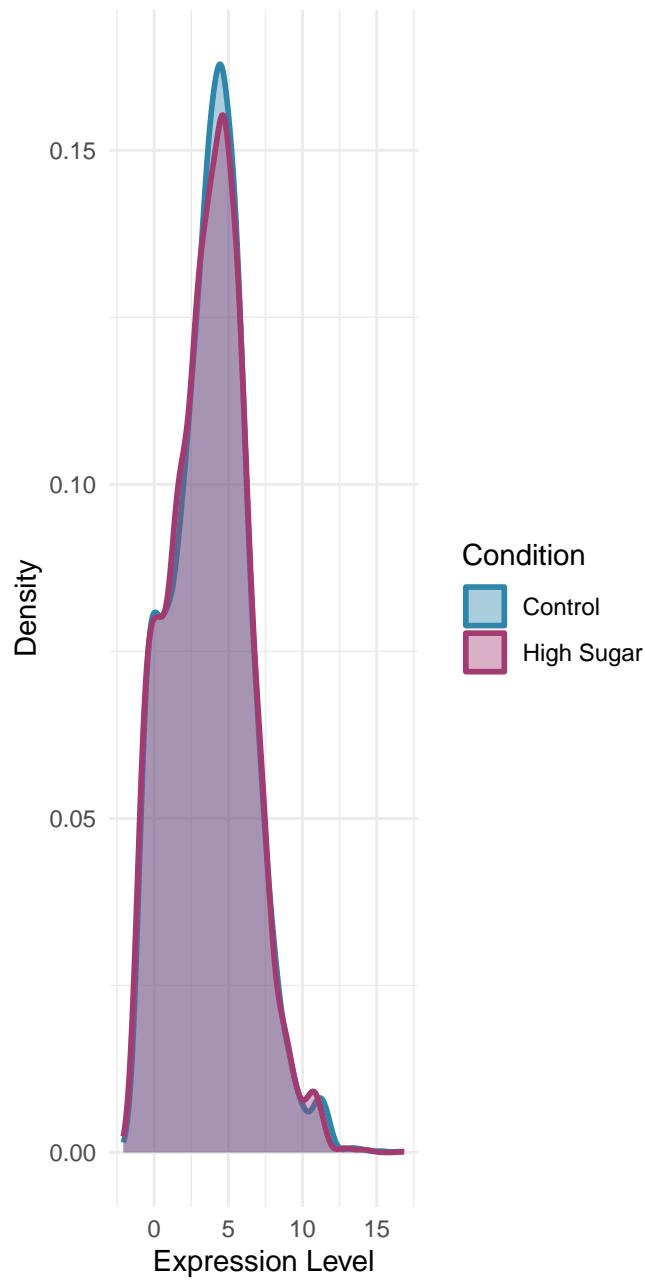
### Mean Expression



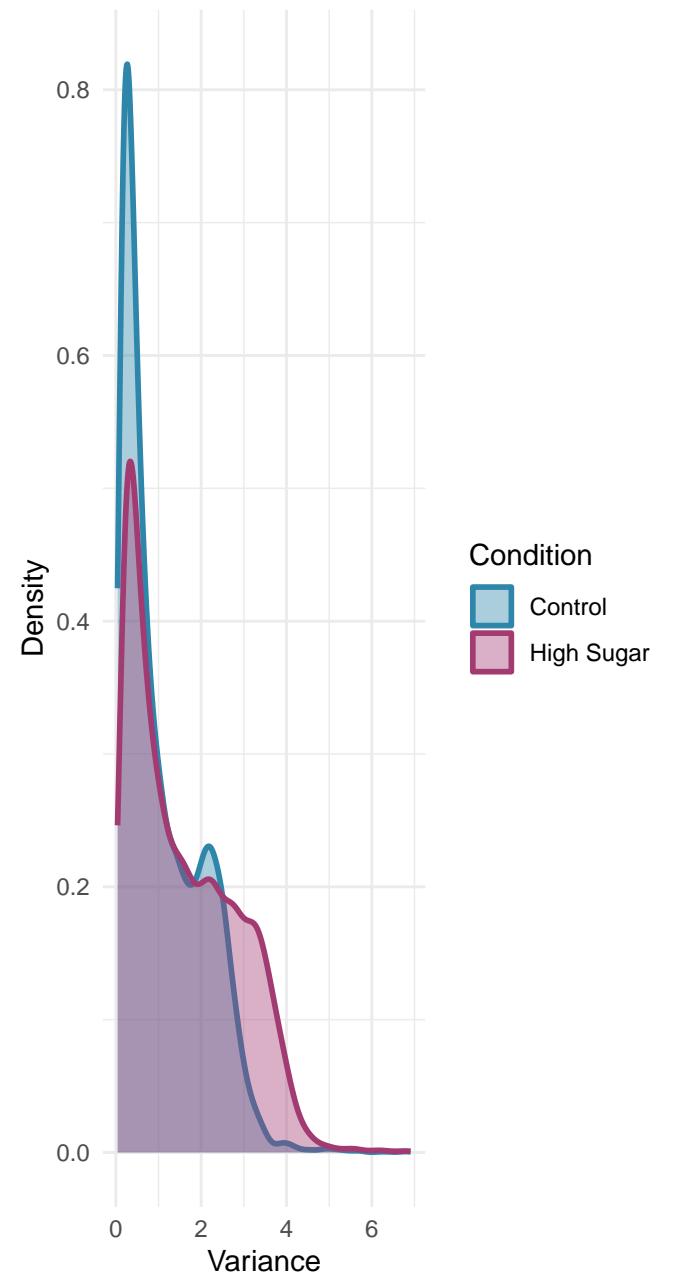
### Expression Variance

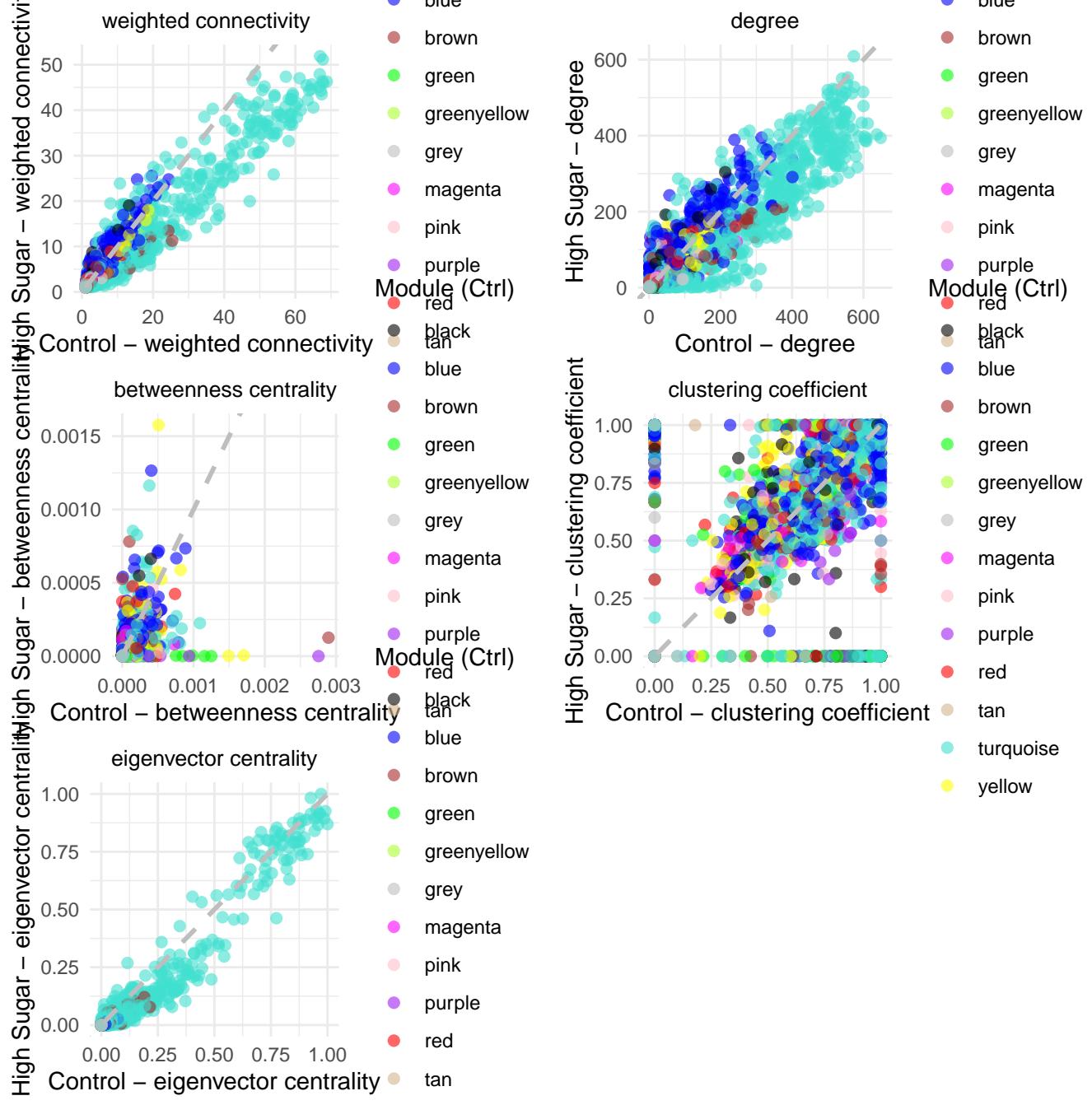


### Mean Expression Distribution (Overall)

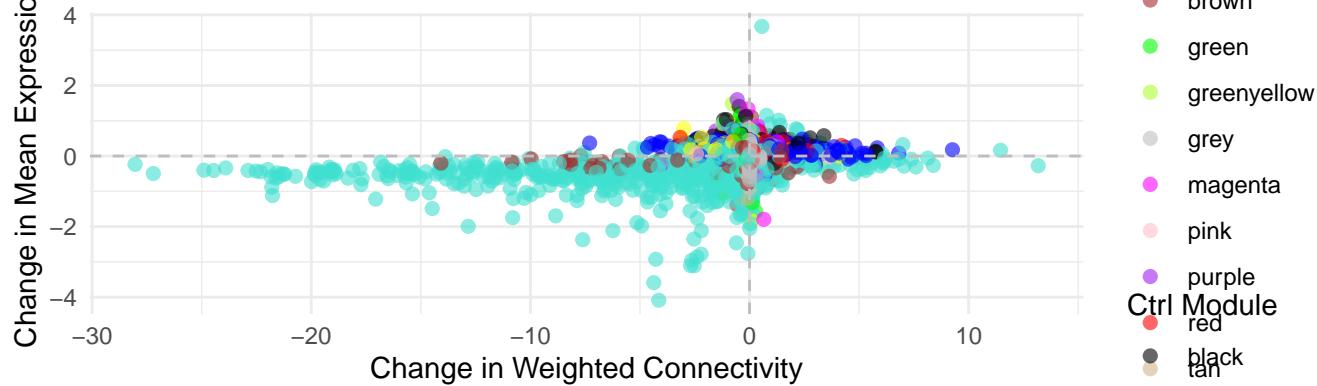


### Expression Variance Distribution (Overall)

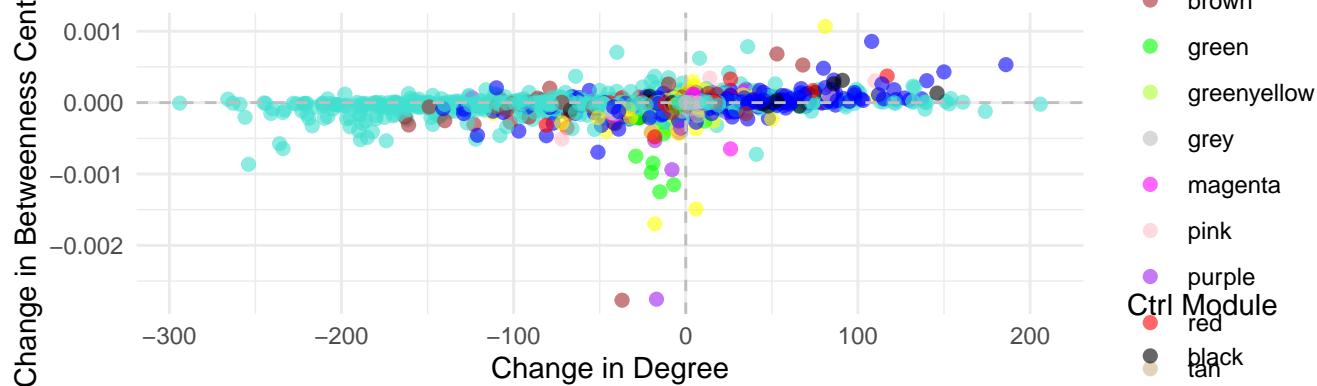




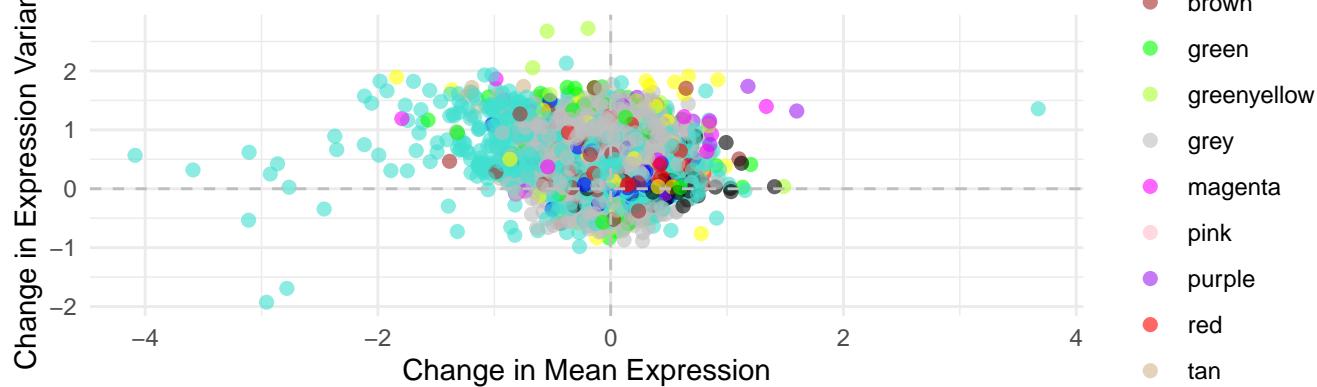
### Connectivity vs Expression Changes (Ctrl → Treat)



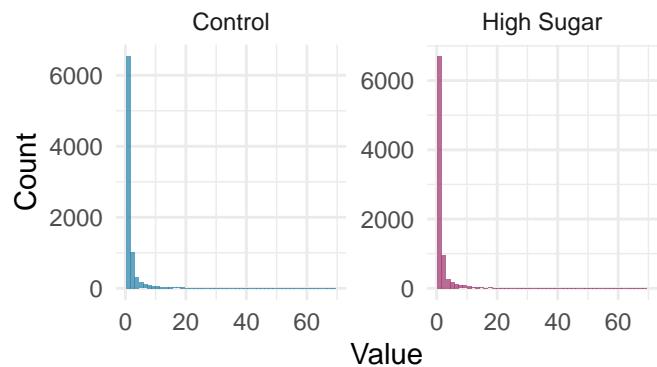
### Degree vs Betweenness Changes (Ctrl → Treat)



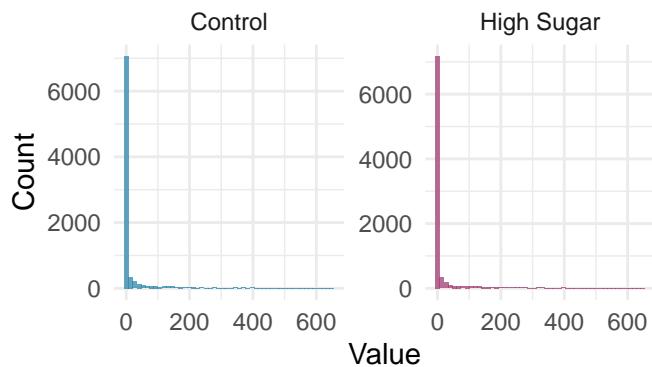
### Expression vs Variance Changes (Ctrl → Treat)



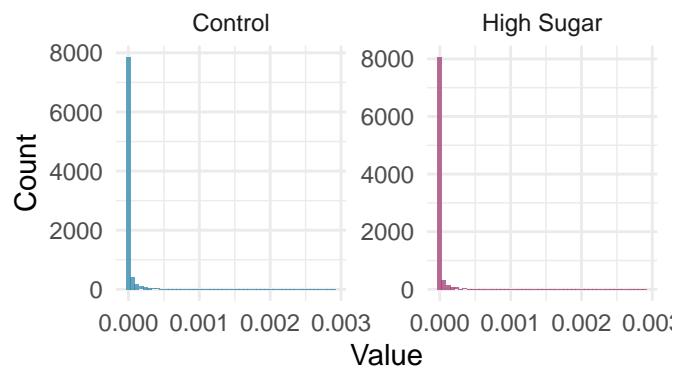
## weighted connectivity



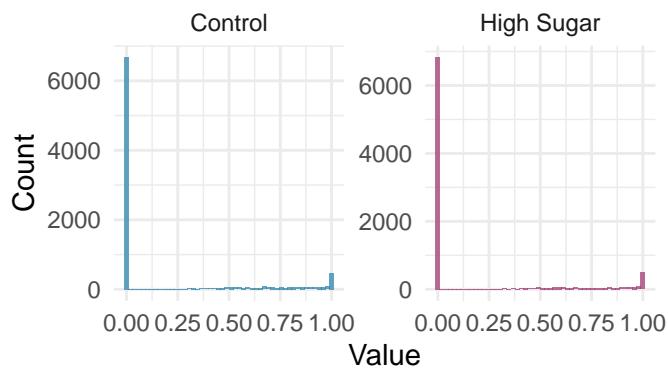
## degree



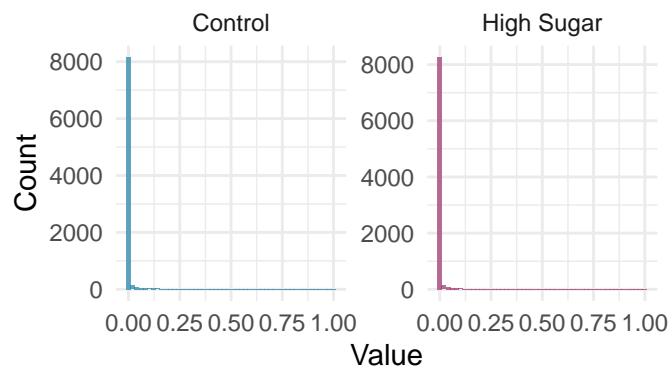
## betweenness centrality



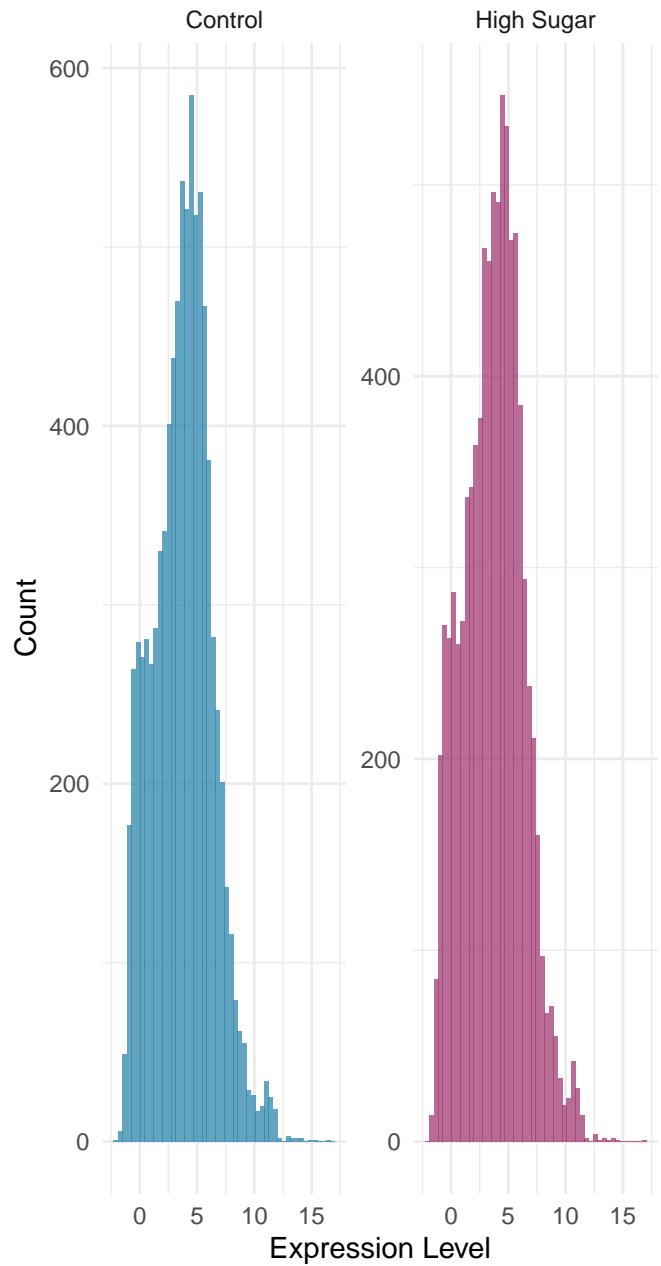
## clustering coefficient



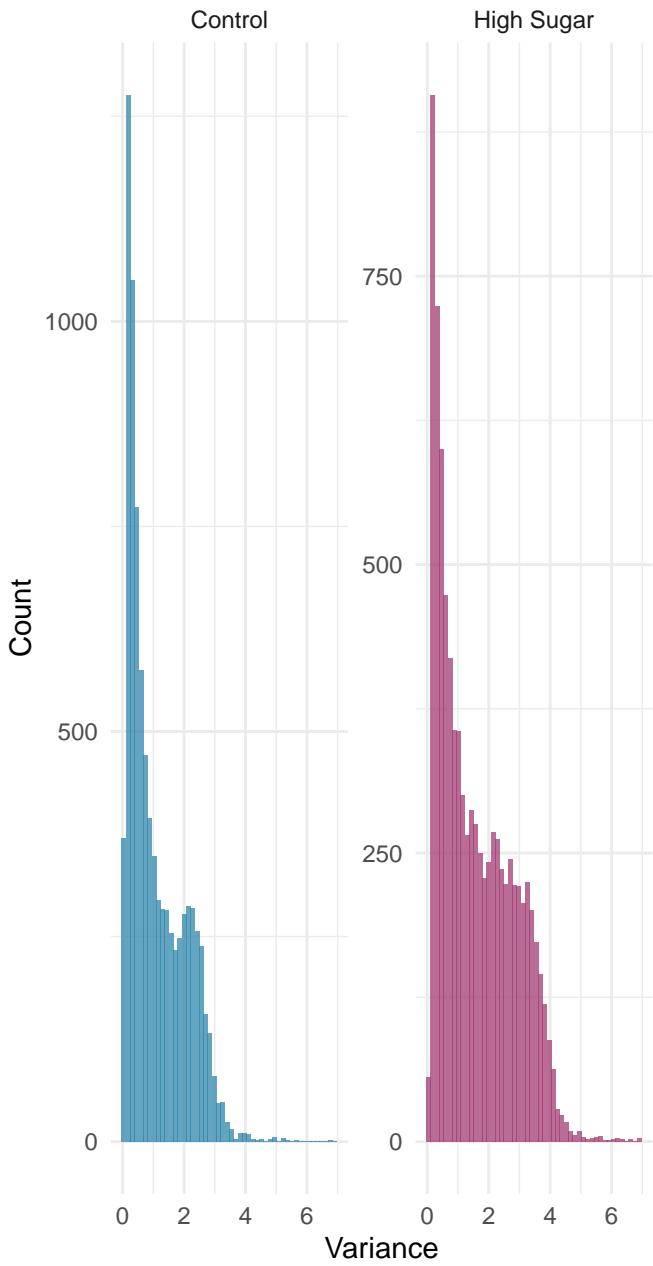
## eigenvector centrality



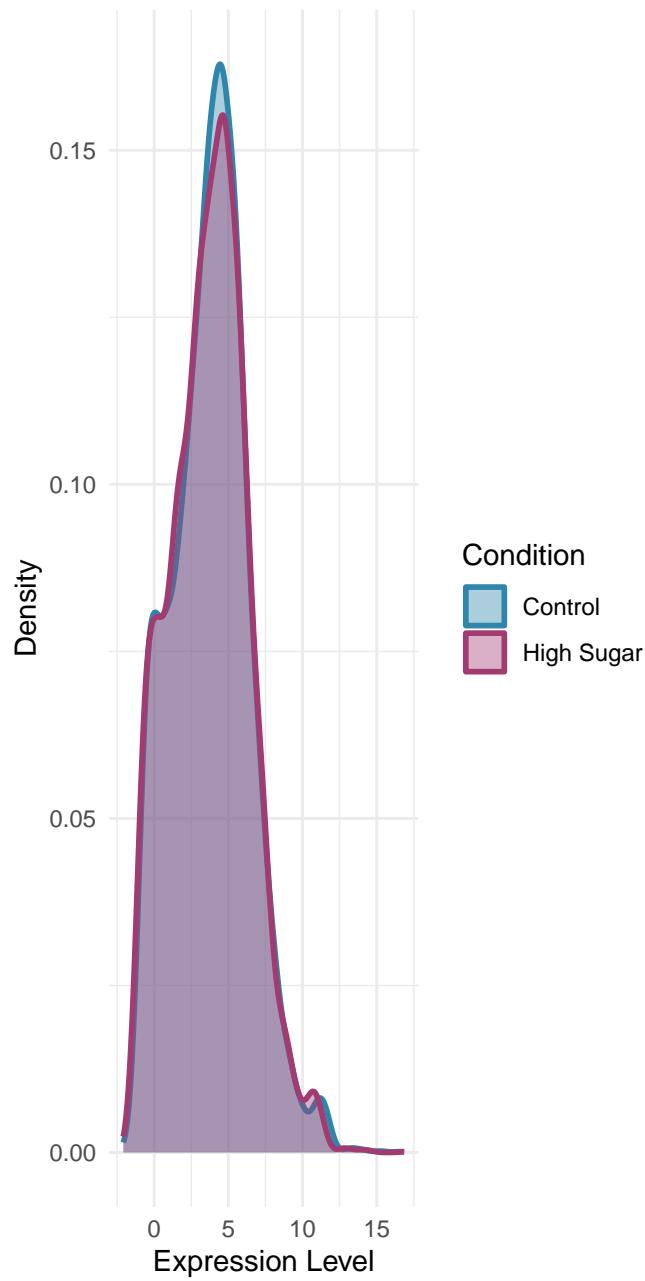
### Mean Expression



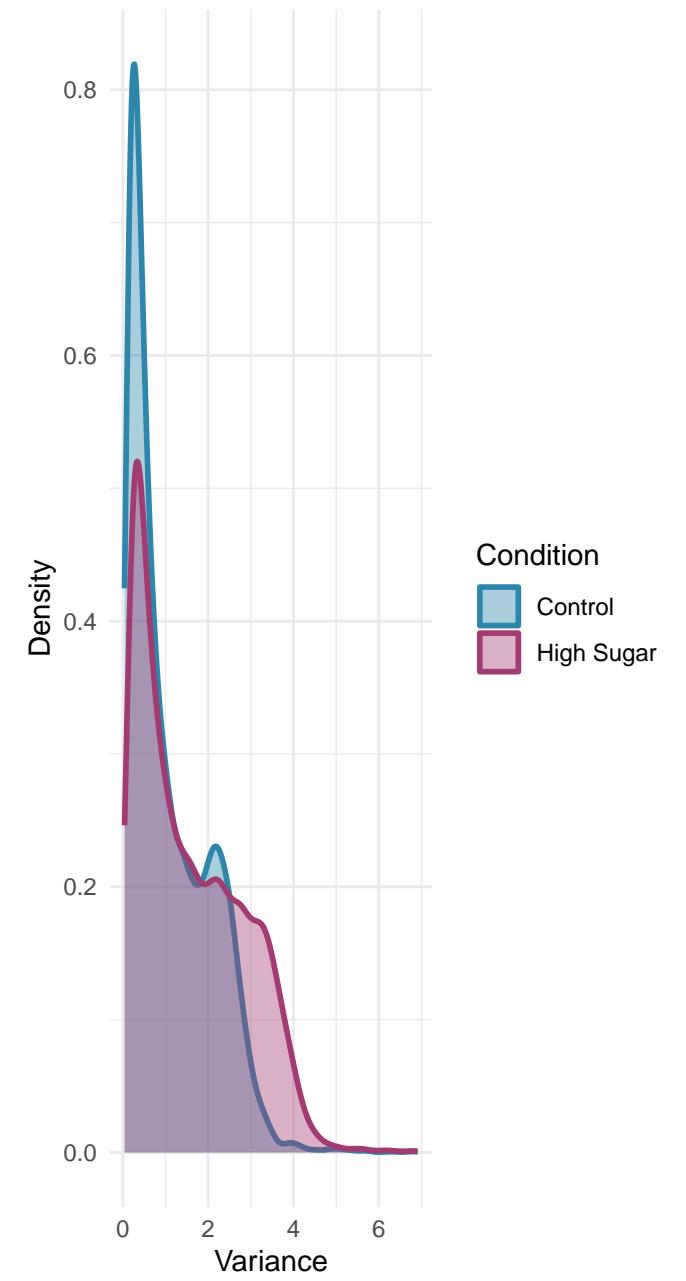
### Expression Variance

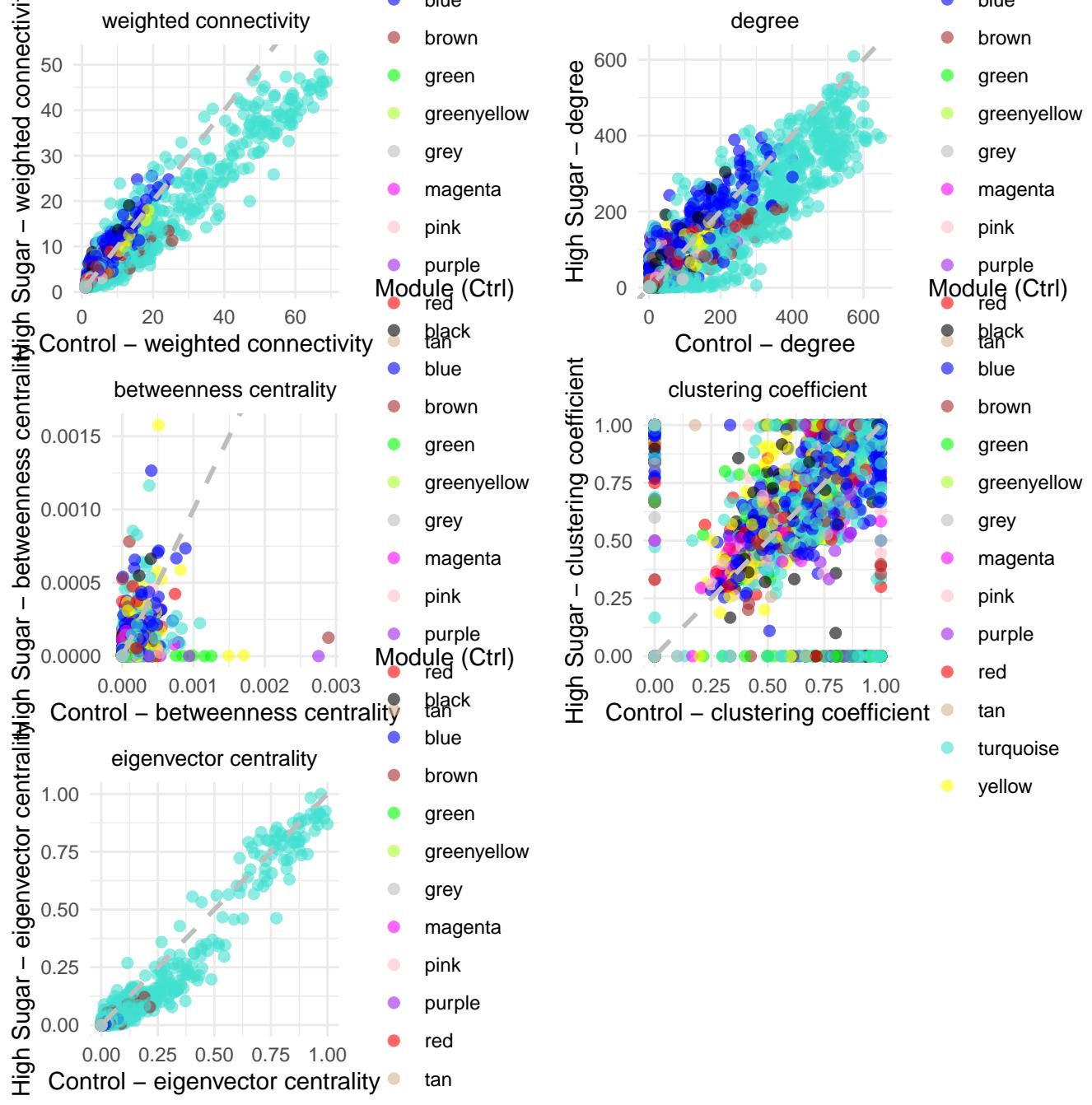


### Mean Expression Distribution (Overall)

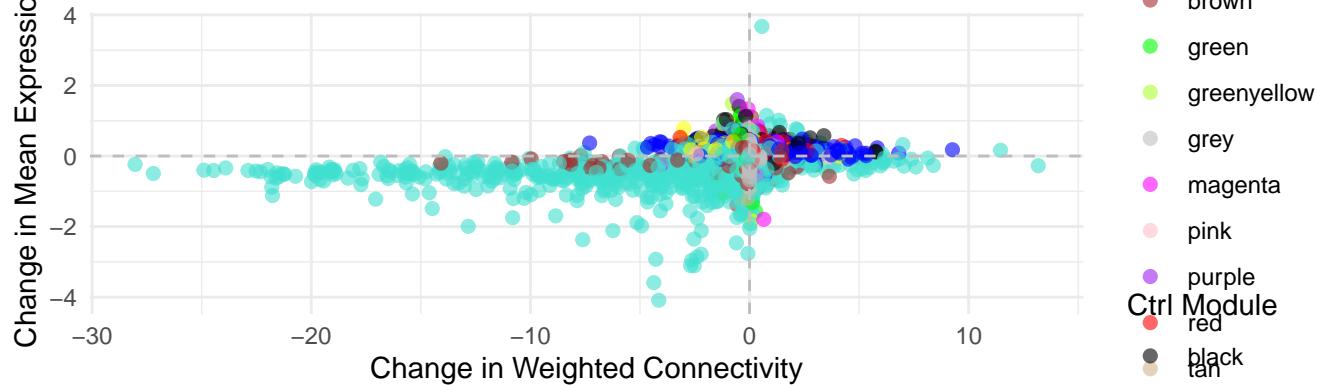


### Expression Variance Distribution (Overall)

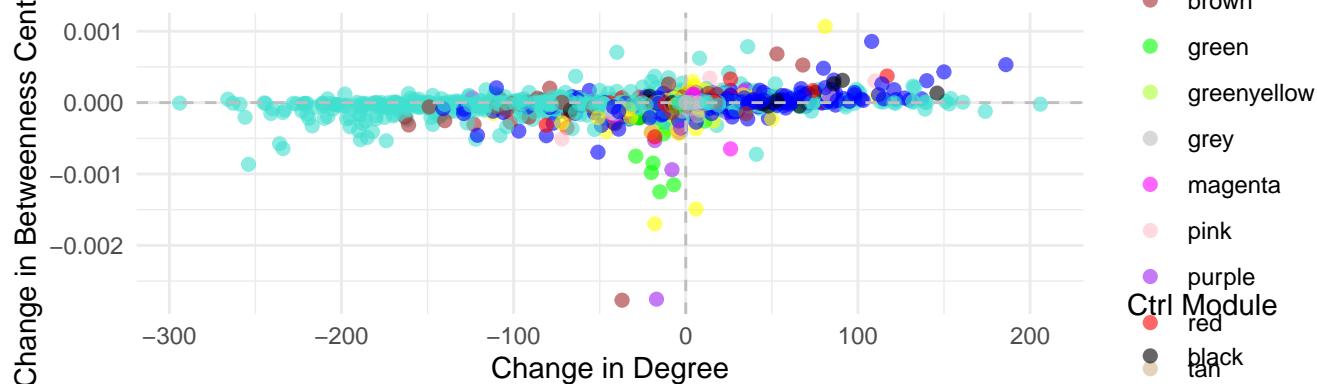




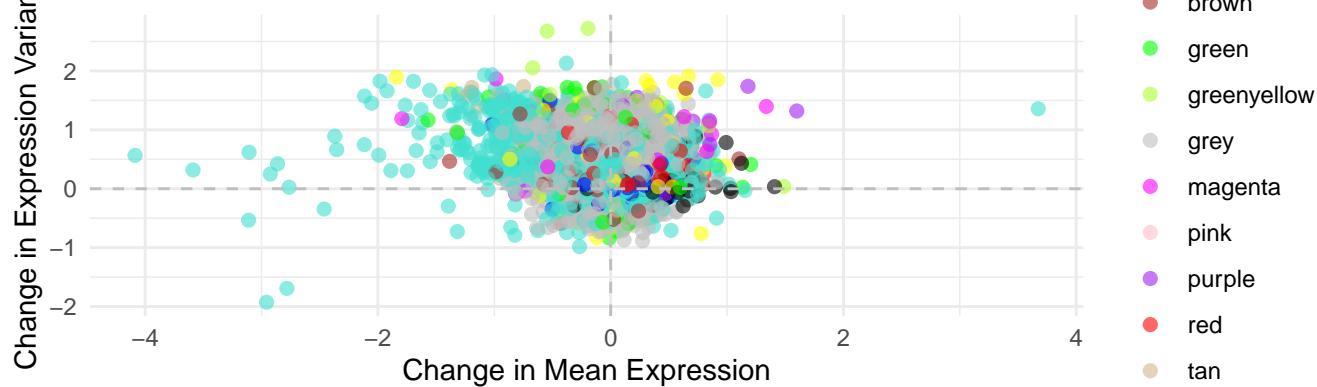
### Connectivity vs Expression Changes (Ctrl → Treat)



### Degree vs Betweenness Changes (Ctrl → Treat)



### Expression vs Variance Changes (Ctrl → Treat)



# Top 50 Genes by Connectivity Change (Z-scored deltas)

