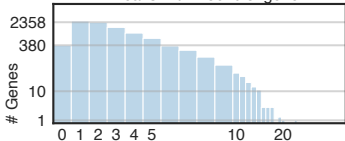
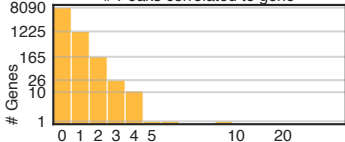


# Peaks within 50kb of gene



# Peaks correlated to gene



# Eigenpeaks correlated to gene

