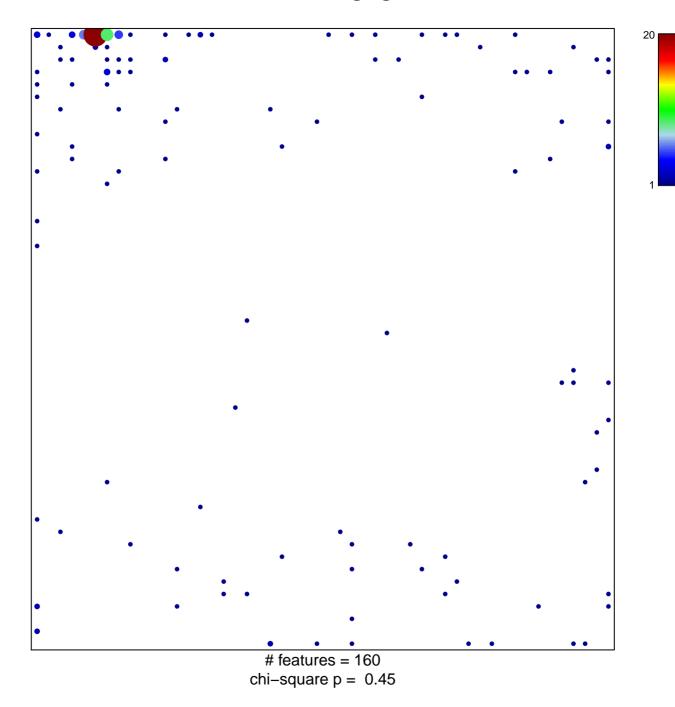
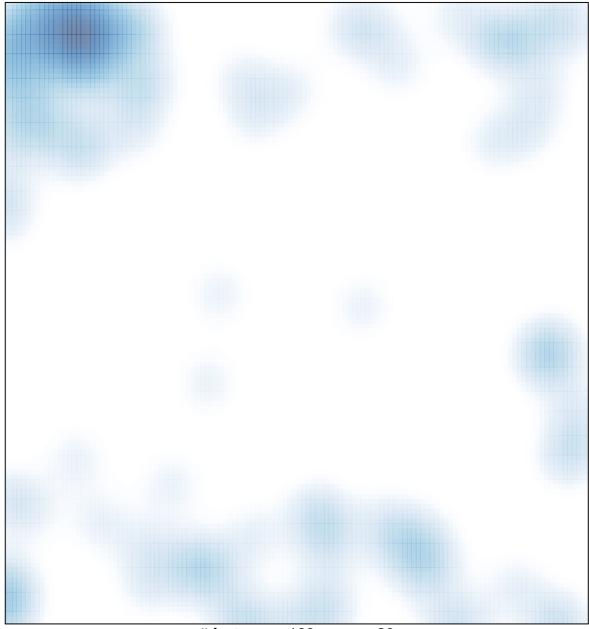
chromosome segregation



## chromosome segregation



# features = 160 , max = 20