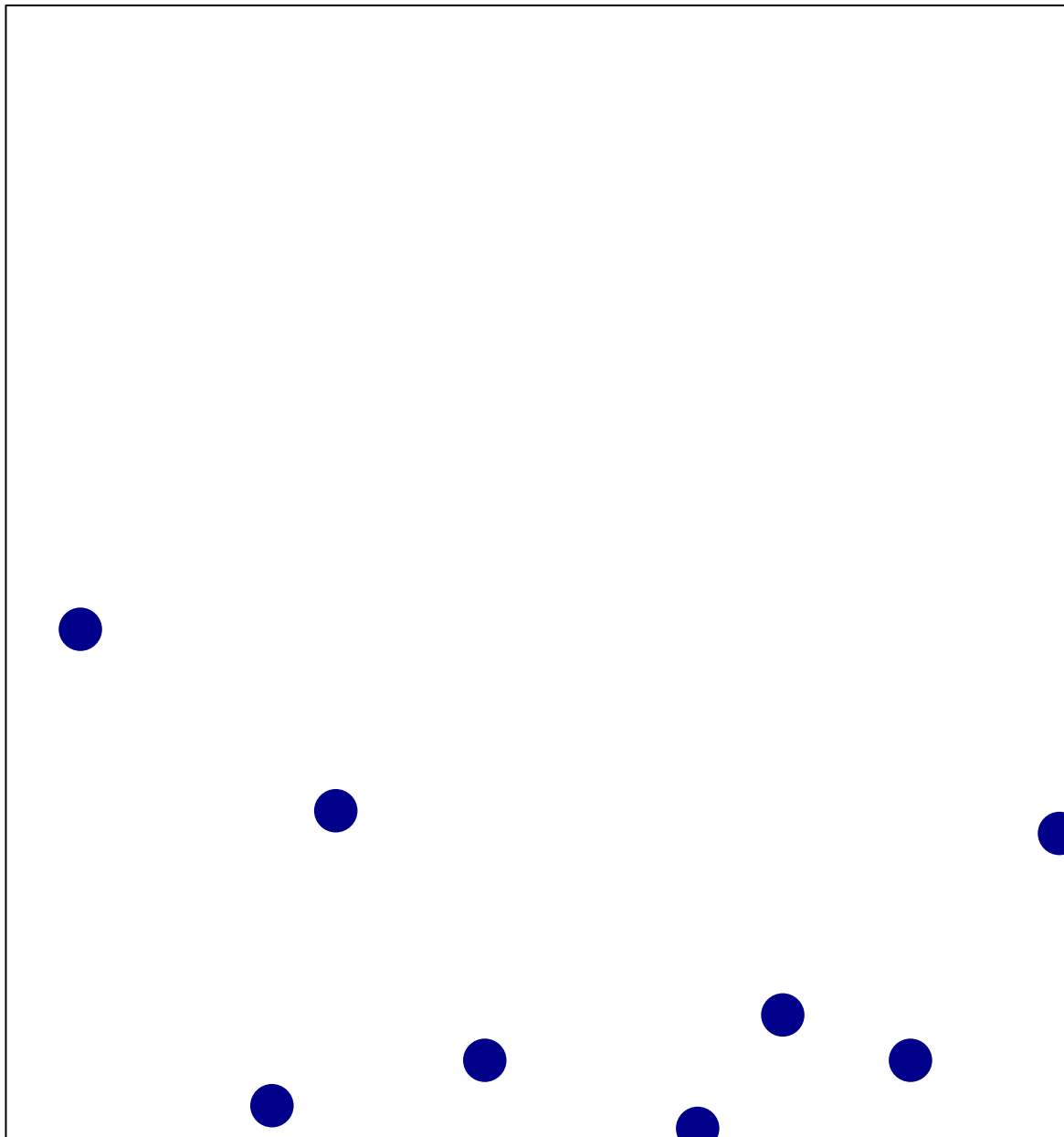
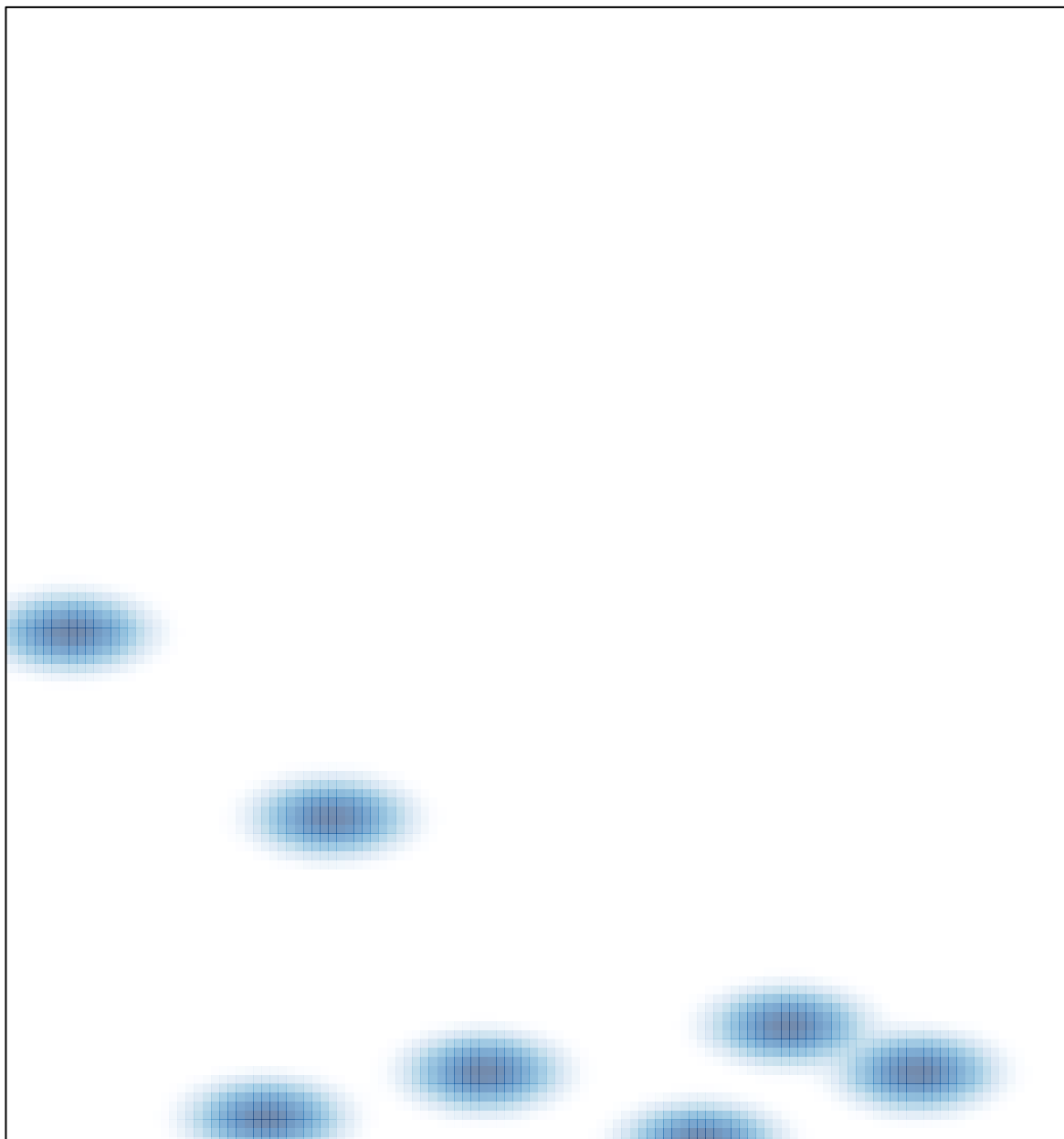


# FIGUEROA\_AML\_METHYLATION\_CLUSTER\_4\_DN



# features = 8  
chi-square p = 0.84

# FIGUEROA\_AML\_METHYLATION\_CLUSTER\_4\_DN



# features = 8 , max = 1