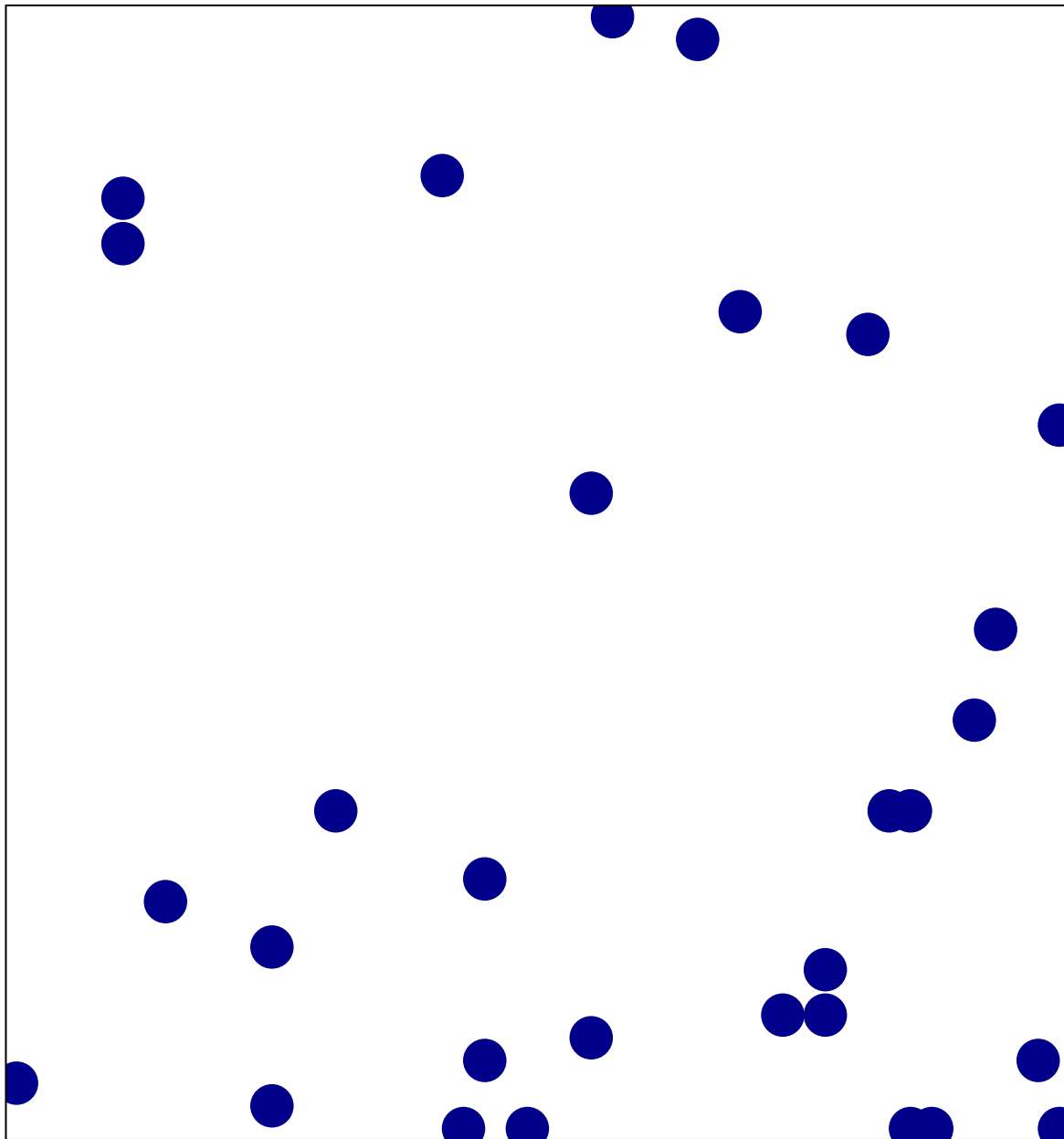
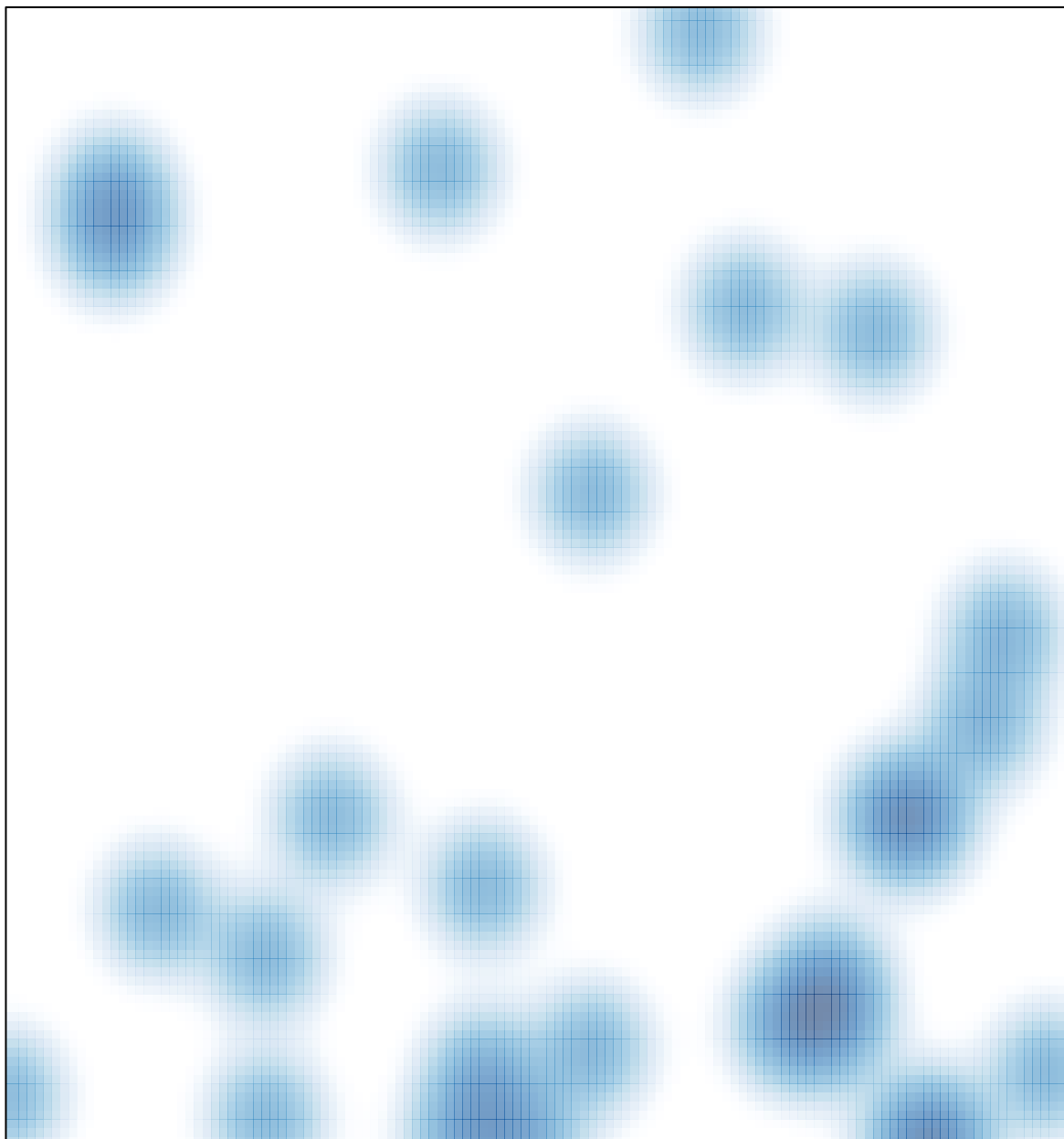


# FIGUEROA\_AML\_METHYLATION\_CLUSTER\_3\_DN



# features = 30  
chi-square p = 0.83

# FIGUEROA\_AML\_METHYLATION\_CLUSTER\_3\_DN



# features = 30 , max = 1