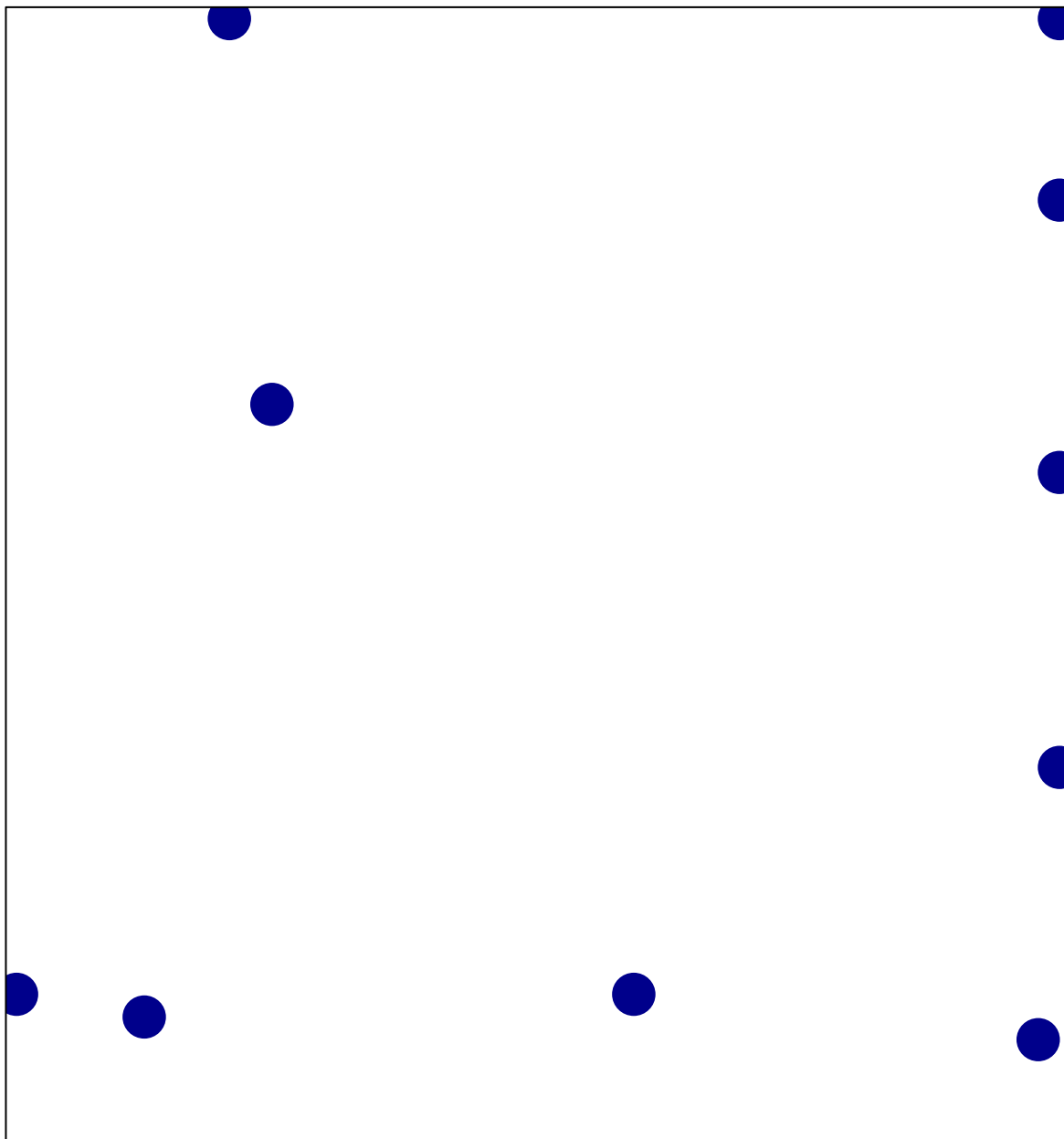
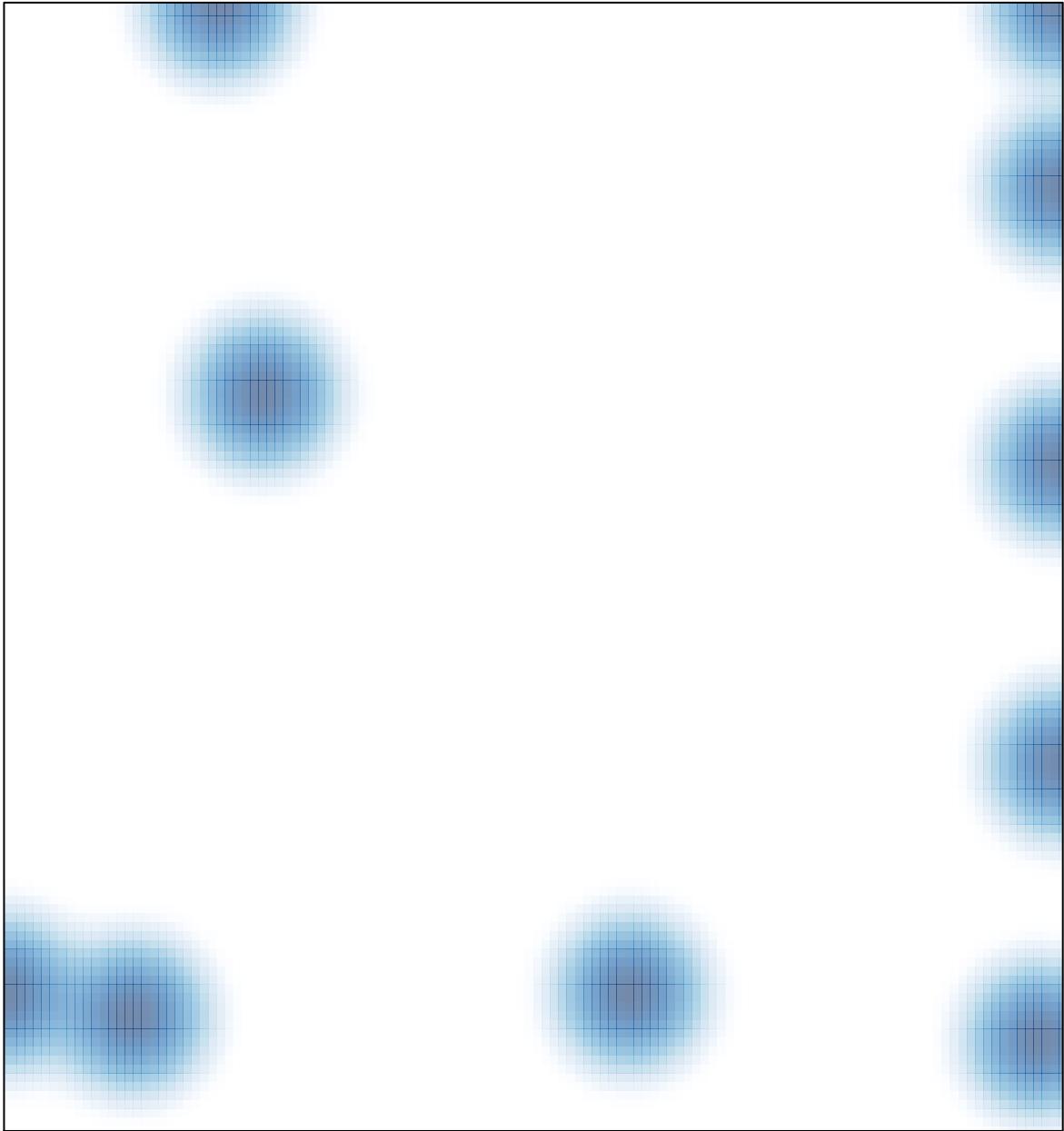


# HELLER\_HDAC\_TARGETS\_SILENCED\_BY\_METHYLATION\_UP



# features = 10  
chi-square p = 0.8

# MLER\_HDAC\_TARGETS\_SILENCED\_BY\_METHYLATIO



# features = 10 , max = 1