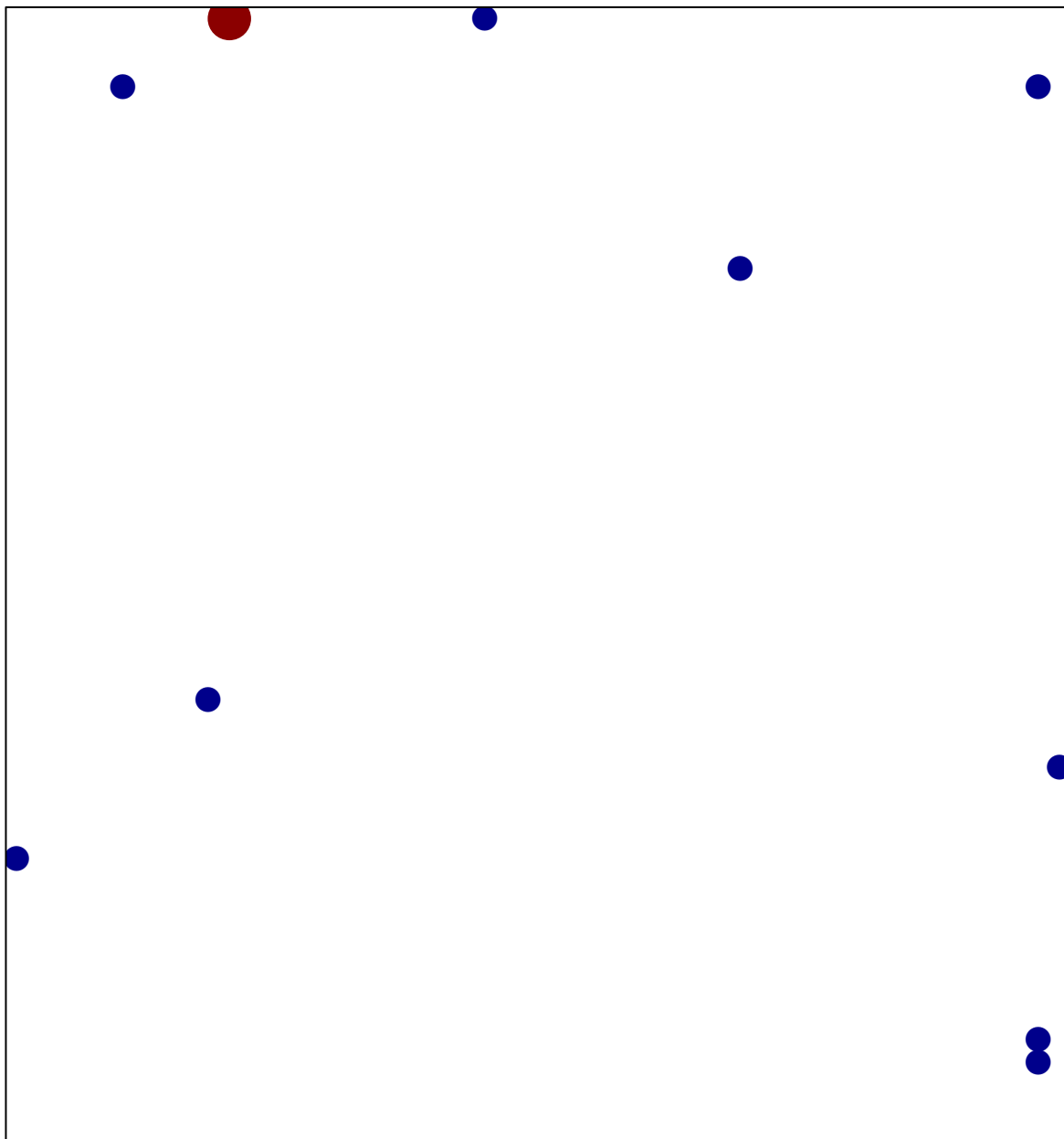
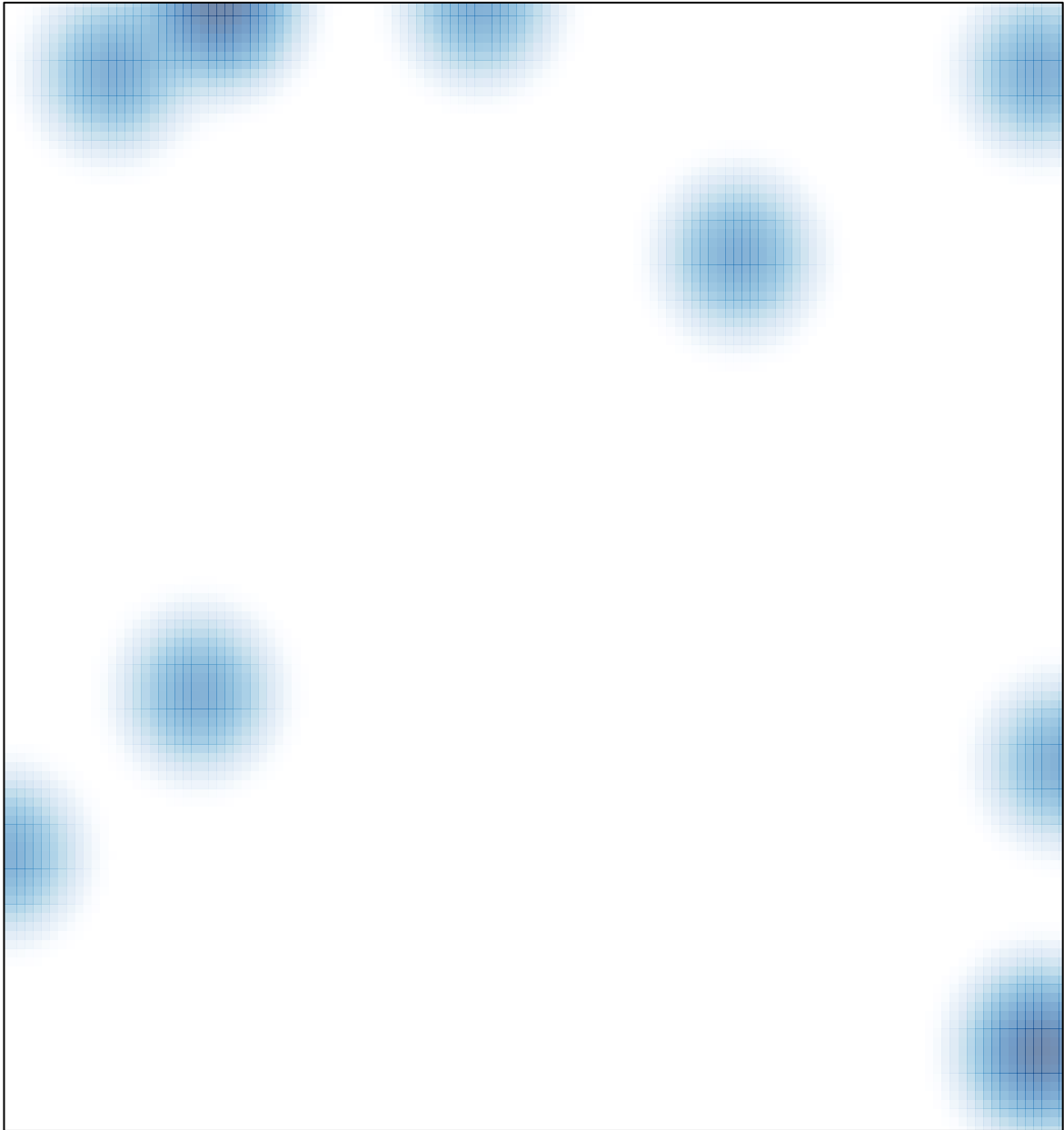


HELLER_HDAC_TARGETS_SILENCED_BY_METHYLATION_DN



features = 11
chi-square p = 0.81

LER_HDAC_TARGETS_SILENCED_BY_METHYLATIO



features = 11 , max = 2