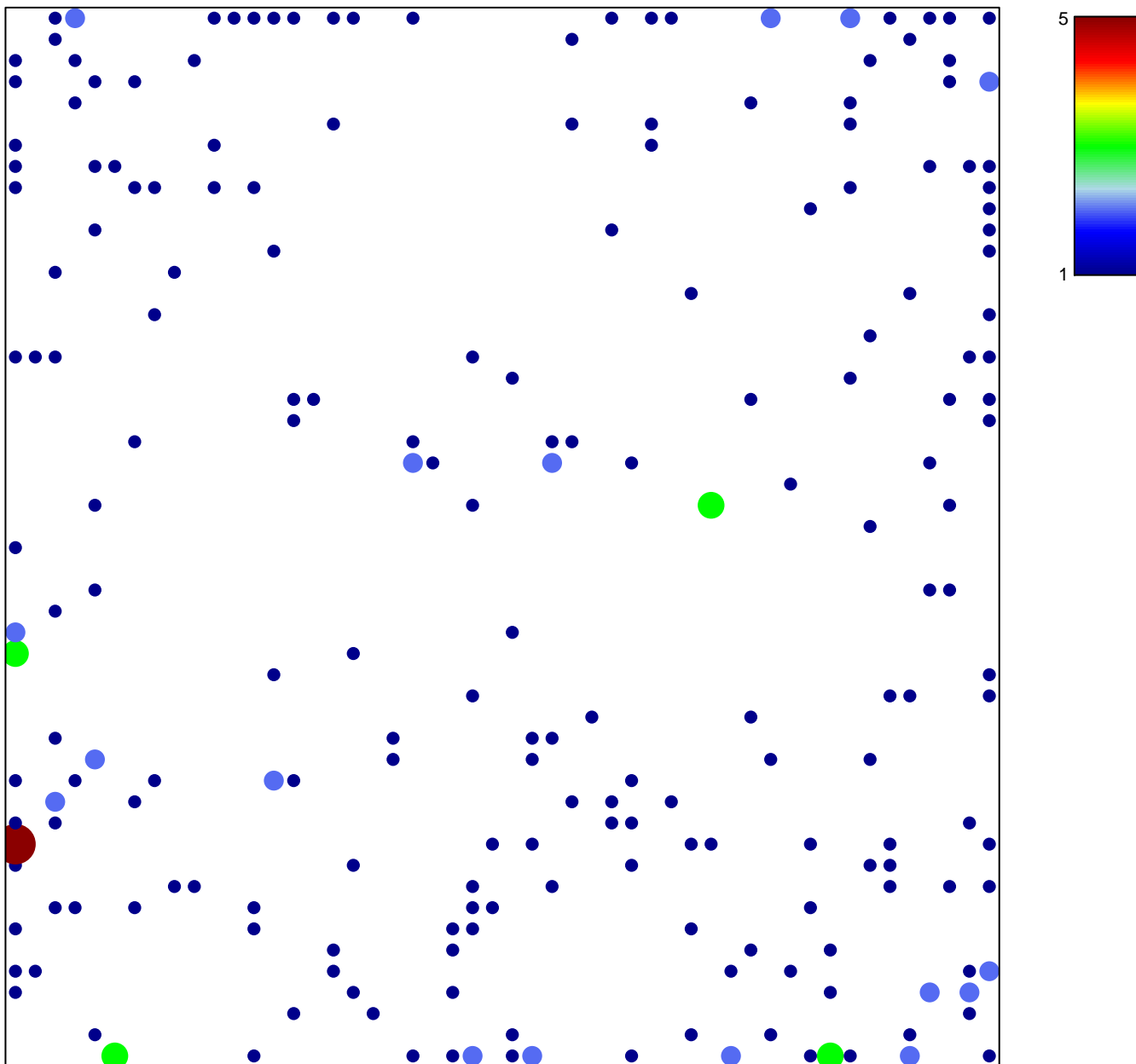
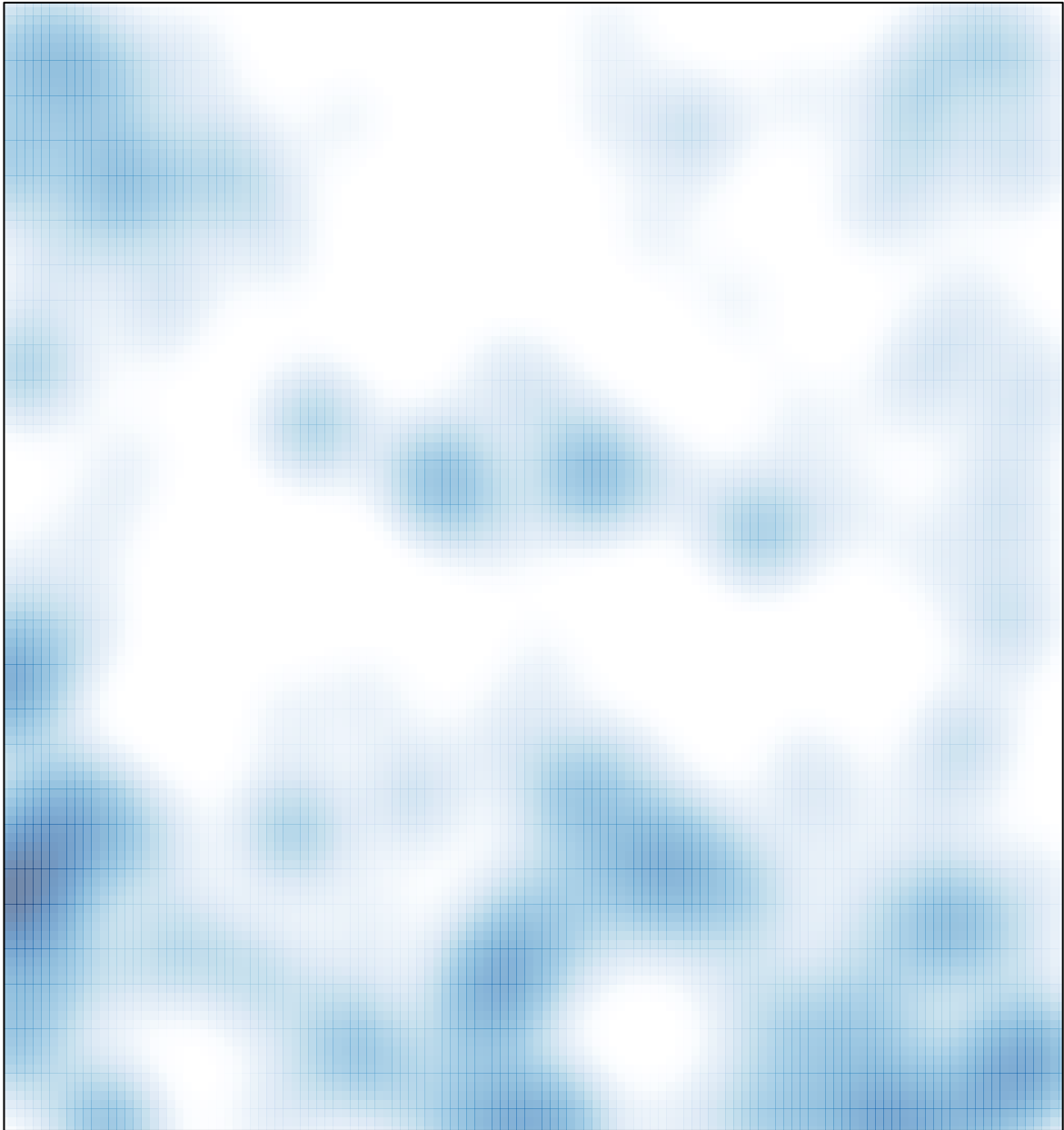


HELLER_HDAC_TARGETS_SILENCED_BY_METHYLATION_DN



features = 241
chi-square p = 0.71

LER_HDAC_TARGETS_SILENCED_BY_METHYLATIO



features = 241 , max = 5