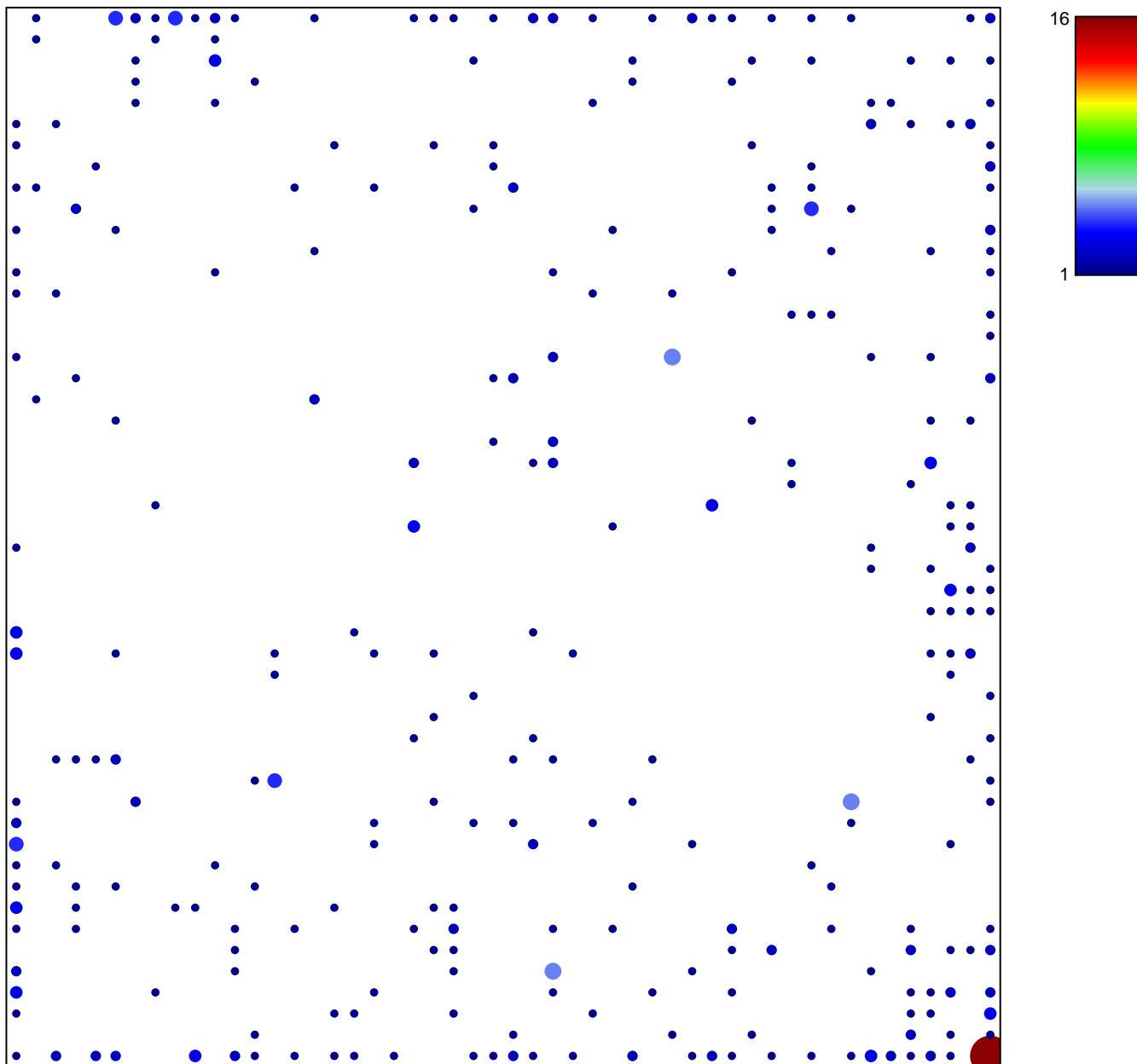
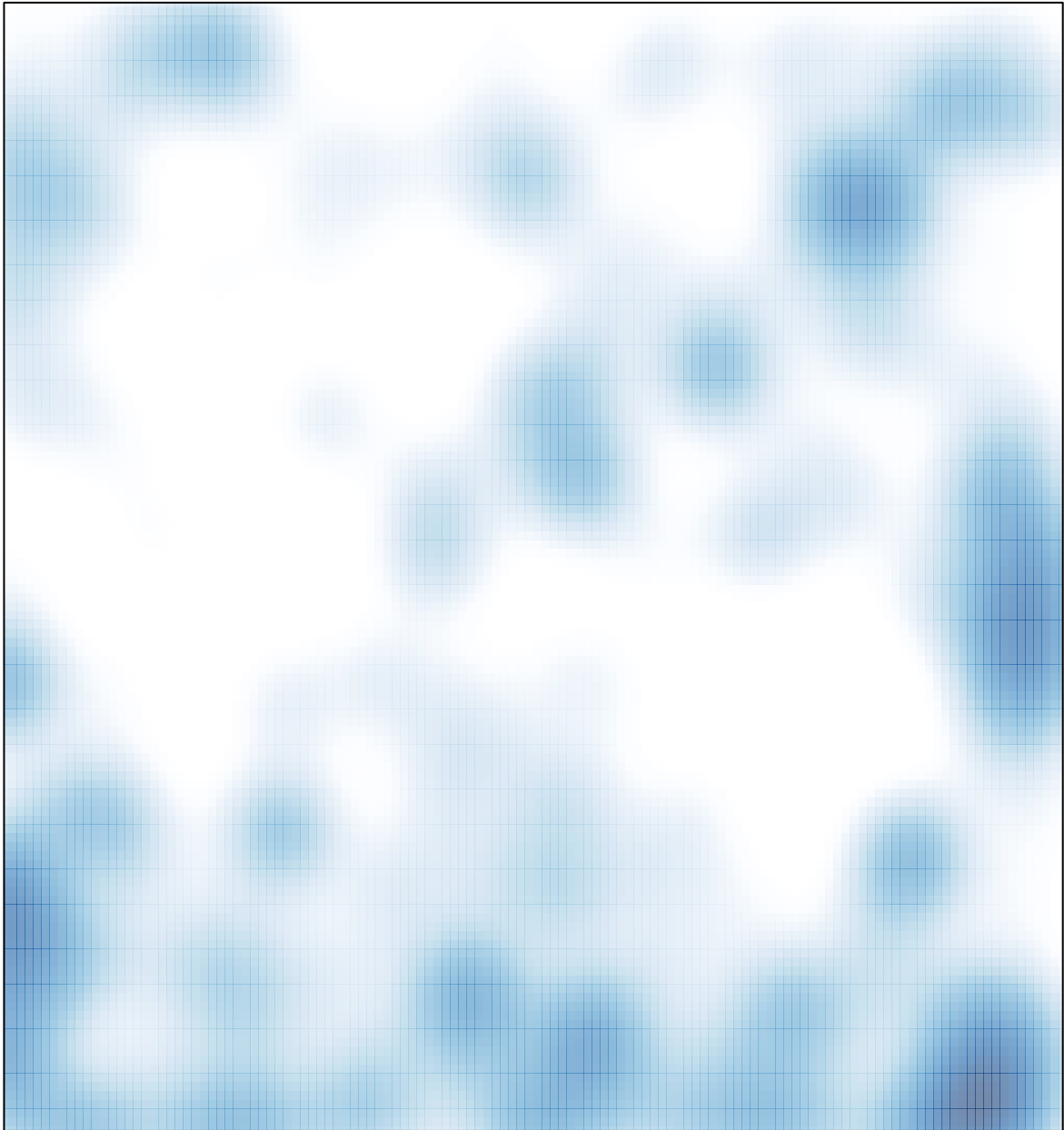


HELLER_HDAC_TARGETS_SILENCED_BY_METHYLATION_UP



features = 400
chi-square p = 0.56

MLER_HDAC_TARGETS_SILENCED_BY_METHYLATIO



features = 400 , max = 16