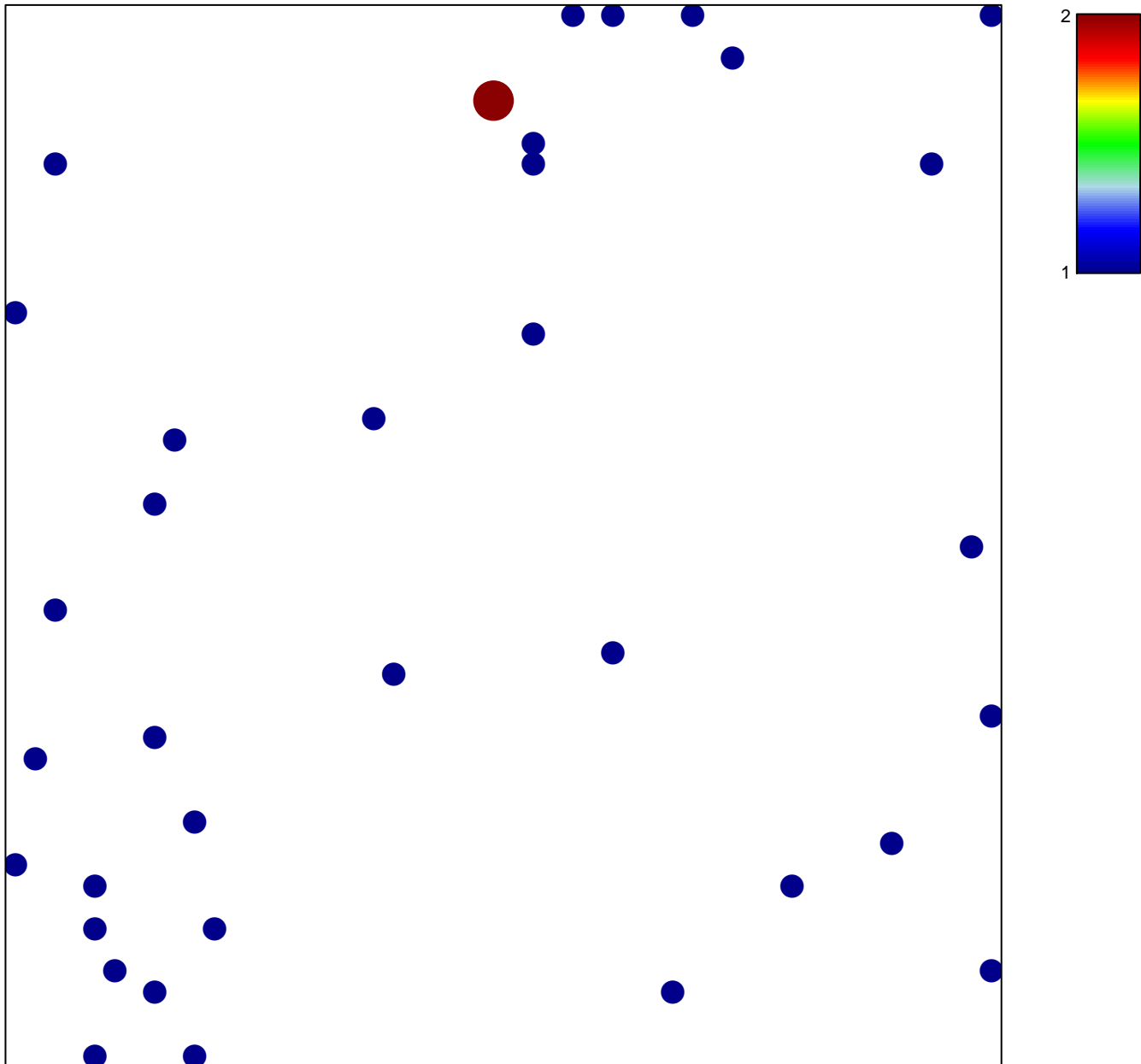
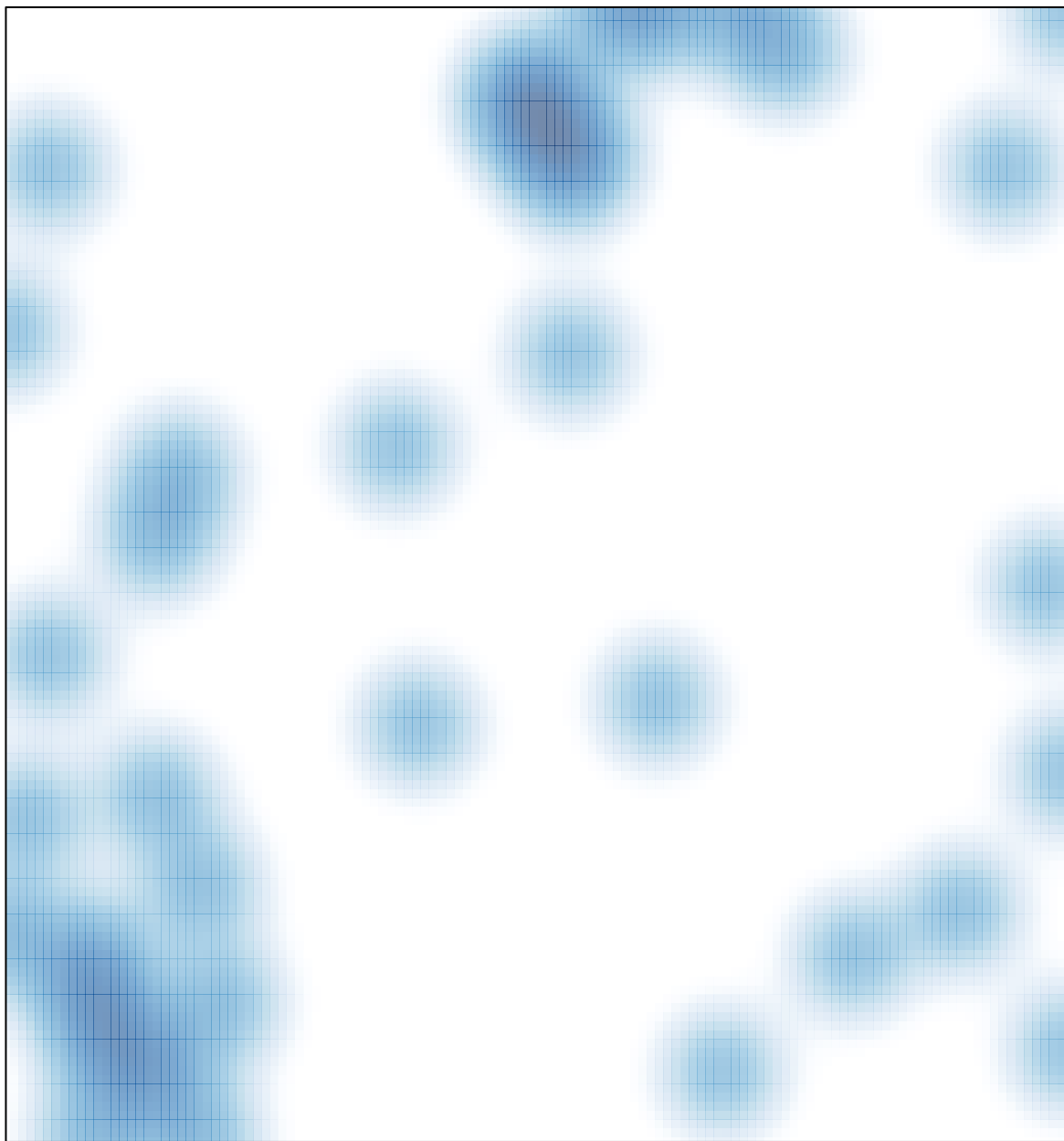


# positive regulation of proteasomal ubiquitin-dependent protein catabolic process



# features = 36  
chi-square p = 0.83

# ation of proteasomal ubiquitin-dependent protein ca



# features = 36 , max = 2