# BARRIER_CANCER_RELAPSE_TUMOR_SAMPLE_UP
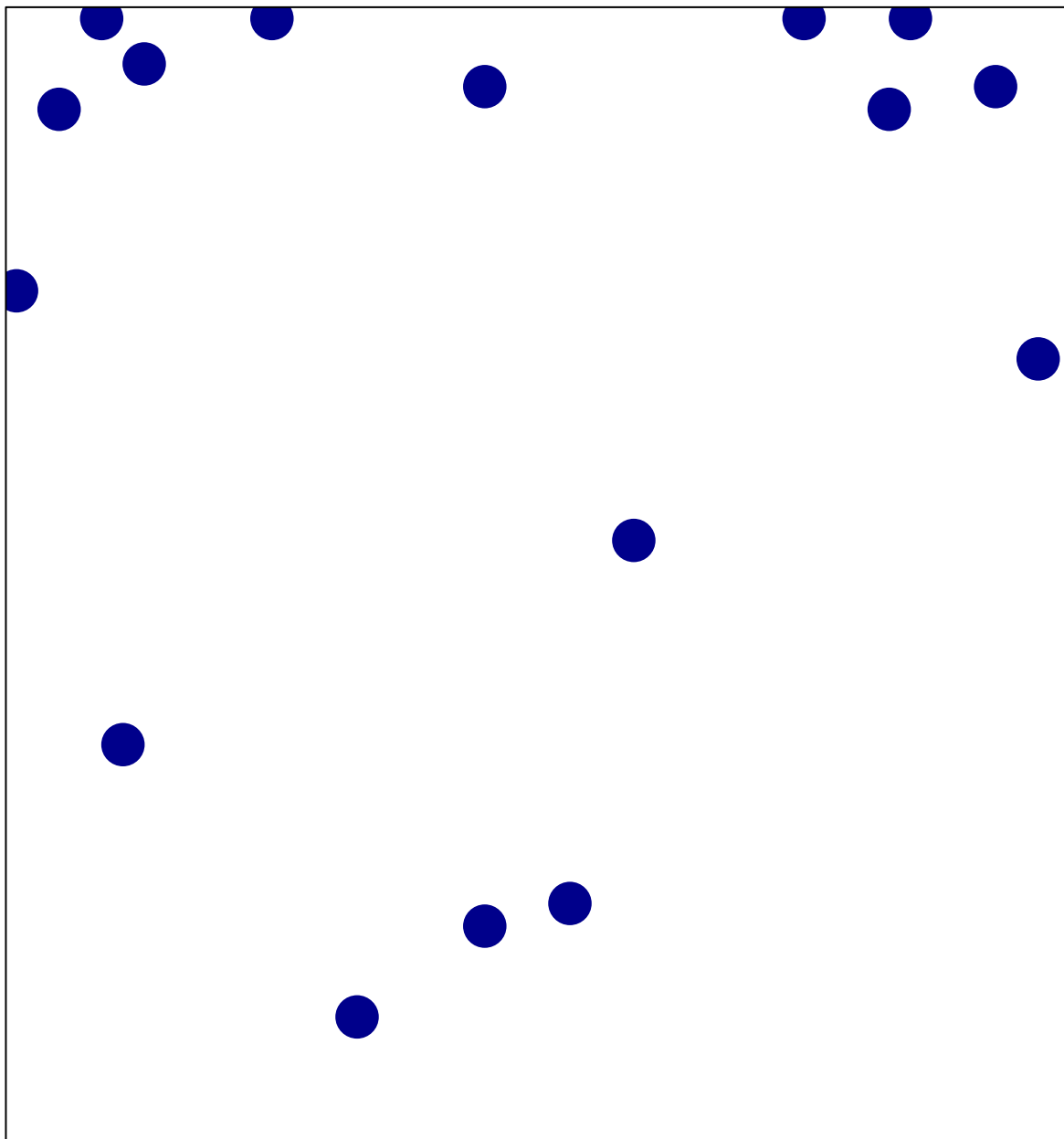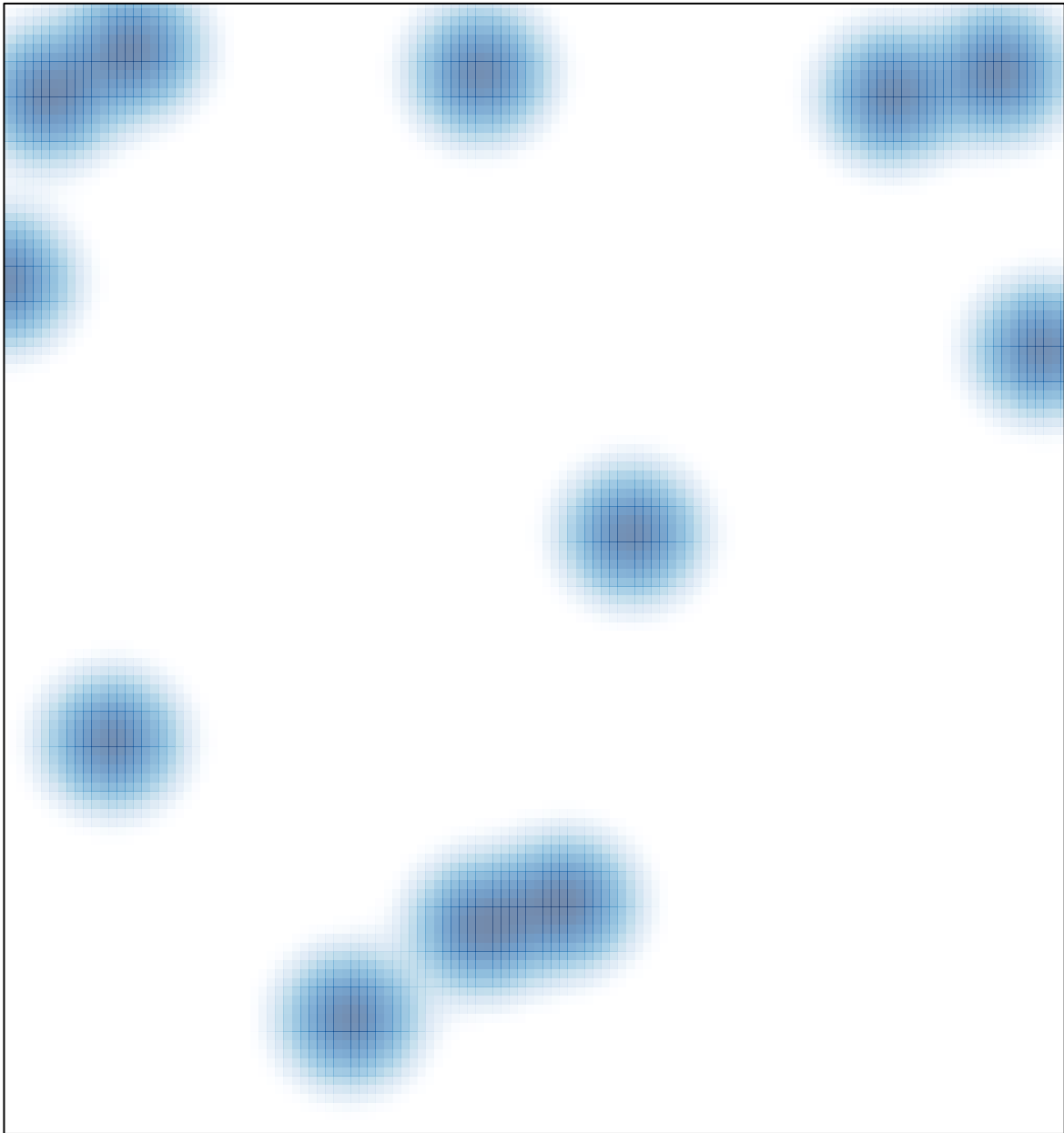


# features = 16
chi−square p =  0.82

# BARRIER_CANCER_RELAPSE_TUMOR_SAMPLE_U



# features = 16 , max = 1