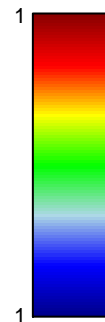
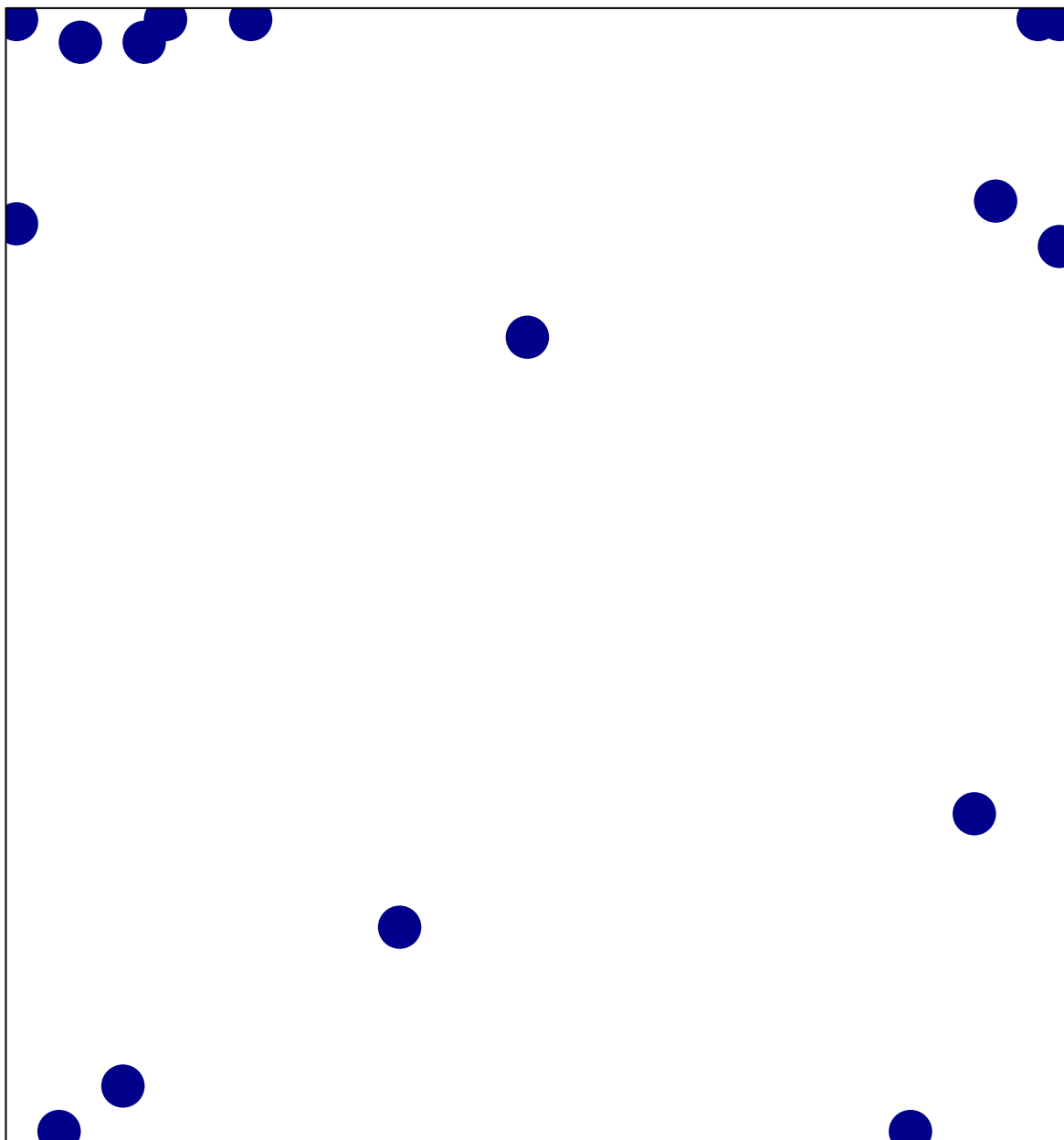
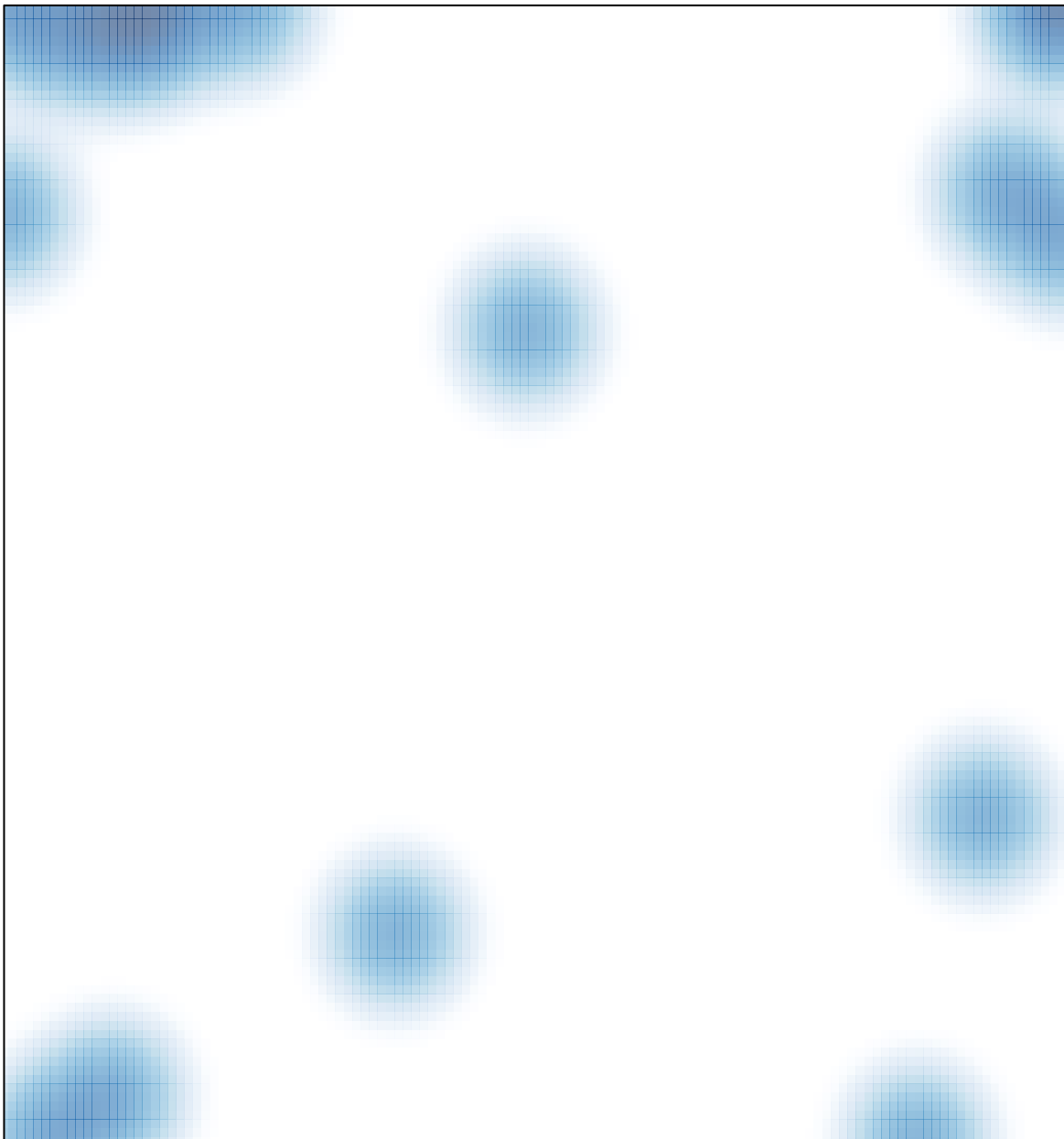


ENGELMANN_CANCER_PROGENITORS_DN



features = 16
chi-square p = 0.8

ENGELMANN_CANCER_PROGENITORS_DN



features = 16 , max = 1