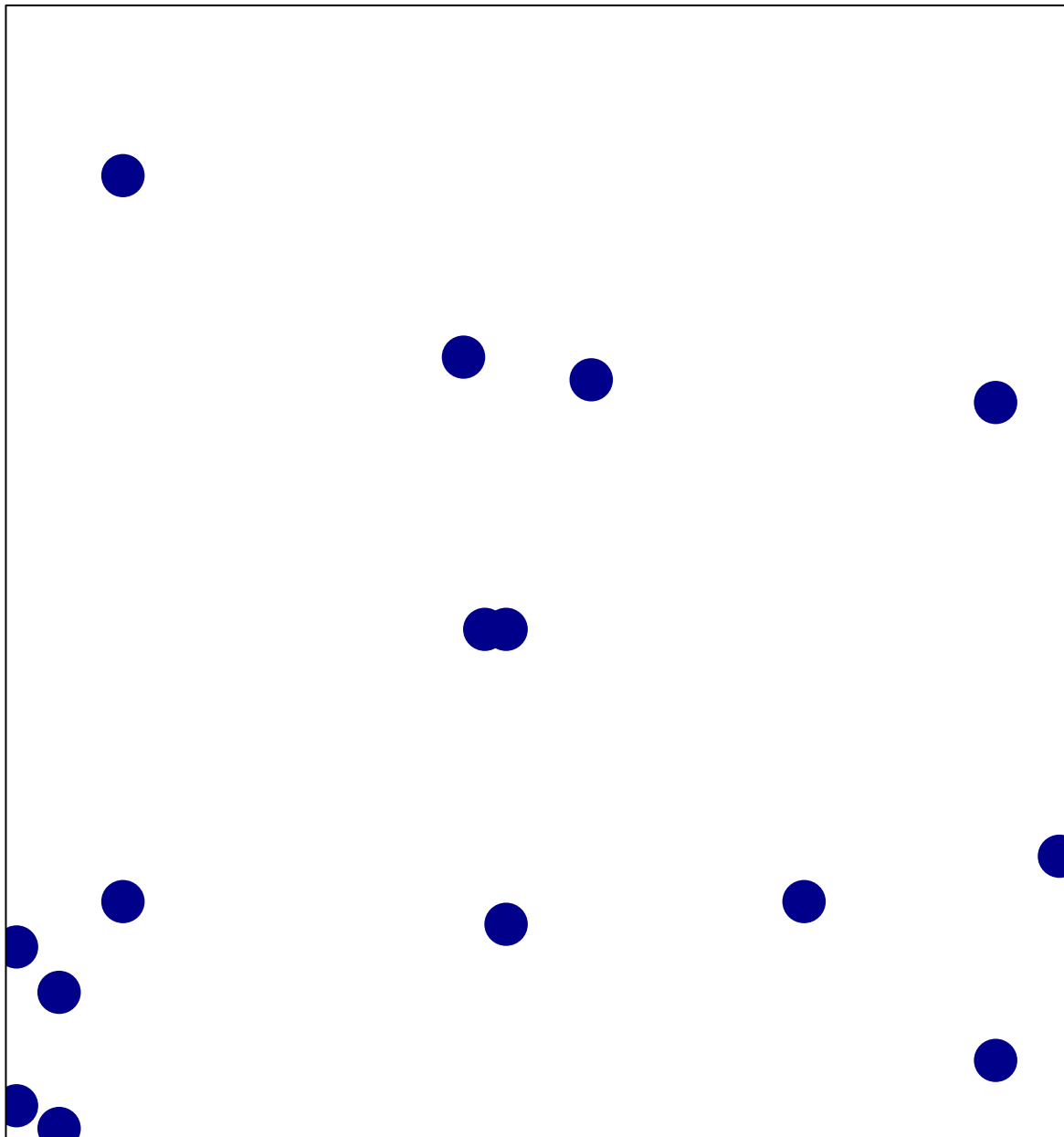


CORRE_MULTIPLE_MYELOMA_UP



features = 15
chi-square p = 0.83

CORRE_MULTIPLE_MYELOMA_UP



features = 15 , max = 1