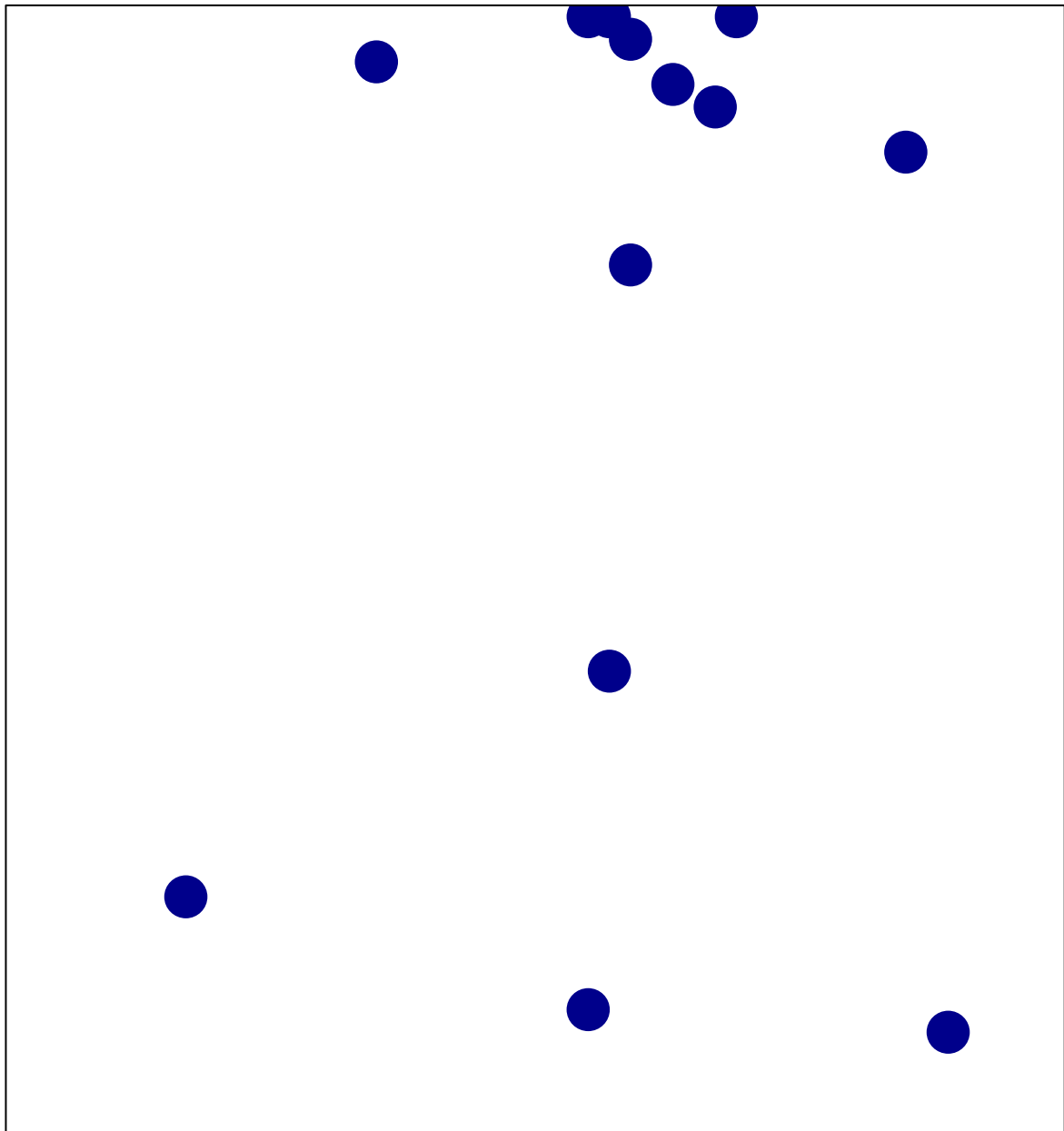


# SAKAI\_TUMOR\_INFILTRATING\_MONOCYTES\_DN



# features = 13  
chi-square p = 0.83

# SAKAI\_TUMOR\_INFILTRATING\_MONOCYTES\_DN



# features = 13 , max = 1