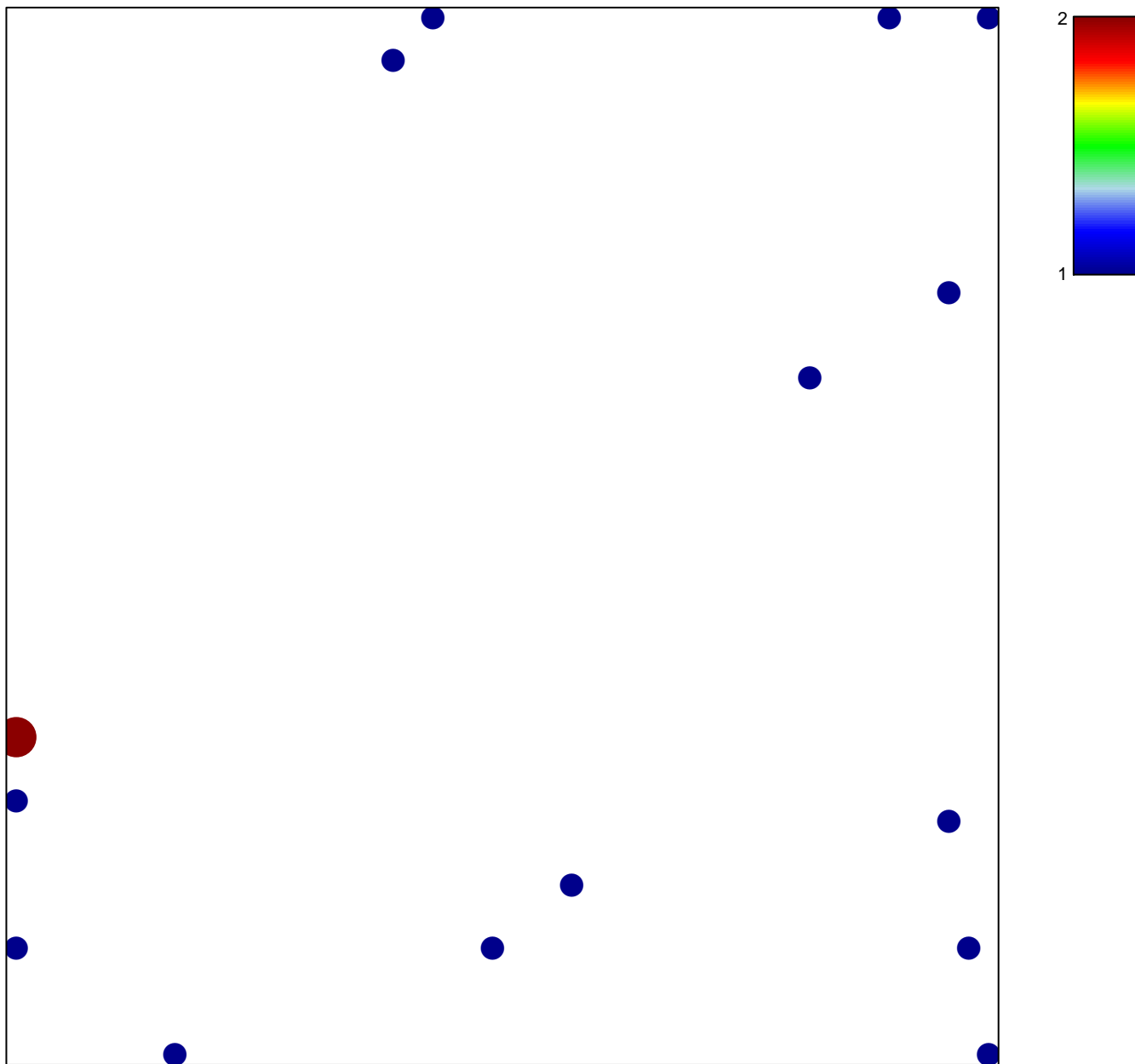


# monocyte differentiation



# features = 16  
chi-square p = 0.8

# monocyte differentiation



# features = 16 , max = 2