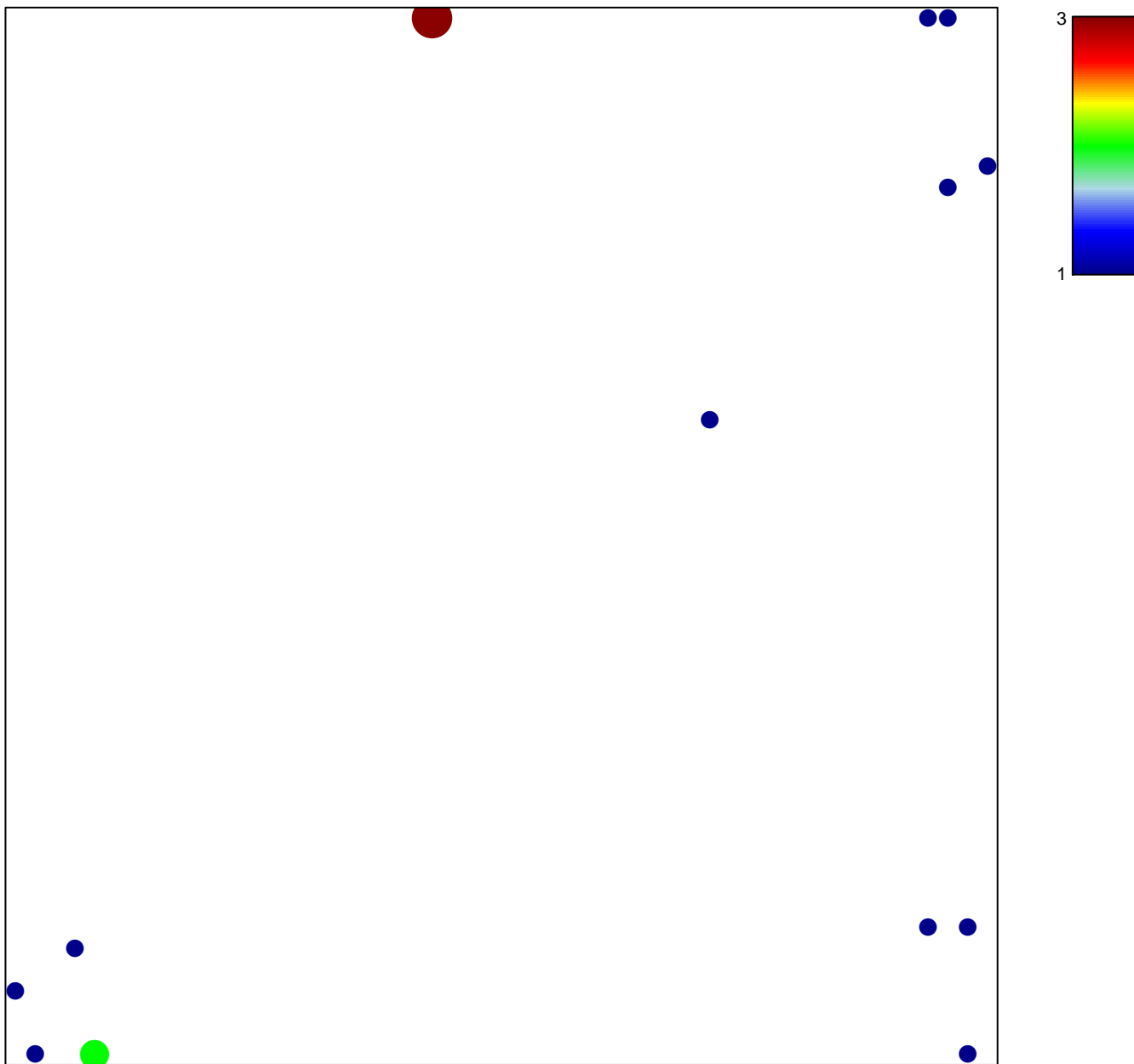


# WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_DN



# features = 16  
chi-square p = 0.77

# WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_DN



# features = 16 , max = 3