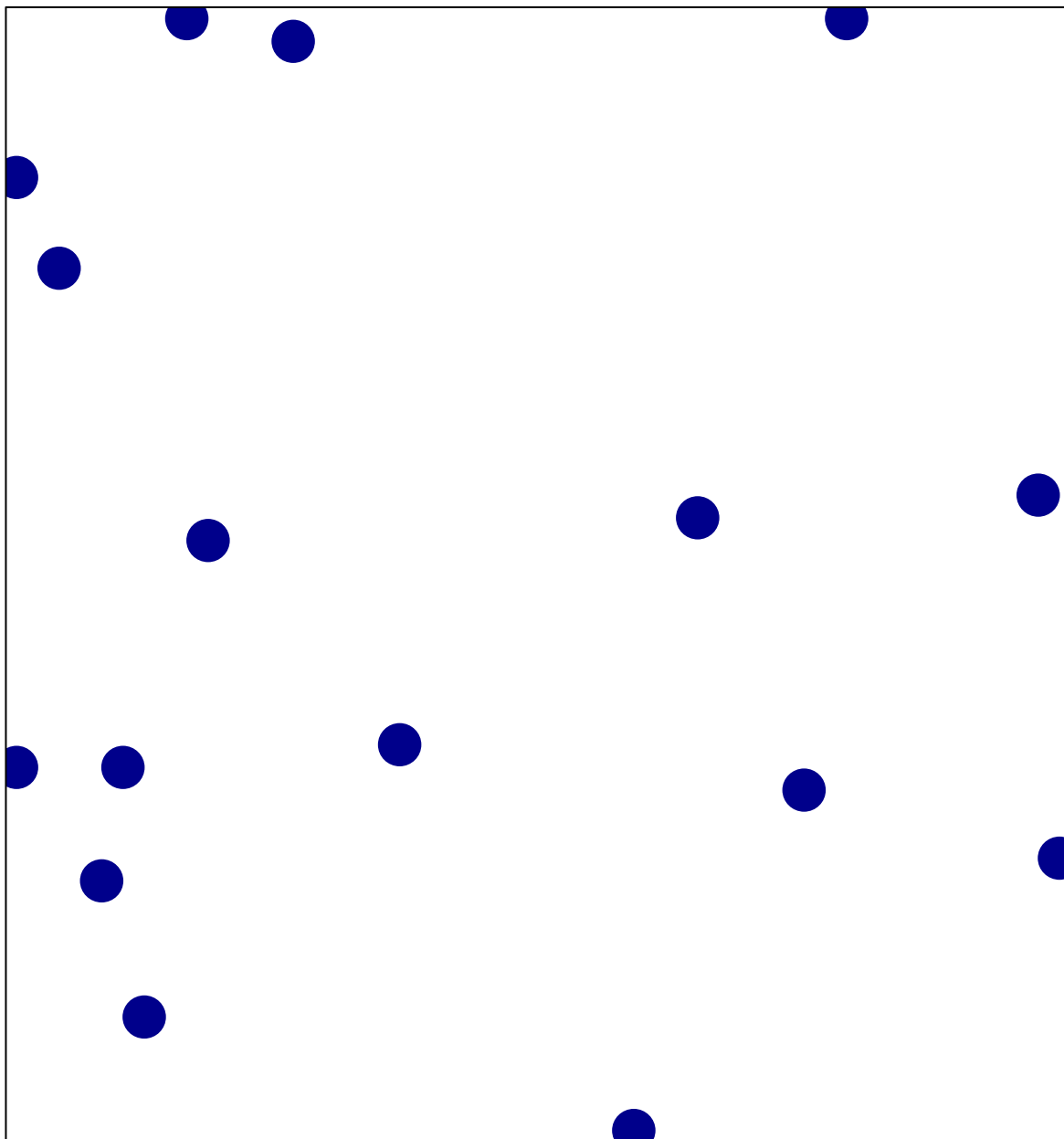
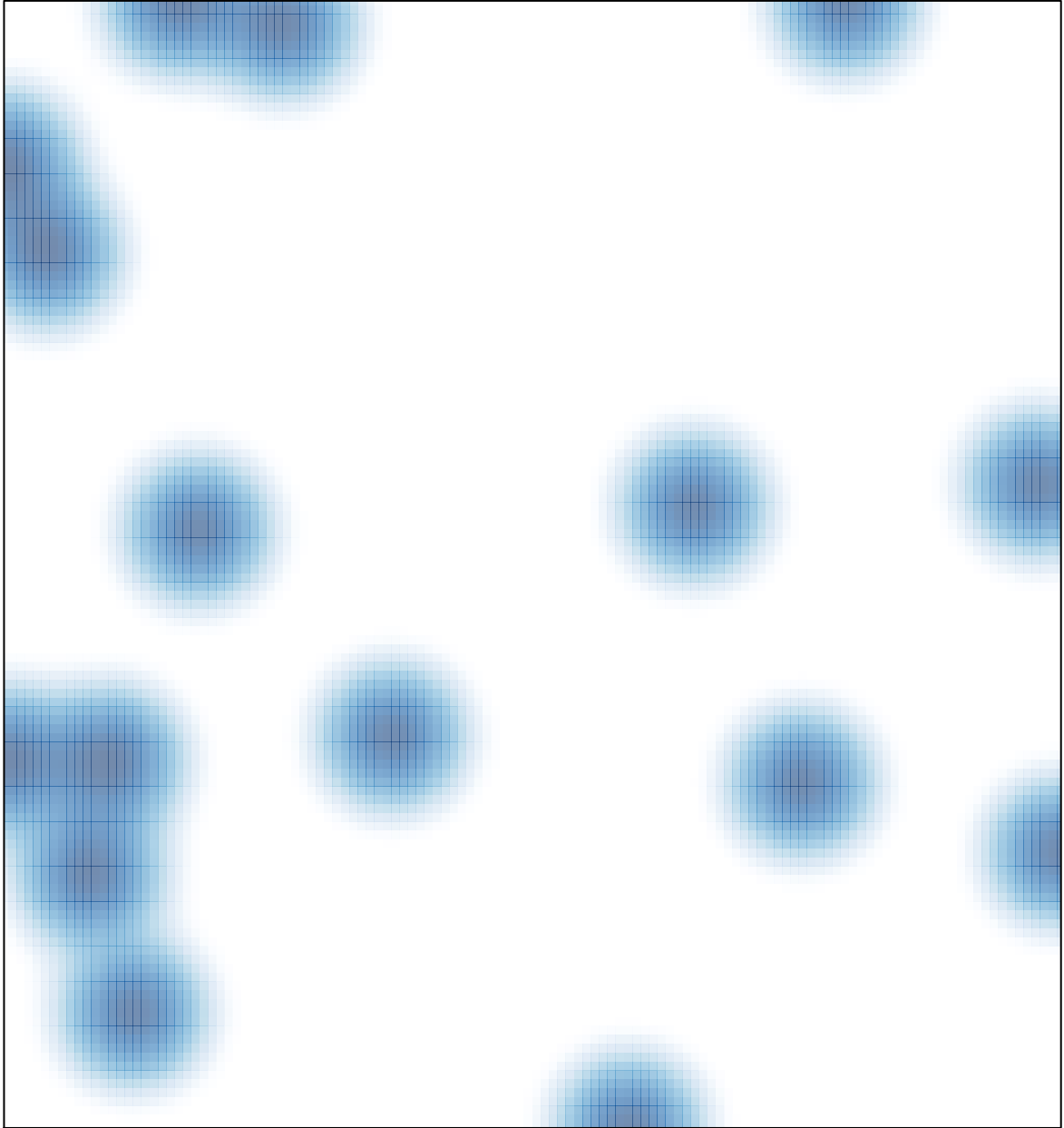


# WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_UP



# features = 16  
chi-square p = 0.85

# WATTEL\_AUTONOMOUS\_THYROID\_ADENOMA\_UP



# features = 16 , max = 1