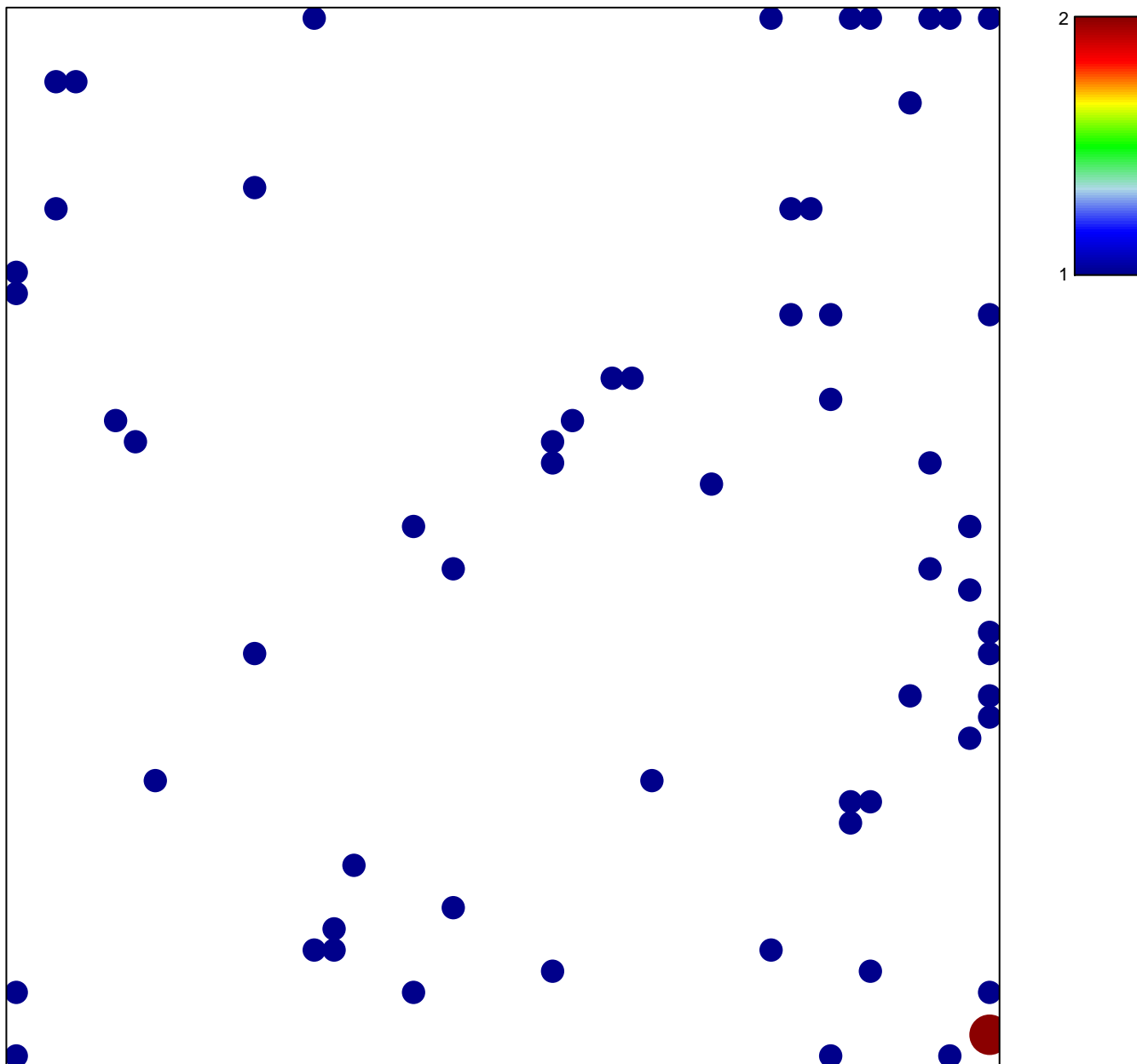
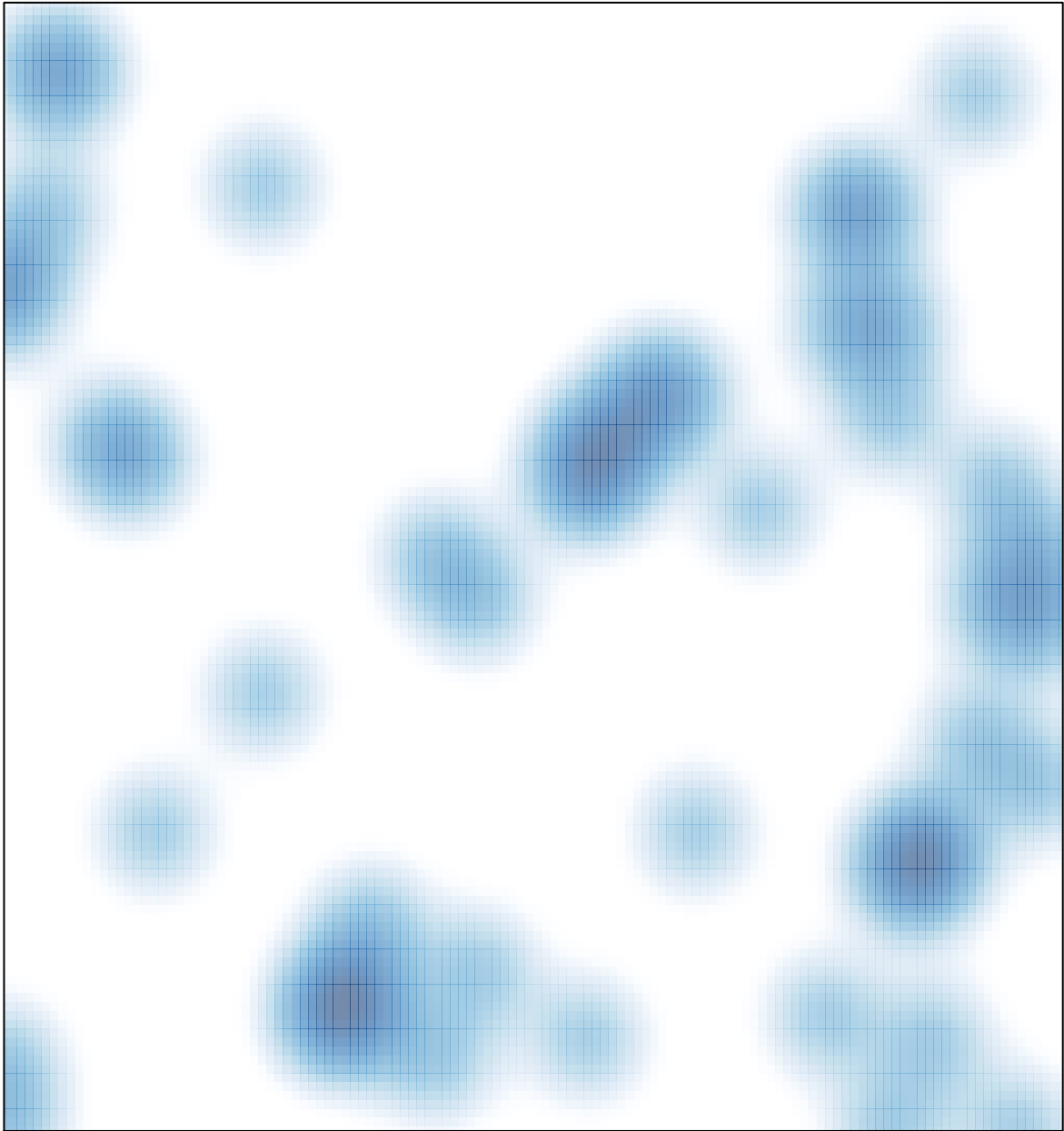


# JECHLINGER\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSITION\_UP



# features = 62  
chi-square p = 0.8

# INGER\_EPITHELIAL\_TO\_MESENCHYMAL\_TRANSIT



# features = 62 , max = 2