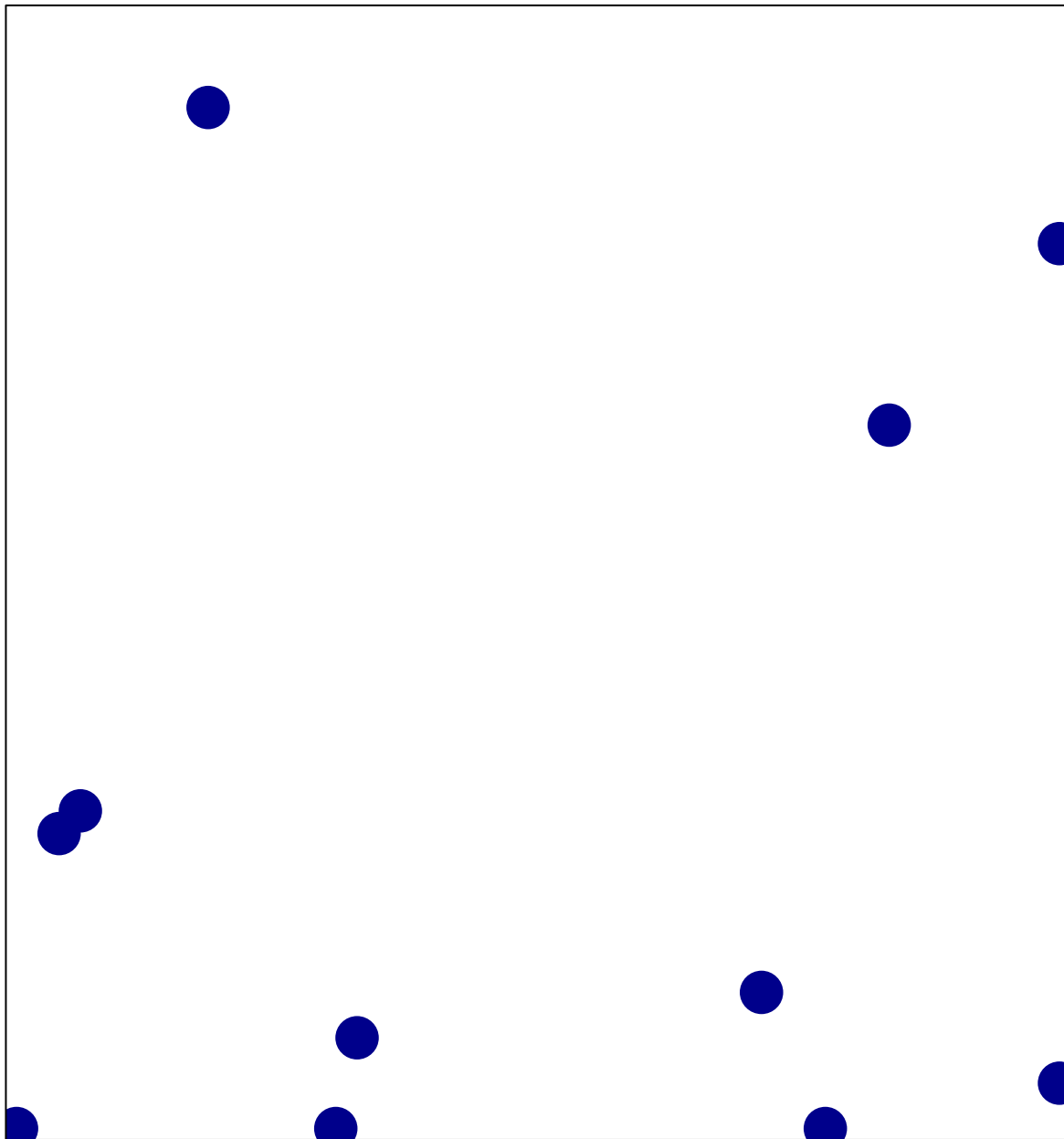
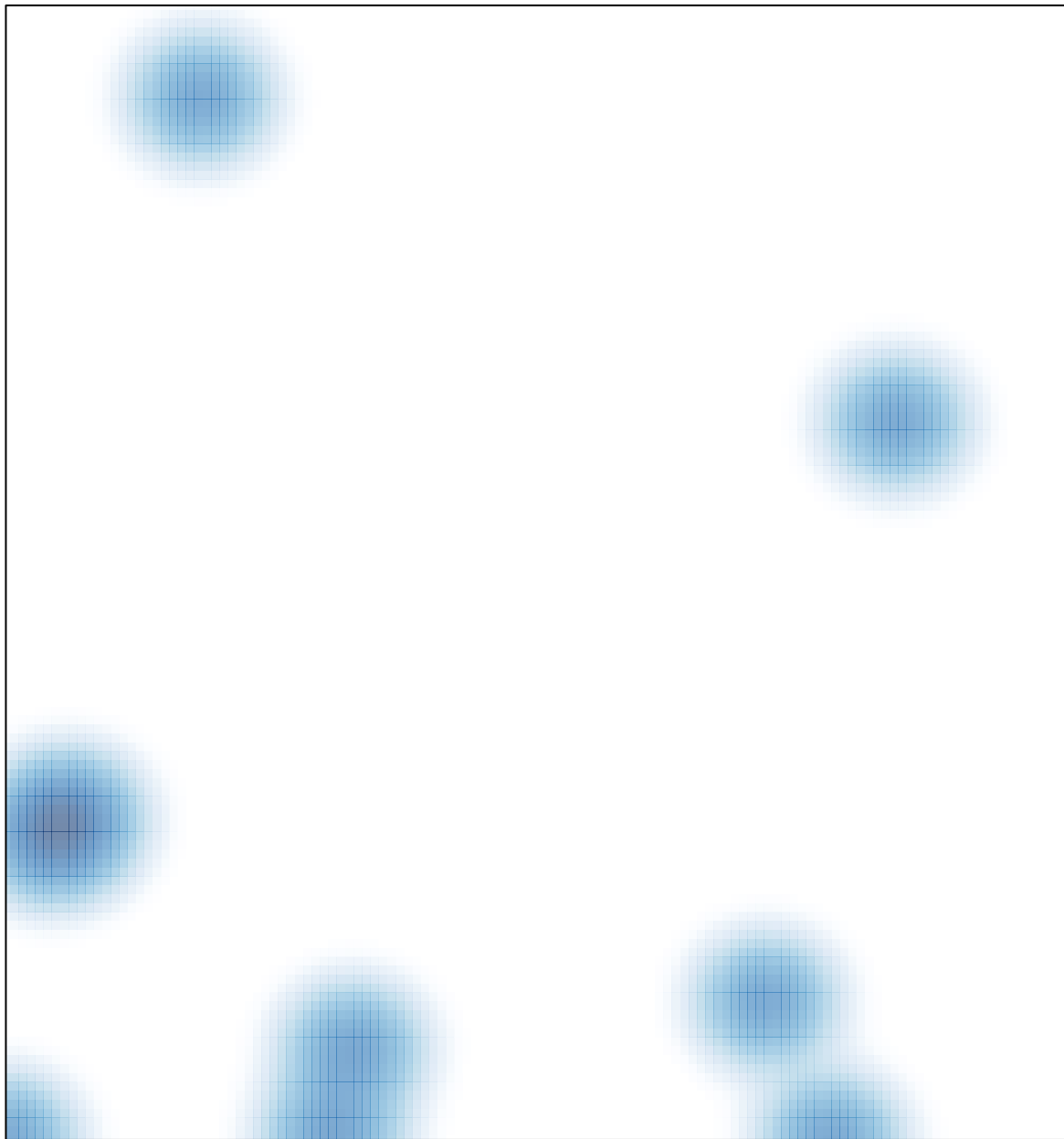


# WILLSCHER\_GBM\_LTSwt\_proteomics-C\_UP



# features = 11  
chi-square p = 0.83

# WILLSCHER\_GBM\_LTSwt\_proteomics-C\_UP



# features = 11 , max = 1