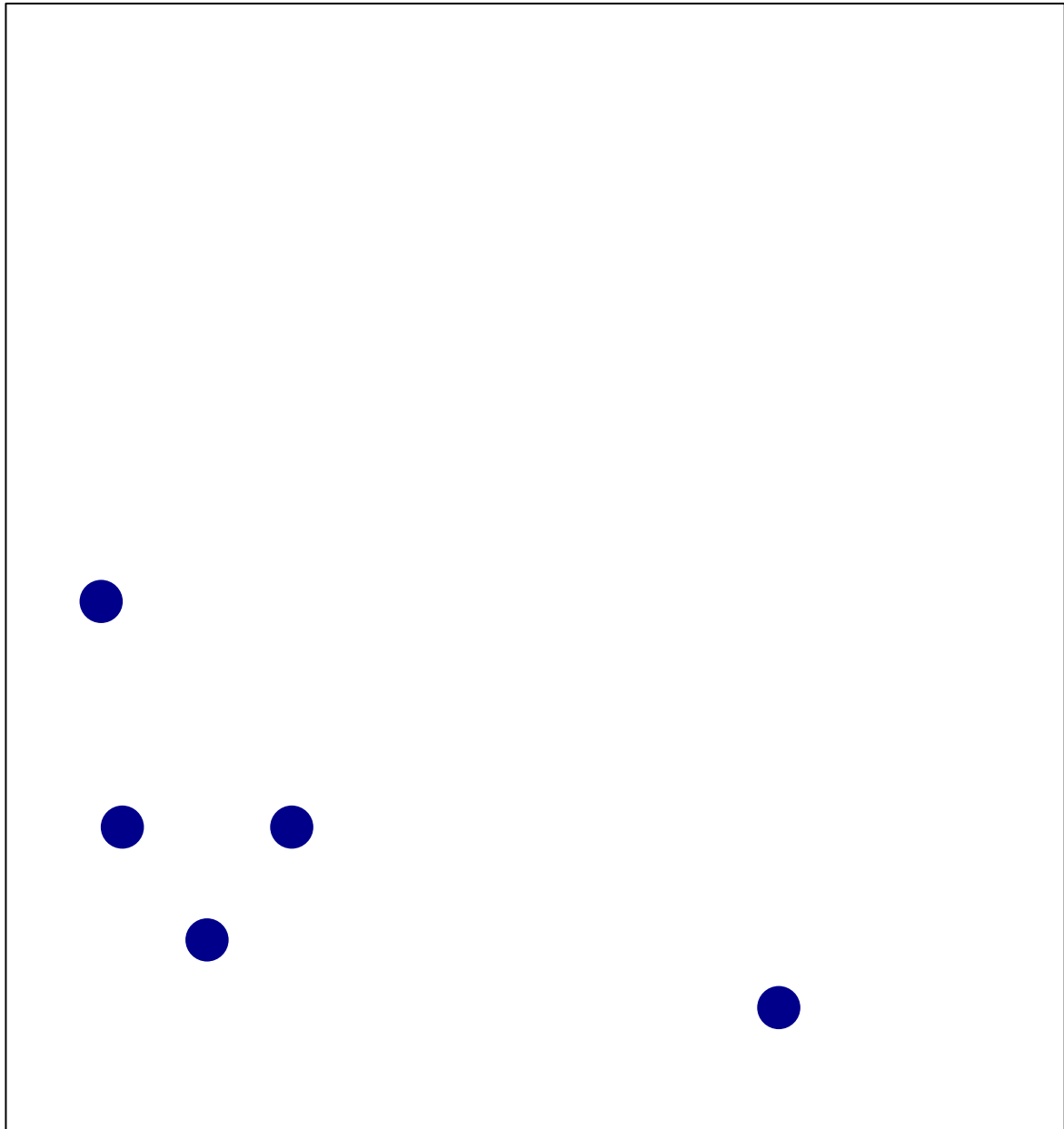
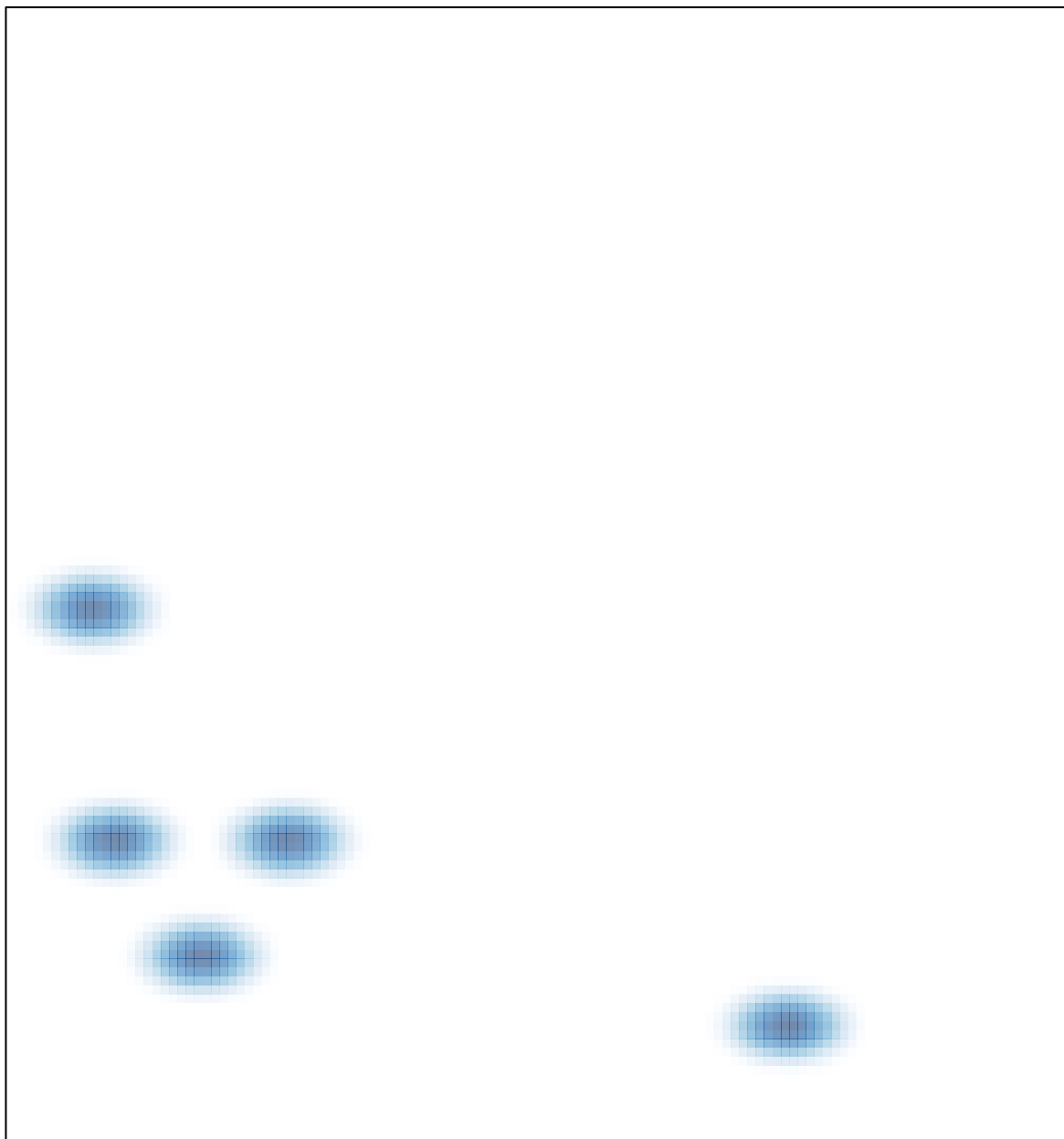


# WILLSCHER\_GBM\_STSwt\_proteomics-N\_UP



# features = 5  
chi-square p = 0.83

# WILLSCHER\_GBM\_STSwt\_proteomics-N\_UP



# features = 5 , max = 1