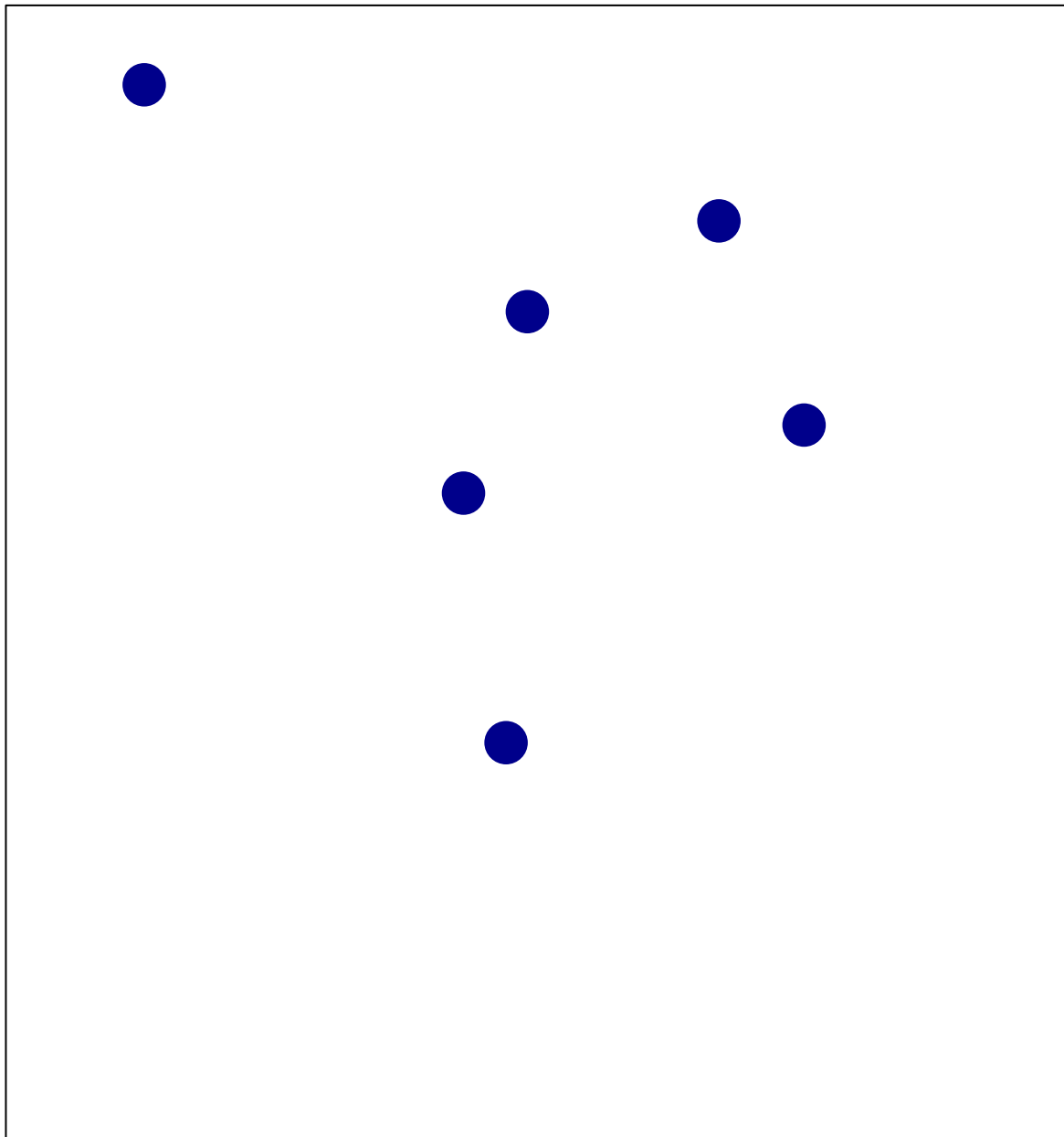
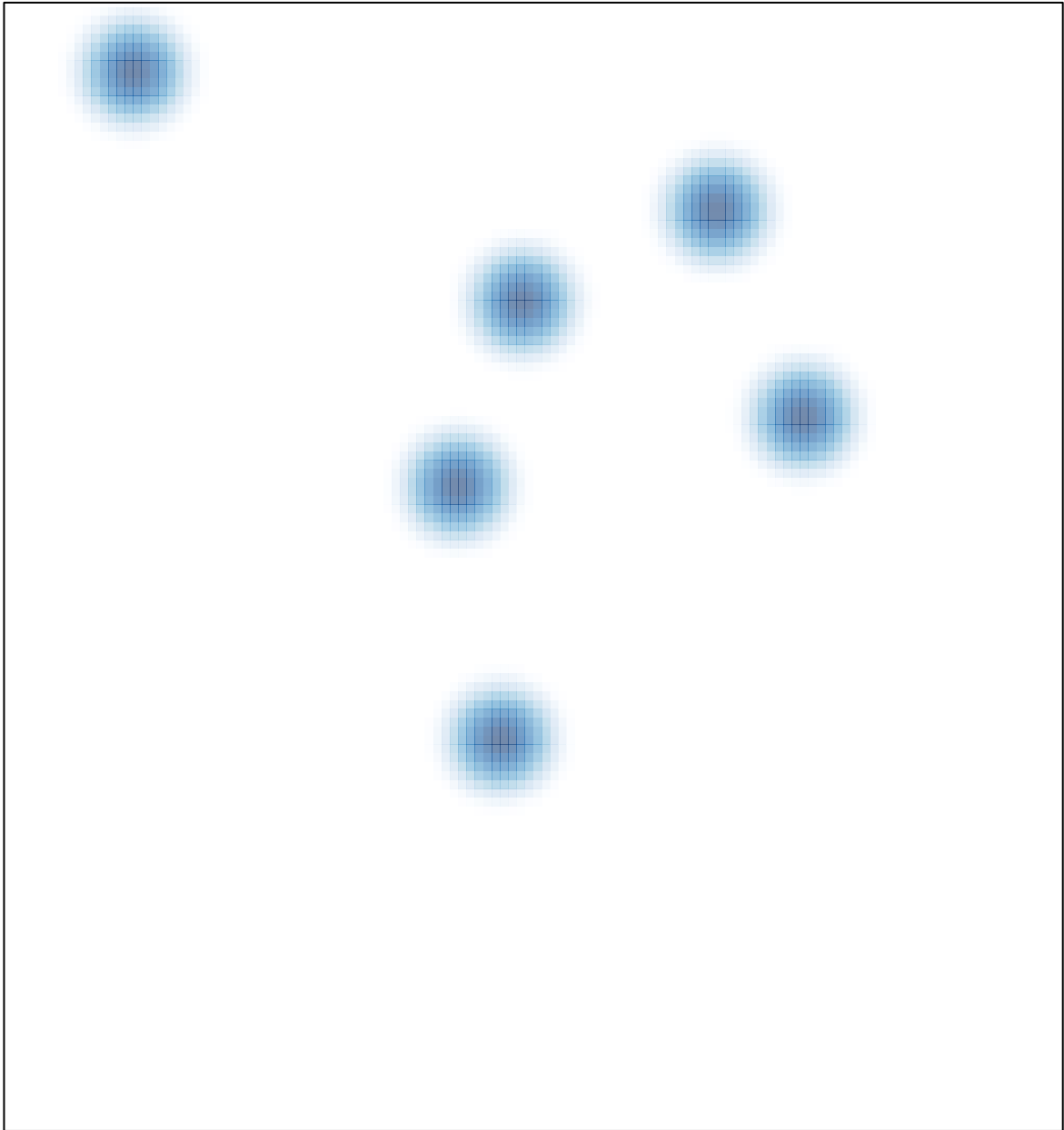


# KINNEY\_DNMT1\_METHYLATION\_TARGETS



# features = 6  
chi-square p = 0.85

# KINNEY\_DNMT1\_METHYLATION\_TARGETS



# features = 6 , max = 1