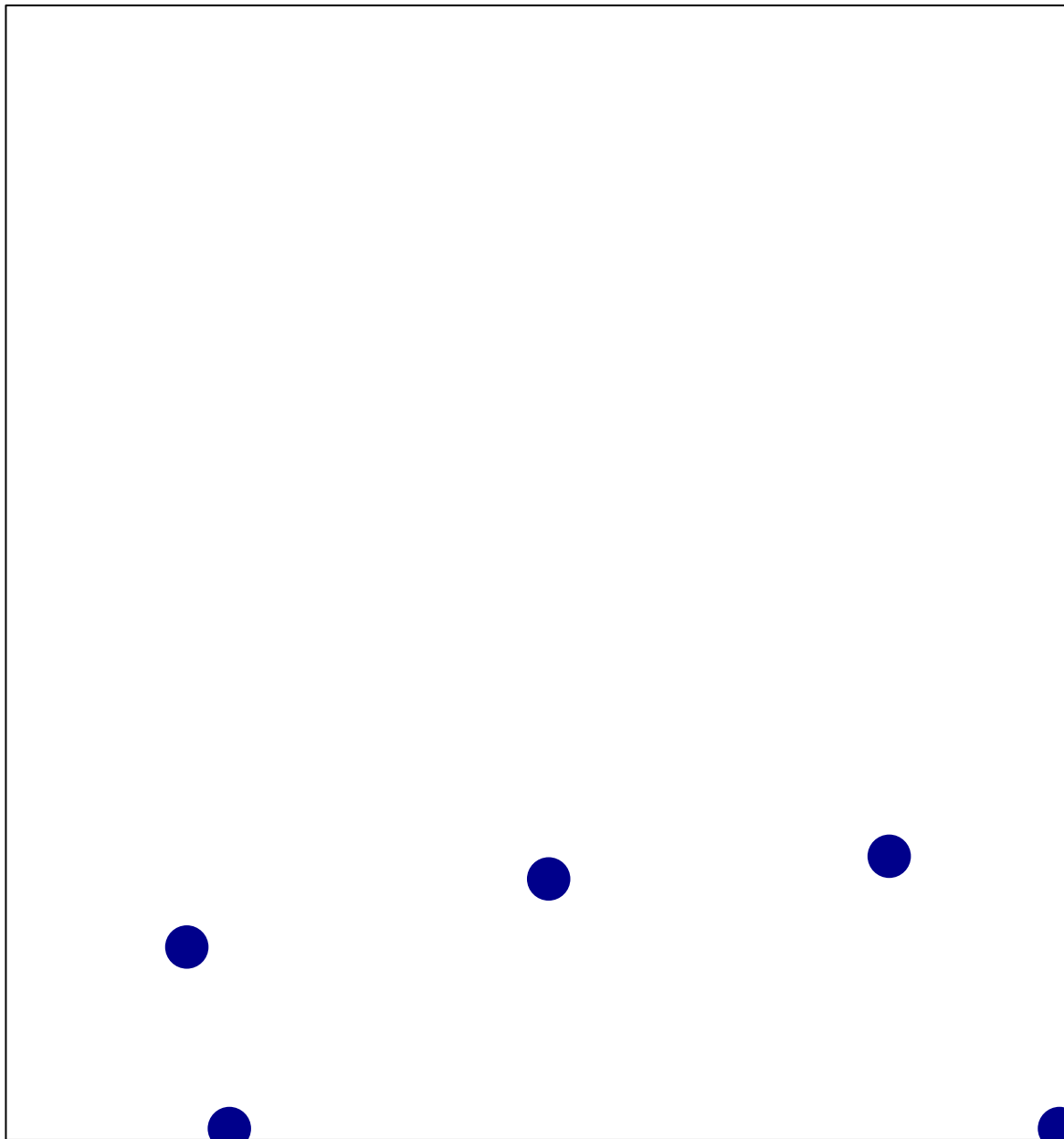
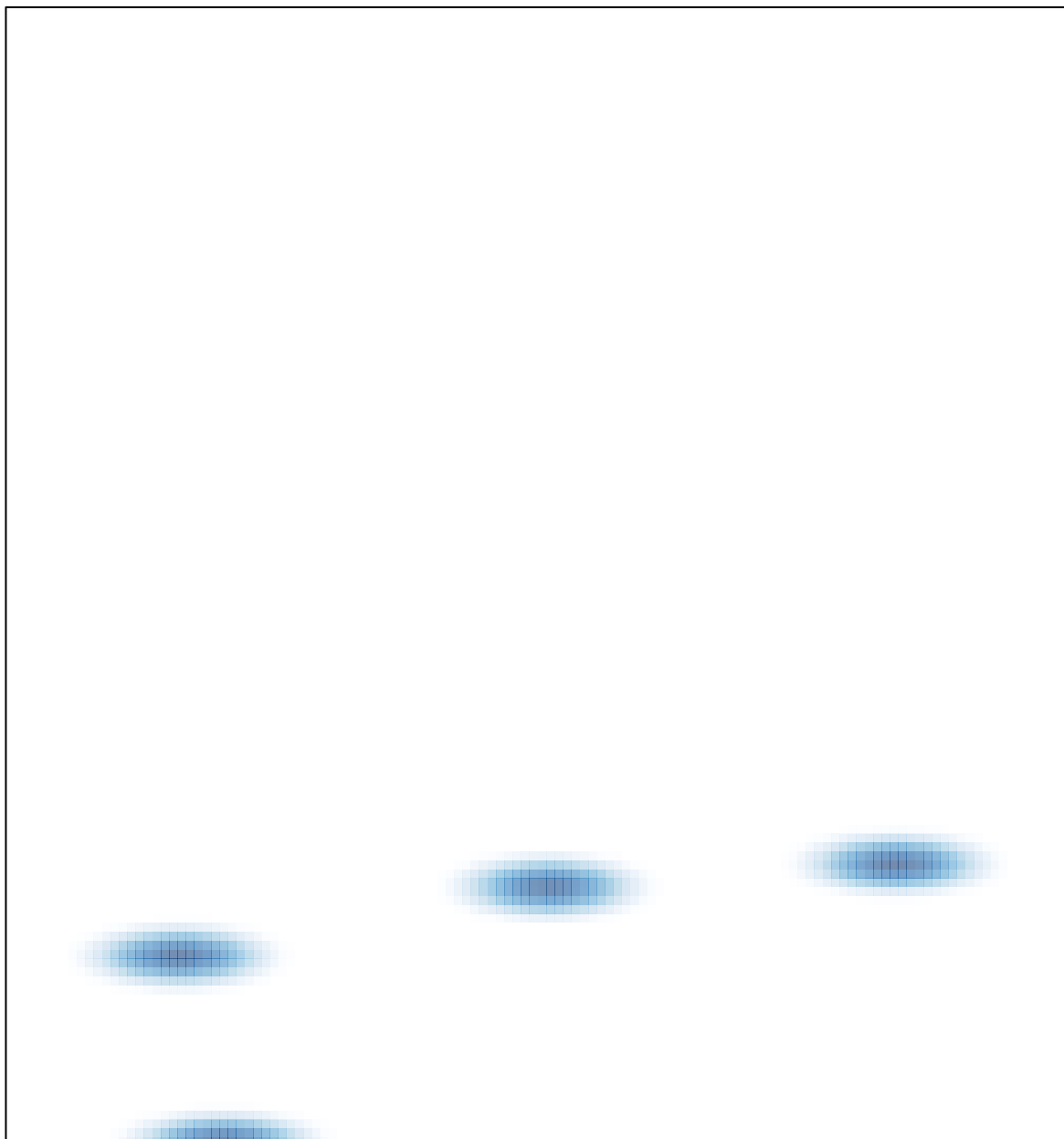


# NOUSHMEHR\_GBM\_GERMLINE\_MUTATED



# features = 5  
chi-square p = 0.83

# NOUSHMEHR\_GBM\_GERMLINE\_MUTATED



# features = 5 , max = 1