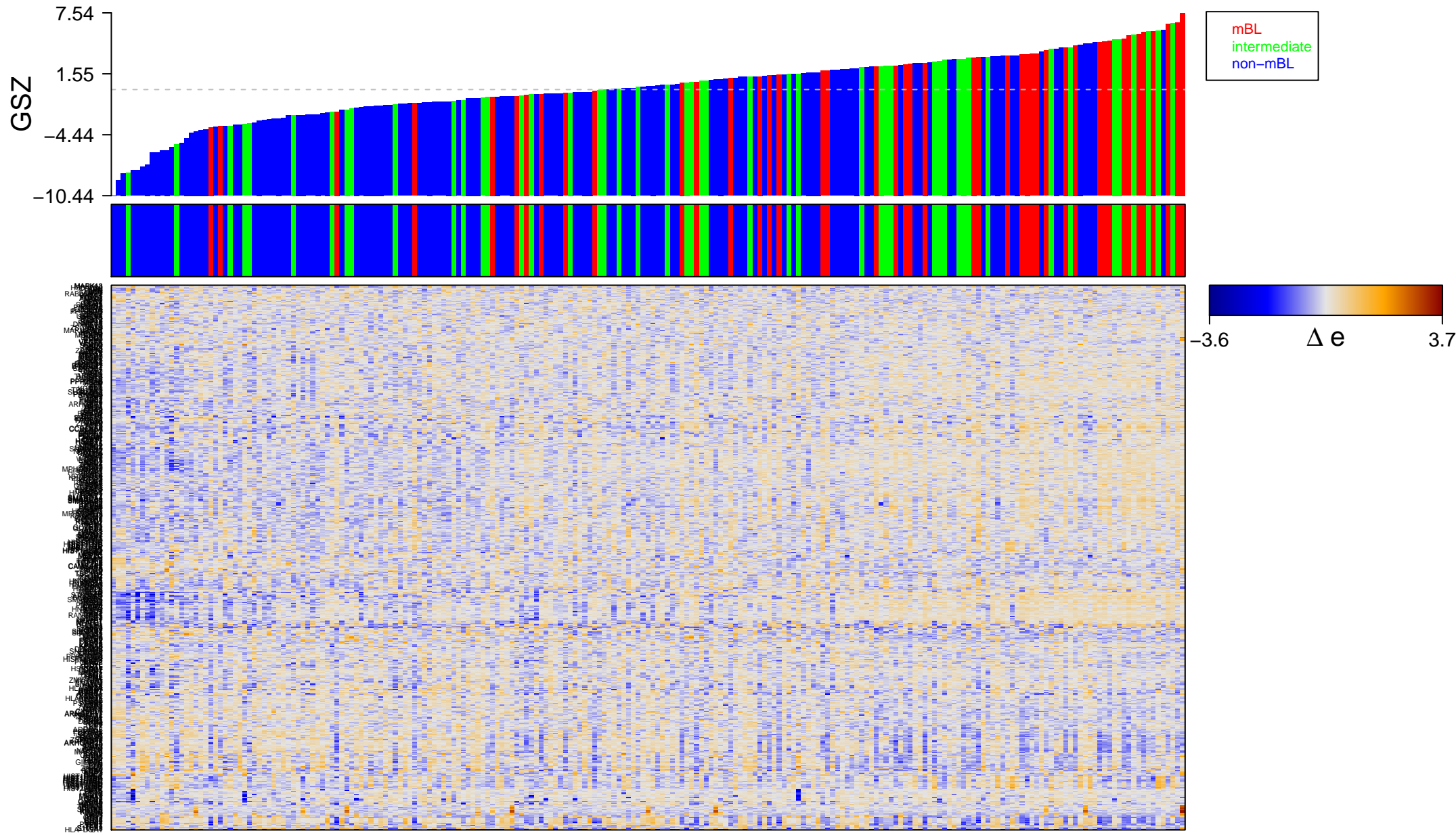
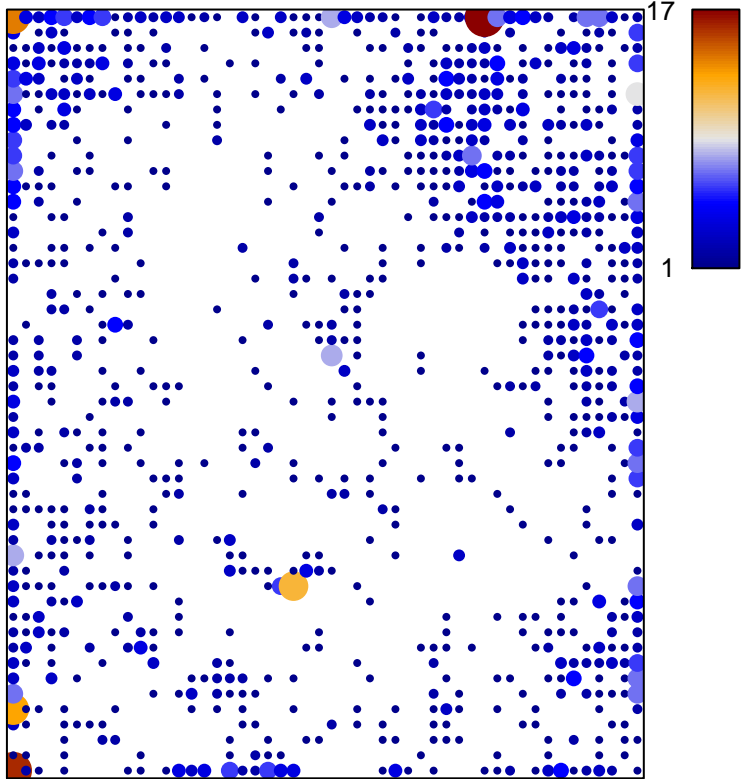
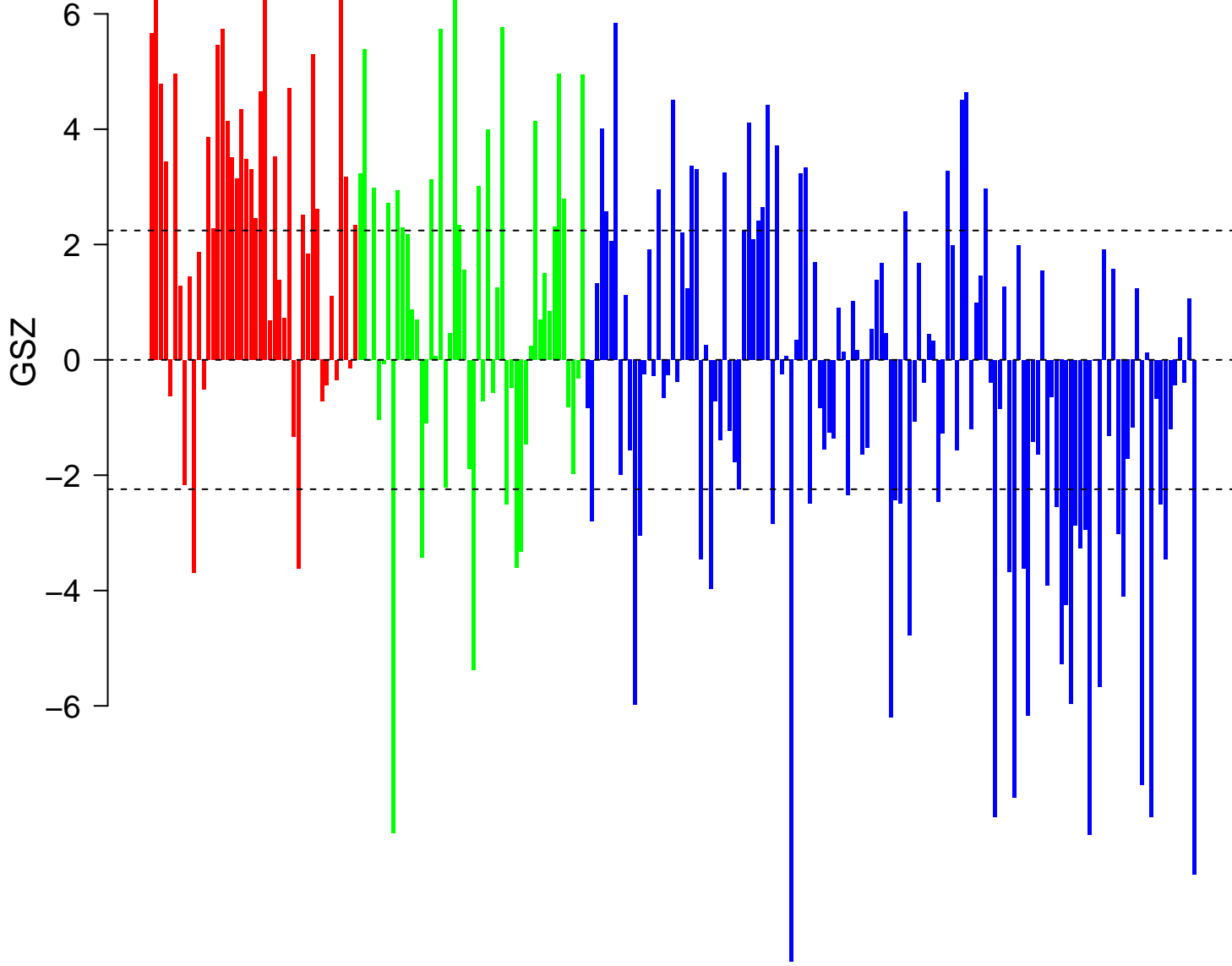


# PROTEINATLAS\_tonsil



# PROTEINATLAS\_tonsil



# features = 1755



# PROTEINATLAS\_tonsil

