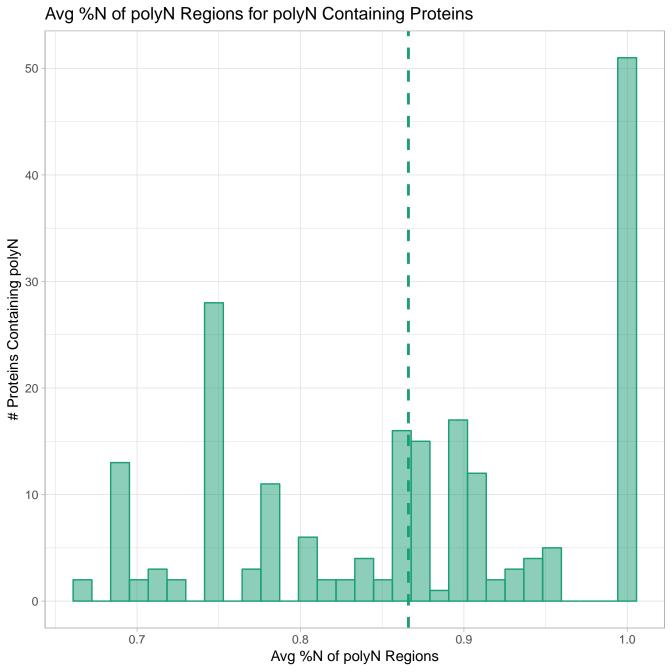
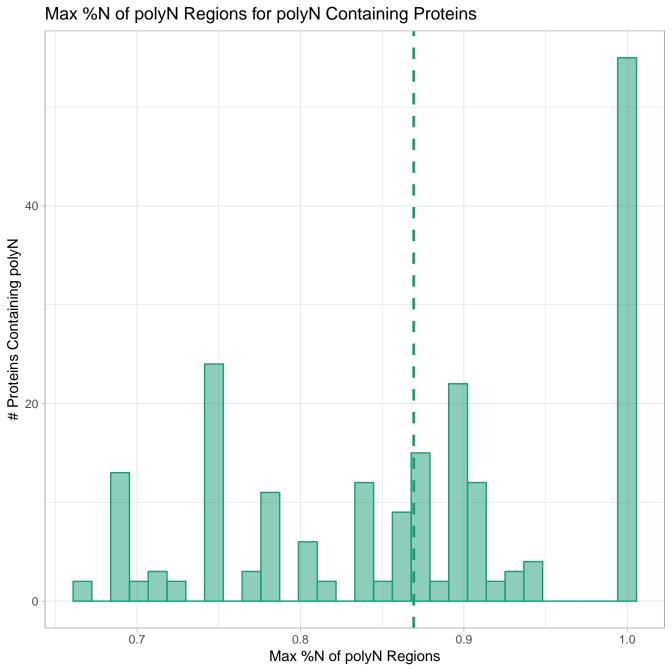
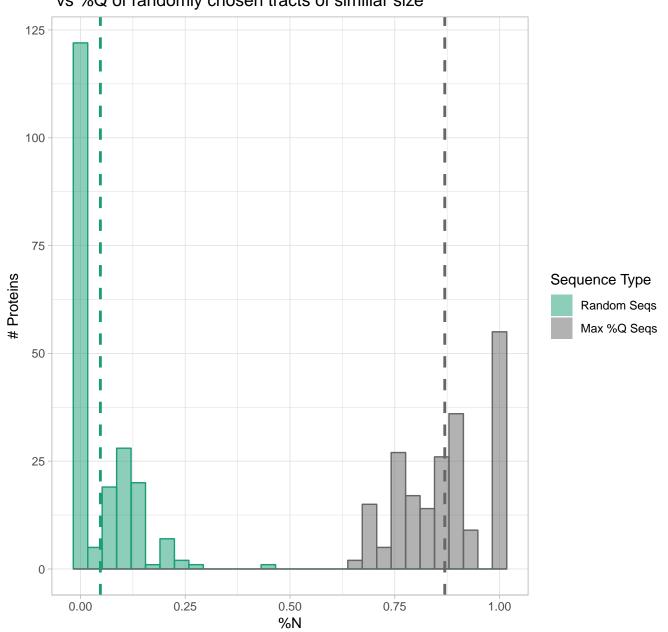


%N for 100000 50AA Random Seqs Histogram 30000 20000 # of Randomly Choosen Pepitides 10000 0 0.2 0.4 0.0 0.6 %N (Random 50AA Seqs)

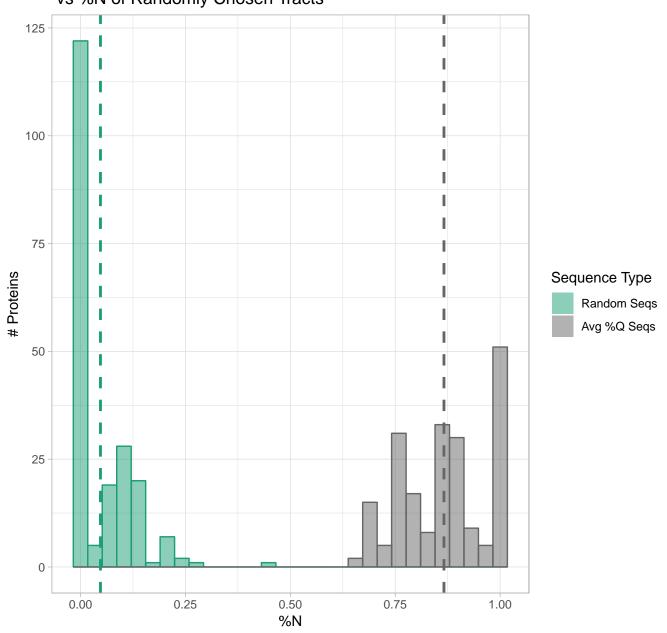




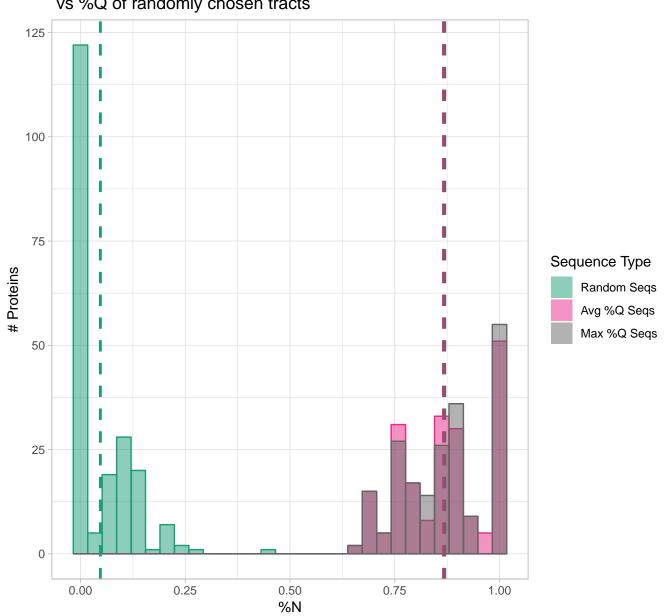
Max %N of polyN Regions for polyN Containing Proteins vs %Q of randomly chosen tracts of similiar size

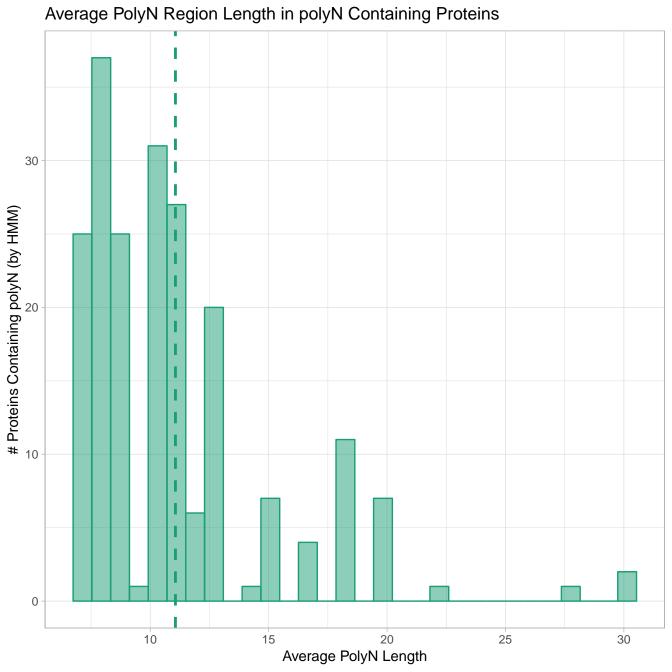


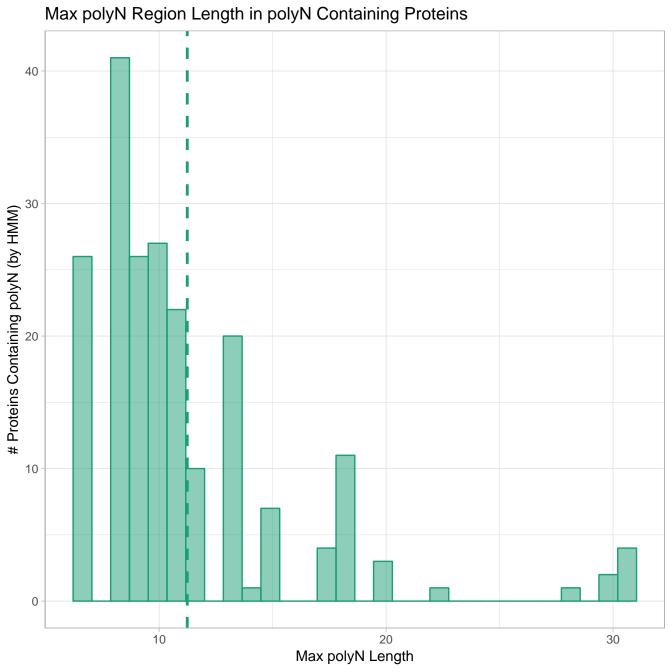
Avg %N of PolyN Regions for polyN Containing Proteins vs %N of Randomly Chosen Tracts



Avg %N of polyN Regions vs Max %N of polyN Regions vs %Q of randomly chosen tracts







Number of polyN Regions in polyN Containing Proteins 150 # of N Proteins 100 50 0 0.5 1.5 1.0 2.0 2.5 Number of polyN Regions