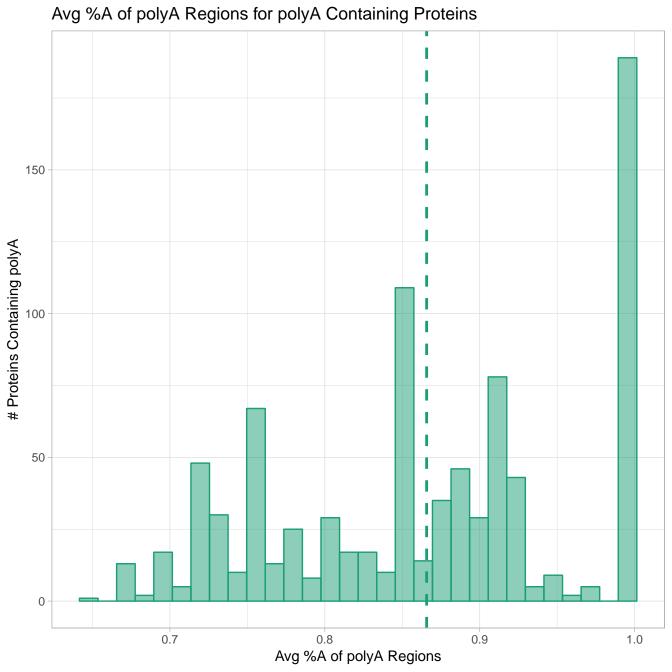
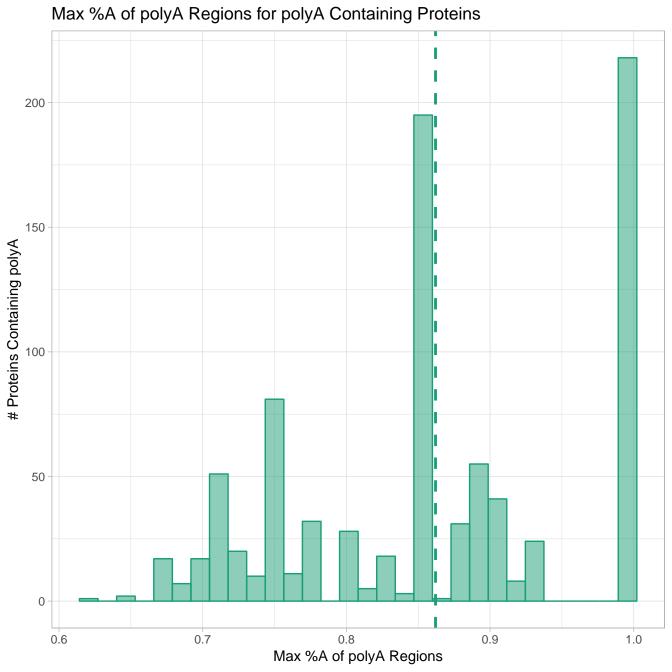
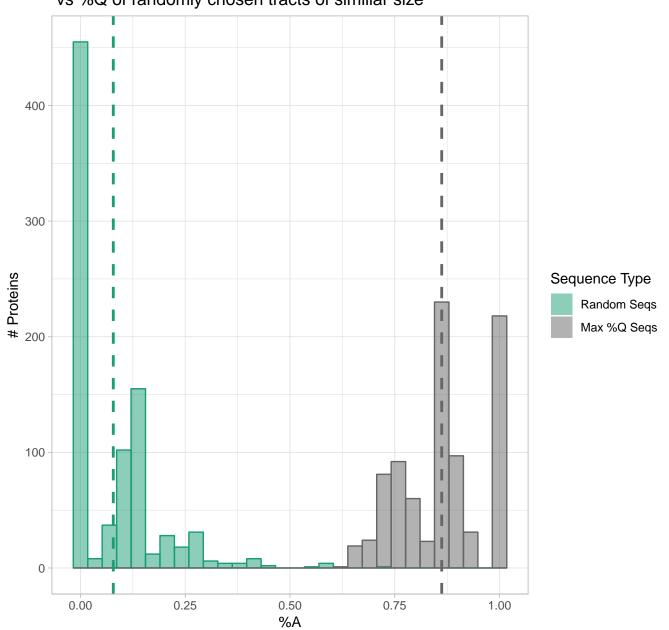


%A for 100000 50AA Random Seqs Histogram 20000 15000 # of Randomly Choosen Pepitides 10000 5000 0 0.0 0.2 0.4 0.6 %A (Random 50AA Seqs)

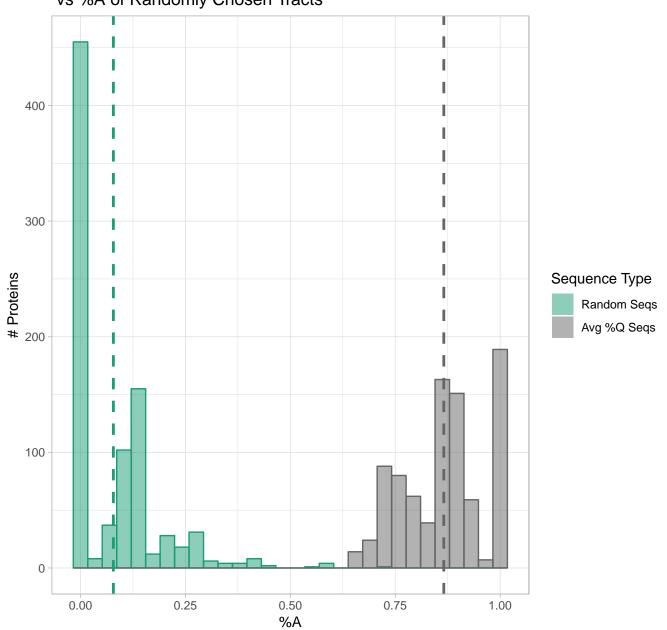




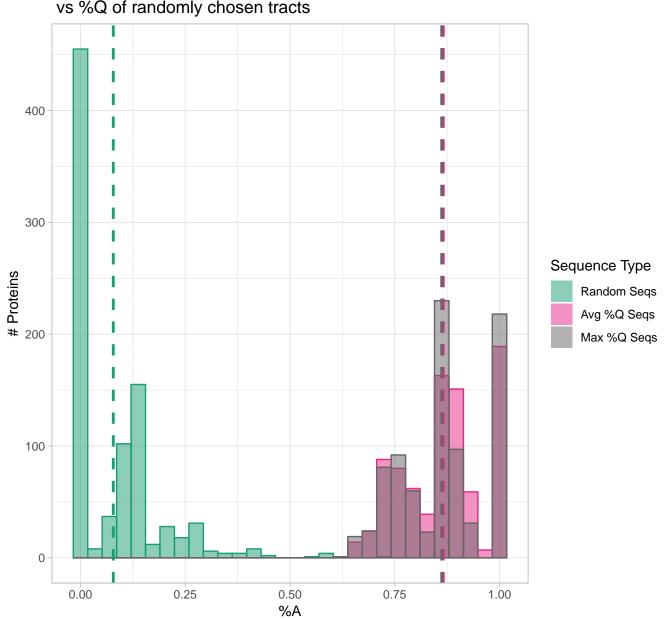
Max %A of polyA Regions for polyA Containing Proteins vs %Q of randomly chosen tracts of similiar size

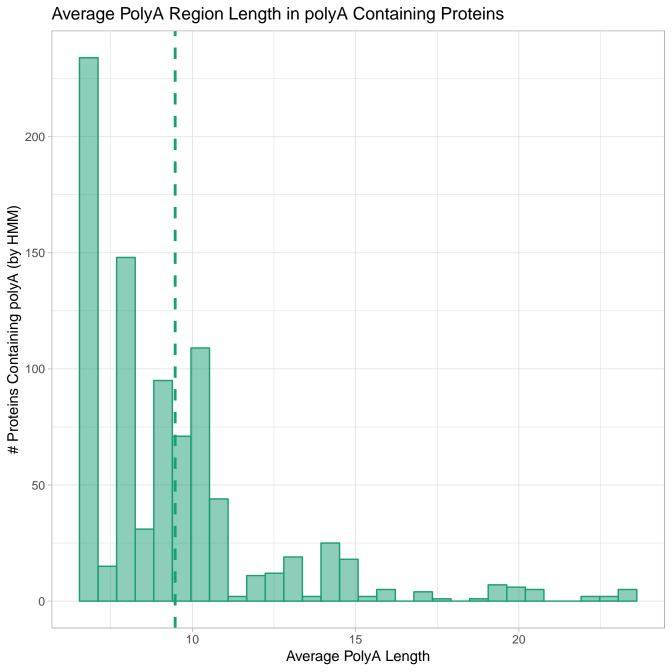


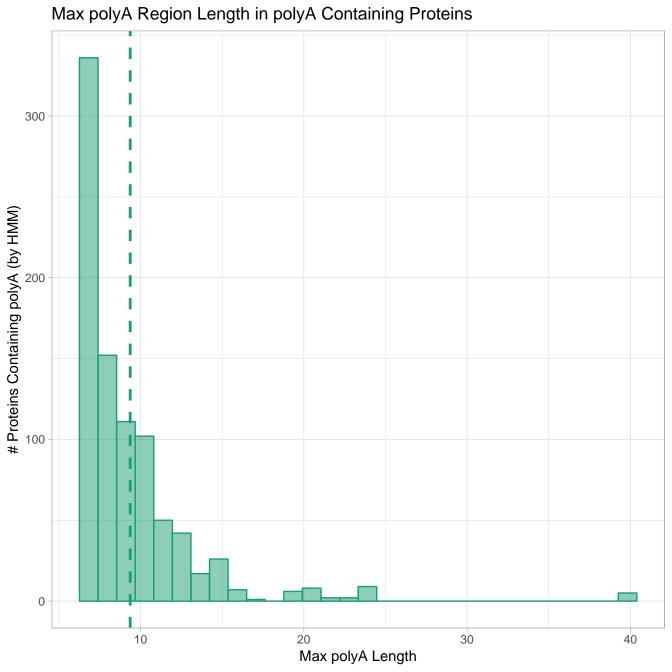
Avg %A of PolyA Regions for polyA Containing Proteins vs %A of Randomly Chosen Tracts



Avg %A of polyA Regions vs Max %A of polyA Regions vs %Q of randomly chosen tracts







Number of polyA Regions in polyA Containing Proteins # of A Proteins Number of polyA Regions