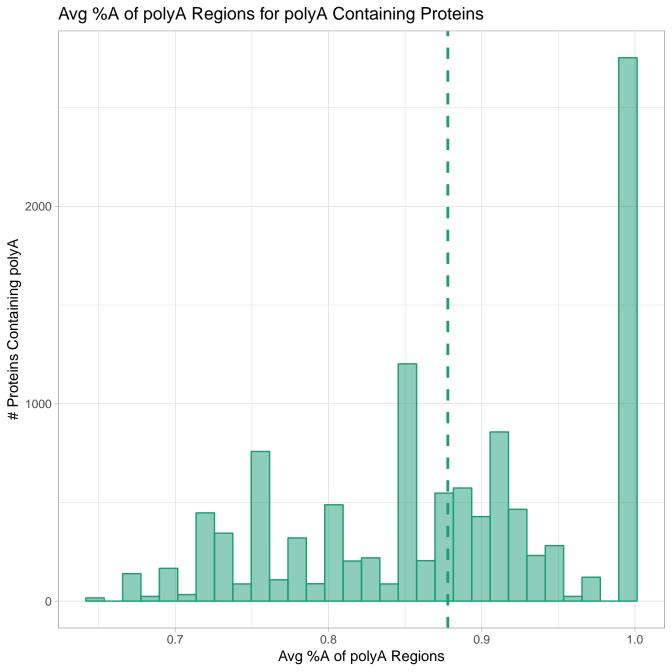
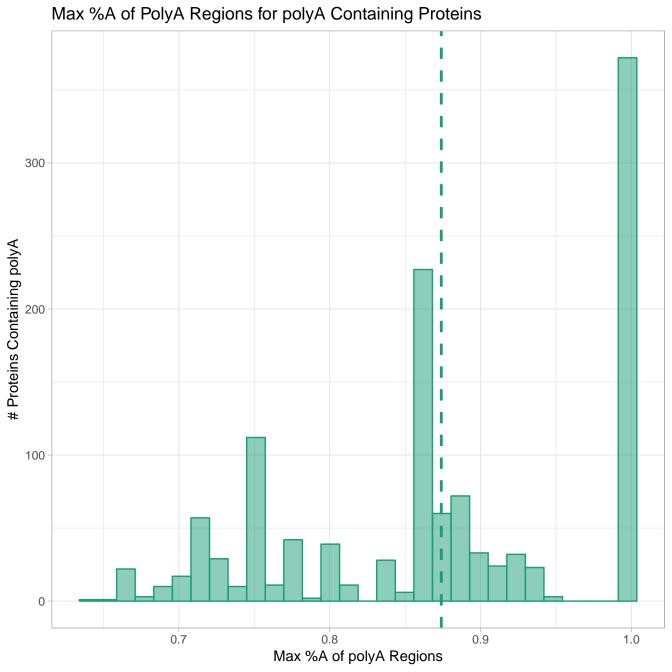
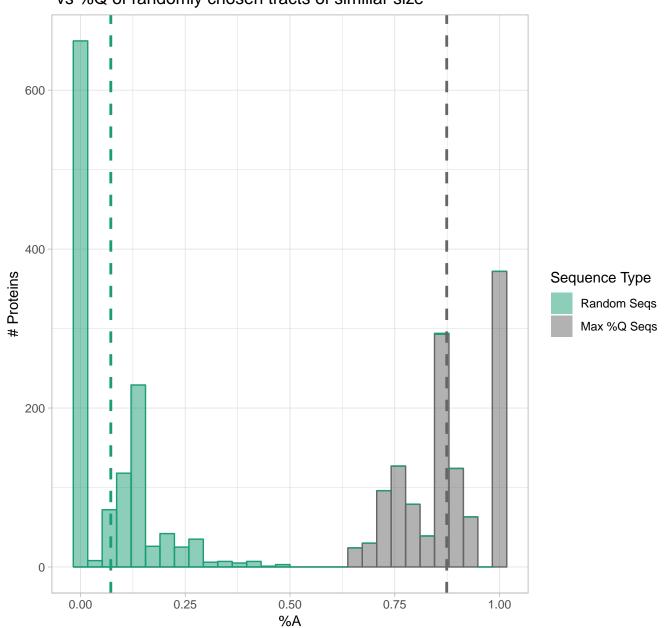


%A for 100000 50AA Random Seqs Histogram 20000 15000 # of Randomly Choosen Pepitides 10000 5000 0 0.2 0.0 0.4 %A (Random 50AA Seqs)





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 600 400 Sequence Type Random Seqs Avg %Q Seqs 200

Proteins

0

0.00

0.25

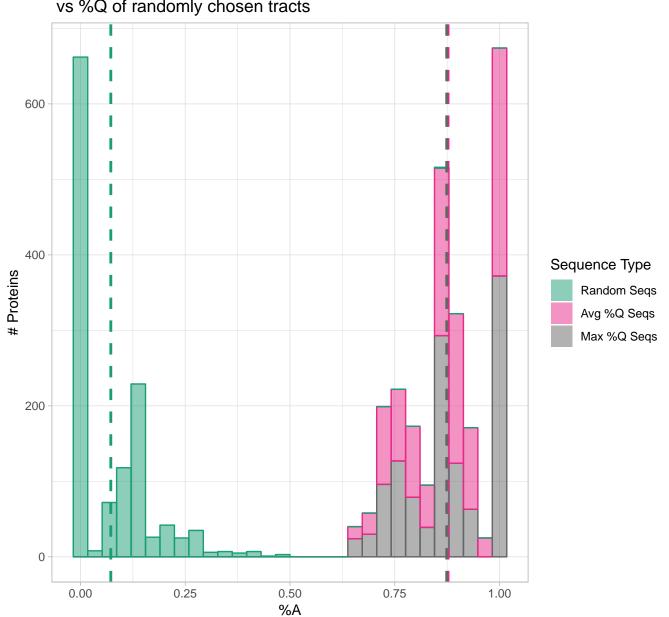
0.50

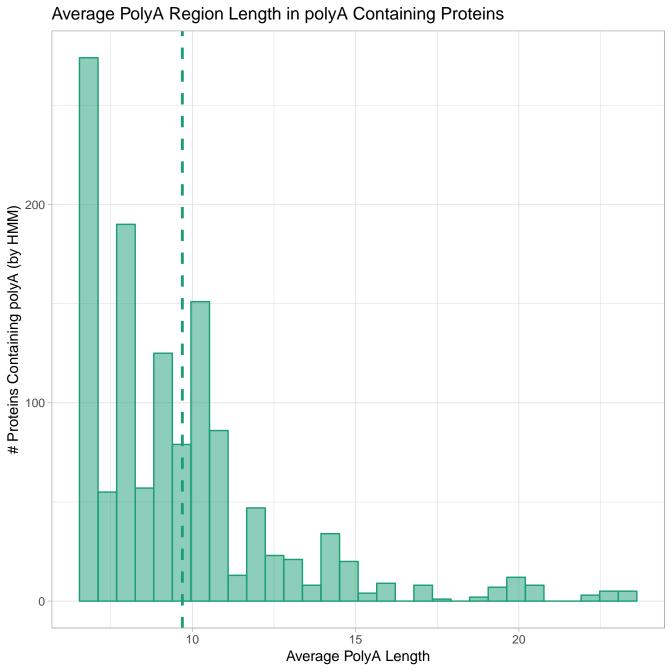
%A

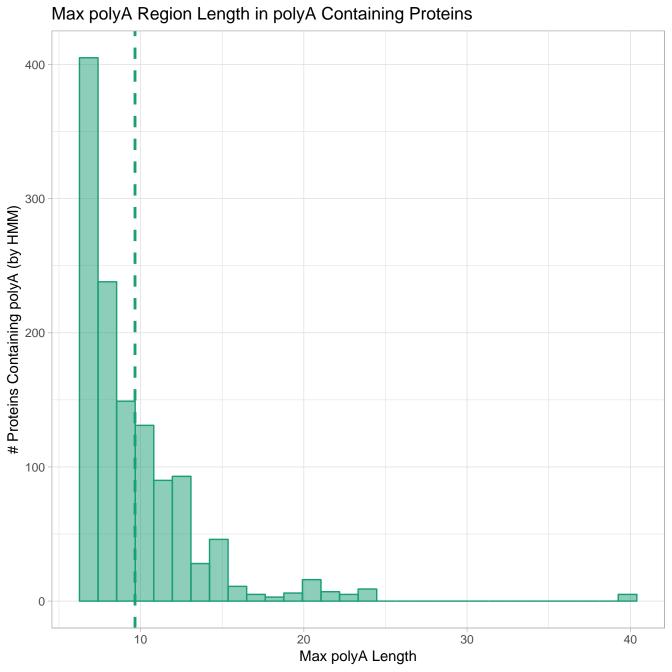
0.75

1.00

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 800 600 # of A Proteins 200 0 2.5 5.0 7.5 Number of polyA Regions