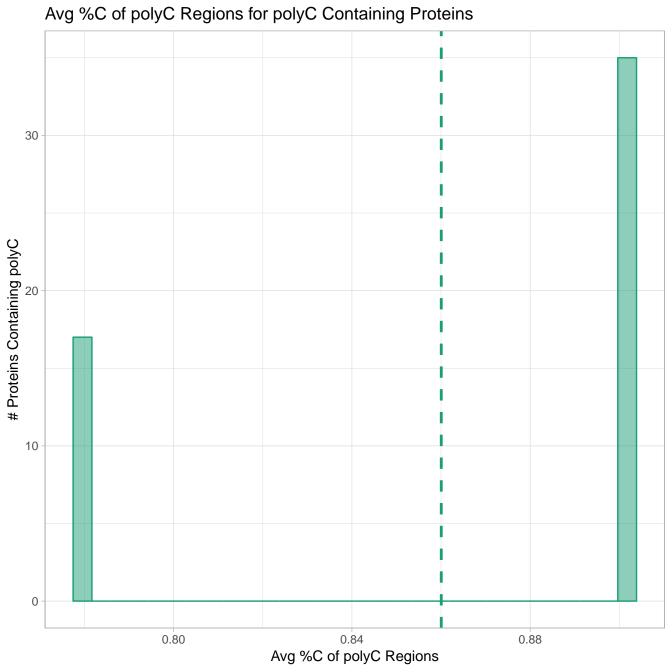
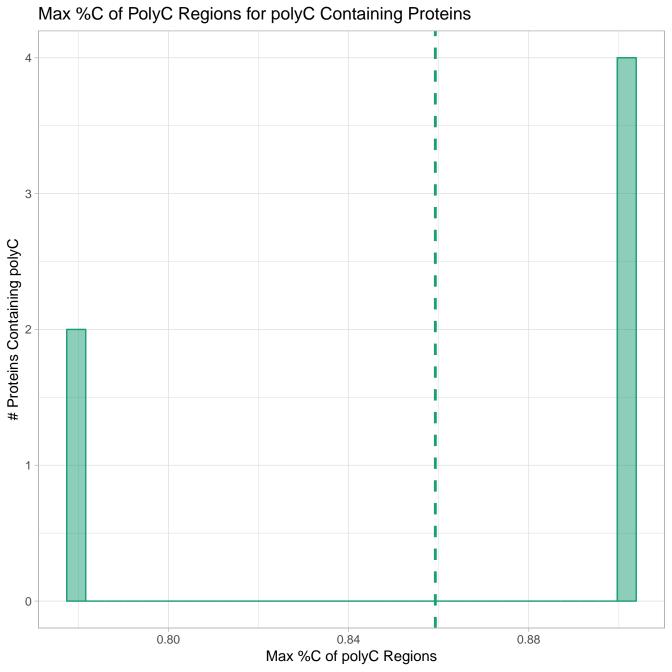
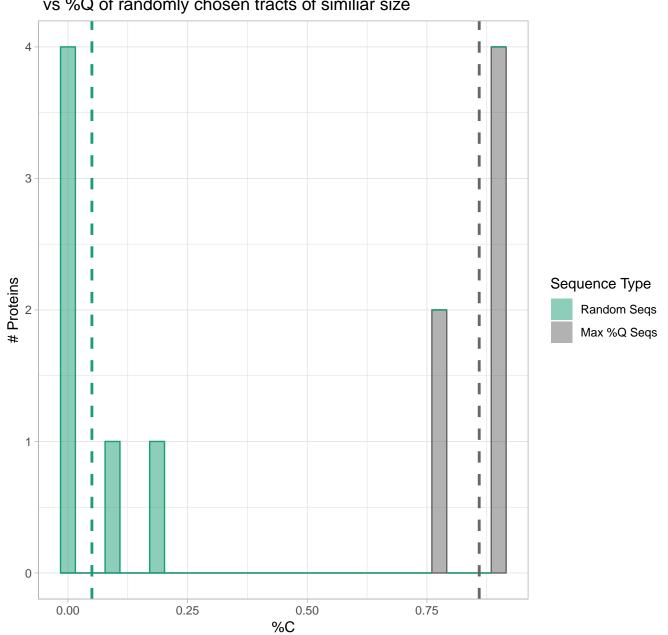


%C for 100000 50AA Random Seqs Histogram 40000 # of Randomly Choosen Pepitides 30000 20000 10000 0 0.00 0.05 0.10 0.15 0.20 0.25 %C (Random 50AA Seqs)



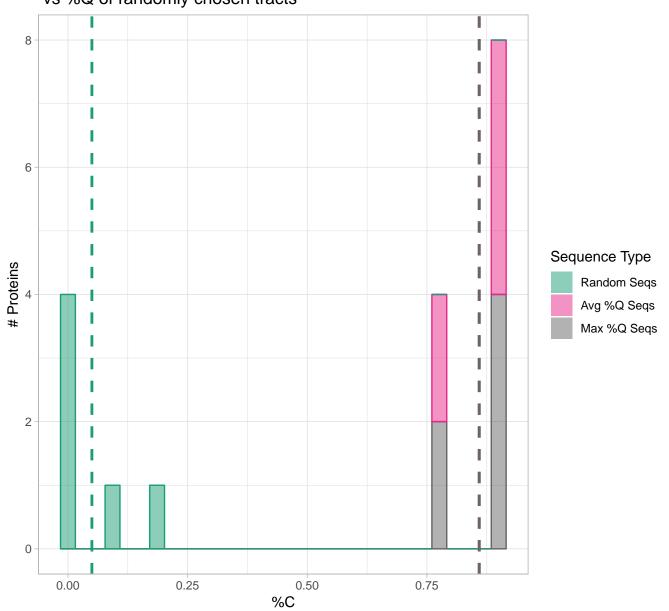


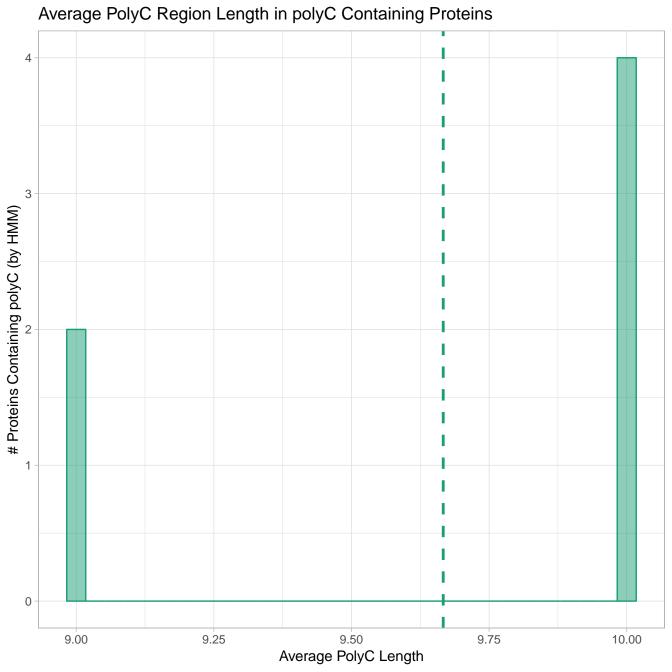
Max %C of PolyC Regions for polyC Containing Proteins vs %Q of randomly chosen tracts of similiar size

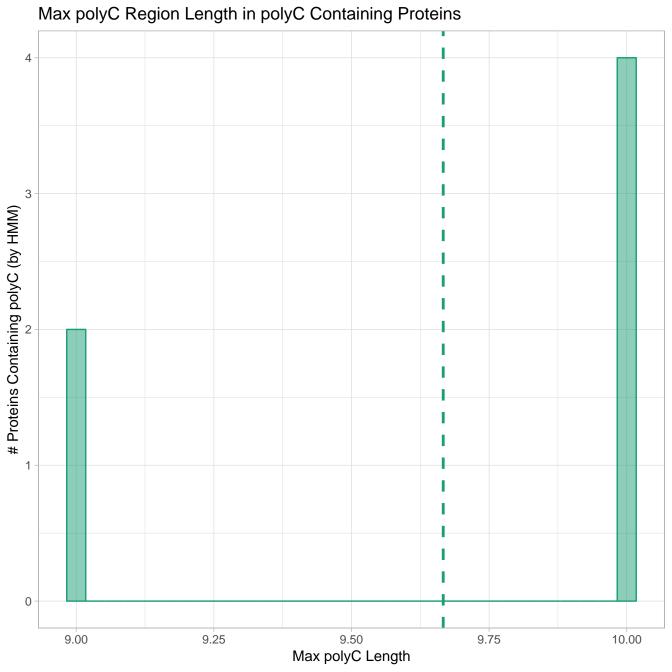


Avg %C of PolyC Regions for polyC Containing Proteins vs %C of Randomly Chosen Tracts 4 3 # Proteins Sequence Type Random Seqs Avg %Q Seqs 0 0.00 0.25 0.50 0.75 %C

Avg %C of polyC Regions vs Max %C of polyC Regions vs %Q of randomly chosen tracts







Number of polyC Regions in polyC Containing Proteins 6 # of C Proteins 2 0 1.00 0.75 1.25 Number of polyC Regions