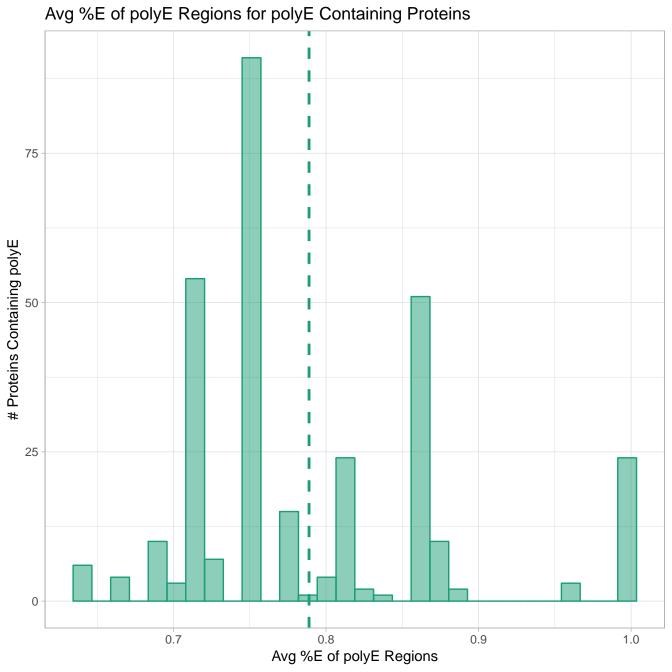
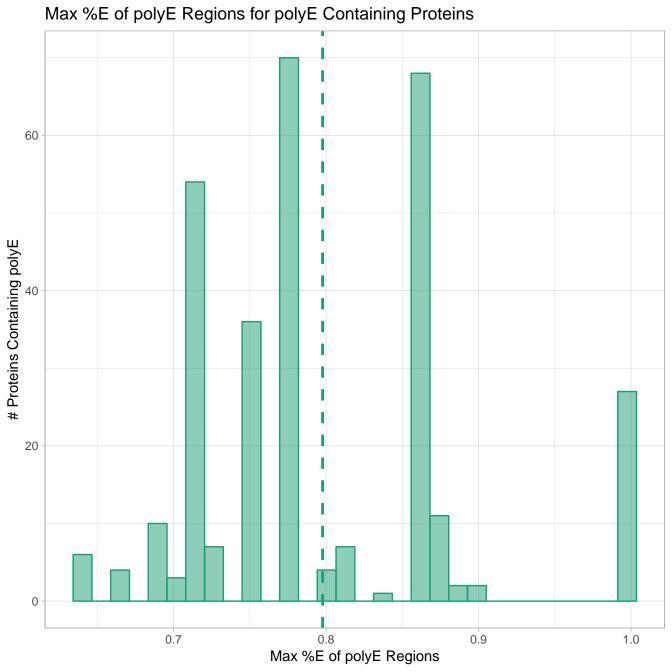
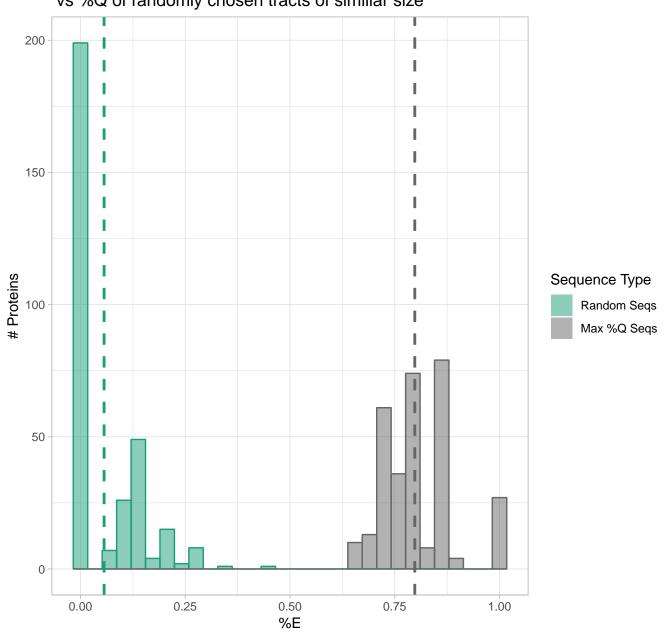


%E for 100000 50AA Random Seqs Histogram 20000 15000 # of Randomly Choosen Pepitides 10000 5000 0 0.2 0.0 0.1 0.3 0.4 %E (Random 50AA Seqs)

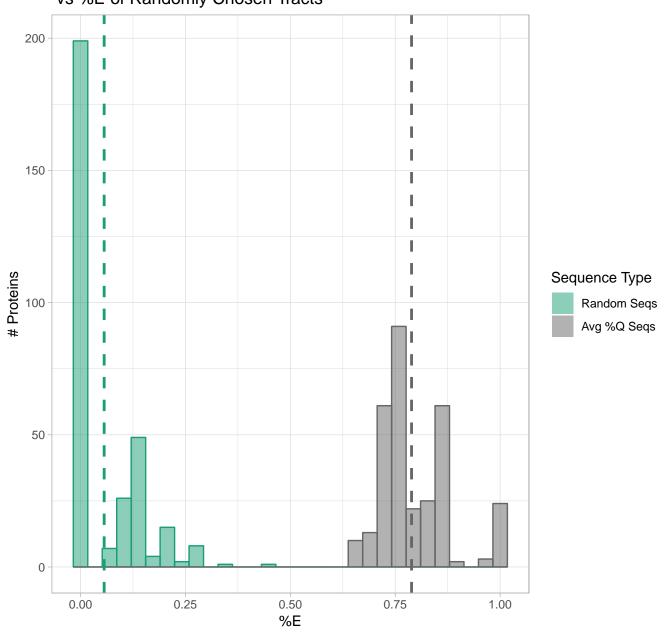




Max %E of polyE Regions for polyE Containing Proteins vs %Q of randomly chosen tracts of similiar size



Avg %E of PolyE Regions for polyE Containing Proteins vs %E of Randomly Chosen Tracts



Avg %E of polyE Regions vs Max %E of polyE Regions vs %Q of randomly chosen tracts

