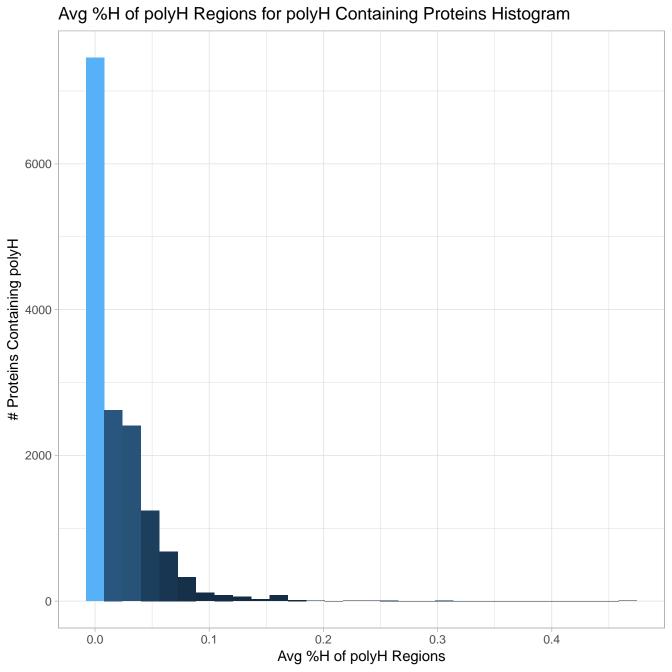


%H for 100000 50AA Random Seqs Histogram 30000 # of Randomly Choosen Pepitides 0 0.2 0.0 0.1 0.3 0.4 0.5 %H (Random 50AA Seqs)



Max %H of PolyH Regions for polyH Containing Proteins Histogram 4000 3000 # Proteins Containing polyH 2000 1000 0 0.2 0.0 0.1 0.3 0.4 Max %H of polyH Regions

