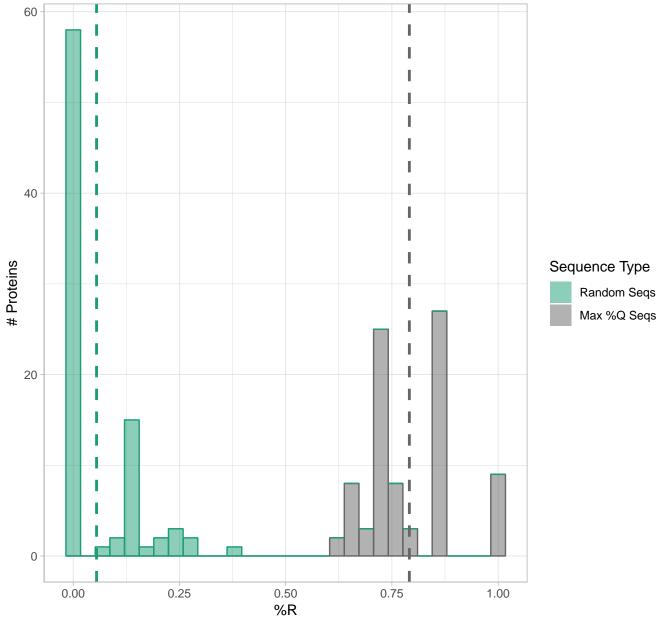


Max %R of PolyR Regions for polyR Containing Proteins vs %Q of randomly chosen tracts of similiar size 60 40 Sequence Type Random Seqs Max %Q Seqs



Avg %R of PolyR Regions for polyR Containing Proteins vs %R of Randomly Chosen Tracts 60 40 Sequence Type # Proteins Random Seqs Avg %Q Seqs 20 0 0.00 0.25 0.50 0.75 1.00 %R

Avg %R of polyR Regions vs Max %R of polyR Regions vs %Q of randomly chosen tracts

