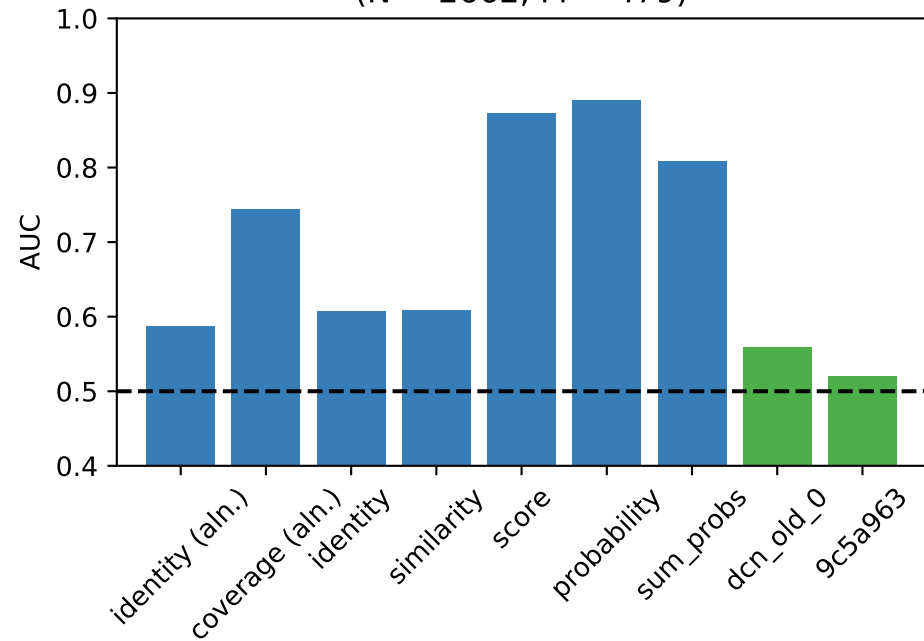


Predicting SCOP fold - combined  
(N = 2662, M = 479)



Predicting SCOP fold - per protein  
(N = 822, M = 149)

