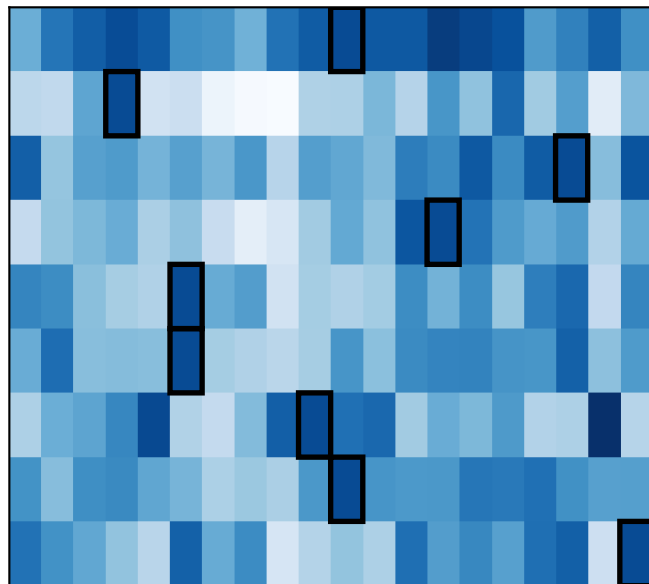


# Amino acid

A I L M F V P G W R H K D E N Q S T Y C

Position within peptide

1  
2  
3  
4  
5  
6  
7  
8  
9



0

-2

-4

-6

-8

-10

-12

-14

HCNN prediction,  $\Delta\log P$