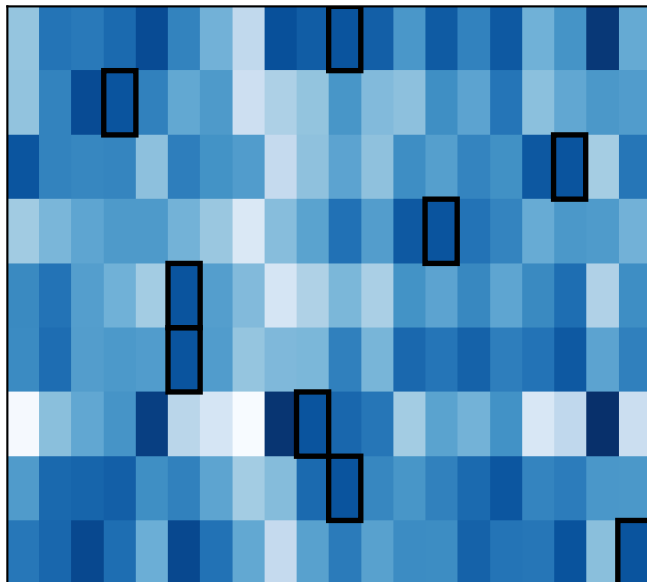


Amino acid

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



HCNN prediction, $\Delta\log P$

2
0
-2
-4
-6
-8
-10
-12
-14