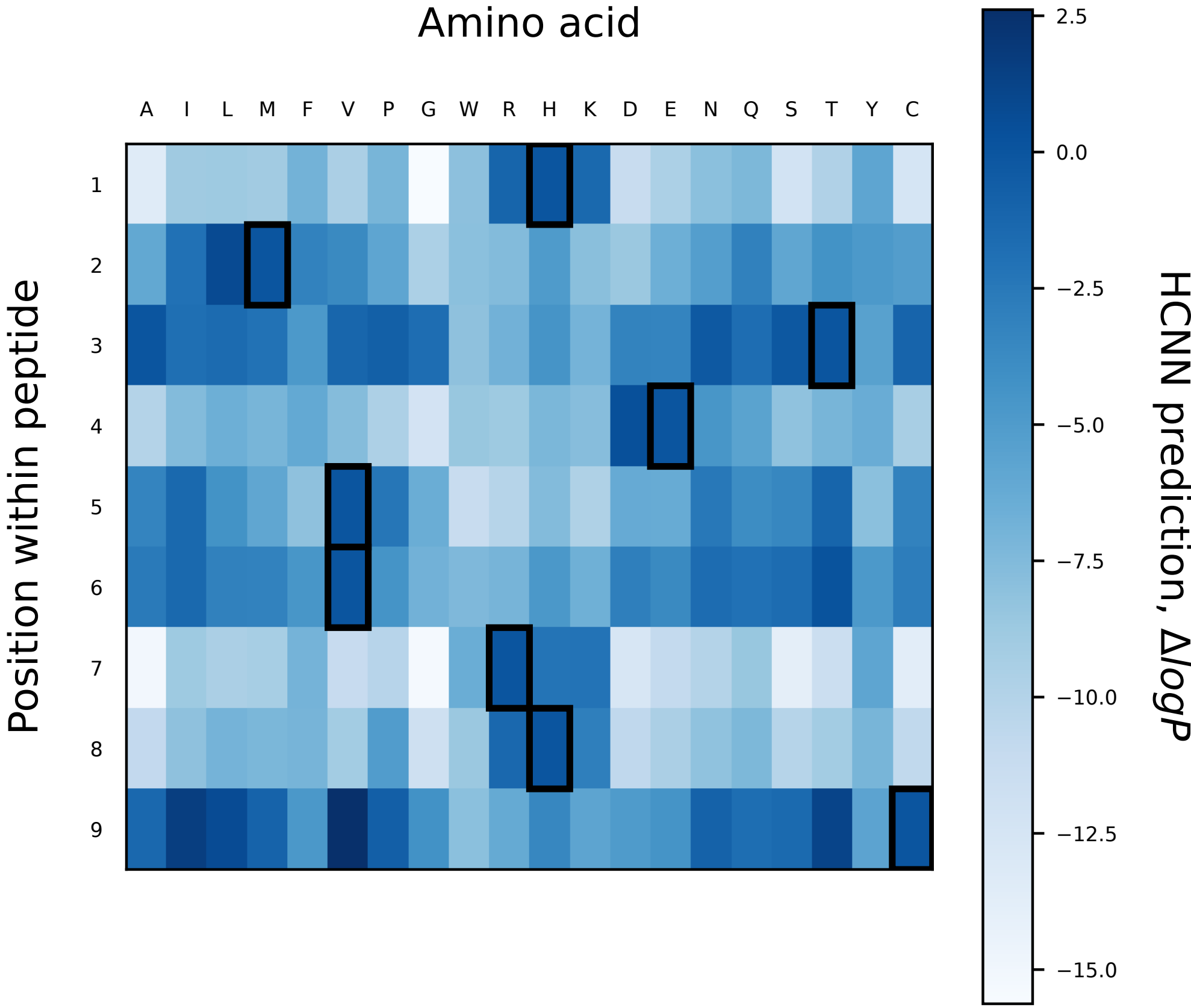
Amino acid
A I L M F V P G W R H K D E N Q S T Y C
Position within peptide
1
2
3
4
5
6
7
8
9
2.5
0.0
−2.5
−5.0
−7.5
−10.0
−12.5
−15.0
HCNN prediction, ΔlogP