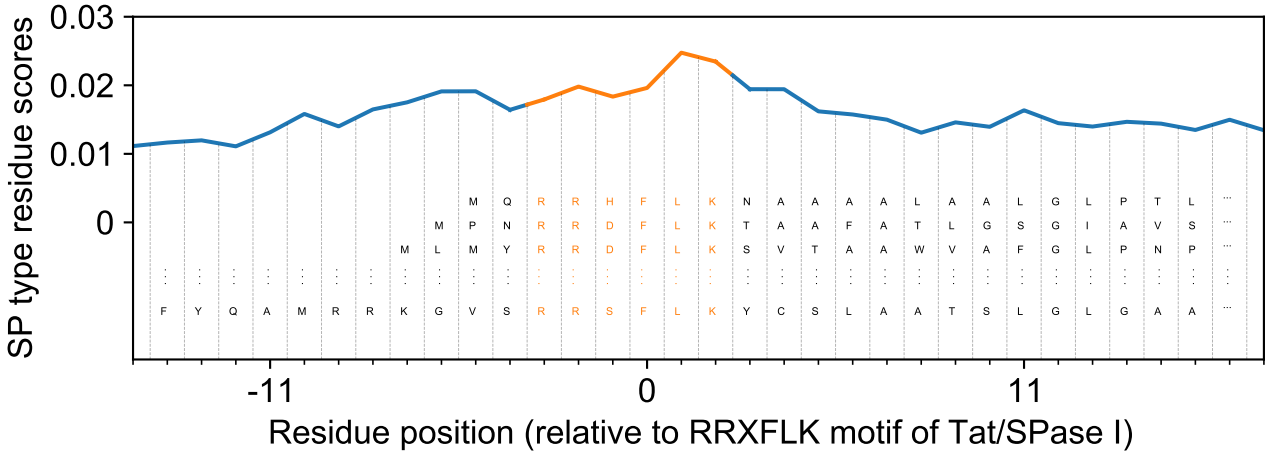
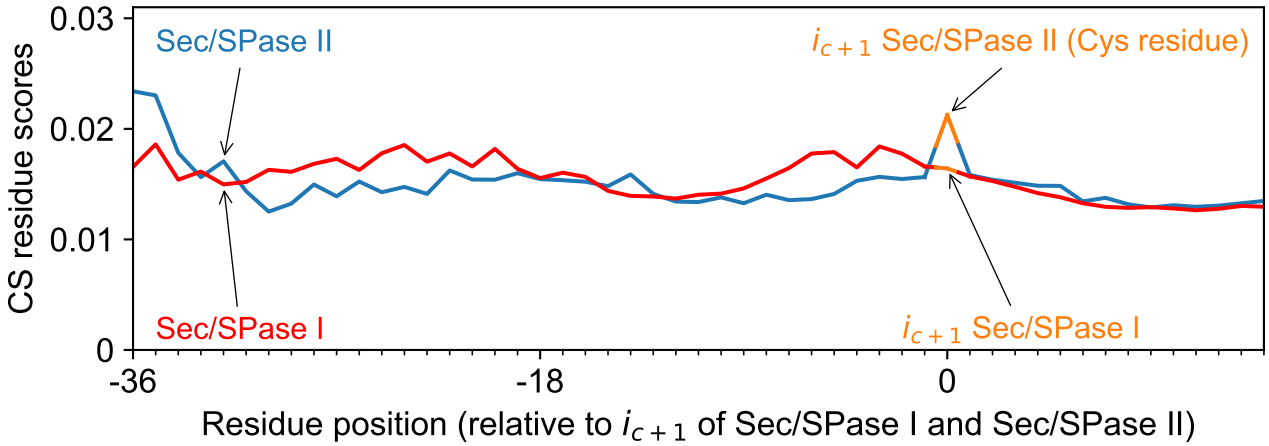

SP type residue scores
0.03
0.02
0.01
0
-11
0
11
Residue position (relative to RRXFLK motif of Tat/SPase I)


CS residue scores
0.03
0.02
0.01
0
Sec/SPase II
Sec/SPase I
i_{c+1} Sec/SPase II (Cys residue)
i_{c+1} Sec/SPase I
-36
-18
0
Residue position (relative to i_{c+1} of Sec/SPase I and Sec/SPase II)