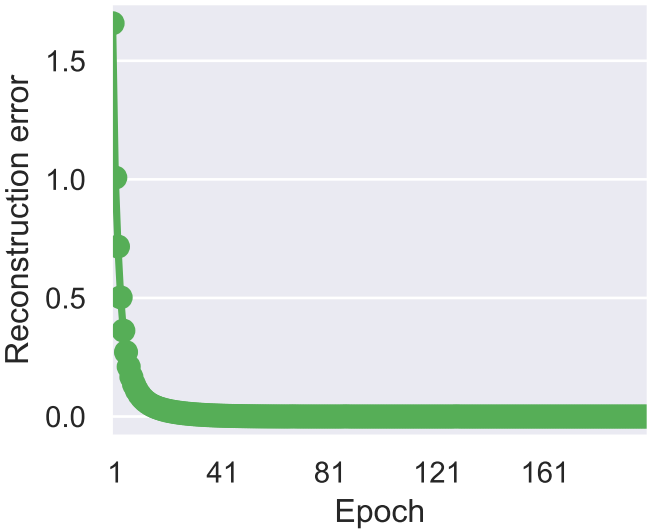


One-hot encoder with arch 399-700-Relu-64 on test set

Reconstruction error during training



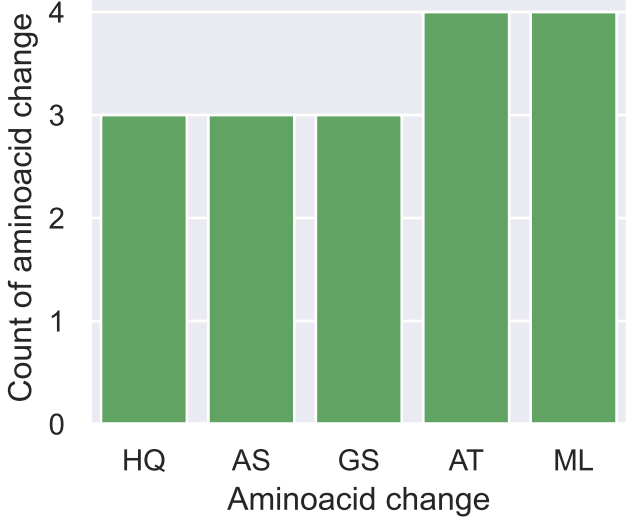
Average reconstruction error on test set:
CELoss: 0.01382

From 7057 sequences with right seq len:7056

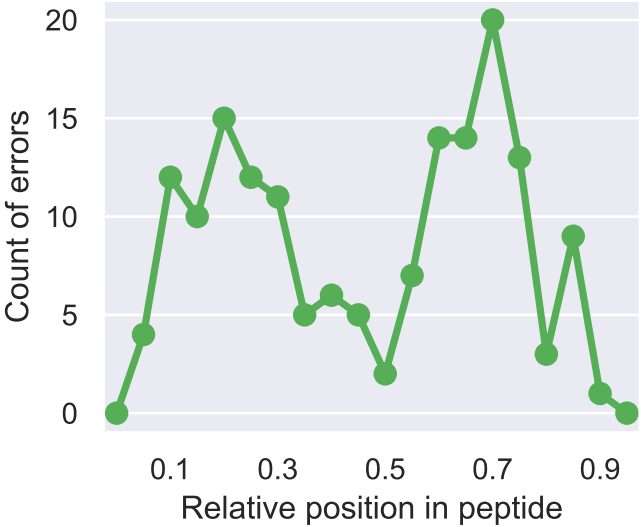
	precision	recall	f1-score	support
A	1.00	1.00	1.00	10584
C	1.00	1.00	1.00	7086
D	1.00	1.00	1.00	2883
E	1.00	1.00	1.00	5978
F	1.00	1.00	1.00	10832
G	1.00	1.00	1.00	10228
H	0.99	0.99	0.99	1108
I	1.00	1.00	1.00	1312
K	0.99	0.99	0.99	668
L	1.00	1.00	1.00	4306
M	0.99	0.98	0.99	383
N	1.00	1.00	1.00	2962
P	1.00	1.00	1.00	2900
Q	1.00	1.00	1.00	6643
R	1.00	1.00	1.00	3198
S	1.00	1.00	1.00	15916
T	1.00	1.00	1.00	6553
V	1.00	0.99	0.99	1826
W	0.99	0.98	0.99	685
Y	1.00	1.00	1.00	6028

accuracy		1.00	102079
macro avg	1.00	1.00	1.00 102079
weighted avg	1.00	1.00	1.00 102079

Aminoacid changes (5 most common)



Count of errors by position



Count of errors by position (in %)

