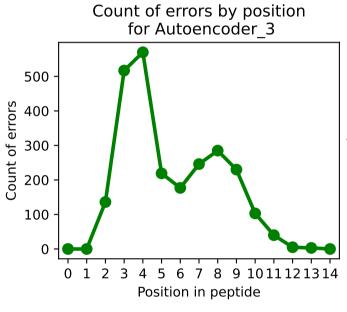
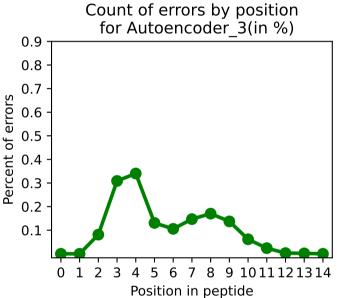


Average reconstruction error on test set:

MSE: 0.00373 RMSE: 0.06104

Architecture: 300->200 Learning rate: 0.001





Autoencoder_3

-0.4 -0.8 -0.8 -1.0 -1.2 -1.4 -1.6 -

5 6 7 8 9 10 Position in peptide

3 4

9 10 11 12 13

Average blossum62 score by position for