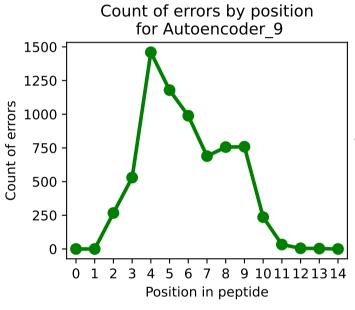


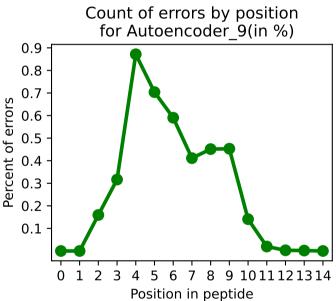
Average reconstruction error on test set:

MSE: 0.00849 RMSE: 0.09215

Architecture: 300->2000->1000->200

Learning rate: 0.001





Autoencoder\_9

-0.4 -0.6 -0.8 -0.8 -1.0 -1.2 -1.6 -

5 6 7 8 9 10 Position in peptide

3

9 10 11 12 13

Average blossum62 score by position for