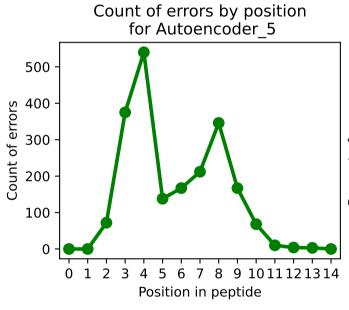


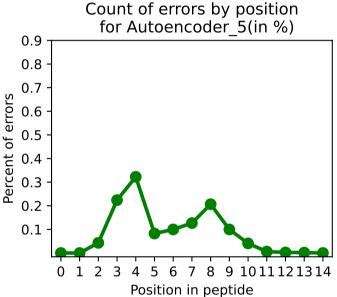
Average reconstruction error on test set:

MSE: 0.00223 RMSE: 0.04718

Architecture: 300->700->200 without L2

Learning rate: 0.001





Autoencoder 5 0.0 Average blossum62 score -0.5 -1.0 -1.5 -2.0

3 4 9

8 Position in peptide

10 11 12 13

Average blossum62 score by position for