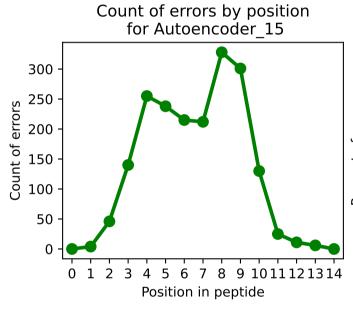


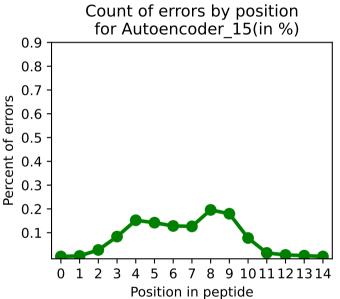
Average reconstruction error on test set:

MSE: 0.00288 RMSE: 0.05365

Architecture: 300->700->200 without L2 sigmoid

Learning rate: 0.001





0.5 - 900 0.0 - 0.5 - 0.0 - 0.5 - 0.0 - 0.5 - 0.0 - 0.5 - 0.0 - 0.5 - 0.0 - 0.5 - 0.

2 3

Average blossum62 score by position for Autoencoder\_15

9 10 11 12 13

8

Position in peptide