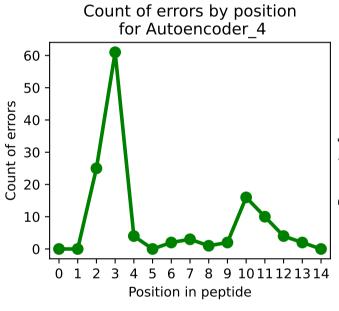


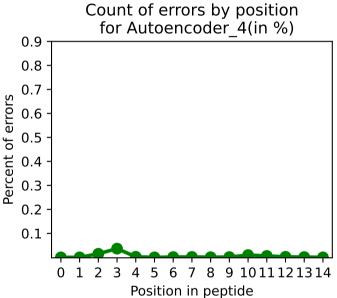
Average reconstruction error on test set:

MSE: 0.00018 RMSE: 0.01351

Architecture: 300->300 without L2

Learning rate: 0.001





Average blossum62 score by position for Autoencoder\_4 0.0 Average blossum62 score 0.5 -1.0-1.5-2.0 -2.5 **-**3.0 9 10 11 12 13 2 3 4 8 Position in peptide