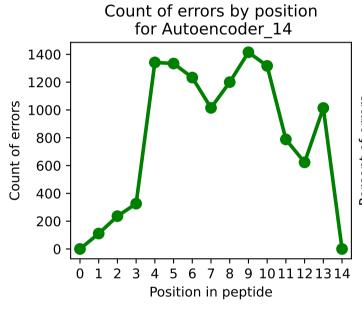


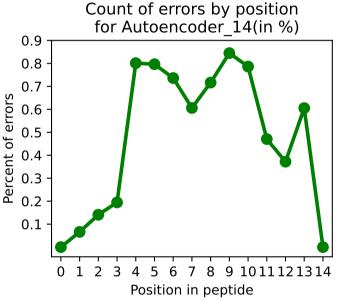
Average reconstruction error on test set:

MSE: 0.01904 RMSE: 0.13797

Architecture: 300->700->200 with big L2

Learning rate: 0.001





Average blossum62 score by position for Autoencoder 14 1.5 Average blossum62 score 1.0 0.5 0.0 -0.5 -1.0-1.5 -2.09 10 11 12 13 2 3 6 7 8 Position in peptide