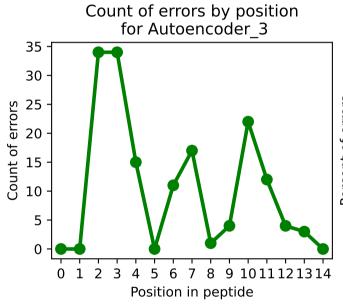


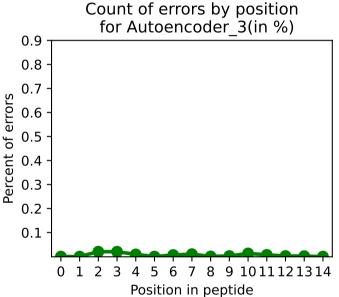
Average reconstruction error on test set:

MSE: 0.00031 RMSE: 0.01765

Architecture: 300->200 without L2

Learning rate: 0.001





Average blossum62 score by position for Autoencoder 3 0.0 Average blossum62 score -0.5 -1.0-1.5 -2.0-2.59 3 10 11 12 13 2 7 8 Position in peptide