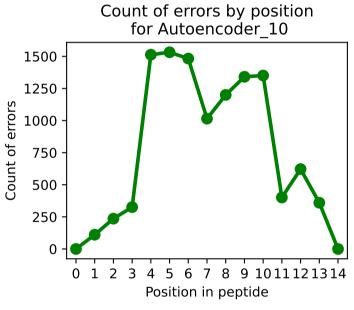


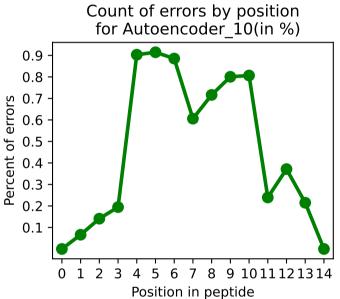
Average reconstruction error on test set:

MSE: 0.01334 RMSE: 0.11551

Architecture: 300->2000->200 without L2

Learning rate: 0.001





Average blossum62 score by position for Autoencoder 10 1.0 Average blossum62 score 0.5 0.0 -0.5 -1.0-1.5-2.0 3 9 10 11 12 13 2 6 7 8 Position in peptide