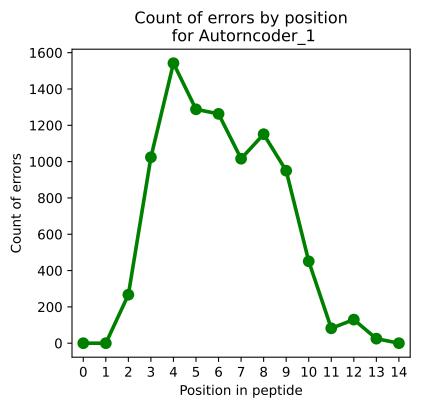
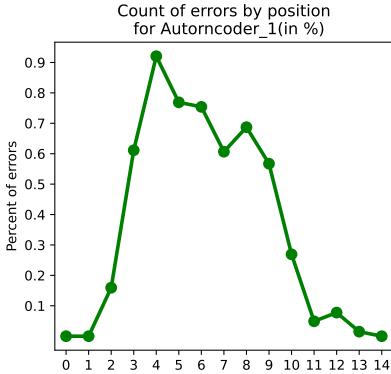


Average reconstruction error on test set:

MSE: 0.01191 RMSE: 0.10912

Architecture: 315->800->600->200





Position in peptide

Average blossum62 score by position for Autoencoder 1

