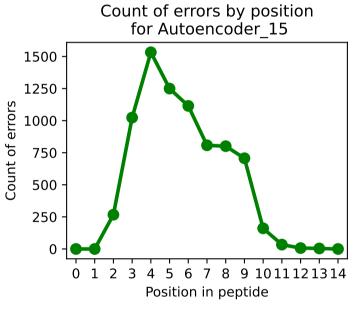


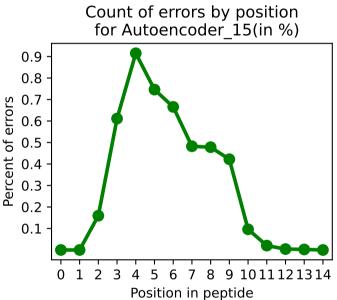
Average reconstruction error on test set:

MSE: 0.00992 RMSE: 0.0996

Architecture: 300->700->200 without L2 sigmoid

Learning rate: 0.001





Average blossum62 score by position for Autoencoder\_15

-0.6
-0.8
-0.8
-1.0
-1.2
-1.4
-0.6
-1.6
-1.8
-2.0
-2 3 4 5 6 7 8 9 10 11 12 13

Position in peptide