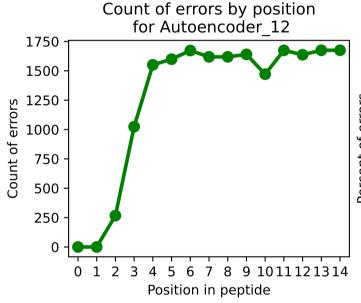


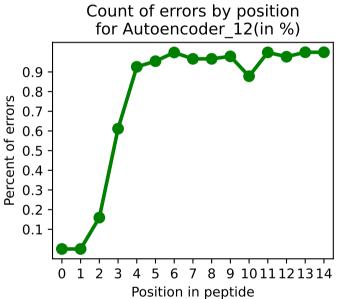
Average reconstruction error on test set:

MSE: 0.0225 RMSE: 0.14999

Architecture: 300->700->200 without L2

Learning rate: 0.01





Average blossum62 score by position for Autoencoder\_12

0.0

-0.5
-0.5
-1.0
-2 3 4 5 6 7 8 9 10 11 12 13 14

Position in peptide