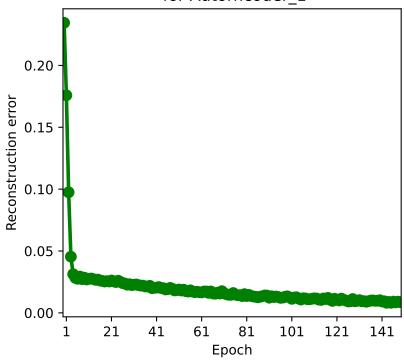
Reconstruction error during training for Autorncoder_1

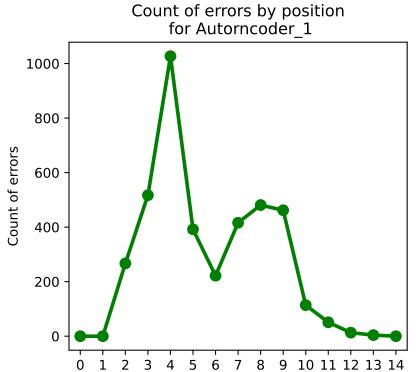


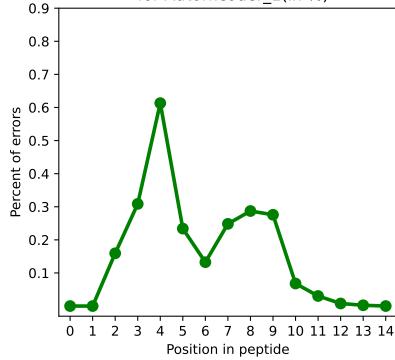
Average reconstruction error on test set:

Count of errors by position for Autorncoder_1(in %)

MSE: 0.00544 RMSE: 0.07375

Architecture: 315->200





Average blossum62 score by position for Autoencoder 1

Position in peptide

