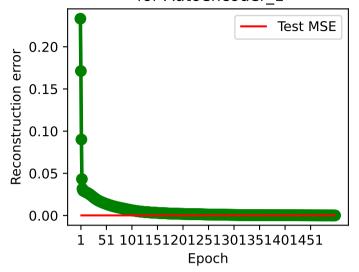
Reconstruction error during training for Autoencoder 1



Architecture: 300->150 without L2

Average reconstruction error on test set:

Learning rate: 0.001

Κ

MSE: 0.00017 RMSE: 0.01299

> recall f1-score precision support 1.00 1.00 1.00 2570 C 1.00 1.00 1683 1.00 D 1.00 739 1.00 1.00 Ε 1.00 1.00 1.00 1381 F 1.00 1.00 1.00 2561 G 1.00 1.00 1.00 2649 Н 1.00 1.00 1.00 315

0.99 0.99 0.99 295 0.99 0.98 0.97 170 0.99 1.00 0.99 1089 М 0.99 0.97 0.98 117

1.00 1.00 784 Ν 1.00 1.00 Ρ 759 1.00 1.00 1.00 1.00 1.00 1654 0.99 1.00 1.00 773

Q R S T 1.00 1.00 1.00 3924 1.00 1.00 1.00 1637 V 0.99 1.00 0.99 469 W 1.00 0.96 0.98 176

1.00

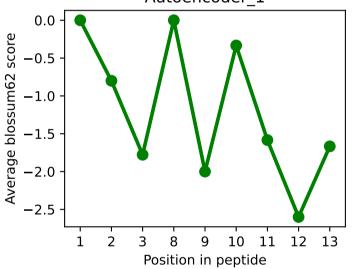
1380

1.00 25125 accuracy macro avg 1.00 0.99 0.99 25125 weighted avg 1.00 1.00 1.00 25125

1.00

0.99

Average blossum62 score by position for Autoencoder 1



Count of errors by position for Autoencoder 1 17.5 15.0 Count of errors 12.5 10.0 7.5 5.0 2.5 0.0 3 5 6 8 9 10 11 12 13 14 2 4 7 Position in peptide

