Combining Word Embeddings of Protein Sequences with Evolutionary Information for Secondary Structure Prediction Using Deep Neural Networks

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Introduction to Bachelor‘s thesis

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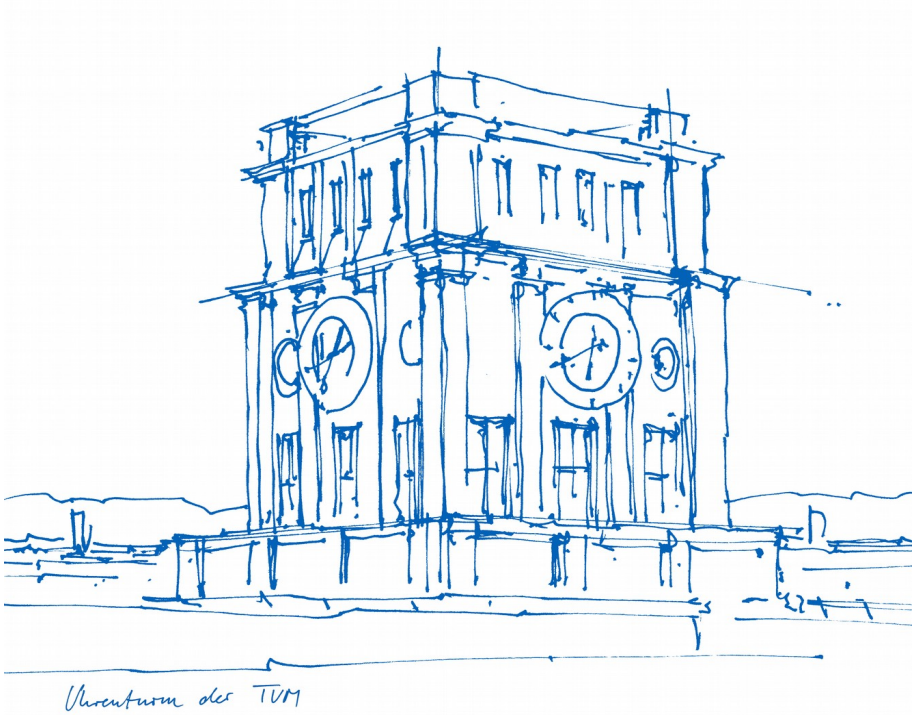


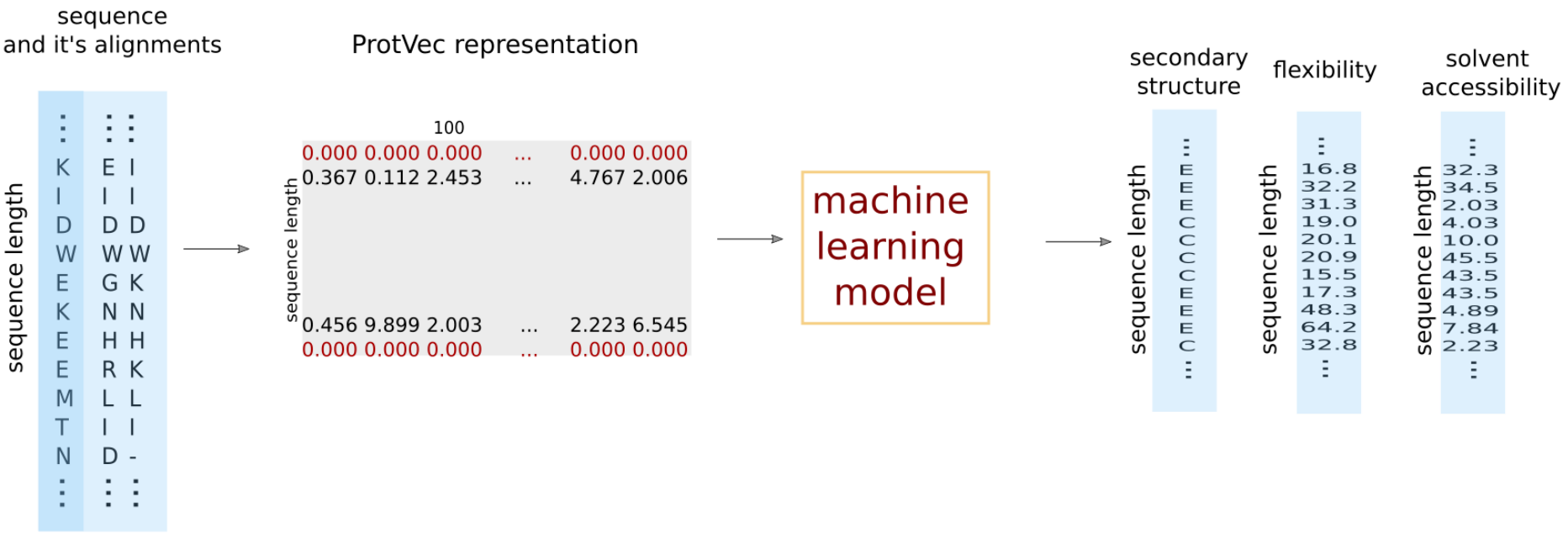
Overview

- Project introduction
- Dataset
- Data preprocessing
- Convolutional neural networks
- Results so far
- Outlook

Project Introduction

- Secondary structure prediction with neural networks
- Integrate evolutionary information in input
- Multi target prediction
  - Secondary structure
  - Solvent accessibility
  - Flexibility
- ProtVec representation on residue basis

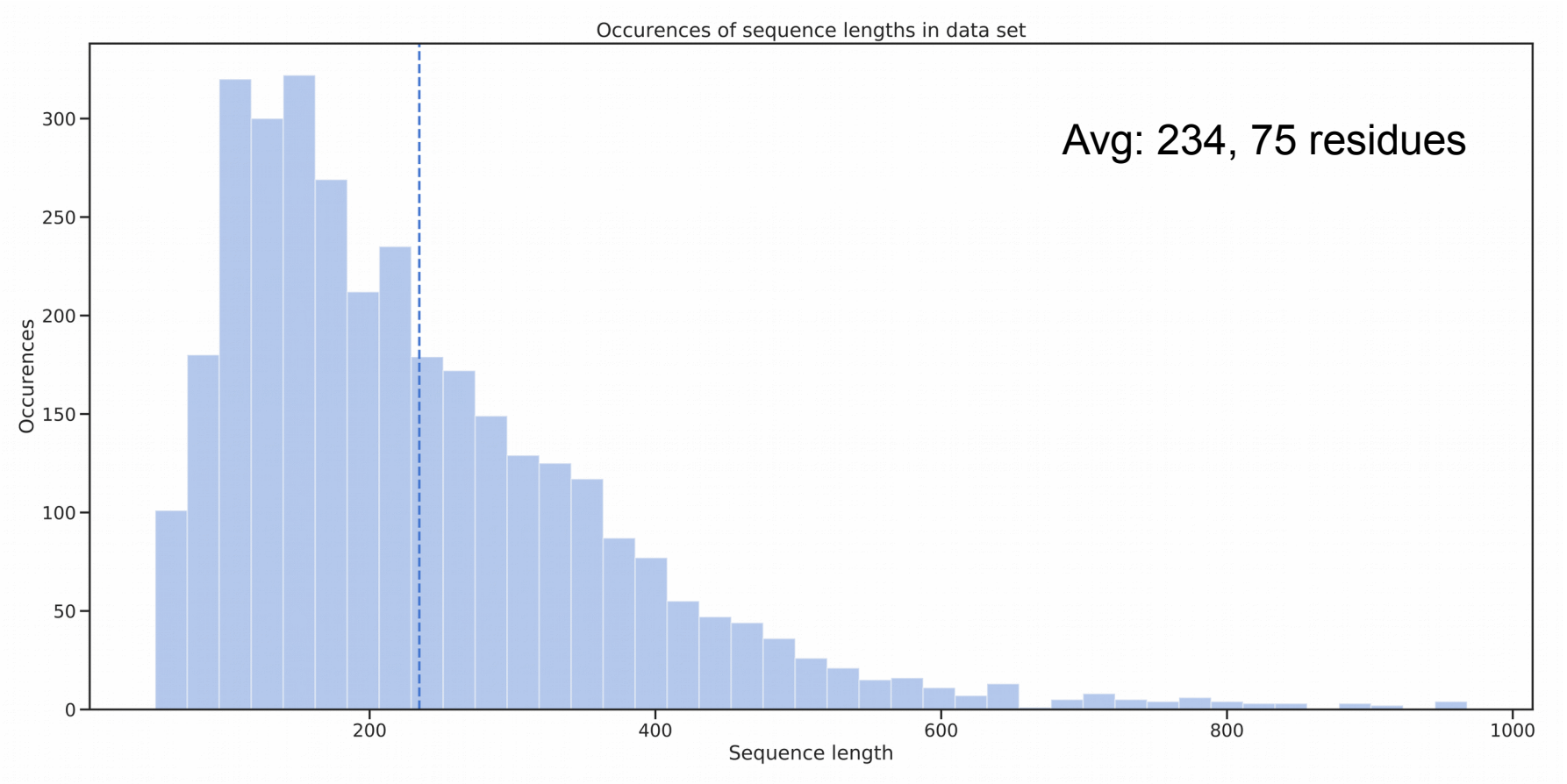
Project Introduction



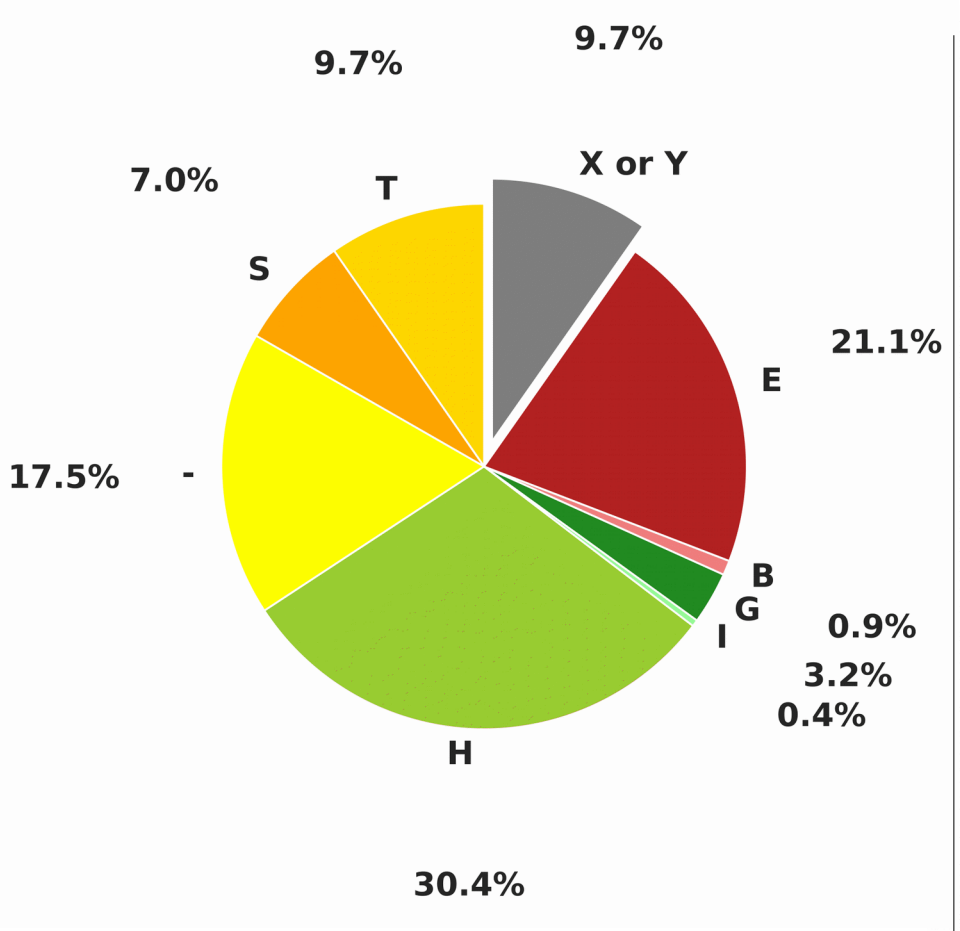
Dataset

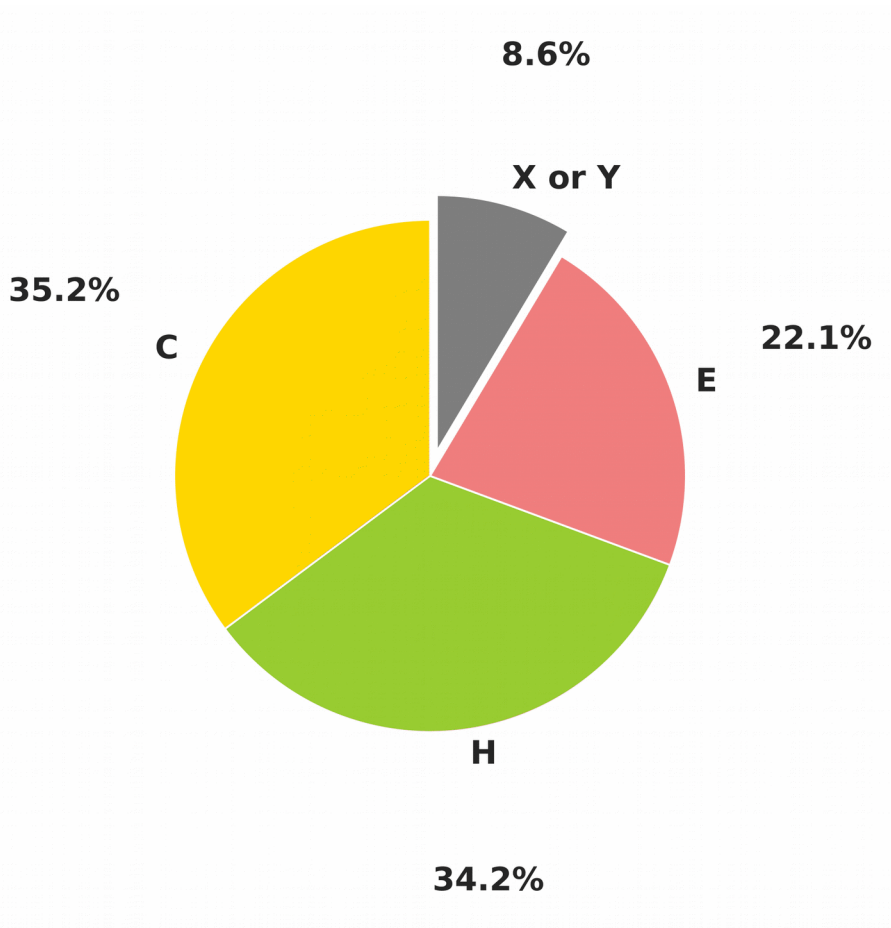
- 3313 protein sequences from PDB
- According structure information, solvent accessibility and flexibility from DSSP algorithm
- Aligned sequences for each protein sequence

Dataset – Sequence Lengths

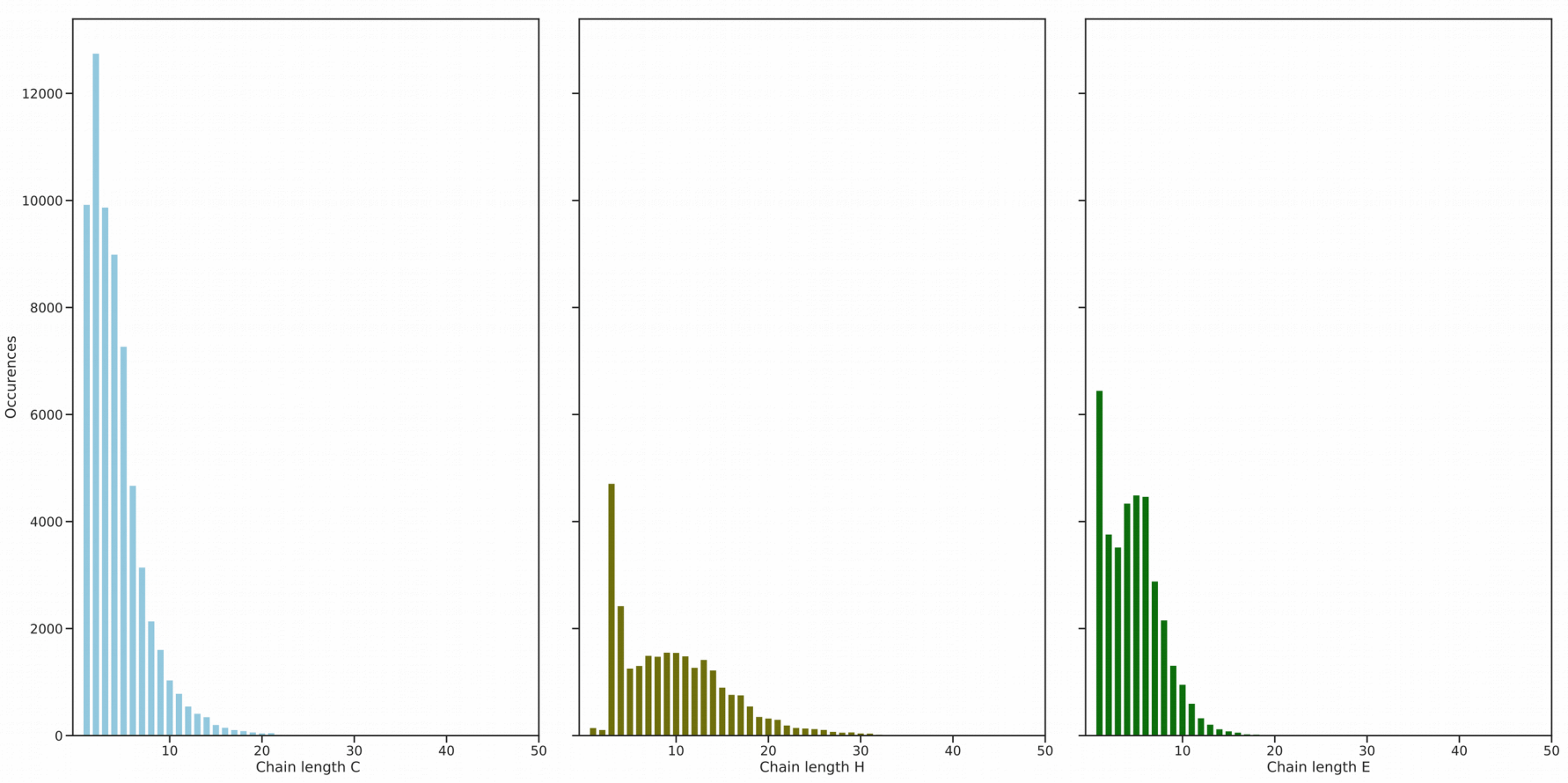


Dataset – DSSP States





Dataset – DSSP-3 States Chains



Avg: 4,2

Avg: 9,8

Avg: 4,7

Dataset – Training, Validation, Test Set

- 20% of the samples → test set (649 samples)
- Remaining 80% → train and validation set (2092 and 529 samples)
- Random selection, assured equal distribution

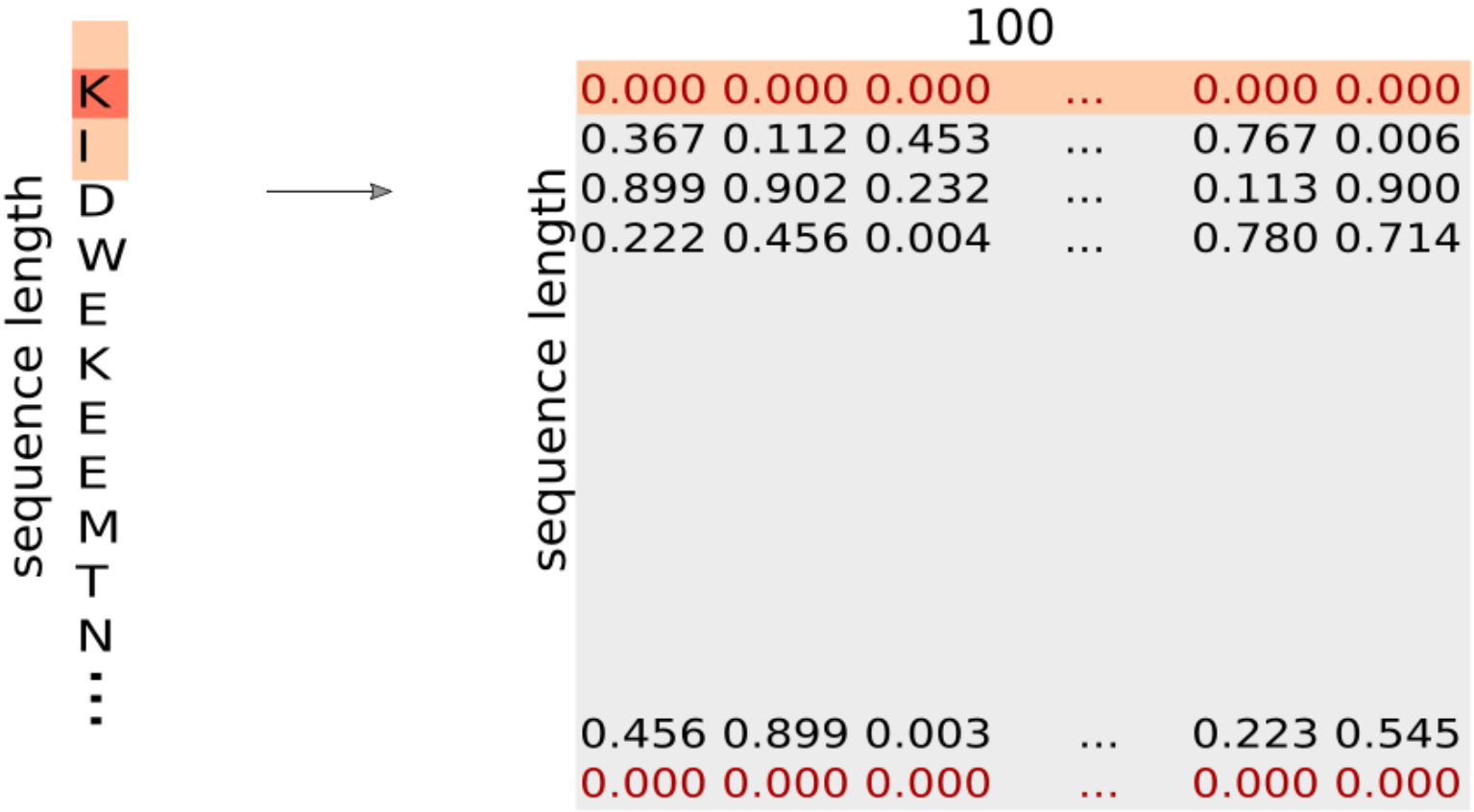
Implementation So Far

- Training and testing pipeline implemented
- Simple CNN used
  - 3 layers
  - Kernels: 1, 7, 15
  - NLL loss + log_softmax
- Secondary structure prediction without evolutionary information

Data Preprocessing

- Protein sequences represented through word embeddings
- ProtVec
  - Vector embeddings learned through Word2Vec
  - Vector for each possible n-gram, fixed length
- 3-grams, length 100

Data Preprocessing



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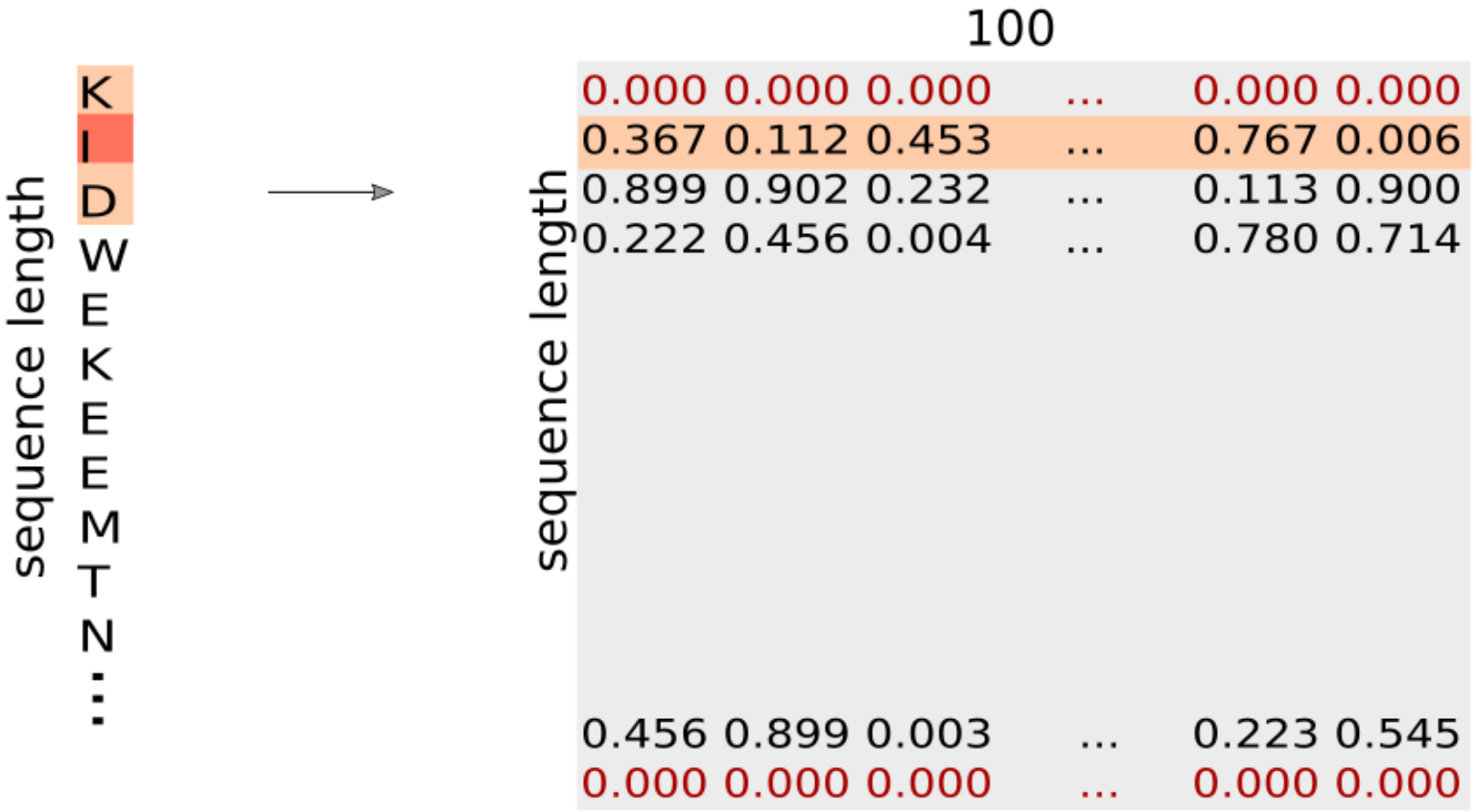

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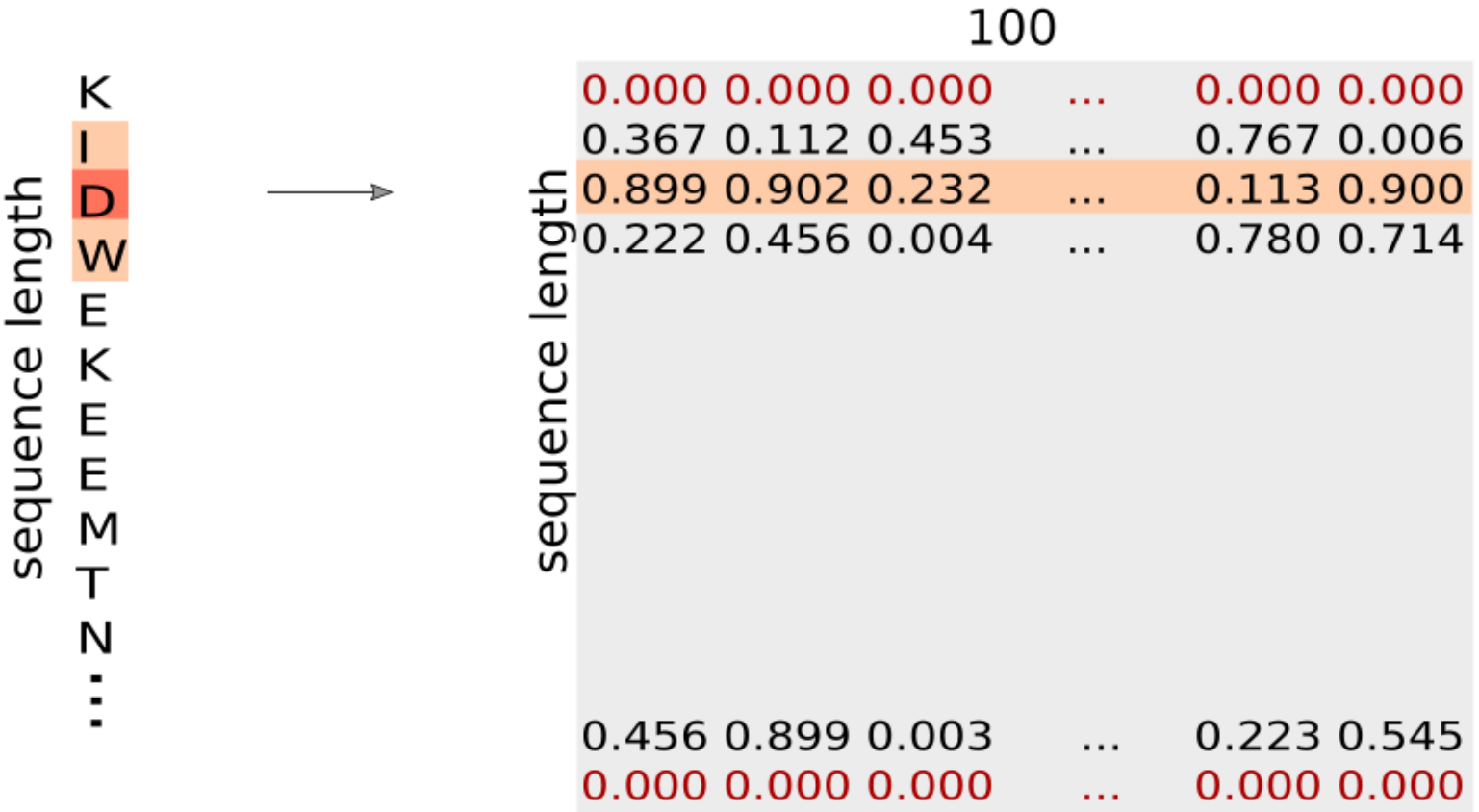
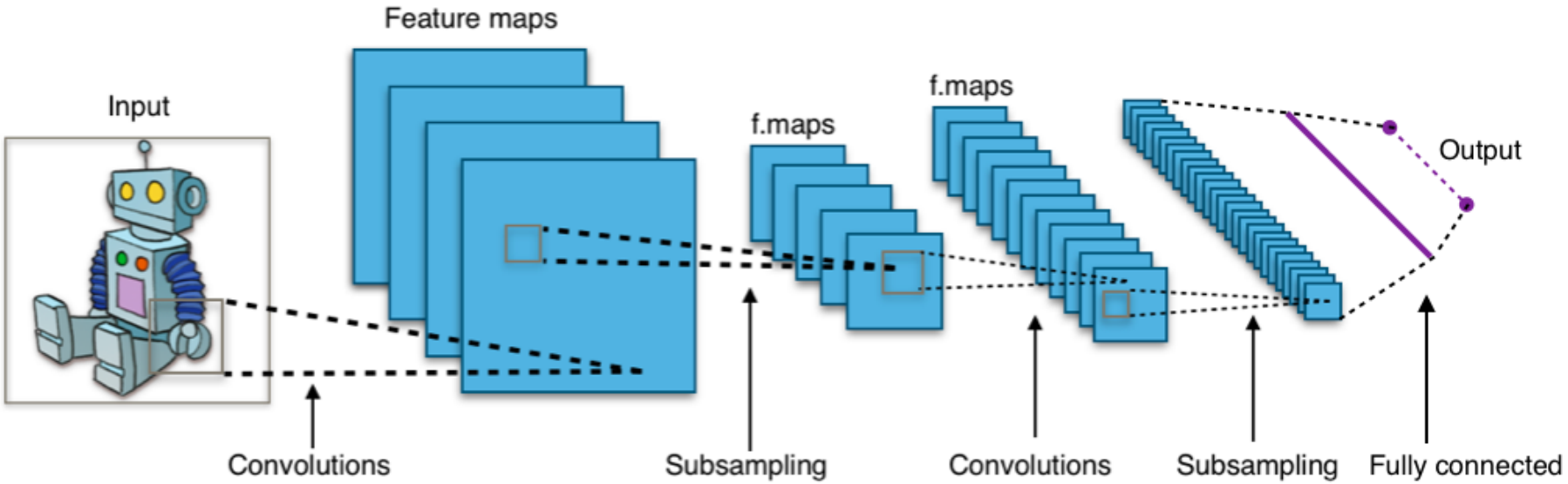
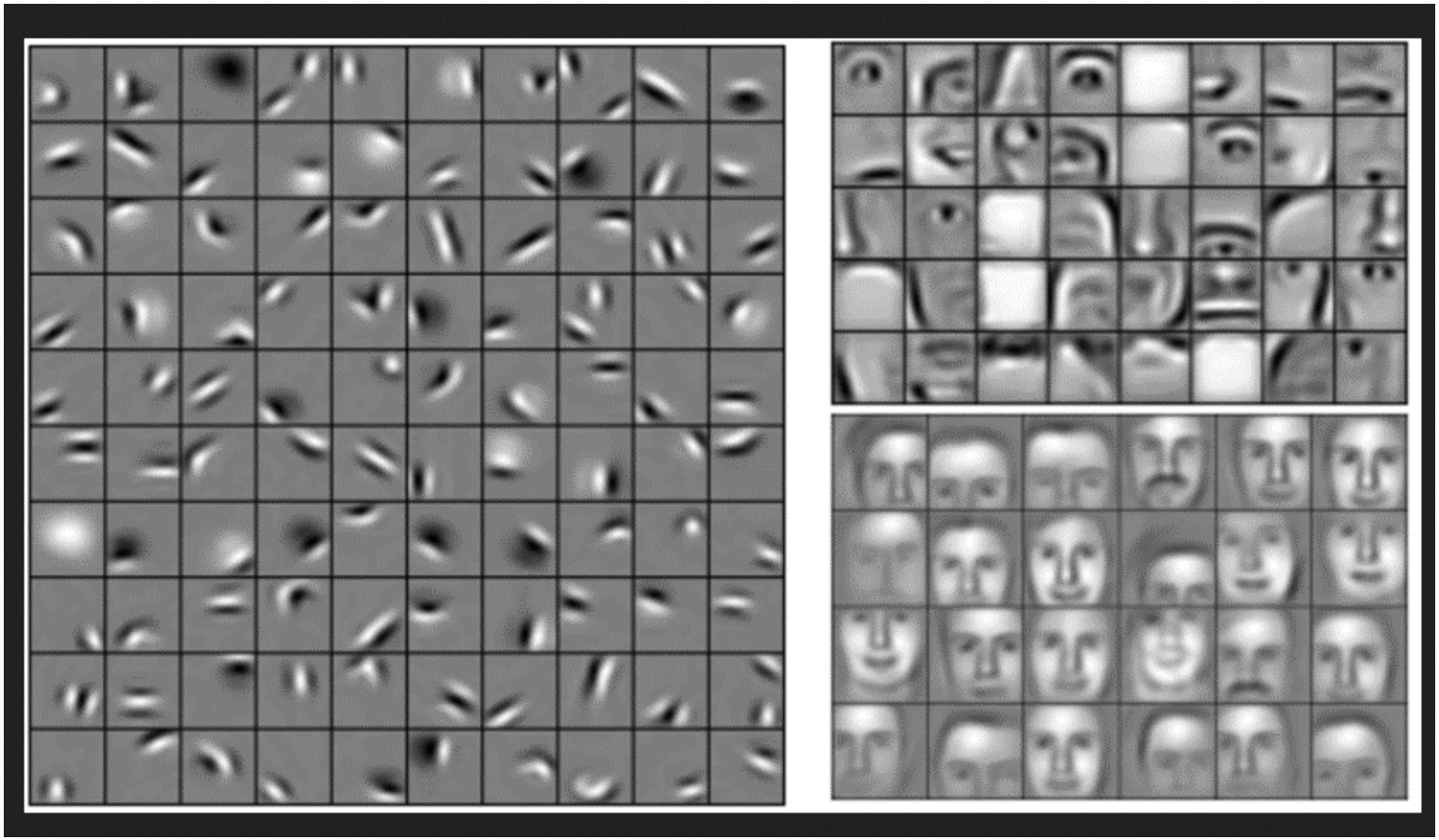


Convolutional Neural Networks



https://de.wikipedia.org/wiki/Convolutional_Neural_Network

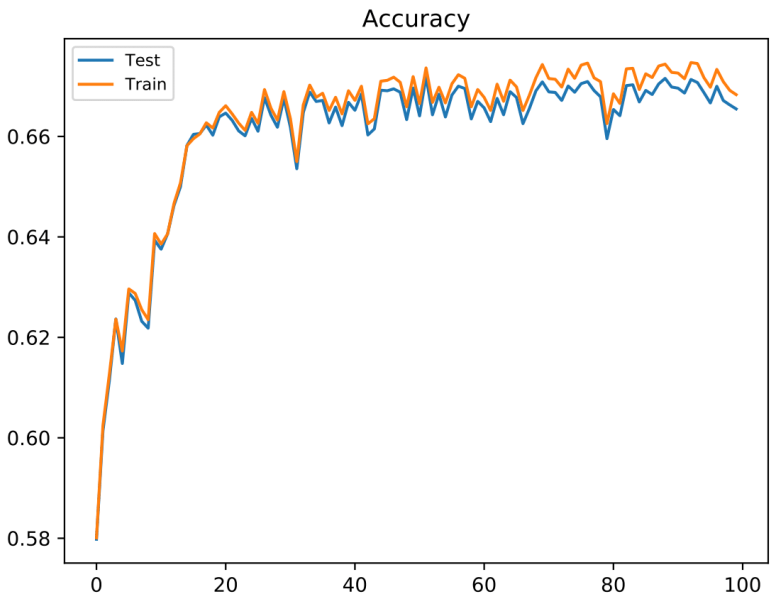
Convolutional Neural Networks



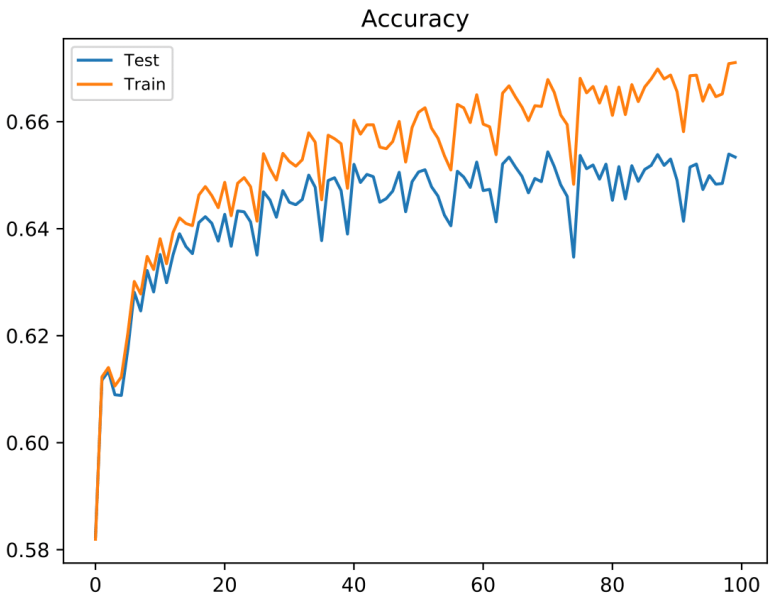
https://brohrer.github.io/images/cnn18.png

Results So Far – 1hot & ProtVec Accuracy

1hot

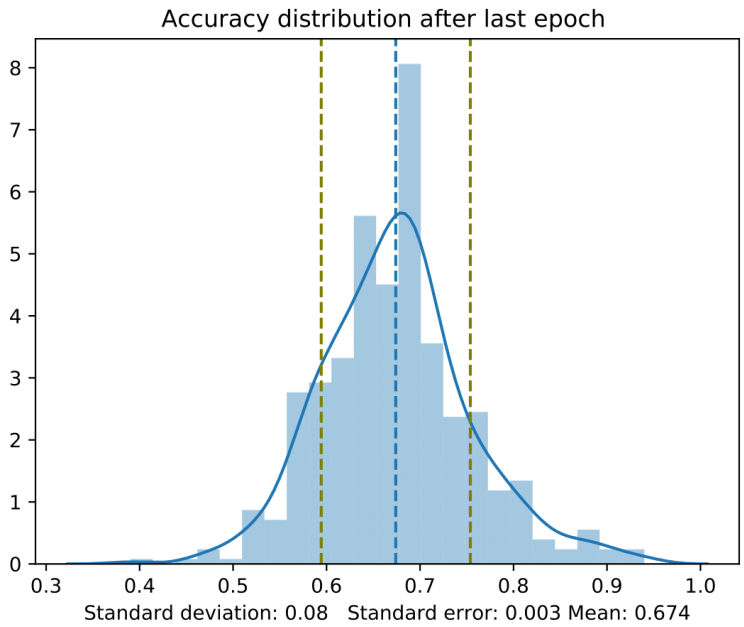


ProtVec

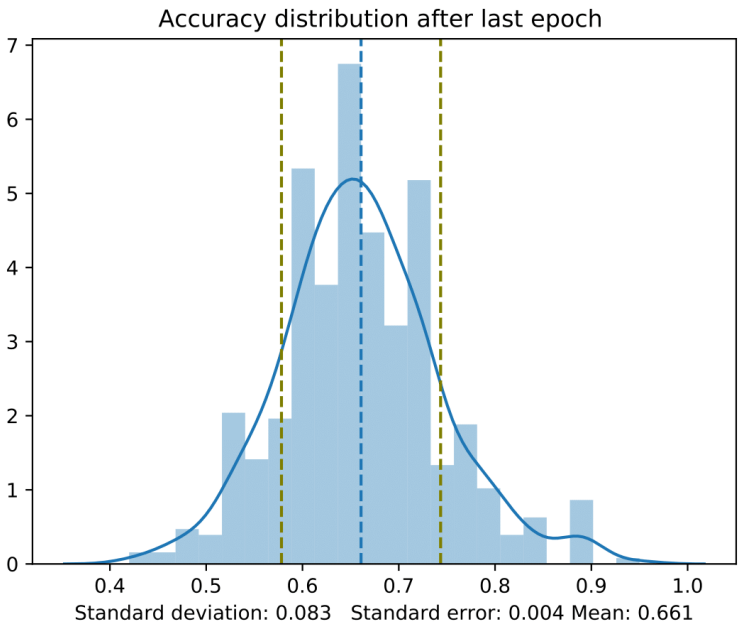


Results So Far – 1hot & ProtVec Accuracy

1hot



ProtVec



Outlook

- Incorporate evolutionary information
- Flexibility and solvent accessibility prediction
- Tune hyperparameters
  - Kernel sizes
  - Regularization
  - Non-linear activation functions e.g. SeLU
- LSTM networks
  - Capable of learning long-term dependencies

Thank you for your attention!