## **Assignment 3**

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## Steps performed for classification of B-cell epitopes and non-epitopes using deep learning.

- 1. Initially for given training and testing set p-features are obtained by computing dipeptide sequences. Tripeptide sequences are also calculated but are not used further as they are not giving higher scores.
- 2. Calculated dipeptides for training and testing dataset are stored in a csv file called dipep train.csv and dipep test.csv.
- 3. Columns with zero values are removed.
- 4. A feed forward neural network has been applied by feeding 400 features as input to the perceptron.
- 5. Activation function used for input layer and hidden layer is hyperbolic tangent(tanh) and for output layer activation function used is sigmoid. Use of other activation functions were not giving a good score.
- 6. The model is then trained and fitted with different batch sizes and epochs values.
- 7. The two highest scores are obtained by setting batch size=100 and epochs =100 with the score of 0.78292. The output of this model is stored in a file named output ANN.csv.
- 8. The score obtained by setting batch size=100 and epochs=150 is 0.76015. This output is stored in a file named final output1.csv.
- 9. Both python file and notebook file are submitted.

## Filenames given in folder are:

trainset.data: Training dataset downloaded from Kaggle.

testset.data: Testing dataset downloaded from Kaggle.

**dipep train.csv:** Training dataset obtained by calculating dipeptide residue sequences

**dipep test.csv:** Testing dataset obtained by calculating dipeptide residue sequences.

**output** ANN.csv: Output file obtained by setting batch size=100 and epochs =100.

**Final output1.csv:** Output file obtained by setting batch size=100 and epochs =150.

## To run the python notebook file:

- 1. Open google colab in your browser
- 2. Open a new jupyter notebook and upload all the training and testing files given in the folder.
- 3. Upload the notebook file named BDMH\_A3\_ANN provided inside the folder.
- 4. Run one by one each cell until the output file is generated.
- 5. Download the output file generated and upload it to the submit prediction page of Kaggle link provided in the assignment page.
- 6. Submit prediction to see the obtained score.