

# Master in Artificial Intelligence

## Advanced Human Language Technologies

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task



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# Outline

## 1 Neural Networks NERC

## 2 General Structure

## 3 Detailed Structure

- Learner
- Classifier
- Required functions

## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

## Session 2 - NERC using neural networks

### Assignment

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

Write a python program that parses all XML files in the folder given as argument and recognizes and classifies drug names.

The program must use a neural network approach.

```
$ python3 ./ml-NER.py data/Devel/
```

```
DDI-DrugBank.d278.s0|0-9|Enoxaparin|drug
```

```
DDI-DrugBank.d278.s0|93-108|pharmacokinetics|group
```

```
DDI-DrugBank.d278.s0|113-124|eptifibatide|drug
```

```
DDI-MedLine.d88.s0|15-30|chlordiazepoxide|drug
```

```
DDI-MedLine.d88.s0|33-43|amphetamine|drug
```

```
DDI-MedLine.d88.s0|49-55|cocaine|drug
```

```
DDI-MedLine.d88.s1|82-95|benzodiazepine|drug
```

```
...
```

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## 2 General Structure

## 3 Detailed Structure

- Learner
- Classifier
- Required functions

## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

# General Structure

The general structure is basically the same than for the traditional ML approach:

- Two programs: one learner and one classifier.
- The learner loads the training (Train) and validation (Devel) data, formats/encodes it appropriately, and feeds the model with the data plus its ground truth.
- The classifier loads the test data, formats/encodes it in the same way that was used in training, and feeds it to the model to get a prediction.

In the case of NN, we don't need to extract features (though we do need some encoding)

# Outline

## 1 Neural Networks NERC

## 2 General Structure

## 3 Detailed Structure

- Learner
- Classifier
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## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

# Outline

## 1 Neural Networks NERC

## 2 General Structure

## 3 Detailed Structure

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- Classifier
- Required functions

## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Learner

Core task

# Learner - Main program

```
1 def learn(traindir, validationdir, modelname) :
2     '''
3     learns a NN model using traindir as training data, and validationdir
4     as validation data. Saves learnt model in a file named modelname
5     '''
6     # load train and validation data in a suitable form
7     traindata = load_data(traindir)
8     valdata = load_data(validationdir)
9
10    # create indexes from training data
11    max_len = 100
12    idx = create_indexes(traindata, max_len)
13
14    # build network
15    model = build_network(idx)
16
17    # encode datasets
18    Xtrain = encode_words(traindata, idx)
19    Ytrain = encode_tags(traindata, idx)
20    Xval = encode_words(valdata, idx)
21    Yval = encode_tags(valdata, idx)
22
23    # train model
24    model.fit(Xtrain, Ytrain, validation_data=(Xval, Yval))
25
26    # save model and indexes, for later use in prediction
27    save_model_and_indexes(model, idx, modelname)
```

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Learner

Core task



# Outline

## 1 Neural Networks NERC

## 2 General Structure

## 3 Detailed Structure

- Learner
- **Classifier**
- Required functions

## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure  
Classifier

Core task

# Classifier - Main program

```
1 def predict(modelname, datadir, outfile) :
2     '''
3     Loads a NN model from file 'modelname' and uses it to extract drugs
4     in datadir. Saves results to 'outfile' in the appropriate format.
5     '''
6
7     # load model and associated encoding data
8     model, idx = load_model_and_indexes(modelname)
9     # load data to annotate
10    testdata = load_data(datadir)
11
12    # encode dataset
13    X = encode_words(testdata, idx)
14
15    # tag sentences in dataset
16    Y = model.predict(X)
17    # get most likely tag for each word
18    Y = [[idx['tags'][np.argmax(y)] for y in s] for s in Y]
19
20    # extract entities and dump them to output file
21    output_entities(testdata, Y, outfile)
22
23    # evaluate using official evaluator.
24    evaluation(datadir,outfile)
```

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Classifier

Core task

# Outline

## 1 Neural Networks NERC

## 2 General Structure

## 3 Detailed Structure

- Learner
- Classifier
- Required functions

## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Required functions

Core task

# Required functions - load\_data

1 **def** load\_data(datadir)

- **Used by:** Learner, Classifier
- **Input:** Receives a directory containing XML files.
- **Output:** Parses XML files in given directory, tokenizes each sentence, extracts ground truth BIO tags for each token, and returns the dataset as a dictionary. Dictionary keys are the sentence id, and values are the list of token tuples (word, start, end, ground truth).

■ **Example:**

```
>>> load_data('data/Train')
{'DDI-DrugBank.d370.s0': [(('as', 0, 1, '0'), ('differin', 3, 10, 'B-brand'),
                           ('gel', 12, 14, '0'), ..., ('with', 343, 346, '0'),
                           ('caution', 348, 354, '0'), ('.', 355, 355, '0'))],
 'DDI-DrugBank.d370.s1': [(('particular', 0, 9, '0'), ('caution', 11, 17, '0'),
                           ('should', 19, 24, '0'), ..., ('differin', 130, 137, 'B-brand'),
                           ('gel', 139, 141, '0'), ('.', 142, 142, '0'))],
 ...
}
```

Use XML parsing and tokenization functions from previous exercises

# Required functions - create\_indexes

1 **def** create\_indexes(datadir, max\_length)

■ **Used by:** Learner

■ **Input:** Receives a dataset produced by load\_data, and the maximum length in a sentence

■ **Output:** Creates a set of words seen in the data and a set of BIO tags. Enumerates those sets, assigning a unique integer to each element. Returns these mappings in a single dictionary, with an additional entry for the given max\_length value.

■ **Example:**

```
>>> create_indexes(traindata)
{'words': {'<PAD>':0, '<UNK>':1, '11-day':2, 'murine':3, 'criteria':4,
          'stroke':5, ... 'carbidopa-levodopa':8510, 'generation':8511,
          'terfenadine*': 8512 },
 'tags': {'<PAD>':0, 'B-group':1, 'B-drug_n':2, 'I-drug_n':3, 'O':4,
          'I-group':5, 'B-drug':6, 'I-drug':7, 'B-brand':8, 'I-brand':9},
 'maxlen' : 100 }
```

Add a <PAD> code to both 'words' and 'tags' indexes, with value 0. Add also an <UNK> code to 'words' with value 1. The coding of the rest of the words or tags is arbitrary.

# Required functions - build\_network

```
1 def build_network(idx) :
2     '''
3     Used by: Learner
4     Input: Receives the index dictionary with the encodings of words and
5           tags, and the maximum length of sentences.
6     Output: Returns a compiled Keras neural network
7     '''
8     # sizes
9     n_words = len(idx['words'])
10    n_tags = len(idx['tags'])
11    max_len = idx['maxlen']
12
13    # create network layers
14    inp = Input(shape=(max_len,))
15    ## ... add missing layers here ... #
16    out = # final output layer
17
18    # create and compile model
19    model = Model(inp,out)
20    model.compile() # set appropriate parameters (optimizer, loss, etc)
21
22    return model
```

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Required functions

Core task

# Required functions - build\_network

```
1  def build_network(idx) :
```

- LSTMs are useful for sequence tagging tasks such as NER.
- You will need to add one Embedding layer after the input, that is where the created indexes will become handy.
- You can base your model in these examples: [1],[2],[3],[4],[5],[6]  
*Note*: some instructions may require to be adapted, depending on your Keras version.  
*Note*: you don't need to follow the **whole** example, only the network construction part.

# Required functions - encode\_words

```
1  def encode_words(dataset, idx) :
```

- **Used by:** Learner, Classifier
- **Input:** Receives a dataset produced by `load_data`, and the index dictionary produced by `create_indexes`
- **Output:** Returns the dataset as a list of sentences. Each sentence is a list of integers, corresponding to the code of each **word** in the sentence. If the word is not in the index, the code for <UNK> is used. If the sentence is shorter than `max_len` it is padded with the code for <PAD>.
- **Example:**

```
>>> encode_words(traindata,idx)
[ [6882 1049 4911 ... 0 0 0]
  [2290 7548 8069 ... 0 0 0]
  ...
  [5964 5183 3519 ... 0 0 0]
  [2002 6582 7518 ... 0 0 0] ]
```



# Required functions - encode\_tags

```
1  def encode_tags(dataset, idx) :
```

- **Used by:** Learner
- **Input:** Receives a dataset produced by `load_data`, and the index dictionary produced by `create_indexes`
- **Output:** Returns the dataset as a list of sentences. Each sentence is a list of integers, corresponding to the code of the **BIO tag** for each word. If the sentence is shorter than `max_len` it is padded with the code for `<PAD>`.

- **Example:**

```
>>> encode_tags(traindata,idx)
[ [ [6] [9] [6] ... [0] [0] [0] ]
  [ [6] [6] [6] ... [0] [0] [0] ]
  ...
  [ [6] [8] [6] ... [0] [0] [0] ]
  [ [6] [6] [6] ... [0] [0] [0] ] ]
```

*Note:* The shape of the produced list may need to be adjusted depending on the architecture of your network and the kind of output layer you use.

# Required functions - Model saving and loading

```
1  def save_model_and_indexs(model, idx, filename) :
```

- **Used by:** Learner
- **Input:** Receives a trained model, an index dictionary, and a string.
- **Output:** Stores the model in a file named `filename.nn`, and the indexs in a file named `filename.idx`

```
1  def load_model_and_indexs(filename) :
```

- **Used by:** Classifier
- **Input:** Loads a model from `filename.nn`, and the indexs from `filename.idx`.
- **Output:** Returns the loaded model and indexs

*Note:* Use Keras `model.save` and `keras.models.load_model` functions to save/load the model.

*Note:* Use your preferred method (pickle, plain text, etc) to save/load the index dictionary.

# Required functions - output\_entities

```
1  def output_entities(dataset, preds, outfilename)
```

- **Used by:** Classifier
- **Input:** Receives a dataset produced by `load_data`, and the corresponding tags predicted by the model.
- **Output:** Prints the detected entities in file *outfilename* in the appropriate format for the evaluator: one line per entity, fields separated by '|', field order: *id*, *offset*, *name*, *type*.
- **Example:**

```
>>> output_entities(dataset, preds, filename)
DDI-DrugBank.d283.s4|14-35|bile acid sequestrants|group
DDI-DrugBank.d283.s4|99-104|trikor|group
DDI-DrugBank.d283.s5|22-33|cyclosporine|drug
DDI-DrugBank.d283.s5|196-208|fibrate drugs|group
DDI-DrugBank.d283.s4|14-35|bile acid sequestrants|group
DDI-DrugBank.d283.s5|220-225|trikor|group
...
```

**Note:** Most of this function can be reused from NER-ML exercise.

# Required functions - evaluation

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Required functions

Core task

```
1  def evaluation(datadir, outfile)
```

- **Used by:** Classifier
- **Input:** Receives a directory with ground truth data, and a file with entities extracted by the model
- **Output:** Runs the official evaluator and gets the results

*Note:* Reuse this function from previous exercises

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## 2 General Structure

## 3 Detailed Structure

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- Classifier
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## 4 Core task

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

# Build a good NN-based drug NERC

**Strategy:** Experiment with different NN architectures and possibilities.

Some elements you can play with:

- Embedding dimension
- Number of LSTM units
- Used optimizer
- Number and kind of layers
- Adding a CRF layer after the LSTM
- Using lowercased and/or non lowercased word embeddings
- Initializing embeddings with available pretrained model
- Using extra input (e.g. suffix embeddings, prefix embeddings, PoS embeddings, ...)
- ...

# Build a good NN-based drug NERC

## Warnings:

- Neural Network training uses randomization, so different runs of the same program will produce different results. For repeatable results, use a random seed.
- During training, Keras reports accuracy on training set and on validation set. Those values are usually over 90%. However, this is due to the fact that most of the words have tag “0” (non-drug). 90% accuracies correspond to  $F_1$  values around 25%. To get a reasonable  $F_1$ , accuracy must reach about 97%. To precisely evaluate how your model is doing, do not rely on reported accuracy: run the classifier on the Development set and use the evaluator.

# Exercise Goals

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

## Goal 5:

Get an overall  $F_1$  score of at least 0.70 on **Devel** dataset.  
(65% is acceptable for a lower grade)



# Deliverables

Neural  
Networks  
NERC

General  
Structure

Detailed  
Structure

Core task

Write one report (max about 5 pages) describing:

- Used architecture
- Performed experiments, tried/discarded/selected options.

The report must include:

- Code for the `build_network` function
- Output of the evaluator on **Devel** and **Test** datasets.

The report must be a PDF file, or a Jupyter notebook (no need that it is are executable, use it only as a presentation support)