

Master in Artificial Intelligence

Advanced Human Language Technologies

Machine
Learning DDI

Relation
Extraction

General
Structure

Resources

Detailed
Structure

Core task

Goals &
Deliverables



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Outline

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Task 2.1 - DDI using machine learning

Assignment

The main program parses input XML file and classifies drug-drug interactions between pairs of drugs, using a machine learning classification algorithm.

```
$ python3 ./ml-DDI.py data/devel/ > result.out
DDI-DrugBank.d398.s0|DDI-DrugBank.d398.s0.e0|DDI-DrugBank.d398.s0.e1|effect
DDI-DrugBank.d398.s0|DDI-DrugBank.d398.s0.e0|DDI-DrugBank.d398.s0.e2|effect
DDI-DrugBank.d211.s2|DDI-DrugBank.d211.s2.e0|DDI-DrugBank.d211.s2.e5|mechanism
...
```

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Relation Extraction

- Relation Extraction is a NLP task, frequently required in Information Extraction applications.
- The goal of the task is to extract relations between entities (previously detected), expressed in the text.
E.g.: `is_CEO_of(Person,Organization)`:
 - *Steve Jobs* was the chairman, the chief executive officer (CEO), and a co-founder of *Apple Inc.*, ...
 - During his career at *Microsoft*, *Bill Gates* held the positions of chairman, chief executive officer (CEO), president and chief software architect.
 - *Mark Zuckerberg* is known for co-founding *Facebook, Inc.* and serves as its chairman, chief executive officer, and controlling shareholder.

Relation Extraction

Other examples:

- **Medical domain:**

- `caused_by(diagnose, drug)`
 - `prescribed_for(drug, diagnose)`
 - `drug_interaction(drug, drug)`

- **Legal domain:**

- `is_suing(Person/Org, Person/Org)`
 - `is_representing(Person, Person/Org)`
 - `is_sentenced_for(Person/Org, Crime)`
 - `is_sentenced_to(Person/Org, Penalty)`

- **Business/Economy:**

- `is_CEO_of(Person, Organization)`
 - `absorbed_by(Organization, Organization)`

- etc.

Relation Extraction

Relation Extraction can be approached as a classical ML classification task, where:

- The objects to be classified are a **text fragment** (sentence, paragraph...) plus a **pair of target entities** in it.
- Each object (*text, entity1, entity2*) is encoded as a feature vector.
- The output class is either **None**, or one **relation type** chosen among a *predefined list*.

Informative enough features are crucial to get good results.

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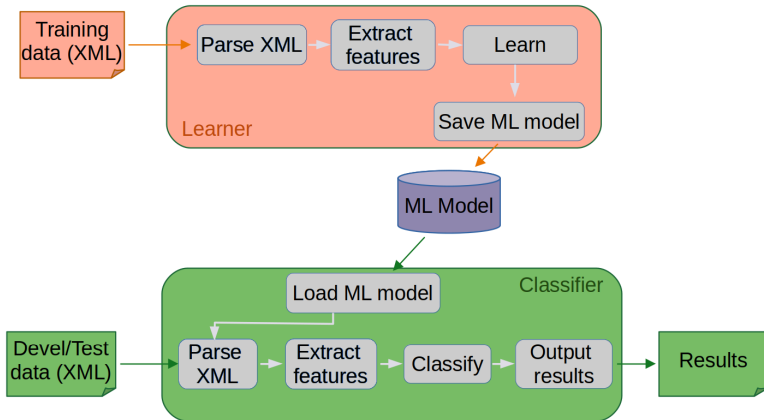
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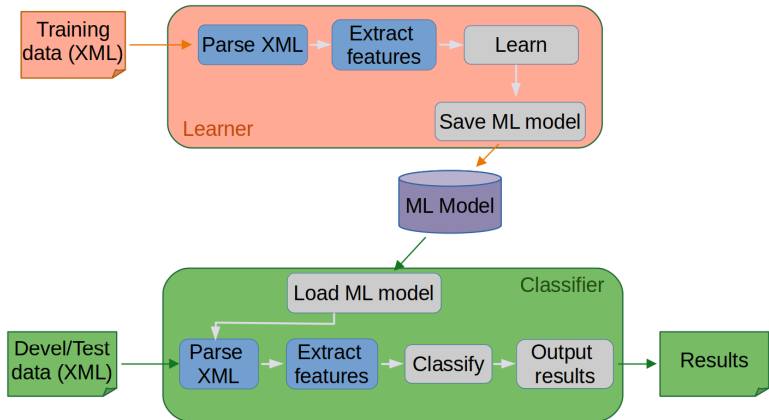
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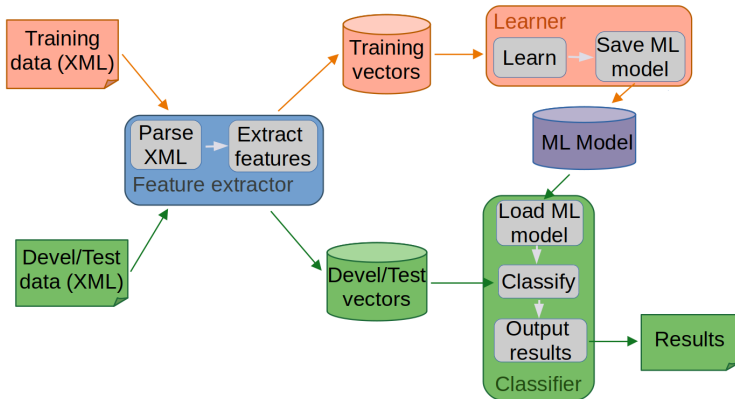
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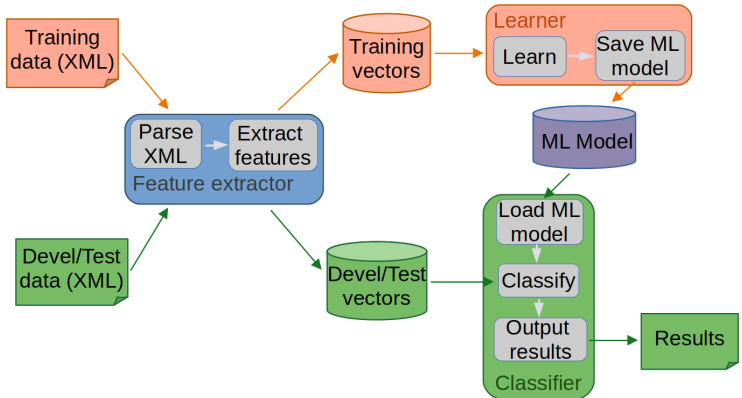
Extracting features is a costly operation, which we do not want to repeat for every possible experiment or algorithm parametrization.

General Structure



Feature extraction process is performed once, out of learning or predicting processes.

General Structure



Feature extraction process is performed once, out of learning or predicting processes.

Thus, we need to write not a single program, but three different components: feature extractor, learner, and classifier.

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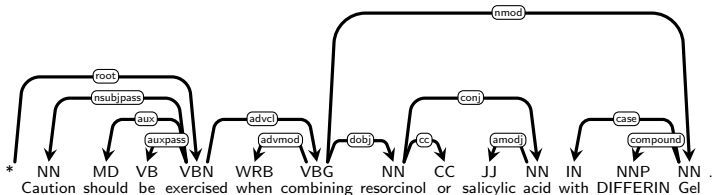
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We will use SpaCy NLP suite, which integrates a [tokenizer](#), a [part-of-speech](#) tagger, and a [dependency parser](#).

- Install [Spacy](#)
- Output is always a list of tokens.
Dependency tree information is inside each token.
Check the [API](#) for details.

Resources - Spacy results

```
nlp = spacy.load("en_core_web_trf")
text = 'Caution should be exercised when combining resorcinol or
        salicylic acid with DIFFERIN Gel'
analysis = nlp(text)
for tk in analysis :
    print(tk.text)    # word form
    print(tk.lemma_)  # lemma
    print(tk.pos_)    # coarse part-of-speech (NOUN, VERB, ADJ, ...)
    print(tk.tag_)    # coarse part-of-speech (NN, NNS, VB, VBZ, VBD, ..)
    print(tk.head_)   # parent token
    print(tk.dep_)    # dependency function wrt parent
    for c in tk.children : ... # iterator over token children nodes
    for c in tk.ancestors : ... # iterator over ancestor chain
etc.
```



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Feature Extractor

The feature extractor:

- Independent program, separated from learner and classifier
- Receives as argument the XML file to encode.
- Prints the feature vectors to given output file

```
$ python3 ./extract_features.py data/devel devel.feats
$ more devel.feats
```

```
DDI-DrugBank.d339.s0 DDI-DrugBank.d339.s0.e0 DDI-DrugBank.d339.s0.e1 null lib=elevated
wib=Elevated lpib=elevated_JJ la2=level wa2=levels lpa2=level_NNS la2=have wa2=have
lpa2=have_VBP la2=be wa2=been lpa2=be_VBN la2=report (...) lpa2=concomitantly_RB path1=NNP
path2=NNP\dep\nsubjpass\compound path=NNP\dep\nsubjpass\compound
DDI-DrugBank.d339.s0 DDI-DrugBank.d339.s0.e0 DDI-DrugBank.d339.s0.e2 null lib=elevated
wib=Elevated lpib=elevated_JJ wib=experience lpib=experience_NN la2=be wa2=is lpa2=be_VBZ
la2=administer wa2=administered lpa2=administer_VBN la2=concomitantly wa2=concomitantly
lpa2=concomitantly_RB path1=NNP path2=NNP\dep\advcl\advcl\nsubjpass
path=NNP\dep\advcl\advcl\nsubjpass
DDI-DrugBank.d339.s0 DDI-DrugBank.d339.s0.e1 DDI-DrugBank.d339.s0.e2 mechanism
lb1=carbamazepine wb1=Carbamazepine wb1=Elevated lpb1=elevated_JJ lib=level wib=levels
lpib=level_NNS lib=have wib=have lpib=have_VBP lib=be wib=been lpib=be_VBN lib=report
wib=reported lpib=report_VBN lib=postmarket wib=postmarketing lpib=postmarket_VBG
lib=experience wib=experience lpib=experience_NN la2=be wa2=is lpa2=concomitantly_RB
path1=compound\nsubjpass_VBN path2=VBN\advcl\advcl\nsubjpass
path=compound\nsubjpass_VBN\advcl\advcl\nsubjpass
...
```

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Feature extraction for NLP

- In most **ML applications**, the feature space is finite and known (e.g. credit scoring, medical diagnose prediction, churn prevention, fraud detection, etc).
- Also, most of the used features are numerical or categorical (income, age, sex, colestherol level, number of receipts returned, etc.)
- Thus, in these ML applications, feature vectors are usually *exhaustive* lists of pairs feature-value.

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Feature extraction for NLP

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BUT...

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Feature extraction for NLP

- In most **ML applications**, the feature space is finite and known (e.g. credit scoring, medical diagnose prediction, churn prevention, fraud detection, etc).
- Also, most of the used features are numerical or categorial (income, age, sex, colestherol level, number of receipts returned, etc.)
- Thus, in these ML applications, feature vectors are usually *exhaustive* lists of pairs feature-value.

BUT...

- In most **NLP applications**, features are related to appearing words, suffixes, prefixes, lemmas, etc.
- Thus, the feature space is huge.

Feature extraction for NLP

- **Example:** We want to encode in a feature the word form for a token.
- We could use a feature (i.e. a single column) `word_form` with thousands of possible values.
- But no ML algorithm is able to deal with categorical features with that amount of possible values, and not enough data will ever be available to learn the behaviour of that feature.
- Thus, features in NLP are typically *binary-valued* (e.g. `word-form-is-THE`, `word-form-is-SOME`, `word-form-is-DOG`, etc.).
That is, a column for each possible value, with boolean value.
- This creates feature spaces with tens or hundreds of thousands of features, but each feature has only two possible values. This is manageable by many ML algorithms.

Feature extraction for NLP

- Thus, in NLP applications, feature vectors are usually *intensive* lists of strings (i.e. listing the names for features with value true, and omitting those with value false), and are stored as *sparse vectors*.
- But, feature spaces of hundreds of thousands of dimensions can not be stored in a table or a matrix.
- However, feature vectors are usually *very sparse*: Each example has only a reduced number of non-zero columns (e.g. even there are thousands of columns like `word-form-is-XXX`, only one of them will be true for a given example).

Sparse vector representation: CSR matrix

- CSR matrix representation consists of storing the coordinates and the value for non-zero elements in the matrix.
- The matrix elements are indexed by their (row,column) coordinates.

$$\begin{pmatrix} 0 & 0 & 3 & 0 & 0 \\ 2 & 0 & 0 & 0 & 0 \\ 0 & 0 & 0 & 0 & 0 \\ 0 & 0 & 1 & 0 & 6 \end{pmatrix} \Rightarrow \begin{pmatrix} \text{row} & \text{col} & \text{val} \\ 0 & 2 & 3 \\ 1 & 0 & 2 \\ 3 & 2 & 1 \\ 3 & 4 & 6 \end{pmatrix}$$

- Each row (of the full matrix) corresponds to a training example, and each column to a feature.
- The CSR matrix is a compressed representation of the same information.
- In the case of DDI, all features are binary (either 0 or 1) so the third column in CSR matrix may be omitted, saving more space.
- A **feature index** is needed to keep the correspondence between feature names and their column number.

Executing the feature extractor

```
$ python3 run.py extract
```

Will process `train.xml` and `devel.xml` and produce `train.feats` and `devel.feats` in directory `predicted`.

If the function `extract_pair_features` is modified (to add or remove features), the datasets must be processed again.

Time to have a look to the `extract_pair_features` and to the `train.feats` file.

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- DDI task is a mere classification, not a sequence prediction.
- For a given sentence and pair of entities in it, the output is just **one** label, not a label per token as in NERC
- Sequence labeling algorithms such as CRFs are overdimensioned, and not straightforward to apply.
- Any classification ML algorithm may be used.
 - Maximum Entropy Classifiers (MEM)
 - Support Vector Machines (SVM)
 - ...

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Learner: MEM (a.k.a. Logistic Regression)

- Install and import scikit-learn
`$ pip install scikit-learn`
- Use provided `train.py` to learn a MEM model.
`$ python3 run.py train MEM`
You may modify learner parameters (loss function, thresholds, learning rates, etc).
- Check performance on development data.
`$ python3 run.py predict MEM`
- scikit-learn implementation of `LogisticRegression` (MEM), admits sparse matrix representations such as CSR (Compressed Sparse Row) matrix.

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Learner: SVM

- Install and import scikit-learn
`$ pip install scikit-learn`
- Use provided train.py to learn a SVM model.
`$ python3 run.py train SVM`
You may modify learner parameters (loss function, thresholds, learning rates, etc).
- Check performance on development data.
`$ python3 run.py predict SVM`
- scikit-learn implementation of SVC (SVM), admits sparse matrix representations such as CSR (Compressed Sparse Row) matrix.

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Classifier

You can apply the learned models to new data and evaluate the results (stored in directory `results`):

```
$ python3 run.py predict MEM devel
$ python3 run.py predict SVM devel
$ python3 run.py predict CRF devel
```

Once a best configuration is selected experimenting with different features, algorithms, and hyperparameters, it can be applied to test data to evaluate generalization ability of the model.

```
$ python3 run.py extract test
$ python3 run.py predict MEM test
$ python3 run.py predict SVM test
$ python3 run.py predict CRF test
```

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Improve provided ML system

Starting point:

- Lemmas-in-between baseline reaches about 25% macro F1.
- Provided ML system reaches about 65%, depending on the algorithm

Task:

- Improve the performance of the provided system by:
 - Adding new features
 - Experimenting with different hyperparameters for the ML algorithms
 - Get insights from training partition. Use devel partition only to evaluate system behaviour.

The goal is not to reach the highest F1 score but to learn through experimenting with different strategies.

Choosing useful features

- Presence of certain *clue verbs* may be indicative of the interaction type.
- Clue verb position (before/inbetween/after) with respect to the target entities.
- Presence of other entities in between.
- Words, lemmas, PoS (or combinations of them) appearing before/inbetween/after the target pair.

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Choosing useful features

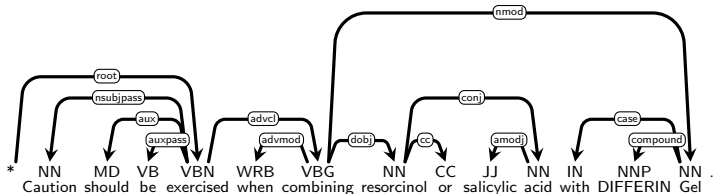
- Presence of certain *clue verbs* may be indicative of the interaction type.
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- Presence of other entities in between.
- Words, lemmas, PoS (or combinations of them) appearing before/inbetween/after the target pair.
- Features encoding information from the syntactic tree.

Choosing useful features

- Presence of certain *clue verbs* may be indicative of the interaction type.
- Clue verb position (before/inbetween/after) with respect to the target entities.
- Presence of other entities in between.
- Words, lemmas, PoS (or combinations of them) appearing before/inbetween/after the target pair.
- **Features encoding information from the syntactic tree.**

Remember: All features are *binary features* (they are either present or not present in a given example) and examples are encoded as *sparse vectors* (lists feature names present in the example)

Choosing useful features



Entities:

e0: resorcinol

e1: salicylic acid

e2: DIFFERIN Gel

Example path features:

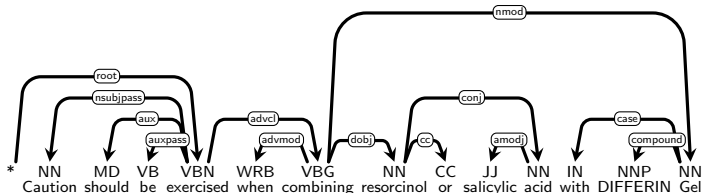
PAIR (e0,e1)

Tree fragment: $e0 \xrightarrow{conj} e1$

(e1 is direct child of e0. The arc is labeled *conj*)

Feature name: **path=conj**>

Choosing useful features



Entities:

e0: resorcinol

e1: salicylic acid

e2: DIFFERIN Gel

Example path features:

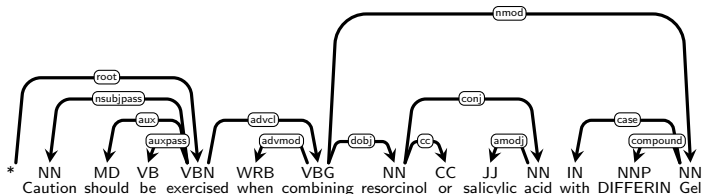
PAIR (e0,e2)

Tree fragment: $e0 \xleftarrow{dobj} \text{combine} \xrightarrow{nmod} e2$

(e0 is direct child of verb "combine" with label *dobj*, and e2 is direct child of the same verb, with label *nmod*)

Feature name: **path=dobj<combine>nmod**

Choosing useful features



Entities:

e0: resorcinol

e1: salicylic acid

e2: DIFFERIN Gel

Example path features:

PAIR (e1,e2)

Tree fragment: $e1 \xleftarrow{conj} \text{resorcinol} \xleftarrow{doobj} \text{combine} \xrightarrow{nmod} e2$

(e1 is *conj* child of "resorcinol", which is under verb "combine" with label *doobj*, and e2 is direct child of the same verb, with label *nmod*)

Feature name: **path=conj<doobj<combine>nmod**

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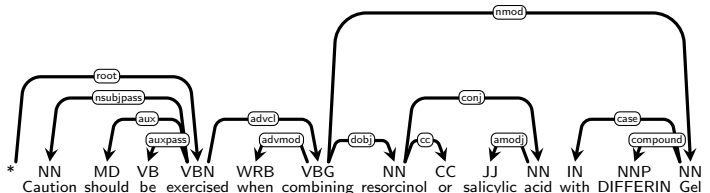
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Choosing useful features



Entities:

e0: resorcinol

e1: salicylic acid

e2: DIFFERIN Gel

Example path features:

PAIR (e1,e2)

Tree fragment: $e1 \xleftarrow{conj} \text{resorcinol} \xleftarrow{dobj} \text{combine} \xrightarrow{nmod} e2$

(e1 is *conj* child of "resorcinol", which is under verb "combine" with label *dobj*, and e2 is direct child of the same verb, with label *nmod*)

Also possible: `path=conj<ENTITY/dobj<combine>nmod`

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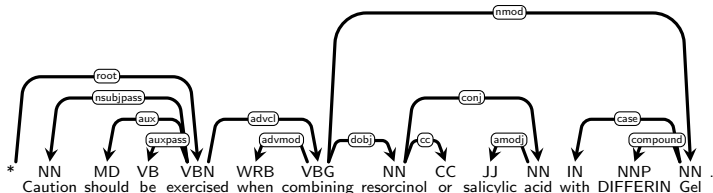
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Choosing useful features



Entities:

e0: resorcinol

e1: salicylic acid

e2: DIFFERIN Gel

Example path features:

PAIR (e1,e2)

Tree fragment: $e1 \xleftarrow{conj} \text{resorcinol} \xleftarrow{dobj} \text{combine} \xrightarrow{nmod} e2$

(e1 is *conj* child of "resorcinol", which is under verb "combine" with label *dobj*, and e2 is direct child of the same verb, with label *nmod*)

Also possible: `path=dobj*<combine>nmod`

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Choosing useful features

Path features may be build in different ways, encoding different information about the tree

- Node words
- Node lemmas
- Node PoS
- Edge labels
- Edge direction
- Direct/indirect dependencies
- ... or any combination of these ...

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Path features may be build in different ways, encoding different information about the tree

- Node words
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- ... or any combination of these ...

Remember: All features are *binary features* (they are either present or not present in a given example) and examples are encoded as *sparse vectors* (lists of present feature names)

Choosing useful features

Some feature possibilities to explore:

- Features detailing the position of words in the sentence (e.g. before E1, between E1 and E2, after E2)
- More general (or more specific) path features (with/without all elements in the path, lemma/tag of each, etc)
- Features encoding specific tree patterns (and maybe their combination with certain verbs)
- Information about the LCS (PoS, lemma, ...)
- Features considering lists of relevant verbs.
- Type of entities in the pair
- Presence of a third entity (in the sentence, in between the target pair, in the path connecting the pair in the tree, ...)
- ...

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Exercise Goals

What you should do:

- Extend your feature extractor. Pay special attention to features encoding **syntactic information**.
- Experiment with different algorithms and parameterizations.
- Keep track of tried features and parameter combinations, and results produced by each.

What you should **NOT** do:

- Use neural network learners. We'll do that later on the course.
- Get insights about errors from **devel** dataset. You have **train** for that. **devel** is only used to evaluate the performance of a given configuration.
- Select features or hyperparameters based on system performance on **test** dataset. You have **devel** for that. **test** is only used to evaluate model generalization ability once the best configuration has been chosen.

Deliverables

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At the end of the DDI task, you will need to deliver a single report on the work carried out on DDI-ML, DDI-NN, and DDI-LLM systems

So, during the development and experimentation on DDI-ML:

- keep track of tried/discarded features
- keep track of used algorithms and parameterizations.
- Record obtained results in the different experiments, and compile the information you'll later need to elaborate the report.