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# **Quoll Documentation**

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Quoll is the name of carnivorous marsupials living in Australia, New Guinea and Tazmania. It is also a python library for running NLP pipelines, based on the [LuigiNLP](#) workflow system. Quoll takes care of the sequence of tasks that are common to basic Machine Learning experiments with textual input: preprocessing, feature extraction, vectorizing, classification and evaluation. Provided that you prepare instances and labels, all these tasks can be ran as a pipeline, in one go. Quoll is built on top of several applications in the [LaMachine](#) Software distribution ([Ucto](#), [Frog](#), [Colibri-core](#), [PyNLPI](#)), as well as popular python packages ([Numpy](#), [Scipy](#) and [Scikit-learn](#)).

Quoll has the following advantages:

- Can run full supervised machine learning pipeline with one command.
- Stores intermediate output of the pipeline.
- Maintains a full log of your experiments.
- Offers various options at each stage of the pipeline.
- If part of the pipeline is already completed, will continue from that point.
- Experiments with different settings can be distinguished based on filenames.



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## The Featurize module

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The Featurize module is the second module in the pipeline, taking care of feature extraction from the output of the [Preprocess](#) module. It makes use of [ColibriCore](#) to count features, and its output forms the input to the [Vectorize](#) module.

### 1.1 Input

*If the input to `Preprocess_` (.txt or .txt\_dir) is given as `inputfile`, this module is ran prior to the Featurize module.*

- inputfile**
  - The featurize module takes preprocessed documents as input. They can come in four formats:
    1. Extension **.tok.txt** - File with tokenized text documents on each line.
    2. Extension **.tok.txt\_dir** - Directory with tokenized text documents (files ending with **.tok.txt**).
    3. Extension **.frog.json** - File with frogged text documents.
    4. Extension **.frog.json\_dir** - Directory with frogged text documents (files ending with **.frog.json**).

### 1.2 Options

- featuretypes**
  - Specify the types of features to extract
  - Options: **tokens, lemmas, pos**
  - lemmas and pos only apply to input with extension *.frog.json* and *.frog.json\_dir*
  - multiple options can be given with quotes, divided by a space (for example: *'tokens lemmas pos'*)

- String parameter; default: **tokens**
- ngrams**
  - Specify the length of the N-grams that you want to include
  - Ngram values should be given within quotes, divided by a space (for example: '1 2')
  - String parameter; default: **'1 2 3'**
- blackfeats**
  - In order to exclude words from the feature space, specify them here
  - Each feature to be excluded should be given within quotes, divided by a space (for example: 'do re mi')
  - Each ngram within the feature space that includes any of the given blackfeats will be removed
  - String parameter; default: **False**
- lowercase**
  - Choose to lowercase all text and lemma features
  - Boolean parameter; default: **False**
- minimum-token-frequency**
  - Option to delete all features that occur below the given threshold
  - Recommended to set to 5 or 10 when applying tfidf or infogain weighting in the Vectorize module
  - Integer parameter; default: **1**

## 1.3 Output

**.features.npz** Binary file in Numpy format, storing the extracted features per document in sparse format

**.vocabulary.txt** File that stores the index of each feature

## 1.4 Overview

--inputfile	--featuretypes	--ngrams	--blackfeats	--lowercase	--minimum-token-frequency	Output
docs.tok.txt	tokens	'1 2 3'	False	True	2	<ul style="list-style-type: none"> <li>• docs.tokens.n_1_2_3.min2</li> <li>• docs.tokens.n_1_2_3.min2</li> </ul>
dos.frog.jsindir	'tokens lemmas pos'	1	'koala kangaroo'	False	10	<ul style="list-style-type: none"> <li>• docs.tokens.lemmas.pos.n</li> <li>• docs.tokens.lemmas.pos.n</li> </ul>



## 1.5 Examples of command line usage

**Extract word Ngrams from tokenized text document, lowercasing them and stripping away token Ngrams that occur less than 5 times**

```
$ luiginlp Featurize --module quoll.classification_pipeline.modules.featurize --inputfile docs.tok.txt --lowercase --minimum-token-frequency 5
```

**Extract lemma and pos Ngrams from directory with frogged texts**

```
$ luiginlp Featurize --module quoll.classification_pipeline.modules.featurize --inputfile docs.frog.jsondir --featuretypes 'lemmas pos'
```

**Frog text document, extract text and pos features and strip away any feature with the word 'snake'**

```
$ luiginlp Featurize --module quoll.classification_pipeline.modules.featurize --inputfile docs.txt --frogconfig /mylama-chinedir/share/frog/nld/frog.cfg --featuretypes 'tokens pos' --blackfeats snake
```



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## The Vectorize module

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The Vectorize module is the third module in the pipeline, taking care of weighting and pruning features based on characteristics in the training data. In contrast to the preliminary **Featurize\_** and **Preprocess** modules, this module requires separate train and test input.

### 2.1 Input

*If the input to Preprocess\_ (.txt or .txt dir) or Featurize (.tok.txt, .tok.txt dir, .frog.json or .frog.json dir) is given as input to traininstances or testinstances, this module is ran prior to the Vectorize module.*

- traininstances**
  - Traininstances takes featurized or vectorized documents as input. They can come in the following formats:
    1. Extension **.features.npz** - Output of the Featurize module.
    2. Extension **.vectors.npz** - Output of the Vectorize module (can be used for vectorizing test documents).
    3. Extension **.csv** - File with feature vectors formatted as comma-separated-values (useful when applying feature extraction that is not accomodated by quoll). When working with '.csv'-input, a file with featurenames should be created that has the same path and name as the '.csv'-file, replacing .csv with .featureselection.txt.
- trainlabels**
  - Extension **.labels** - File with a label per line, should be as many instances as traininstances, where the position of the label corresponds to the instance on the same position.
- testinstances**
  - Like traininstances, test instances takes featurized or vectorized documents as input. They can come in the following formats:
    1. Extension **.features.npz** - Output of the Featurize module.
    2. Extension **.csv** - File with feature vectors formatted as comma-separated-values (useful when applying feature extraction that is not accomodated by

quoll). Can only be used when the input to traininstances is of the same format; the number of columns should be as many as for the traininstances input. Like

## 2.2 Options

*Options for Preprocess\_ and Featurize\_ also apply and are effective in combination with the inputfiles for these modules.*

- weight**
  - Specify the feature weighting
  - Does not work in combination with a ‘.csv’-file
  - Options: **frequency**, **binary**, **tfidf**, **\*\*infogain**
  - For tfidf or infogain, it is recommended to set minimum feature frequency to 5 or 10 in the Featurize module
  - String parameter; default: **frequency**
- prune**
  - Specify the number of features to maintain after pruning
  - Does not work in combination with a ‘.csv’-file
  - Pruning is done by ranking features based on their feature weight (total count of a feature is taken in case of ‘frequency’ and ‘binary’ weighting)
  - Integer parameter; default: 5000
- balance**
  - Choose to balance the number of train instances
  - The number of instances for each class label is decreased to the instance count of the least frequent class
  - Can help in case of strong class skewness
  - Boolean parameter; default: **False**
- delimiter**
  - Specify the delimiter by which columns in the ‘.csv’-file are separated
  - Only applies to ‘.csv’-files
  - String parameter; default: ,
- scale**
  - Option to normalize feature values to the same scale
  - Only applies to ‘.csv’-file
  - Useful in combination with some classifiers, if the features in the ‘.csv’-file are from different sources and have a wide range of values
  - Boolean parameter; default: **False**

## 2.3 Output

**.balanced.features.npz** Balanced instances Only applied when ‘balance’ is chosen

**.balanced.labels** Labels related to balanced instances Only applies when ‘balance’ is chosen

**.balanced.vocabulary** Vocabulary related to balanced instances

## 2.4 Overview

–inputfile	–featuretypes	–ngrams	–blackfeats	–lowercase	–minimum-token-frequency	Output
docs.tok.txt	tokens	‘1 2 3’	False	True	2	<ul style="list-style-type: none"> <li>docs.tokens.n_1_2_3.min2</li> <li>docs.tokens.n_1_2_3.min2</li> </ul>
docs.frog.jsongdir	‘tokens lemmas pos’	1	‘koala kangaroo’	False	10	<ul style="list-style-type: none"> <li>docs.tokens.lemmas.pos.n_1_2_3.min2</li> <li>docs.tokens.lemmas.pos.n_1_2_3.min2</li> </ul>

## 2.5 Examples of command line usage

**Extract word Ngrams from tokenized text document, lowercasing them and stripping away token Ngrams that occur less than 5 times**

```
$ luiginlp Featurize --module quoll.classification_pipeline.modules.featurize --inputfile docs.tok.txt --lowercase --minimum-token-frequency 5
```

**Extract lemma and pos Ngrams from directory with frogged texts**

```
$ luiginlp Featurize --module quoll.classification_pipeline.modules.featurize --inputfile docs.frog.jsongdir --featuretypes 'lemmas pos'
```

**Frog text document, extract text and pos features and strip away any feature with the word ‘snake’**

```
$ luiginlp Featurize --module quoll.classification_pipeline.modules.featurize --inputfile docs.txt --frogconfig /mylama-chinedir/share/frog/nld/frog.cfg --featuretypes 'tokens pos' --blackfeats snake
```



## CHAPTER 3

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### Indices and tables

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- `genindex`
- `modindex`
- `search`