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# Orange3 Text Mining Documentation

*Release 0.1.1*

**Biolab**

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<b>1</b>	<b>Widgets</b>	<b>1</b>
<b>2</b>	<b>Scripting Reference</b>	<b>3</b>
<b>3</b>	<b>Indices and tables</b>	<b>5</b>



## 1.1 Preprocess

The description of the tokenizer goes here.



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## Scripting Reference

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### 2.1 Preprocessor

```
class orangecontrib.text.preprocess.Preprocessor (incl_punct=False, lowercase=True,  
stop_words='english', trans=None,  
min_df=1)
```

Holds pre-processing flags and other information, about stop word removal, lowercasing, text morphing etc.(the options are set via the Preprocess widget).

```
Preprocessor.__init__ (incl_punct=False, lowercase=True, stop_words='english', trans=None,  
min_df=1)
```

#### Parameters

- **incl\_punct** (*boolean*) – Determines whether the tokenizer should include punctuation in the tokens.
- **lowercase** (*boolean*) – If set, transform the tokens to lower case, before returning them.
- **stop\_words** (*'english' or list or None*) – Determines whether stop words should(“english”), or should not(None) be removed. If this is list, it should contain stop-words.
- **trans** – An optional pre-processor object to perform the morphological transformation on the tokens before returning them.

#### Returns

**class** *orangecontrib.text.preprocess.Preprocessor*





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## Indices and tables

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



## Symbols

`__init__()` (orangecontrib.text.preprocess.Preprocessor  
method), 3

## P

Preprocessor (class in orangecontrib.text.preprocess), 3