
uparma-py Documentation

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Contents

1	uparma-py - the Python framework for the uparma project	1
2	Indices and tables	5

uparma-py - the Python framework for the uparma project

UPARMA - Universal Parameter Mapper for Proteomics

1.1 Installation

Use pip to install, use virtualenv for maximum convenience:

```
user@localhost$ pip install -r requirements
```

1.2 Testing

Use pytest to run tests:

```
user@localhost$ pytest
```

1.2.1 Uparma - The Universal parameter mapper class in Python

class uparma.uparma.**UParma** (*refresh_jsons=False, source_style='ursgal_style_1'*)
Universal Parameter Mapper Class

Keyword Arguments

- **refresh_jsons** (*bool*) – indicates if uparma jsons should be pulled from central repo (<https://github.com/uparma/uparma-lib>) or not. Note that if jsons are not available in uparma folder, option is overridden and set to True.
- **source_style** (*str*) – Convenience when mapper is used to map from the same source style every time, in which case self.convert can be used instead of self.translate

convert (*param_dict*, *target_style=None*)

Convenient wrapper to translate params with the source style defined during init.

Calls self.translate with `source_style = self.source_style`

translate (*param_dict*, *source_style=None*, *target_style=None*)

Translate `param_dict` from source style into target style.

Keyword Arguments

- **param_dict** (*dict*) – dict containing parameter and value in a given style
- **source_style** (*str*) – style of the input format
- **target_style** (*str*) – style to which the parameters should be translated to.

Returns

dict with the translated key and values for the input dict with additional information in `self.details` (see below).

Return type `translated_params` (dict-like)

For example an input in ursgal style:

```
{
    "precursor_mass_tolerance_unit": "ppm",
    "min_pep_length" : 8
}
```

can be converted to msgfplus style, yielding:

```
{
    '-minLength' : 8,
    '-t' : 'ppm'
}
```

the return object is a dict-like structure which holds additional detailed information accessible via `self.details`. For the example above, `self.details` looks like:

```
{
    'min_pep_length': {
        'source_key': 'min_pep_length',
        'source_style': 'ursgal_style_1',
        'source_value': 8,
        'target_key': '-minLength',
        'target_style': 'msgfplus_style_1',
        'target_value': 8
    },
    'precursor_mass_tolerance_unit': {
        'source_key': 'precursor_mass_tolerance_unit',
        'source_style': 'ursgal_style_1',
        'source_value': 'ppm',
        'target_key': '-t',
        'target_style': 'msgfplus_style_1',
        'target_value': 'ppm'
    }
}
```

1.2.2 Examples

Get parameter information

`get_param_info.main(parameter)`

Get uparma information for a specific parameter name. Description, value translation and uparma id is extracted.

usage:

`./get_param_info.py <parameter_name>`

E.g.:

`./get_param_info.py 'spectrum, parent monoisotopic mass error units'`

The example provides the name of the monoisotopic mass error units of the X!Tandem engines. This name is mapped back to translation styles matching this name.

CHAPTER 2

Indices and tables

- [genindex](#)
- [modindex](#)
- [search](#)

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C

`convert()` (`uparma.uparma.UParma` method), 1

M

`main()` (in module `get_param_info`), 3

T

`translate()` (`uparma.uparma.UParma` method), 2

U

`UParma` (class in `uparma.uparma`), 1