# Improving \$ click\_library

Architecture of large projects in bioinformatics final project

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## What is \$ click\_

Command Line Interface Creation Kit

- python package for creating command line interfaces
- arbitrary nesting of commands
- automatic help page generation
- supports lazy loading of subcommands at runtime

## Click program example

import click

```
@click.command()
@click.option('--count', default=1, help='Number of greetings.')
@click.option('--name', prompt='Your name',
        help='The person to greet.')
def hello(count, name):
  """Simple program that greets NAME for a total of COUNT times."""
  for x in range(count):
    click.echo(f"Hello {name}!")
if __name__ == '__main__':
  hello()
```

## Output

\$ python hello.py --count=3

Your name: John

Hello John!

Hello John!

Hello John!

\$ python hello.py --help Usage: hello.py [OPTIONS]

Simple program that greets NAME for a total of COUNT times.

#### Options:

- --count INTEGER Number of greetings.
- --name TEXT The person to greet.
- --help Show this message and exit.

## Parameter Types

#### Arguments

- arguments can do less than options
- accept an arbitrary number of arguments
- convenient to use

#### Options

- automatic prompting for missing input
- act as flags (boolean or otherwise)
- option values can be pulled from environment variables, arguments can not
- options are fully documented in the help page, arguments are not

## Why choose click?

Click vs Argparse, Docopt etc.

- Click does not just parse, it also dispatches to the appropriate code
- Click has strong information available for all parameters and commands,
- Click has a strong understanding of what types are, and it can give the user consistent error messages
- Click has enough meta information available for its whole program to evolve without forcing developers to adjust their programs

## Click as a useful tool for bioinformaticians

#### **Example of usage**

- The program that connects to the BLAST based on the given sequences
- Returns the result of program operation as a file in .fasta format

```
@click.command()
@click.argument("sequence_file")
@click.argument("output")
@click.option("-i", "--identity", "ident", default=0.9,
              help="Minimal percent identity used during searching the NCBI database, Input range: <0, 1>. Default: 0.9.
@click.option("-e", "--e_value", "e", default=10e-10,
              help="E-value used during searching the NCBI database. Default: 10e-10.")
    run(sequence_file: str, output: str, ident: float, e: float):
    ....
                                                                            Usage: extend.py [OPTIONS] SEQUENCE_FILE OUTPUT
    sequence_file: Path to a fasta format file with a protein sequence.
                                                                               sequence_file: Path to a fasta format file with a protein sequence.
    output: Path to a fasta format output file.
                                                                              output: Path to a fasta format output file.
                                                                            Options:
                                                                               -i, --identity FLOAT Minimal percent identity used during searching the
                                                                                                    NCBI database. Input range: <0, 1>. Default: 0.9.
                                                                               -e, --e_value FLOAT
                                                                                                    E-value used during searching the NCBI database.
                                                                                                    Default: 10e-10.
                                                                                                    Show this message and exit.
                                                                               --help
```

## Our ideas for improvement

- help parameter for @click.argument decorator to preserve an UI consistency
- 2. Parameter for **@click.argument** decorator to give a possibility of creating a Python argument differed from an argument name displayed in a help message Allows to avoid shadowing built-in names, e.g. when argument has to be named "filter"

## @click.arguemnt decorator

#### **Basic:**

```
@click.command()
@click.argument('filename')
def touch(filename):
    """Print FILENAME."""
    click.echo(filename)
$ touch foo.txt
foo.txt
```

#### Variadic:

```
@click.command()
@click.argument('src', nargs=-1)
@click.argument('dst', nargs=1)
def copy(src, dst):
  """Move file SRC to DST."""
  for fn in src:
     click.echo(f"move {fn} to folder {dst}")
```

\$ copy foo.txt bar.txt my\_folder move foo.txt to folder my\_folder move bar.txt to folder my\_folder

## Challenges

- Massive documentation
- Learning about the existing code
- The improvement could not affect the current user scripts

davidism Merge pul	l request pallets#2301 from p 13 days	ago 🕲 <b>2,186</b>
.github	Bump actions/cache from 3.0.2 to 3.0.3	13 days ago
artwork	Initial commit	8 years ago
docs	Update quickstart.rst	2 months ago
examples	Merge remote-tracking branch 'origin/8	12 months ago
requirements	update requirements	2 months ago
src/click	Merge branch '8.1.x'	2 months ago
tests	disallow use of is_flag and multiple in opt	2 months ago
.editorconfig	add EditorConfig	2 years ago
gitignore	delete directory .DS_Store (pallets#1938)	13 months ago
.pre-commit-confi	update requirements	2 months ago
.readthedocs.yaml	pin os and python version in rtd build	6 months ago
CHANGES.rst	Merge branch '8.1.x'	2 months ago
CODE_OF_CONDU	Create CODE_OF_CONDUCT.md	3 years ago
CONTRIBUTING.rst	Improve the contributing guide	13 months ago
LICENSE.rst	standardize license	3 years ago
MANIFEST.in	add typing annotations	14 months ago
README.rst	update pip link	10 months ago
setup.cfg	Merge branch '8.0.x'	5 months ago
🖺 setup.py	install importlib_metadata on Python < 3.8	13 months ago
🖰 tox.ini	drop Python 3.6	7 months ago

### GitHub workflow

- 1. First time setup
  - Configure git
  - Clone main repository
  - Create a virtualenv
  - Upgrade pip and setuptools
  - Install the development dependencies
  - Install the pre-commit hooks
- 2. Start coding
  - Push your commits to your fork on GitHub
- 3. Run the tests
- 4. Run the test coverage
  - Generate the report
- 5. Build the docs

#### How to contribute to Click

Thank you for considering contributing to Click!

#### **Support questions**

Please don't use the issue tracker for this. The issue tracker is a tool to address bugs and feature requests in Click itself. Use one of the following resources for questions about using Click or issues with your own code:

- The #get-help channel on our Discord chat: https://discord.gg/pallets
- The mailing list flask@python.org for long term discussion or larger issues.
- Ask on Stack Overflow. Search with Google first using: site:stackoverflow.com python click {search term, exception message, etc.}

#### **Reporting issues**

Include the following information in your post:

- Describe what you expected to happen.
- If possible, include a minimal reproducible example to help us identify the issue. This also helps check that the issue is not with your own code.
- Describe what actually happened. Include the full traceback if there was an exception.
- List your Python and Click versions. If possible, check if this issue is already fixed in the latest releases or the latest code in the repository.

#### Submitting patches

If there is not an open issue for what you want to submit, prefer opening one for discussion before working on a PR. You can work on any issue that doesn't have an open PR linked to it or a maintainer assigned to it. These show up in the sidebar. No need to ask if you can work on an issue that interests you.

## Results

#### First assumption

Before After

```
@click.command()
@click.command()
                                                           @click.argument('argument', help="It's a feature!")
@click.argument('argument')
@click.option('-o', "--option", "opt", help='Option help text')
                                                           @click.option('-o', "--option", "opt", help='Option help text')
def run(argument, opt):
                                                           def run(argument, opt):
                                                               print(argument, opt)
   argument: Argument help text.
   print(argument, opt)
                                                            Usage: example.py [OPTIONS] ARGUMENT
Usage: example.py [OPTIONS] ARGUMENT
                                                            Arguments:
                                                              ARGUMENT It's a feature! [required]
  argument: Argument help text.
                                                            Options:
Options:
                                                              -o, --option TEXT Option help text
  -o, --option TEXT Option help text
                                                              --help
                                                                                   Show this message and exit.
                      Show this message and exit.
  --help
```

### Results

First assumption cd.

```
@click.command()
@click.argument('argument', help="It's a feature!", hidden=True)
@click.option('-o', "--option", "opt", help='Option help text')
idef run(argument, opt):
   print(argument, opt)
Usage: example.py [OPTIONS] ARGUMENT
Options:
  -o, --option TEXT Option help text
                   Show this message and exit.
  --help
```

# Results Second assumption

Before

```
@click.command()
@click.argument('argument')
@click.option('-o', "--option", "opt", help='Option help text')
idef run(argument, opt):
   argument: Argument help text.
   print(argument, opt)
Usage: example.py [OPTIONS] ARGUMENT
  argument: Argument help text.
Options:
  -o, --option TEXT Option help text
  --help Show this message and exit.
```

#### After

```
@click.command()
@click.argument('argument', "arg")
@click.option('-o', "--option", "opt", help='Option help text')
idef run(arg, opt):
   print(arg, opt)
Usage: example.py [OPTIONS] ARGUMENT
Options:
  -o, --option TEXT Option help text
                      Show this message and exit.
  --help
```

## **Tests**

### PASSED!

platform linux Python 3.9.7, pytest-7.1.2, pluggy-1.0.0 rootdir: /home/macius/PycharmProjects/click/click, configfile: setup.cfg, testpaths: tests plugins: anyio-2.2.0 collected 598 items	
tests/test_arguments.py	
tests/test_basic.py	
tests/test_chain.pyx	
tests/test_command_decorators.py	
tests/test_commands.py	
tests/test_compat.py .	
tests/test_context.py	[ 26%
tests/test_custom_classes.py	[ 27%
tests/test_defaults.py	[ 27%
tests/test_formatting.py	[ 30%
tests/test_imports.py	[ 30%
tests/test_info_dict.py	[ 34%
tests/test_normalization.py	[ 34%
tests/test_options.py	[ 33%
tests/test_parser.py tests/test_shell_completion.py	
tests/test_termui.pysssssssssssssssssssssssssssssss	[ 01/ [ 74%
tests/test_testing.py	[ 78%
tests/test_types.py	F 84%
tests/test_utils.py	[ 15% [ 17% [ 18% [ 22% [ 22% [ 26% [ 27% [ 30% [ 30% [ 34% [ 34% [ 34% [ 53% [ 55% [ 61% [ 74% [ 78% [ 78%

### References

https://click.palletsprojects.com/en/8.1.x/