Software Documentation

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Disclaimer: All pictures used are from random searches of the web and for educational purpose. They might be subject to specific licenses and should be checked before using further

| | tter how good your softvole will not use it. | vare is, because if the | documentation is not | good |
|---------------|--|-------------------------|----------------------|------|
| -Daniele Proc | cida | | | |
| | | | | |

Documentation: Outline

Introduction

Coding Style

Documentation

Best practices

Reflection



Intro: how many keystrokes do you have left?

Let's assume you'll live to at least **90**. That's **70** years left. We'll assume you spend **half your working day** typing full speed. That's **4** hours a day of nothing but typing. There's an average of **5** letters in a word in English.

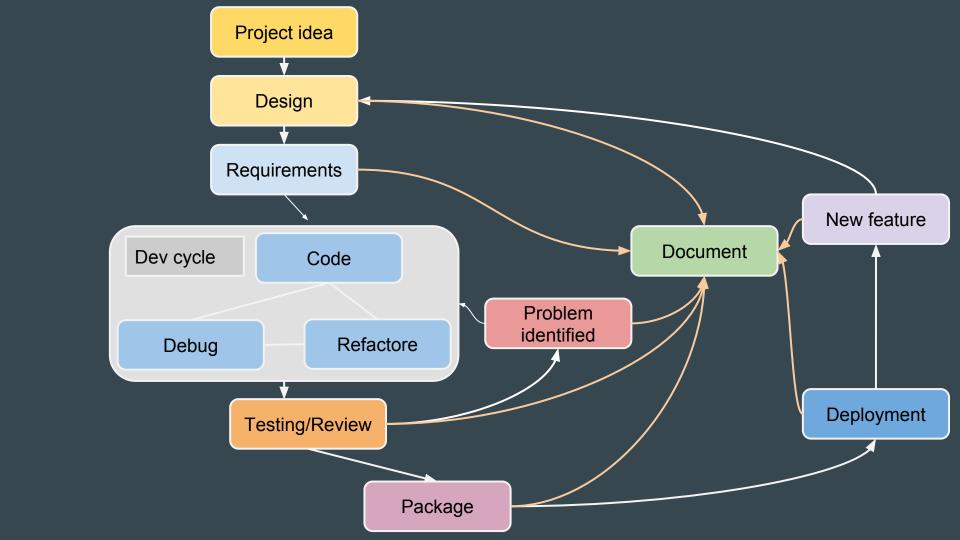
That means, in your remaining lifetime you have these left in your hands:

- **80,639,999** Keystrokes Left
- **575,999** Tweets Left
- 26 Novels Left
- 161 Computer Programs Left
- 8,063 Love Letters Left

Or

403,199 Emails to your boss left

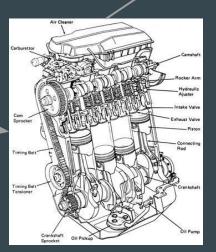
where do you want the gift of your keystrokes to go?

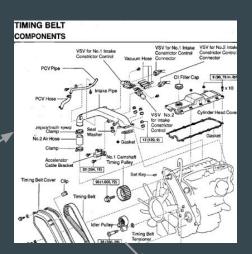


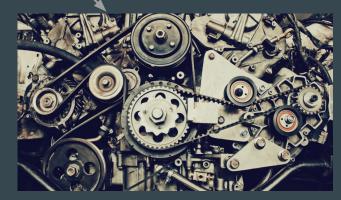
Introduction: levels of detail











Introduction: Target your Audience

Code documentation

Detailed information on how things work

2

Technical

API documentation



Architecture

How the software works in greater detail, and how different components interact



Manual

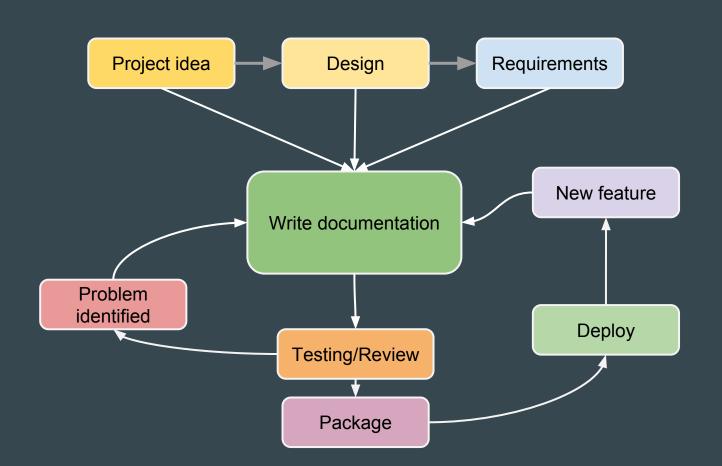
High level usage of software, eg: installation or compiling



Website

To attract uses





Coding Style: Standards & guidelines

Pep8: Style Guide for Python Code

Pep20: The Zen of Python

Google Python Style Guide

python-guide.org style guide

Docstring Conventions



Coding Style: Find your personal style

Coding Style: Valid does not mean good

```
x = \{ 'a': 37, 'b': 42, \}
'c':927}
y = 'hello ''world'
z = 'hello '+'world'
a = 'hello {}'.format('world')
class foo ( object ):
  def f
       (self ):
           37*-+2
   return
  def q(self, x, y=42):
     return y
def f (a):
 return 37+-+a[42-x : y**3]
```

```
x = \{ 'a': 37, 'b': 42, 'c': 927 \}
y = 'hello ' 'world'
z = 'hello ' + 'world'
a = 'hello {}'.format('world')
class foo (object):
    def f(self):
        return 37 * -+2
    def g(self, x, y=42):
        return y
def f(a):
    return 37 + -+a[42 - x:y**3]
```

Coding Style: Hands on pep8 checking

Use the tool on some of your python code:

http://pep8online.com

What errors/warnings do you see?

Do you agree with the errors?

Documentation: What are useful comments

```
# add 1 to counter
i += 1
# Calculate area of circle
area = pi*radius*radius
#cannot be bothered with this right now
# state = set state( input1, current state())
state = 1
// Here be dragons. Thou art forewarned
images.sort(ImageSprite::image path comparator,5,3);
```

Documentation: Python docstring

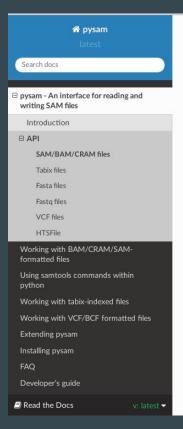
#function help:
help(max)

```
def function with pep484 type annotations(param1: int, param2: str) -> bool:
     """Example function with PEP 484 type annotations.
     Args:
         param1 (int): The first parameter.
         param2 (str): The second parameter.
     Returns:
         The return value. True for success, False otherwise.
         ValueError: If `param2` is equal to `param1`.
   11 11 11
# Class help:
import string
help(string)
```



Documentation: Integrated documentation (sphinx)

```
test documentation master file
word counter: count words in files
:Author: Kim Brugger
:Date: |today|
:Version: |version|
Word counter is a python package/module ...
.. toctree::
   :maxdepth: 1
   :caption: Contents:
```



access to genomic coordinates.

find introns(self, read iterator)

Return a dictionary {(start, stop): count} Listing the intronic sites in the reads (identified by 'N' in the cigar strings), and their support (= number of reads).

read_iterator can be the result of a .fetch(...) call. Or it can be a generator filtering such reads. Example samfile.find_introns((read for read in samfile.fetch(...) if read.is_reverse)

find_introns_slow(self, read_iterator)

Return a dictionary $\{(\text{start, stop}): \text{ count}\}\ \text{Listing the intronic sites in the reads (identified by 'N' in the cigar strings), and their support (= number of reads).}$

read_iterator can be the result of a .fetch(...) call. Or it can be a generator filtering such reads. Example samfile.find_introns((read for read in samfile.fetch(...) if read.is_reverse)

get_index_statistics(self)

return statistics about mapped/unmapped reads per chromosome as they are stored in the index.

Returns: list - 'mapped', 'unmapped' and 'total'.

Return type: a list of records for each chromosome. Each record has the

attributes 'contig',

Best practices: Repository readme file

Project Title

- Overview
- Getting Started
 - Prerequisites
 - Installing
 - Testing the installation
- Deployment
 - Configuration
 - Security
- Contributing
- Versioning
- Authors
 - Contributors
 - Acknowledgments
- <u>Licence</u>



Best practices: Distribution of documentation

readthedocs.com

Github.io

https://sourceforge.net/

Dedicated project webpage



Best practices: Look at Examples (20 min):

https://github.com/pysam-developers/pysam

https://github.com/pallets/flask

https://github.com/django/django

https://github.com/pandas-dev/pandas

https://github.com/google/yapf

Documentation: reflection & thoughts

What is your current practice?

How could you improve it?

How do you find the documentation of other projects?

Any good practices?



Various resources

https://softdev4research.github.io/4OSS-lesson/

https://semver.org/

https://choosealicense.com/licenses/

https://sphinxcontrib-napoleon.readthedocs.io/en/latest/example_google.html

http://www.sphinx-doc.org/en/master/