

---

**cellseg**

***Release 0.1.0***

**Nelson Gonzabato**

**Sep 23, 2021**



**CONTENTS:**

<b>1</b>	<b>cellseg: Multiclass Cell Segmentation</b>	<b>1</b>
<b>2</b>	<b>Development stage</b>	<b>3</b>
<b>3</b>	<b>Introduction</b>	<b>5</b>
<b>4</b>	<b>Installation</b>	<b>7</b>
<b>5</b>	<b>Usage</b>	<b>9</b>
5.1	Script mode . . . . .	9
5.2	Programming mode . . . . .	9
<b>6</b>	<b>cellseg</b>	<b>11</b>
6.1	data module . . . . .	11
6.2	model module . . . . .	11
6.3	utils module . . . . .	12
<b>7</b>	<b>cellseg: Multiclass Cell Segmentation</b>	<b>13</b>
<b>8</b>	<b>Contributing to cellseg</b>	<b>15</b>
<b>9</b>	<b>Indices and tables</b>	<b>17</b>
	<b>Python Module Index</b>	<b>19</b>
	<b>Index</b>	<b>21</b>



## **CELLSEG: MULTICLASS CELL SEGMENTATION**



## DEVELOPMENT STAGE

- ☒ Read Tiff Images
- ☒ Read Non Tiff Images
- ☒ Write Data Transformers and Loaders
- ☐ Write functional model plus scripts
- ☐ Modify model weights/layers
- ☐ Read stacked tiff images/videos





## INTRODUCTION

`cellseg` is a PyTorch (`torch`) based deep learning package aimed at multiclass cell segmentation.



## INSTALLATION

```
pip install cellseg
```

Or if you want to build from source

```
git clone git@github.com:Nelson-Gon/cellseg.git
cd cellseg
python setup.py install
```



## 5.1 Script mode

### View images

```
python -m cellseg -d data/train -t "image" -n 4 -s 512
```

### To get help

```
python -m cellseg --help
#usage: __main__.py [-h] -d IMAGE_DIRECTORY -s IMAGE_SIZE -t TARGET -n NUMBER
#
#optional arguments:
#  -h, --help            show this help message and exit
#  -d IMAGE_DIRECTORY, --image-directory IMAGE_DIRECTORY
#                        Path to image directory containing images and
#                        masks/labels
#  -s IMAGE_SIZE, --image-size IMAGE_SIZE
#                        Size of images
#  -t TARGET, --target TARGET
#                        Target images to show
#  -n NUMBER, --number NUMBER
#                        Number of images to show
```

## 5.2 Programming mode

### Importing relevant modules

```
from cellseg.data import DataProcessor
from cellseg.model import CellNet
from cellseg.utils import DataProcessor, show_images
```

### Creating a model object

```
my_model = CellNet()
```

### Load training data

```
train_data = DataProcessor(image_dir="data/train/images", label_dir="data/train/images
↪", image_suffix="tif")
```

### View loaded images or masks

```
show_images(train_data, number = 8, target="image")
```

## **Training**

## CELLSEG

## 6.1 data module

```
class cellseg.data.DataProcessor (image_dir, label_dir, target_size=(512, 512), im-
                                age_suffix='tif')
    Bases: Generic[torch.utils.data.dataset.T_co]
    transform (image, mask)
```

**Parameters**

- **image** – Image to be transformed.
- **mask** – Label/Mask to be transformed.

**Returns** Transformed images that have been randomly cropped, flipped, and resized to the target size.

## 6.2 model module

```
class cellseg.model.CellNet (input_shape=32, channels=1)
    Bases: torch.nn.modules.module.Module
```

**forward** (*x*)

Defines the computation performed at every call.

Should be overridden by all subclasses.

---

**Note:** Although the recipe for forward pass needs to be defined within this function, one should call the `Module` instance afterwards instead of this since the former takes care of running the registered hooks while the latter silently ignores them.

---

```
training: bool
```

## 6.3 utils module

`cellseg.utils.show_images` (*dataset\_object*, *number=None*, *fig\_size=(20, 20)*, *target='image'*)

### Parameters

- **dataset\_object** – An object of class DataProcessor.
- **number** – Number of images to plot
- **target** – Type of images to show. One of “image” or “mask”, defaults to image
- **fig\_size** – Figure size, defaults to (20, 20)

**Returns** A plot showing images or labels



## CELLSEG: MULTICLASS CELL SEGMENTATION

### Version 0.1.0

- Initialized script mode.
- Optimized imports
- Updated docs to reflect proper `DataProcessor` usage.
- Initial tests.
- `DataProcessor` now errors if image and mask/label lengths differ.
- Dropped thresholding methods. Please use `pyautocv` or any other image processing package of your choice.
- Refactored `show_images` to handle follow current `DataProcessor` logic. Fixed a bug that caused rows and column switch.
- `DataProcessor` now returns a dictionary containing an image, its label, and index.
- Reading stacked tiff images is no longer supported for now.
- `dir_type` was dropped in `DataProcessor`. Only provide a directory to `image_dir` containing images and masks.
- Added data from `cytounet`
- Versioning is now automated.
- Updated docs to show that this is a work in progress.
- Updated docs to use explicit imports.

### version 0.0.0

- Preserve name on PyPI
- Fixed issues with `show_images` showing blank images for masks (labels).
- Fixed issues with `uint16` not working with `PIL`.
- `DataProcessor` can now transform images to a given target size.
- Renamed `DataLoader` class to `DataProcessor` to avoid conflicts with `torch.utils.data.DataLoader`
- Added initial simple CNN model with a single layer
- Added `show_images` in `utils.py` to allow quick visualization of a given number of images from a given stack of images.
- Implemented data loaders.
- Conceptualized project



## CONTRIBUTING TO CELLSEG

This document provides guidelines for contributions to `cellseg`.

### Kinds of contribution

- Typo fixes
- Documentation enhancements
- Pull requests

### Fixing typos and enhancing documentation

To fix typos and/or grammatical errors, please edit the corresponding `.py` or `.md` file that generates the documentation.

Please also update the docs using `sphinx`

### Pull Requests

- Please raise an issue for discussion and reproducibility checks at [issues](#)
- Once the bug/enhancement is approved, please create a Git branch for the pull request.
- Make changes and ensure that builds are passing the necessary checks on Travis.
- Update `changelog.md` to reflect the changes made.
- Do the following:

```
bash scripts/mkdocs.sh #projectnamehere
```

- Releasing

```
bash scripts/release.sh
```

The above does the following:

- Makes dist with `python setup.py sdist` at the very minimum. Ensure everything necessary is included in `Manifest.in`.
- Uploads dist to `test.pypi.org` with `twine upload --repository-url https://test.pypi.org/legacy/ dist/*`
- If everything looks good, asks you to upload to `pypi.org` with `twine upload dist/*`

Please note that the ‘`cellseg`’ project is released with a [Contributor Code of Conduct](#). By contributing to this project, you agree to abide by its terms.

[See also](#) for a guide on Sphinx documentation.



## INDICES AND TABLES

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



## PYTHON MODULE INDEX

### C

`cellseg.data`, [11](#)  
`cellseg.model`, [11](#)  
`cellseg.utils`, [12](#)





## INDEX

### C

`CellNet` (*class in cellseg.model*), 11  
`cellseg.data`  
    module, 11  
`cellseg.model`  
    module, 11  
`cellseg.utils`  
    module, 12

### D

`DataProcessor` (*class in cellseg.data*), 11

### F

`forward()` (*cellseg.model.CellNet method*), 11

### M

module  
    `cellseg.data`, 11  
    `cellseg.model`, 11  
    `cellseg.utils`, 12

### S

`show_images()` (*in module cellseg.utils*), 12

### T

`training` (*cellseg.model.CellNet attribute*), 11  
`transform()` (*cellseg.data.DataProcessor method*),  
    11