



TOPAZ – PICK MORE OF
ANYTHING

TOPAZ: REVOLUTIONIZING PARTICLE PICKING IN CRYO-EM

- **Background:** Cryo-electron microscopy (cryo-EM) is crucial for determining protein structures, but manual particle picking is time-consuming and prone to bias.
- **Topaz:** A particle-picking pipeline using convolutional neural networks (CNNs) trained with a positive-unlabeled (PU) learning method.
- **Advantages:**
 - Efficient and accurate, reducing the need for manual effort.
 - Capable of detecting challenging particles, including small, non-globular, and asymmetric ones.
 - Low false-positive rate and minimal need for post-picking curation.
- **Key Features:** Modular, standalone, free, open-source, and compatible with various cryo-EM software suites.


PERFORMANCE AND IMPACT OF TOPAZ IN CRYO-EM

- **High-Resolution Reconstruction:** Topaz enables high-resolution reconstructions with fewer false positives and better representation of particle views.
- **Case Studies:** Demonstrated effectiveness on datasets with difficult particles, such as the Toll receptor and clustered protocadherin.
- **Integration and Efficiency:** Appion Relion and CryoSparc. Runs efficiently on a single GPU computer.

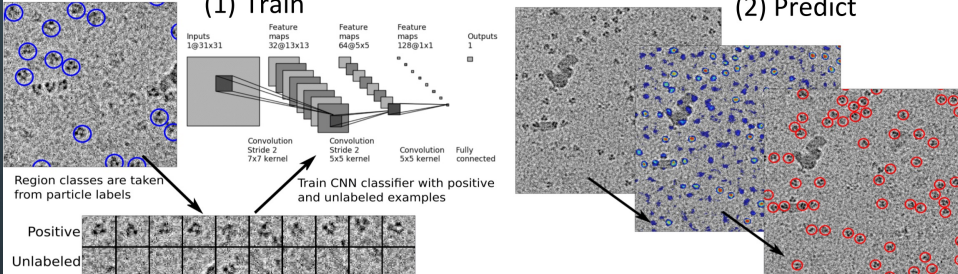
OVERVIEW AND INTERFACE(S)

TOPAZ

Pick more of anything



(1) Train **(2) Predict**

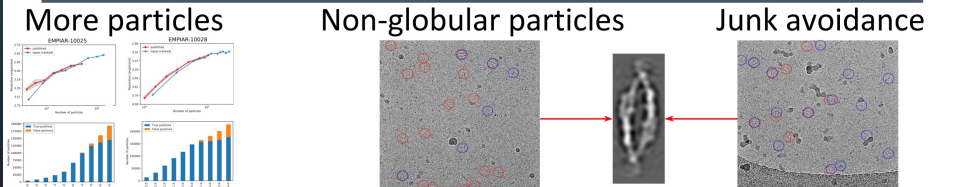


Region classes are taken from particle labels

Train CNN classifier with positive and unlabeled examples


Positive
Unlabeled

More particles **Non-globular particles** **Junk avoidance**



TOPAZ

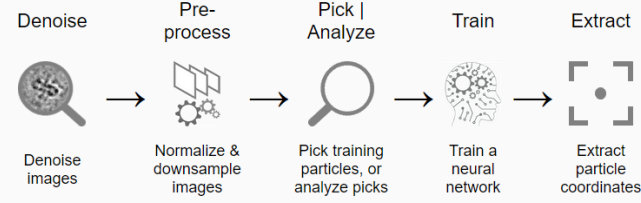
Pick more of anything



Start
Denoise
Pre-process
Pick | Analyze
Train
Extract
More Tools

Welcome!

This GUI allows you to pick particles & make Topaz commands ([watch Topaz presentation](#)). General workflow:



Note: To use pre-trained picking models, first Pre-process then Extract using a pre-trained model (don't Pick or Train).

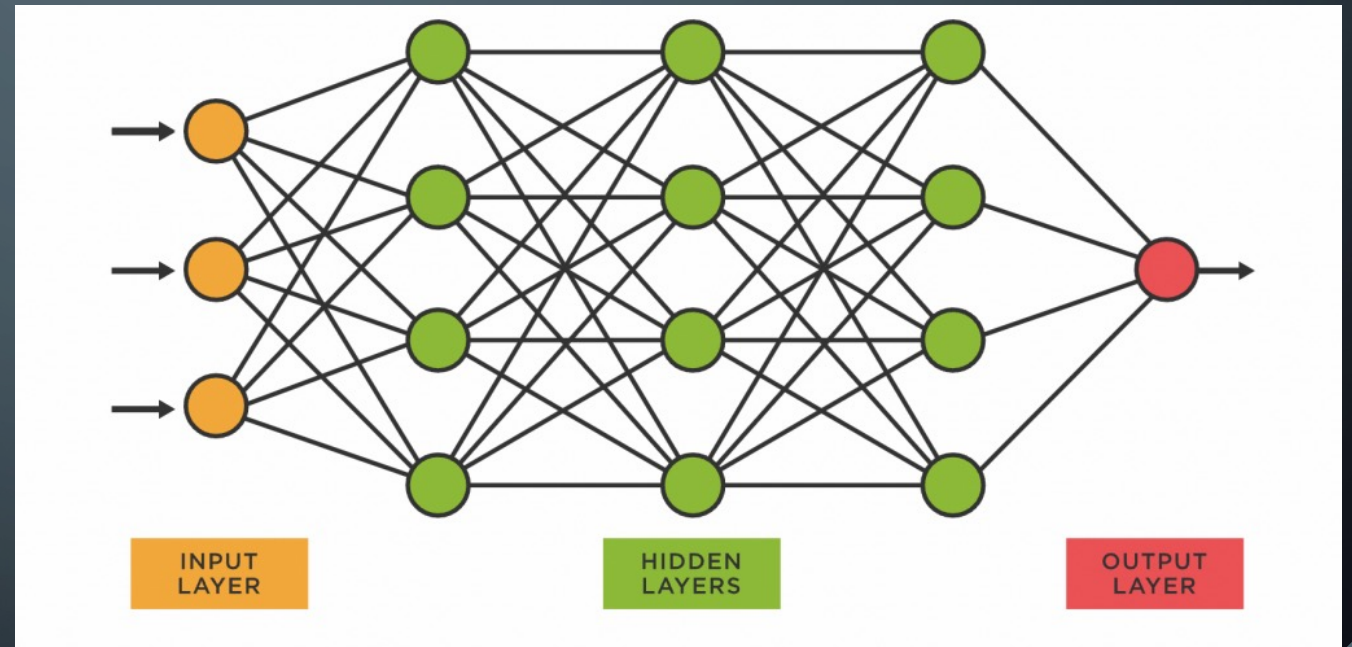
Citations: *Bepler, T, Morin, A, Rapp, M, Brasch, J, Shapiro, L, Noble, AJ, Berger, B (2019). Positive-unlabeled convolutional neural networks for particle picking in cryo-electron micrographs. Nat Meth*

Bepler, T, Kelley, K, Noble, AJ, Berger, B (2020). Topaz-Denoise: general deep denoising models for cryoEM and cryoET. Nat Comm

CNN ARCHITECTURE

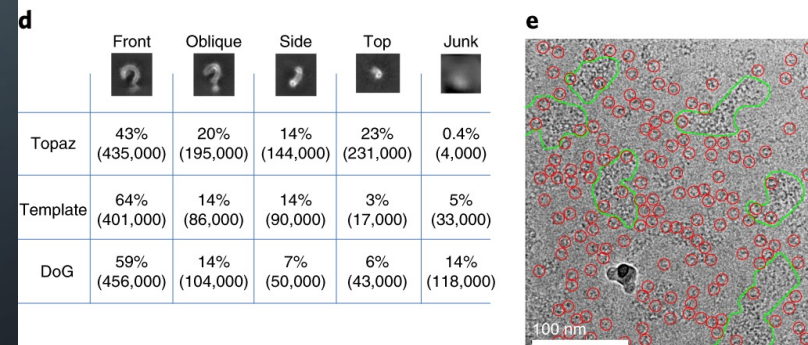
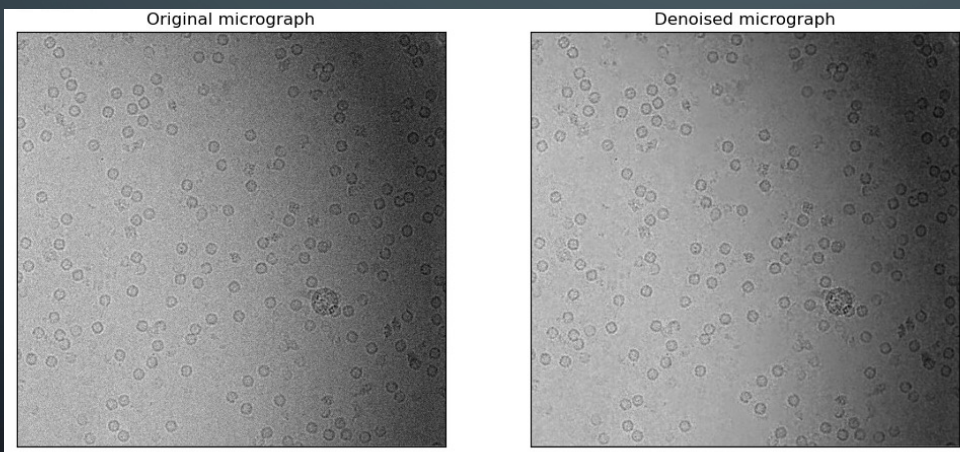
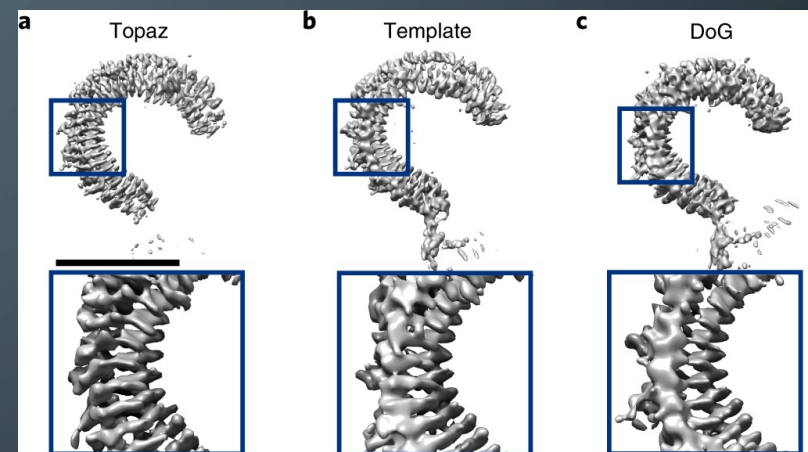
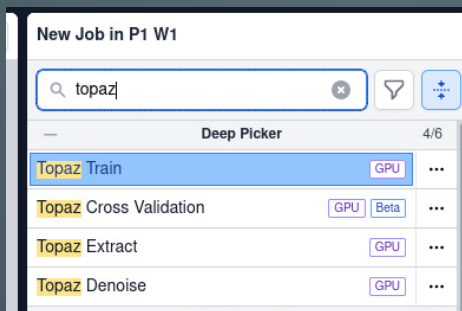
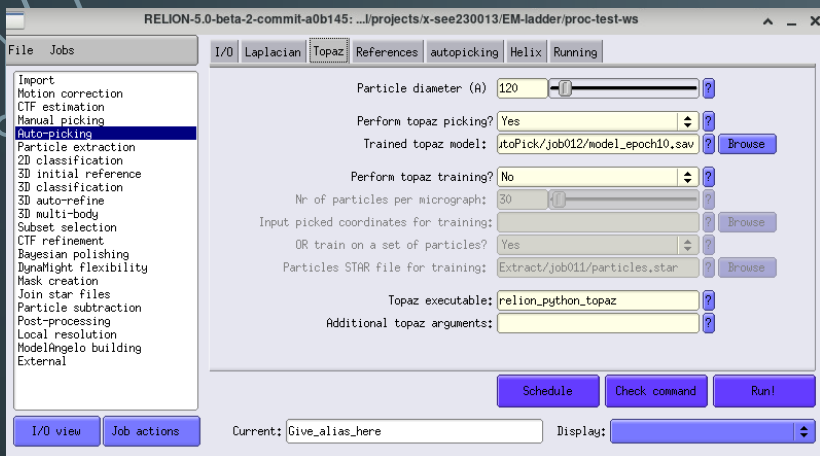
Convolutional neuronal networks are used to analyse multi-dimensional data and present it in human readable form. They operate on reducing multidimensionality to a perceivable dimension.

- **Input Layer:** Represents the features of the dataset, with each neuron corresponding to one feature.
- **Hidden Layers:** Multiple layers of neurons that process input features, learning complex patterns and relationships.
- **Output Layer:** Final layer that provides the prediction or classification result, with each neuron representing a possible output category.



[DEMO](#)

TOPAZ EXAMPLES



TOPAZ ISN'T MAGIC

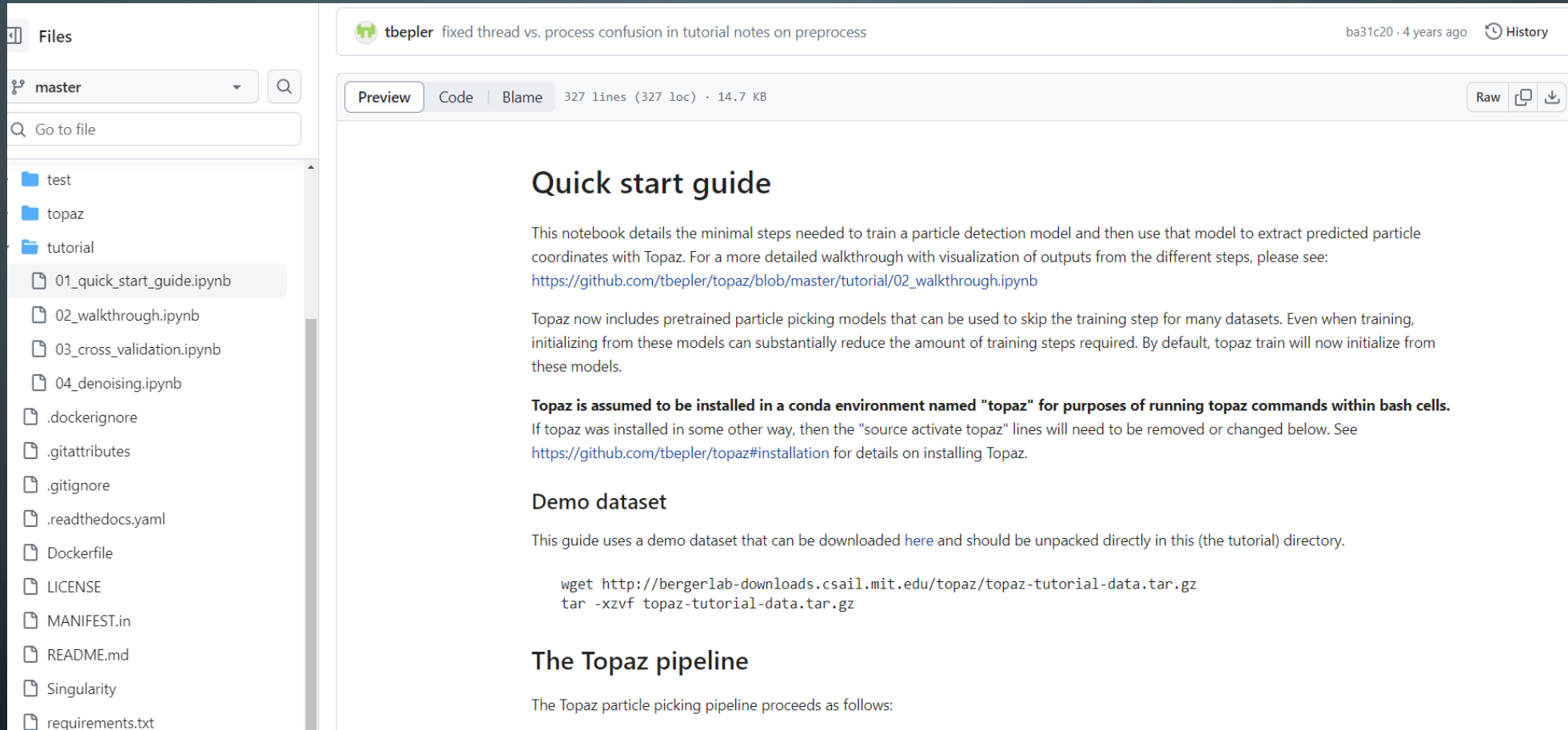
- What Topaz can do

- Denoise the mics to enhance contrast
- Extract particles using own models
- Get trained on your data
- Extract particles using trained models
- Great on single GPU

- What Topaz can't do

- Filter particles for you
- Cannot box ptcls, you have to EXTRACT
- MAGIC

TRY IT YOURSELF



The screenshot shows a GitHub repository interface. On the left is a file explorer for the 'master' branch, listing files like '01_quick_start_guide.ipynb', '02_walkthrough.ipynb', '03_cross_validation.ipynb', '04_denoising.ipynb', and various configuration files. The main area displays a notebook preview titled 'fixed thread vs. process confusion in tutorial notes on preprocess' by user 'tbepler', updated 4 years ago. The notebook content includes a 'Quick start guide' section with a link to a walkthrough, a 'Demo dataset' section with a wget command, and a 'The Topaz pipeline' section.

Files

master

Go to file

- test
- topaz
- tutorial
 - 01_quick_start_guide.ipynb
 - 02_walkthrough.ipynb
 - 03_cross_validation.ipynb
 - 04_denoising.ipynb
- .dockerignore
- .gitattributes
- .gitignore
- .readthedocs.yaml
- Dockerfile
- LICENSE
- MANIFEST.in
- README.md
- Singularity
- requirements.txt

tbepler fixed thread vs. process confusion in tutorial notes on preprocess ba31c20 · 4 years ago History

Preview Code Blame 327 lines (327 loc) · 14.7 KB Raw Copy Download

Quick start guide

This notebook details the minimal steps needed to train a particle detection model and then use that model to extract predicted particle coordinates with Topaz. For a more detailed walkthrough with visualization of outputs from the different steps, please see: https://github.com/tbepler/topaz/blob/master/tutorial/02_walkthrough.ipynb

Topaz now includes pretrained particle picking models that can be used to skip the training step for many datasets. Even when training, initializing from these models can substantially reduce the amount of training steps required. By default, topaz train will now initialize from these models.

Topaz is assumed to be installed in a conda environment named "topaz" for purposes of running topaz commands within bash cells. If topaz was installed in some other way, then the "source activate topaz" lines will need to be removed or changed below. See <https://github.com/tbepler/topaz#installation> for details on installing Topaz.

Demo dataset

This guide uses a demo dataset that can be downloaded [here](#) and should be unpacked directly in this (the tutorial) directory.

```
wget http://bergerlab-downloads.csail.mit.edu/topaz/topaz-tutorial-data.tar.gz
tar -xzvf topaz-tutorial-data.tar.gz
```

The Topaz pipeline

The Topaz particle picking pipeline proceeds as follows:

Install it from: <https://github.com/tbepler/topaz/blob/master/README.md>

https://github.com/tbepler/topaz/blob/master/tutorial/01_quick_start_guide.ipynb