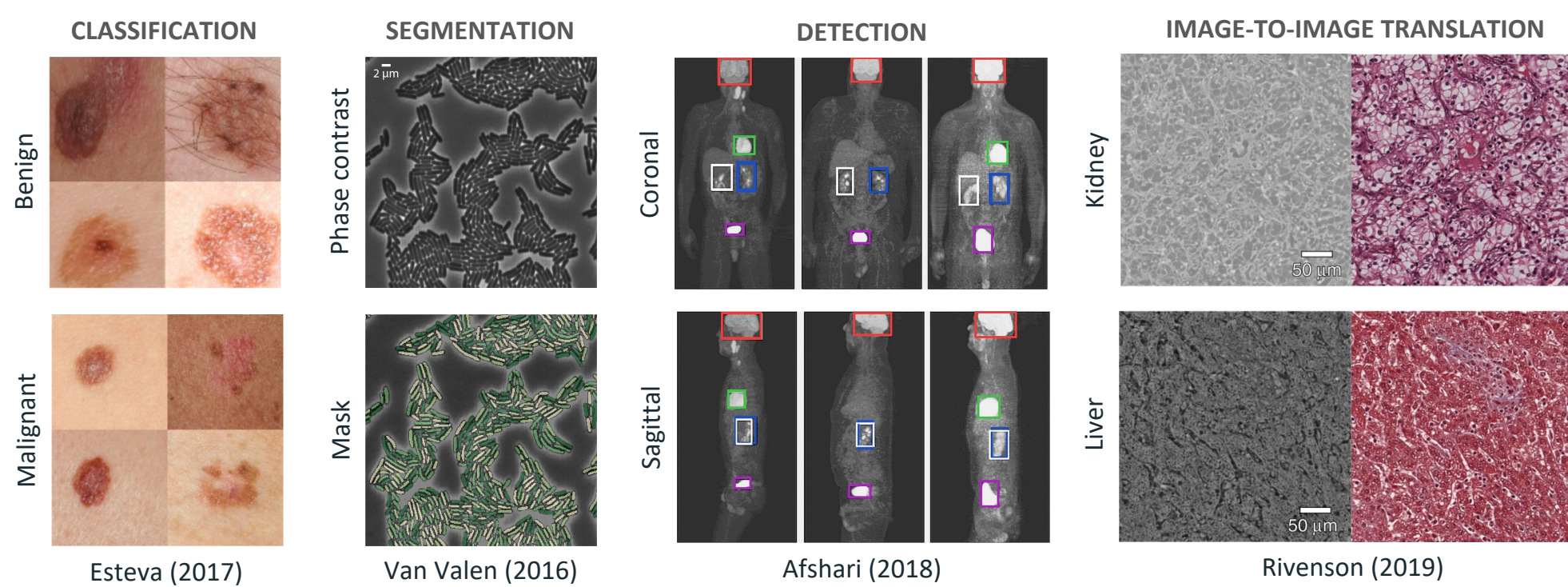


Abstract

Deep learning has transformed many aspects of industrial pipelines recently. Scientists involved in biomedical imaging research are also benefiting from the power of AI to tackle complex challenges (Kim, 2019). Although academic community has widely accepted image processing tools, such as scikit-image, ImageJ, there is still a need for a tool which integrates deep learning into biomedical image analysis. We propose a minimal, but convenient Python package based on PyTorch with common deep learning models, extended by flexible trainers and medical datasets.



Package Structure

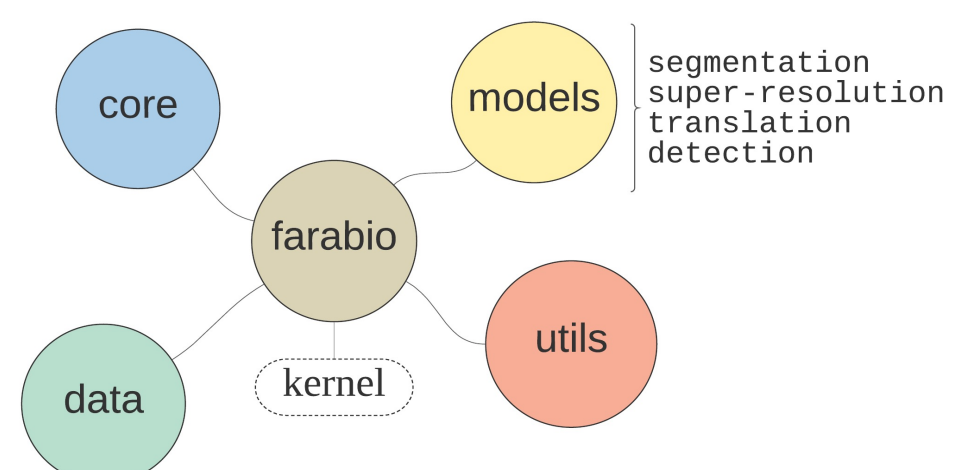


Fig 1. Package structure diagram

farabio package has 4 modules:

- **.core**: base trainer classes
- **.data**: preprocessing, datasets
- **.models**: architectures, trainers
- **.utils**: log, visualization, helpers

Every deep learning model trainer derives from the **core** module. Based on the type, it has either **ConvnetTrainer** or **GanTrainer** as its parent.

farabio incorporates and modifies common deep learning models from open-source code released on GitHub (see the docs)

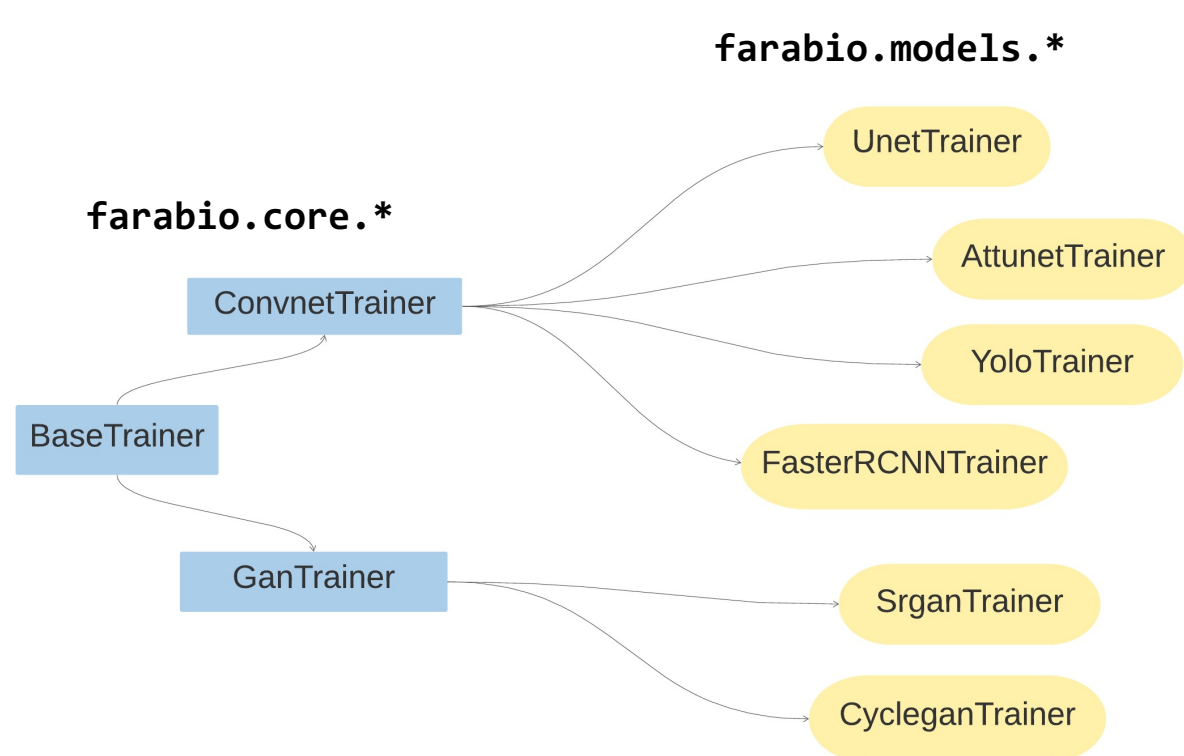


Fig 2. Inheritance diagram of trainers

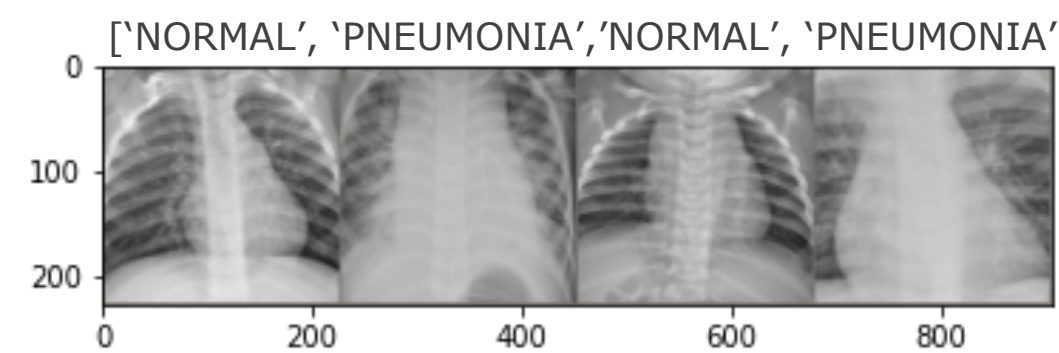
Biomedical Datasets

Several publicly available biomedical datasets which are retrieved from ISBI, MICCAI and Kaggle can be loaded from **farabio.data.biiodatasets** module:

```
import torch, torchvision
from torch.utils.data import DataLoader
from farabio.data.biiodatasets import ChestXray
from farabio.utils.vistools import imshow
```

```
train_dataset = ChestXray(split='train')
train_loader = DataLoader(train_dataset, batch_size=4, shuffle=True)
inputs, classes = next(iter(train_loader))
class_names = train_dataset.classes
```

```
out = torchvision.utils.make_grid(inputs)
imshow(out, title=[class_names[x] for x in classes])
```



Lifecycle of Trainers

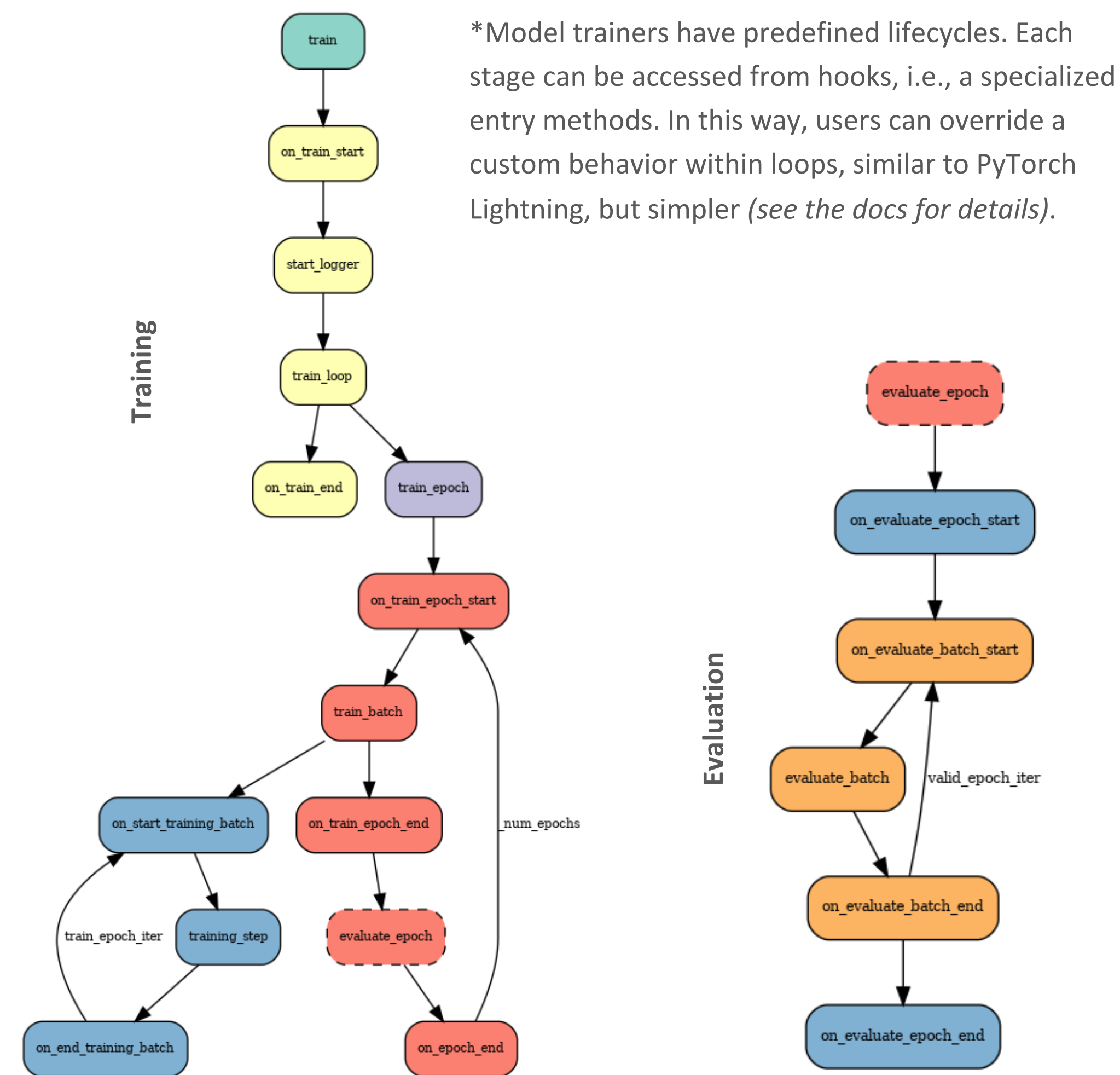


Fig 3. Diagrams of ConvnetTrainer lifecycle in training and evaluation loops

*Model trainers have predefined lifecycles. Each stage can be accessed from hooks, i.e., a specialized entry methods. In this way, users can override a custom behavior within loops, similar to PyTorch Lightning, but simpler (see the docs for details).

Preliminary Results

Each of the existing model trainers is *complete* and functions well with traditional computer vision datasets, such as ImageNet, VOC2012, COCO, Monet2Photo. However, the current priority is to extend these modules to comply with data configurations from **farabio.data.biiodatasets**.



Fig 4. Visualization of FasterRCNNTrainer training the PASCAL VOC 2012 image dataset

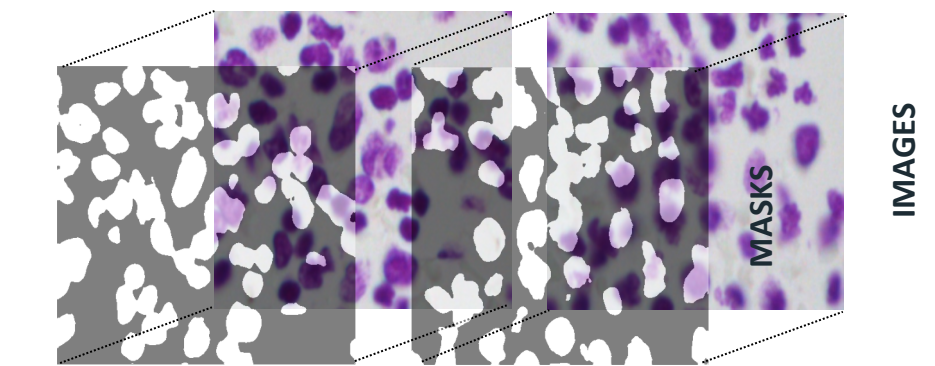


Fig 5. Qualitative results of UnetTrainer with default configurations (see the docs), on segmented nuclei images from 2018 Data Science Bowl, Kaggle

Future Works

In perspective, the vision of **farabio** package is to become an important player of PyTorch ecosystem, with research and academic community of biomedical engineering adopting it widely for different deep learning purposes. This is only possible with robust architectural foundations, rich dataset extensions and openness to new ideas, collaborations, which we aspire to cultivate.



Fig 5. Deep learning for medical imaging and healthcare industry (Dutta, 2020)

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Links

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