





### Software - Research - Education

Readability (train script)

Package or tool (network class)

Pre-defined workflow (data loader)

Manual, documentation or tutorial (docs vs papers)

Test (alternative codebase)

**Promotion** 

Clinical deployment

Where are we heading





# Medical Image Registration

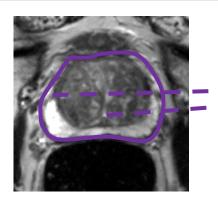


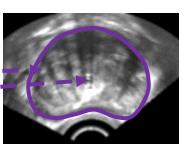
## **Registration applications**

- Multi-modal, e.g. image-guided interventions
- Inter-subject, e.g. atlas-based segmentation
- Medical image analysis, e.g. longitudinal analysis

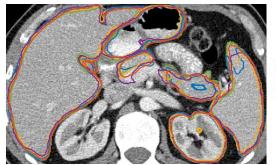
## Why correspondence is a problem

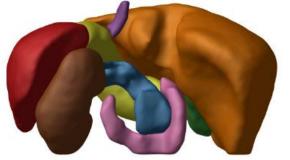
- Different imaging coordinates
- Motion and deformation
- Homology

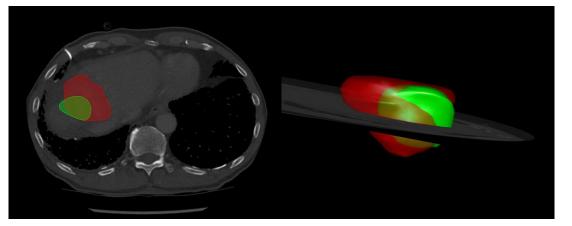














**o** cmic

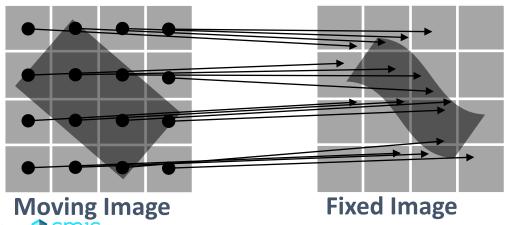
# Medical Image Registration

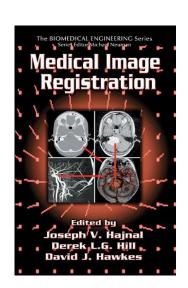


## The three-component formulation

- Regularised transformation models:  $\mathcal{T}(I_A, \theta)$ , e.g. rigid, affine, TPS, FFD ...
- Image similarity measures:  $S(I_A, I_B)$ , e.g. SSD, CC, MI ...
- Optimisation:  $\hat{\theta} = \max_{\theta} \mathcal{S}[I_{Fixed}, \mathcal{T}(I_{Moving}, \theta)]$

e.g. [Rueckert et al 1999] and two-decades research





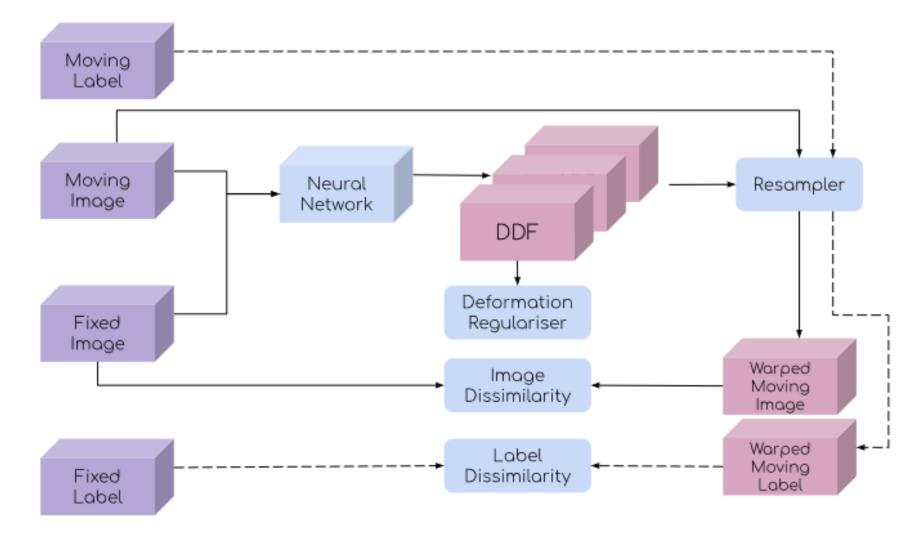
Intensity- and Feature-

based methods





## Medical image registration using deep learning









# Freely available, community-supported open-source tools for medical image registration using deep learning

DeepReg.net

github.com/DeepRegNet/DeepReg

DeepReg.readthedocs.io



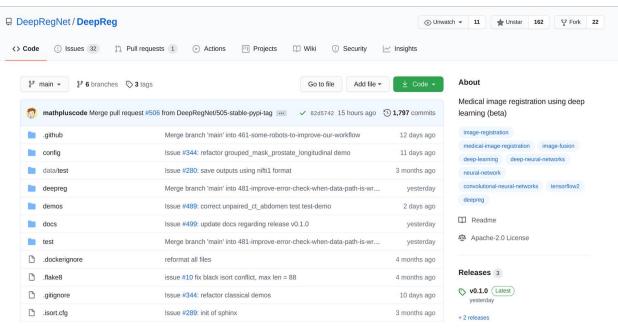


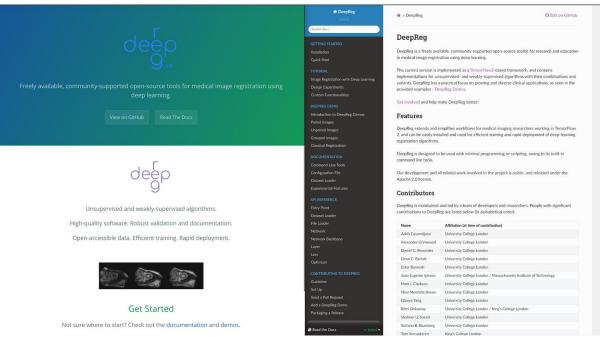




















## **Readability** — the original training.py script

```
import tensorflow as tf
import sys
import random
import time
import labelreg.helpers as helper
 import labelreg.networks as network
import labelreg.utils as util
import labelreg.losses as loss
# 0 - get configs
config = helper.ConfigParser(sys.argv, 'training')
reader_moving_image, reader_fixed_image, reader_moving_label, reader_fixed_label = helper.get_data_readers(
    config['Data']['dir_moving_image'],
    config['Data']['dir_fixed_image'],
    config['Data']['dir_moving_label'],
    config['Data']['dir fixed label'])
ph_moving_image = tf.placeholder(tf.float32, [config['Train']['minibatch_size']]+reader_moving_image.data_shape+[1])
ph_fixed_image = tf.placeholder(tf.float32, [config['Train']['minibatch_size']]+reader_fixed_image.data_shape+[1])
ph_moving_affine = tf.placeholder(tf.float32, [config['Train']['minibatch_size']]+[1, 12])
ph_fixed_affine = tf.placeholder(tf.float32, [config['Train']['minibatch_size']]+[1, 12])
input_moving_image = util.warp_image_affine(ph_moving_image, ph_moving_affine) # data augmentation
input_fixed_image = util.warp_image_affine(ph_fixed_image, ph_fixed_affine) # data augmentation
reg net = network.build network(network type=config['Network']['network type'],
                                minibatch_size=config['Train']['minibatch_size'],
                                image_moving=input_moving_image,
                                image_fixed=input_fixed_image)
```

```
# loss
ph_moving_label = tf.placeholder(tf.float32, [config['Train']['minibatch_size']]+reader_moving_image.data_shape+[1])
ph_fixed_label = tf.placeholder(tf.float32, [config['Train']['minibatch_size']]+reader_fixed_image.data_shape+[1])
input moving label = util.warp image affine(ph moving label, ph moving affine) # data augmentation
input_fixed_label = util.warp_image_affine(ph_fixed_label, ph_fixed_affine) # data augmentation
warped_moving_label = reg_net.warp_image(input_moving_label) # warp the moving label with the predicted ddf
loss_similarity, loss_regulariser = loss.build_loss(similarity_type=config['Loss']['similarity_type'],
                                                   similarity_scales=config['Loss']['similarity_scales'],
                                                   regulariser_type=config['Loss']['regulariser_type'],
                                                   regulariser_weight=config['Loss']['regulariser_weight'],
                                                   label_moving=warped_moving_label,
                                                   label_fixed=input_fixed_label,
                                                   network_type=config['Network']['network_type'],
                                                   ddf=reg net.ddf)
train_op = tf.train.AdamOptimizer(config['Train']['learning_rate']).minimize(loss_similarity+loss_regulariser)
# utility nodes - for information only
dice = util.compute_binary_dice(warped_moving_label, input_fixed_label)
dist = util.compute_centroid_distance(warped_moving_label, input_fixed_label)
```

```
# 3 - training
num_minibatch = int(reader_moving_label.num_data/config['Train']['minibatch_size'])
train_indices = [i for i in range(reader_moving_label.num_data)]
saver = tf.train.Saver(max_to_keep=1)
sess = tf.Session()
sess.run(tf.global_variables_initializer())
for step in range(config['Train']['total_iterations']):
    if step in range(0, config['Train']['total_iterations'], num_minibatch):
        random.shuffle(train indices)
    minibatch_idx = step % num_minibatch
    case_indices = train_indices[
                    minibatch_idx*config['Train']['minibatch_size']:(minibatch_idx+1)*config['Train']['minibatch_size']]
    label_indices = [random.randrange(reader_moving_label.num_labels[i]) for i in case_indices]
    trainFeed = {ph_moving_image: reader_moving_image.get_data(case_indices),
                  ph fixed image: reader fixed image.get data(case indices),
                 ph_moving_label: reader_moving_label.get_data(case_indices, label_indices),
                  ph_fixed_label: reader_fixed_label.get_data(case_indices, label_indices),
                  ph_moving_affine: helper.random_transform_generator(config['Train']['minibatch_size']),
                  ph_fixed_affine: helper.random_transform_generator(config['Train']['minibatch_size'])}
    sess.run(train_op, feed_dict=trainFeed)
```









## **Readability** — the original training.py script

```
import tensorflow as tf
def build_loss(similarity_type, similarity_scales, regulariser_type, regulariser_weight,
               label_moving, label_fixed, network_type, ddf):
    label_similarity = multi_scale_loss(label_fixed, label_moving, similarity_type.lower(), similarity_scales)
    if network_type.lower() == 'global':
        ddf_regularisation = tf.constant(0.0)
        ddf_regularisation = tf.reduce_mean(local_displacement_energy(ddf, regulariser_type, regulariser_weight))
    return tf.reduce_mean(label_similarity), ddf_regularisation
def weighted_binary_cross_entropy(ts, ps, pw=1, eps=1e-6):
    ps = tf.clip_by_value(ps, eps, 1-eps)
    return -tf.reduce_sum(
       tf.concat([ts*pw, 1-ts], axis=4)*tf.log(tf.concat([ps, 1-ps], axis=4)),
        axis=4, keep_dims=True)
def dice_simple(ts, ps, eps_vol=1e-6):
    numerator = tf.reduce_sum(ts*ps, axis=[1, 2, 3, 4]) * 2
    denominator = tf.reduce_sum(ts, axis=[1, 2, 3, 4]) + tf.reduce_sum(ps, axis=[1, 2, 3, 4])+eps_vol
```









## **Readability** – the DeepReg training.py script

```
train(
    gpu="0",
    config_path=config_path,
    gpu_allow_growth=True,
    ckpt_path="",
    log_dir=log_dir,
    exp_name=exp_name,
```

```
import argparse
    from datetime import datetime
    from deepreg.train import train
    name = "paired_mrus_prostate"
    # parser is used to simplify testing
    # please run the script with --full flag to ensure non-testing mode
    # for instance:
12 # python script.py --full
    parser = argparse.ArgumentParser()
14 parser.add_argument(
        "--test",
        help="Execute the script with reduced image size for test purpose.",
        dest="test",
        action="store_true",
20 parser.add_argument(
        help="Execute the script with full configuration.",
        dest="test",
        action="store_false",
25 )
    parser.set_defaults(test=True)
    args = parser.parse_args()
30 print(
        "\n\n\n\n\n"
        "=======\n"
        "The training can also be launched using the following command.\n"
        "deepreg_train --gpu '0' "
        f"--config_path demos/{name}/{name}.yaml "
        f"--log_dir demos/{name} "
        "--exp_name logs_train\n"
        "\n\n\n\n\n"
40 )
42 log_dir = f"demos/{name}"
43 exp_name = "logs_train/" + datetime.now().strftime("%Y%m%d-%H%M%S")
44 config_path = [f"demos/{name}.yaml"]
45 if args.test:
        config_path.append("config/test/demo_paired.yaml")
```







**Readability** – the DeepReg training.py script

```
48 train(
49 gpu="0",
50 config_pat =config_path,
51 gpu_allow_grawth=True,
52 ckpt_path="",
53 log_dir=log_dir,
54 exp_name=exp_name,
```

```
- demos/paired_mrus_prostate/dataset/part09
 moving_image_shape: [69, 69, 64]
method: "ddf" # the registration method, value should be ddf / dvf / conditional
   name: "local" # value should be local / global / unet
  extract_levels: [0, 1, 2, 3]
   label:
     weight: 1.0
     name: "dice"
     name: "bending" # options include "bending", "gradient"
   learning_rate: 1.0e-5
   shuffle_buffer_num_batch: 1 # shuffle_buffer_size = batch_size * shuffle_buffer_num_batch
 epochs: 5000 # number of training epochs
```





## Package or tool

- -> For users who do not code
- -> command line tool
- -> custom config file
- -> RFGISTRY

Feedback from developers, testers, and users

```
@<mark>REGISTRY</mark>.register_model(name="ddf")
class DDFModel(RegistrationModel):
   A registration model predicts DDF.
   When using global net as backbone,
   the model predicts an affine transformation parameters,
   and a DDF is calculated based on that.
   ....
   name = "DDFModel"
   def _resize_interpolate(self, field, control_points):
       resize = layer.ResizeCPTransform(control_points)
       field = resize(field)
       interpolate = layer.BSplines3DTransform(control_points, self.fixed_image_size)
       field = interpolate(field)
       return field
   def build_model(self):
       """Build the model to be saved as self._model."""
       # build inputs
       self._inputs = self.build_inputs()
       moving_image = self._inputs["moving_image"] # (batch, m_dim1, m_dim2, m_dim3)
       fixed_image = self._inputs["fixed_image"] # (batch, f_dim1, f_dim2, f_dim3)
       # build ddf
       control_points = self.config["backbone"].pop("control_points", False)
       backbone_inputs = self.concat_images(moving_image, fixed_image)
       backbone = REGISTRY.build_backbone(
           config=self.config["backbone"],
           default_args=dict(
               image_size=self.fixed_image_size,
               out_channels=3,
               out_kernel_initializer="zeros",
               out_activation=None,
```



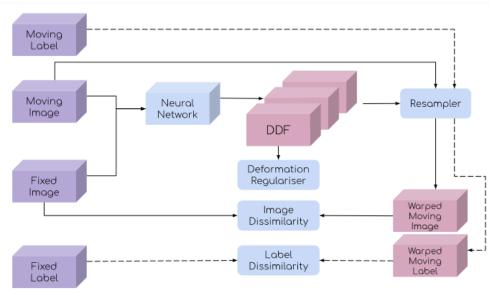




### Pre-defined workflow

Data loader script – for a two-stage, image/label feeding

```
class DataReader:
   def __init__(self, dir_name):
        self.dir_name = dir_name
        self.files = os.listdir(dir_name)
        self.files.sort()
        self.num_data = len(self.files)
        self.file_objects = [nib.load(os.path.join(dir_name, self.files[i])) for i in range(self.num_data)]
       self.num_labels = [self.file_objects[i].shape[3] if len(self.file_objects[i].shape) == 4
                           else 1
                           for i in range(self.num_data)]
        self.data_shape = list(self.file_objects[0].shape[0:3])
   def get_num_labels(self, case_indices):
        return [self.num_labels[i] for i in case_indices]
   def get_data(self, case_indices=None, label_indices=None):
        if case_indices is None:
            case_indices = range(self.num_data)
        # todo: check the supplied label_indices smaller than num_labels
        if label_indices is None: # e.g. images only
           data = [np.asarray(self.file_objects[i].dataobj) for i in case_indices]
        else:
           if len(label_indices) == 1:
               label_indices *= self.num_data
           data = [self.file_objects[i].dataobj[..., j] if self.num_labels[i] > 1
                   else np.asarray(self.file_objects[i].dataobj)
                   for (i, j) in zip(case_indices, label_indices)]
        return np.expand_dims(np.stack(data, axis=0), axis=4)
```



### Software - Research - Education



### Pre-defined workflow

DeepReg data loaders

Possibilities and permutations

Compromises

Readability

Unpaired Images Grouped Images Classical Registration **DOCUMENTATION** Command Line Tools Logging Configuration File □ Dataset Loader

Dataset type

Dataset requirements

□ Paired images

Sampling

Configuration

□ File loader

Nifti

H5

⊕ Unpaired images

⊞ Grouped images

Registry

**Experimental Features** 

**API REFERENCE** 

**Entry Point** 

Dataset Loader

File Loader

Registry

Network

Backbone

Layer

Loss

Optimizer

### **CONTRIBUTING TO DEEPREG**

### Configuration

An example configuration for paired dataset is provided as follows.

```
dataset:
 train:
    dir: "data/test/h5/paired/train" # folder containing data
    format: "h5" # nifti or h5
    labeled: true # true or false
  valid:
    dir: "data/test/h5/unpaired/test"
    format: "h5"
   labeled: true
  test:
    dir: "data/test/h5/unpaired/test"
    format: "h5"
   labeled: true
  type: "paired" # value should be paired / unpaired / grouped
  moving_image_shape: [16, 16, 16] # value should be like [dim1, dim2, dim3]
 fixed_image_shape: [8, 8, 8] # value should be like [dim1, dim2, dim3]
```

where, the configuration can be split into common configurations that shared by all dataset types and specific configurations for paired images:

- · Common configurations
  - dir/train gives the directory containing training data. Same for dir/valid and dir/test.
  - format can only be Nifti or h5 currently.
  - type can be paired, unpaired or grouped, corresponding to the dataset type described above.
  - labeled is a boolean indicating if the data is labeled or not.
- · Paired images configurations
  - moving image shape is the shape of moving images, a list of three integers.
  - fixed\_image\_shape is the shape of fixed images, a list of three integers.

Optionally, multiple dataset directories can be specified, such that the data will be sampled from several directories, for instance:

```
dataset:
 train:
    dir: # folders containing data
      - "data/test/h5/paired/train1"
      - "data/test/h5/paired/train2"
```

#### Nifti

Nifti data are stored in files with suffix .nii.gz . Each file should contain only one 3D or 4D tensor, corresponding to an image or a label.

obs is short for one observation of a data sample - a 3D image volume or a 3D/4D label volume - and the name can be any string.

All image data should be placed under moving\_images/ , fixed\_images/ with respect to the provided directory. The label data should be placed under moving\_labels/, and fixed\_labels/, if available. These are top directories.

File names should be consistent between top directories, e.g.:

- moving\_images/
- obs1.nii.gz
- obs2.nii.gz
- fixed images/
- obs1.nii.gz
- obs2.nii.gz
- o ... moving\_labels/
- obs1.nii.gz
- obs2.nii.gz
- fixed labels/
- obs1.nii.gz obs2.nii.gz

Check test paired Nifti data as an example.

Optionally, the data may not be all saved directly under the top directory. They can be further grouped in subdirectories as long as the data paths are consistent.

#### H5

H5 data are stored in files with suffix .h5. Hierarchical multi-level indexing is not used. Each file should contain multiple key-value pairs and values are

Yipeng Hu



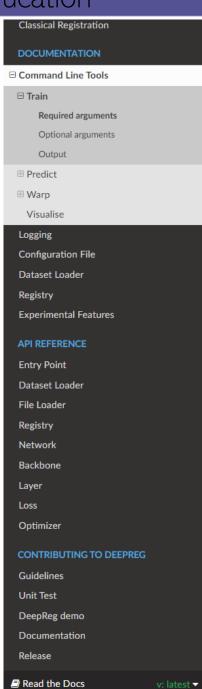






### CLT user manual

(for a standalone software)



### Required arguments

· GPU:

--gpu or -g, specifies the index or indices of GPUs for training.

#### Example usage:

- --gpu "" for CPU only
- --gpu "0" for using only GPU 0
- --gpu "0,1" for using GPU 0 and 1.

#### Configuration:

--config\_path or -c , specifies the configuration file for training.

The path must end with .yaml .

Optionally, multiple paths can be specified, and the configuration will be merged. In case of conflicts, values are overwritten by the last config file defining them.

#### Example usage:

- --config\_path\_config1.yaml for using one single configuration file.
- --config\_path\_config1.yam1 config2.yam1 for using multiple configuration files.

### Optional arguments

· CPU allocation:

--num\_workers , if given, TensorFlow will use limited CPUs.

By default, it uses only 1 CPUs. Setting it to non-positive values will be using all CPUs.

#### Example usage:

- --num\_workers 2 for using at most 2 CPUs.
- GPU memory allocation:

--gpu\_allow\_growth or -gr , if given, TensorFlow will only grow the memory usage as is needed.

By default, it allocates all available GPU memory.

#### Example usage:

- --gpu\_allow\_growth , no extra argument is needed.
- Load checkpoint:

 $\overline{\ \ \ }_{\text{--ckpt\_path}} \text{ or } \overline{\ \ \ }_{\text{-k}} \text{, specifies the path of the saved model checkpoint, so that the training will be}$ 







### Software - Research - Education



### **Tutorials**

(for research, education)

Cluster)

#### **DEEPREG DEMO**

Introduction to DeepReg Demos

Paired Images

**Unpaired Images** 

**Grouped Images** 

Classical Registration

#### **DOCUMENTATION**

Command Line Tools

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#### CONTRIBUTING TO DEEPREG

Guidelines

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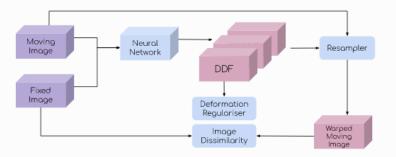
Release

### Learning

Depending on the availability of the data labels, registration networks can be trained with different approaches:

### Unsupervised

When the data label is unavailable, the training can be driven by the unsupervised loss. The loss function often consists of the intensity based loss and deformation loss. The following is an illustration of an unsupervised DDF-based registration network.



### Weakly-supervised

When there is no intensity based loss that is appropriate for the image pair one would like to register, the training can take a pair of corresponding moving and fixed labels (in addition to the image pair), represented by binary masks, to compute a label dissimilarity (feature based loss) to drive the registration.

Combined with the regularisation on the predicted displacement field, this forms a weakly-supervised training. An illustration of an weakly-supervised DDF-based registration network is provided below.

When multiple labels are available for each image, the labels can be sampled during the training iteration, such that only one label per image is used in each iteration of the data set (epoch).









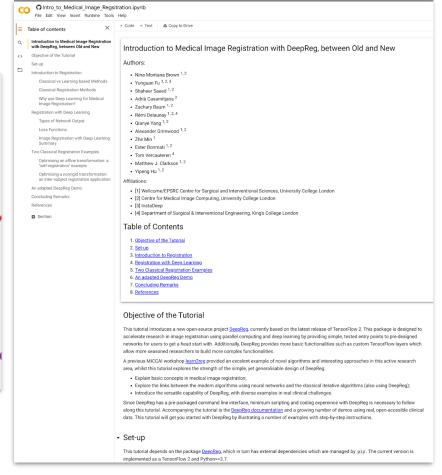
### **Tutorials**

(for research, education)

1st Place Winner in MICCAI Educational Challenge 2020

### MEC 2020 Materials View Teaser Videos For 2020 Finalists • 1st Place! Introduction to Medical Image Registration with DeepReg, Between Old and New (Teaser Video) Nina Montana Brown, Yunquan Fu, Shaheer Saeed, Adrià Casamitjana, Zachary Baum, Rémi Delaunay, Qianye Yang, Alexander Grimwood, Zhe Min, Ester Bonmati, Tom Vercauteren, Matthew J. Clarkson and Yipeng Hu - University College London, InstaDeep & King's College London • 2nd Place! An Introduction to SimpleITK (Teaser Video) Ziv Yaniv, Bradley C. Lowekamp and David T. Chen - National Institute of Allergy and Infectious Diseases, National Institutes of Health & Medical Science and Computing LLC. • 3rd Place! How to Organize a Challenge -- Behind the Scenes (Teaser Video) Annika Reinke, Matthias Eisenmann, Laura Aguilera Saiz, Sinan Onogur, Leonardo Antonio Ayala Menjiyar and Peter Maximilian Full - German Cancer Research Center (DKFZ) & University of Heidelberg • Finalist Guitar and Musical MRI Concepts (Part 2) (Part 3) (Teaser Video) Jerry J. Battista - Western University Finalist Hands on Machine Learning Training -- HaMLeT (Teaser Video) Leon Weninger, Laxmi Gupta and Philipp Gräbel - RWTH Aachen University Finalist The Basic Augmented Reality Demo -- BARD (Teaser Video)

Matt Clarkson, Steve Thompson, Ester Bonmati, Tom Dowrick, Yipeng Hu and Ann Blandford - University College of London











## **Papers**

(for research)



DeepReg: a deep learning toolkit for medical image registration

Yunguan Fu<sup>1, 2, 3</sup>, Nina Montaña Brown<sup>1, 2</sup>, Shaheer U. Saeed<sup>1, 2</sup>, Adrià Casamitjana<sup>2</sup>, Zachary M. C. Baum<sup>1, 2</sup>, Rémi Delaunay<sup>1, 4</sup>, Qianye Yang<sup>1, 2</sup>, Alexander Grimwood<sup>1, 2</sup>, Zhe Min<sup>1</sup>, Stefano B. Blumberg<sup>2</sup>, Juan Eugenio Iglesias<sup>2, 5, 6</sup>, Dean C. Barratt<sup>1, 2</sup>, Ester Bonmati<sup>1, 2</sup>, Daniel C. Alexander<sup>2</sup>, Matthew J. Clarkson<sup>1, 2</sup>, Tom Vercauteren<sup>4</sup>, and Yipeng Hu<sup>1, 2</sup>

1 Wellcome/EPSRC Centre for Surgical and Interventional Sciences, University College London, London, UK 2 Centre for Medical Image Computing, University College London, London, UK 3 InstaDeep, London, UK 4 Department of Surgical & Interventional Engineering, King's College London, London, UK 5 Martinos Center for Biomedical Imaging, Massachusetts General Hospital and Harvard Medical School, Boston, USA 6 Computer Science and Artificial Intelligence Laboratory, Massachusetts Institute of Technology, Boston, USA

**DOI:** 10.21105/joss.02705

#### Software

- Review 🗗
- Repository ♂
- Archive ♂

Editor: Kevin M. Moerman 다

### Summary

Image fusion is a fundamental task in medical image analysis and computer-assisted intervention. Medical image registration, computational algorithms that align different images together (Hill, Batchelor, Holden, & Hawkes, 2001), has in recent years turned the research attention towards deep learning. Indeed, the representation ability to learn from population







C Edit on GitHub

## Docstring, API documentation

Paired Images

**Unpaired Images** 

Grouped Images

Classical Registration

#### **DOCUMENTATION**

**Command Line Tools** 

Logging

Configuration File

Dataset Loader

Registry

**Experimental Features** 

#### **API REFERENCE**

**☐ Entry Point** 

Train

Predict

Warp

Dataset Loader

File Loader

Registry

Network

Backbone

Layer

Loss

Optimizer

#### CONTRIBUTING TO DEEPREG

v: latest ~

Guidelines

**Unit Test** 

DeepReg demo

Documentation

Release

Read the Docs

**Entry Point** 

\* » Entry Point

### Train

Module to train a network using init files and a CLI.

deepreg.train.build\_config(config\_path: Union[str, List[str]], log\_dir: str, exp\_name: str, ckpt\_path: str, max\_epochs: int = -1)  $\rightarrow$  Tuple[Dict, str, str] %

Function to initialise log directories, assert that checkpointed model is the right type and to parse the configuration for training.

Parameters:

- config\_path list of str, path to config file
- log\_dir path of the log directory
- exp\_name name of the experiment
- · ckpt\_path path where model is stored.
- max\_epochs if max\_epochs > 0, use it to overwrite the configuration

Returns:

- config: a dictionary saving configuration
- exp\_name: the path of directory to save logs

deepreg.train.main(args=None)

Entry point for train script.

Parameters: args – arguments

deepreg.train.train(gpu: str, config\_path: Union[str, List[str]], ckpt\_path: str, num\_workers: int = 1, gpu\_allow\_growth: bool = True, exp\_name: str = ", log\_dir: str =  $'logs', max_epochs: int = -1)$ 

Function to train a model.

Parameters:

- gpu which local gpu to use to train.
- config\_path path to configuration set up.
- ckpt\_path where to store training checkpoints.
- num workers number of cpu cores to be used.







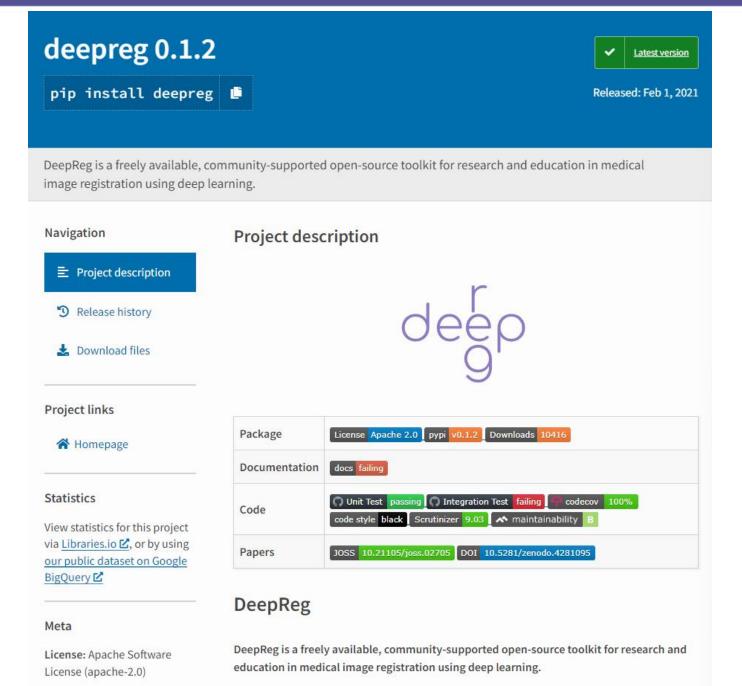


### Installation

Conda

PYPI packaging

Docker











## DeepReg Demos

Reference Paired brain MR-ultrasound registration Paired prostate MR-ultrasound registration Unpaired Images **Grouped Images** Classical Registration DOCUMENTATION Command Line Tools Logging Configuration File Dataset Loader Registry **Experimental Features API REFERENCE Entry Point** Dataset Loader File Loader Registry Network Backbone Layer Loss Optimizer **CONTRIBUTING TO DEEPREG** Guidelines **Unit Test** DeepReg demo Documentation

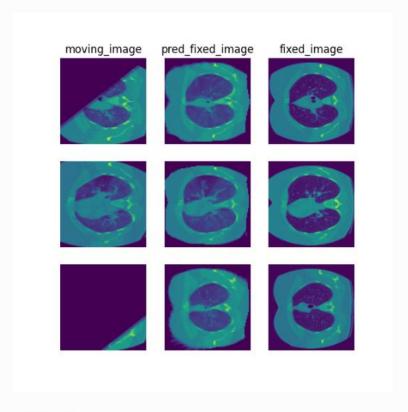
Release

### **Visualise**

The following command can be executed to generate a plot of three image slices from the the moving image, warped image and fixed image (left to right) to visualise the registration. Please see the visualisation tool docs here for more visualisation options such as animated gifs.

```
deepreg_vis -m 2 -i 'demos/paired_ct_lung/logs_predict/<time-stamp>/test/<pair-r
```

Note: The prediction must be run before running the command to generate the visualisation. The <time-stamp> and <pair-number> must be entered by the user.



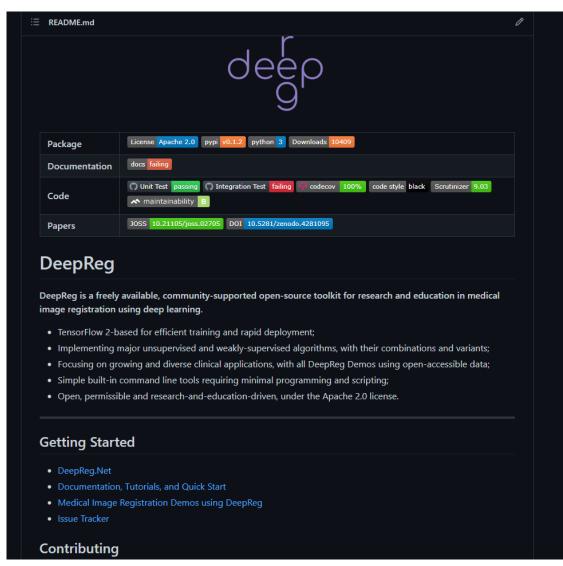


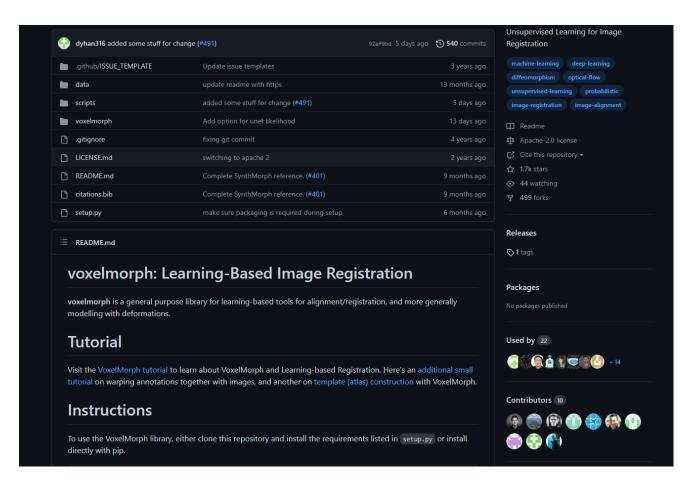






### Tests

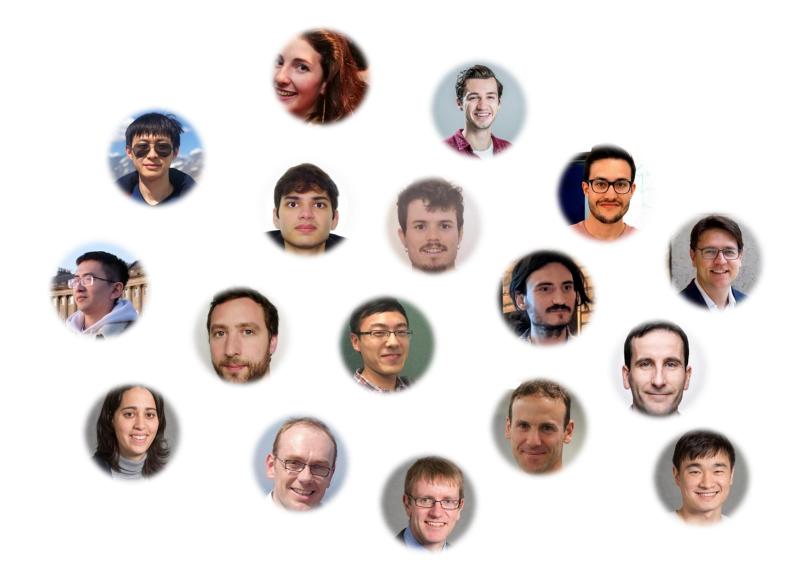








## We have no clue...!





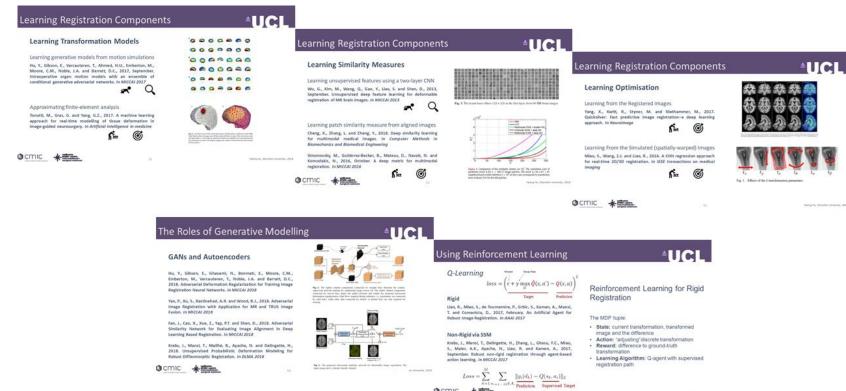




Trained models

Open data sets, e.g. learn2reg challenges

Clinical data pre-processing and curation



Plenty research, no clinical application yet









¥ 6 1 0

As is and achieved?

Next interested PhD students?

MONAI?

Paper-based? e.g.

Yang 2019 MICCAI (DeepReg branch)

Yang 2021 ISBI (using DeepReg)

Yang 2022 TMI (PyTorch)

Baum 2021 MedIA (standalone repo)

Baum 2022 ASMUS (using DeepReg)

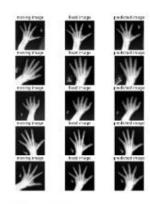
Baum 2022 TMI (using DeepReg)

Shen 2022 MICCAI (PyTorch)

UCL Modules? e.g.

MPHY0041 (DeepReg code)

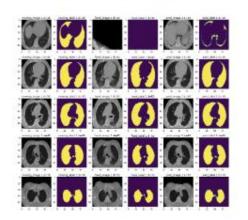
MPHY0043 (MONAl code)

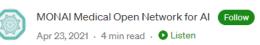


#### 3D intra-subject lung CT registration

The 3D intra-subject lung CT registration tutorial is an example of registration between 3D lung CT images acquired at inspiration and expiration from a single patient. This type of intra-subject registration is helpful in trackling material features of intenset like airways in airflow analysis or compressating motion during radiotherage.

The numerial showcases several features described above, including unsupervised and weakly-supervised losses, deformation regulariser, non-rigid transformation based on DDPs, and 3D volumetric registration using real clinical images. To learn more, check out the privided notebook and the original DeepBeg demo using the same open-accessible data set.





# Monai Explores Learning-Based Medical Image Registration

<u>MONAI</u> has been working closely with <u>DeepReg</u> on learning-based medical image registration using PyTorch. In the latest release, <u>MONAI v0.5.0</u>, we are delighted to provide a set of essential tools for developing registration pipelines.

### Medical Image Registration

<u>Image registration</u>, a process that spatially aligns one image with another, is important to many medical imaging applications. The two images that register are often referred to as moving and fixed images. An image registration algorithm generates a spatial transformation that can transform, or "warp", the moving image onto the fixed image coordinates.

The ability to align two or more images together allows the complementary information acquired from different imagi (multimodal registration, e.g., MR and ultrasound), at diffe (longitudinal registration), or from different patients (inter registration). For example, track which is a part of the patients (interpretation) with from a part of the patients (interpretation).



