# **Submission Instructions**

MOOD 2020

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# **General Instructions**

### Introduction

This document contains instructions on how to **access the data**, how to create a **docker image** and how to **submit it** to the synapse portal to participate in our challenge.

Challenge details can be found at our web pages: <a href="https://www.synapse.org/#!Synapse:syn21343101/wiki/599515">https://www.synapse.org/#!Synapse:syn21343101/wiki/599515</a>

# **Technical Instructions**

### Data access

# Training phase

- 1. Create an account at synapse and register for the challenge: <a href="https://www.synapse.org/#!Synapse:syn21343101/wiki/599515">https://www.synapse.org/#!Synapse:syn21343101/wiki/599515</a>
- 2. Download the data as zip.

# Test phase

Testing data won't be accessible during training. To be part of the challenge, you must **create a docker file and upload it to the synapse platform**. The challenge organizers will perform the complete evaluation. Please follow the instructions in the following sections to install, create and submit docker images.

# Creation of docker images

Please install and use docker for submission: https://www.docker.com/get-started

You can build and use any docker base/ image you like.

There are already a good base docker images to build on for:

- Pytorch: https://hub.docker.com/r/pytorch/pytorch/
- Tensorflow https://www.tensorflow.org/install/docker

For GPU support you may need to install the NVIDIA Container Toolkit: <a href="https://github.com/NVIDIA/nvidia-docker">https://github.com/NVIDIA/nvidia-docker</a>

For a simple example checkout out our Github: https://github.com/MIC-DKFZ/mood

# **Docker preparation**

For the different tasks the docker needs the following scripts:

- Sample-level:
  - /workspace/run\_sample\_brain.sh input\_folder output\_folder (for the brain dataset)
  - /workspace/run\_sample\_abdom.sh input\_folder output\_folder (for the abdominal dataset)
- Pixel-level:
  - /workspace/run\_pixel\_brain.sh input\_folder output\_folder (for the brain dataset)
  - /workspace/run\_pixel\_abdom.sh input\_folder output\_folder (for the abdominal dataset)

The docker has to allow mounting the input folder to /mnt/data and the output folder to /mnt/pred. We will mount the input and output folder and pass them to the run scripts.

During testing, the docker image will be run with the following commands:

#### Sample-level:

```
docker run --gpus all -v "<input folder>/:/mnt/data" -v
"<output folder>:/mnt/pred" docker-image-name
/workspace/run_sample_brain.sh /mnt/data /mnt/pred

docker run --gpus all -v "<input folder>/:/mnt/data" -v
"<output folder>:/mnt/pred" docker-image-name
/workspace/run sample abdom.sh /mnt/data /mnt/pred
```

#### Pixel-level:

```
docker run --gpus all -v "<input folder>/:/mnt/data" -v
"<output folder>:/mnt/pred" docker-image-name
/workspace/run_pixel_brain.sh /mnt/data /mnt/pred

docker run --gpus all -v "<input folder>/:/mnt/data" -v
"<output folder>:/mnt/pred" docker-image-name
/workspace/run pixel abdom.sh /mnt/data /mnt/pred
```

```
In other words, it will mount two folders, one at "/mnt/data" and one at "/mnt/pred".
```

Then, we will run the run scripts therefore, it must be written in such a way that it **will run your model automatically**.

The folder "/mnt/data" will contain the test data set, in the same way as the training data set, with the image dimensions equal to the image dimensions in the training data set.

Your model should process all scans and write the outputs to the given output directory.

#### Please be aware of the different output formats depending on your task:

- <u>Sample-level:</u> For each input file (e.g. input1.nii.gz) create a text file with the same name and an appended ".txt" file-ending (e.g. input1.nii.gz.txt) in the output directory and write out a single (float) score.
- <u>Pixel-level:</u> For each input file (e.g. input1.nii.gz) create a nifti file with the same dimensions and save it under the same (e.g. input1.nii.gz) in the output directory.

For more information have a look at our github example: <a href="https://github.com/MIC-DKFZ/mood">https://github.com/MIC-DKFZ/mood</a>

### Docker submission

# 0. Becoming a certified Synapse user

**Important:** In order to use all docker functionality e.g. <code>docker push</code>, you must be a certified user. You can become a certified user by filling out the following quiz (attention, the url contains a ":" at the end!): <a href="https://www.synapse.org/#!Quiz:">https://www.synapse.org/#!Quiz:</a>

# 0. Test your docker on the Toy data

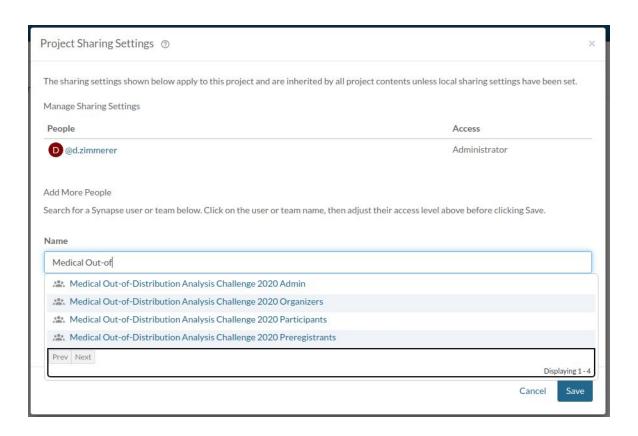
Please test your docker on the test data beforehand. See <a href="https://github.com/MIC-DKFZ/mood">https://github.com/MIC-DKFZ/mood</a>

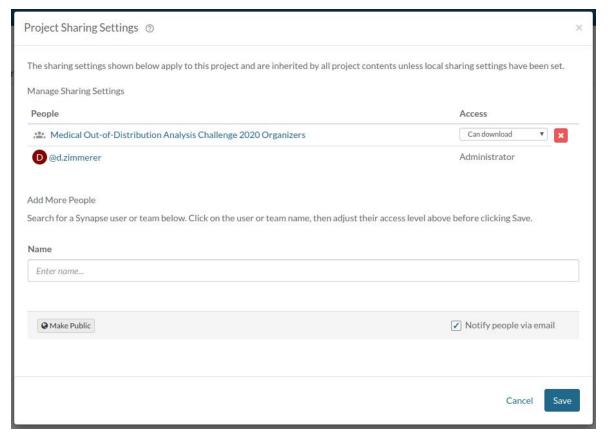
# 1. Create a new project on synapse

- To submit a docker file, you first need to **create a new project** on the synapse platform: (e.g.)

MOOD\_submission\_<task>\_<Your team name>

- Note the Synapse Project ID (e.g. syn20482334).
- Furthermore, once the project has been created, the organizing team of the challenge (*Medical Out of Distribution Analysis Challenge 2020 Organizers*) must be given download permissions to the project (Under *Project Settings* -> *Project Sharing Settings*), as shown in the following screenshots:





# 2. Upload docker-image to synapse

- docker login docker.synapse.org (Enter synapse username and password)

```
(base) rosst@mbi99:~$ sudo docker login docker.synapse.org
[sudo] password for rosst:
Username: schnobi1990
Password:
WARNING! Your password will be stored unencrypted in /home/rosst/.docker/config.
json.
Configure a credential helper to remove this warning. See
https://docs.docker.com/engine/reference/commandline/login/#credentials-store
Login Succeeded
```

- docker tag <imagename\_local>
 docker.synapse.org/<ProjectSynapseID>/<imagename synapse>

```
rosst@mbi99:~$ sudo nvidia-docker tag pytorch_example_final_v2 docker.synapse.org_syn20482334/example_submission rosst@mbi99:~$
```

- Docker push

docker.synapse.org/<SynapseProjectID>/<imagename>:latest

```
(base) rosst@mbi99:-$ sudo docker push docker.synapse.org/syn20482334/example_submission
[sudo] password for rosst:
The push refers to repository [docker.synapse.org/syn20482334/example_submission]
f12f1c6ace83: Preparing
bac58b2a5ace: Preparing
bac58b2a5ace: Layer already exists
92eb77d4bece: Layer already exists
32ae97716e4: Layer already exists
838dfd1aef44: Layer already exists
838dfd1aef44: Layer already exists
80782a85330f: Layer already exists
46d1ee805a8f: Layer already exists
46d1ee805a8f: Layer already exists
46d2e805a8f: Layer already exists
4798cbaaa513: Layer already exists
```

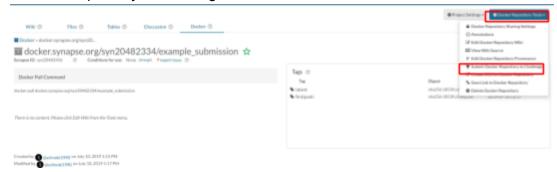
Verify the success of your push

You can verify on the Synapse website if the push was successful (If the upload fails, it might not be a **certified user**. Please refer to the upper paragraph "Becoming a certified user".)

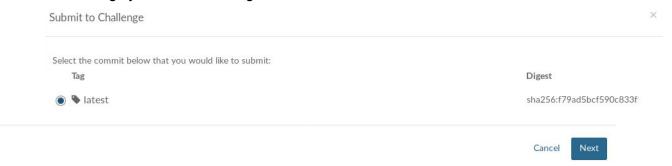
Synapse ID: syn204	82334	⑦ Si	torage Location: Synap	ose Storage		
Wiki ①	File	s ②	Tables ⑦	Discussion ①	Docker ⑦	
docker.synap			334/example_subm	nission		

### 3 . Submit the docker image

1) Click on the docker image. In the next view, select "Docker Repository Tools → Submit Docker Repository to Challenge"



To submit the docker to the challenge, first click on the docker image you want to submit which brings you to the following screen:



For each of the three tasks there is an individual submission queue:

- MOOD 2020 Sample-Level,
- MOOD 2020 Pixel-Level,

Choose the task for which you want to submit and click "Next".

If you plan to participate in **multiple tasks** of the challenge, you need to **submit your corresponding docker to each queue individually**.

Specify if you are entering alone or as a team. The submission pipeline will be checked in regular intervals. You will then be notified whether your submission is invalid (due to wrong format, etc...) or has been accepted.

The committee wishes you much success :-)