# Supplementary Section 3. Installing and populating BIAFLOWS locally

It is possible to install BIAFLOWS on a local server or a desktop computer. This might be useful to manage images locally or to analyse them with existing or custom BIAFLOWS workflows. The procedure is described below and should take less than 30 minutes using a regular Internet connection (UNIX based system recommended).

# Installing a local instance of BIAFLOWS

The procedure described in this section is for Linux Ubuntu but since BIAFLOWS leverages Docker container technologies, it should be possible to install BIAFLOWS on other platforms (not tested). Some specific details related to deployment on Mac OS can be found online.

## 1/ Install requirement

BIAFLOWS runs in Docker containers, so that the only requirement is to install Docker. Check official Docker documentation to install Docker for Ubuntu. In particular, choose Install using the repository, set up the repository and install Docker CE.

## 2/ Retrieve BIAFLOWS installation files

```
mkdir Biaflows/
cd Biaflows/
git clone https://github.com/Neubias-WG5/Biaflows-bootstrap.git
cd Biaflows-bootstrap
```

#### 3/ Configure the local instance

Edit configuration.sh file and if needed, update URLs (CORE\_URL, IMS\_URL, UPLOAD\_URL). Make sure to use URLs that are not already used by other applications (avoid localhost) to prevent conflicts. Add XXX\_URL variable values into the /etc/hosts of the host machine. In the /etc/hosts, add the following lines and don't forget to adapt them with values previously chosen for XXX\_URL variables in the configuration.sh file.

```
127.0.0.1 biaflows
127.0.0.1 biaflows-ims
127.0.0.1 biaflows-upload
127.0.0.1 rabbitmq
```

If needed, update data path variables (IMS\_STORAGE\_PATH ...). All data paths must be valid and mappable in the Docker engine. Create all the directories (**mkdir**) corresponding to the following variables (if they don't exist):

```
IMS_STORAGE_PATH
IMS_BUFFER_PATH
FAST_DATA_PATH
PROXY_CACHE_PATH
```

SOFTWARE\_CODE\_PATH SOFTWARE\_DOCKER\_IMAGES\_PATH JOBS\_PATH SERVER\_SSHKEYS\_PATH

Configure BIAFLOWS\_WORKFLOWS\_METRICS to *true* or *false* depending if you want to perform benchmarking: ground truth annotations are then required for all images. Setting this flag to false is the valid option if you plan to manage and process local images.

### 4/ Initialize the deployment

Generate the installation script with the command: sudo bash init.sh

#### 5/ Deploy the local instance

Run the generated deployment script with the command sudo bash start.sh

### 6/ Check the running instance

When start up is finished, check the application is running in your browser on the URL specified in your CORE\_URL variable (by default: http://biaflows).

Three accounts are created by default (username: admin; password: admin; username: guest; password: guest; username: neubias; password: neubias) with different access rights. Passwords should be updated from the Account page at the top right.

## 7/ Install sample projects (images and ground-truth data)

After BIAFLOWS successfully installed locally, this local instance is still empty of data. All projects available on BIAFLOWS maintained by NEUBIAS can be imported to this local instance.

Get the public and private keys of the admin account (at the end of the Account page). Then, run:

```
cd Biaflows-bootstrap sudo bash ./inject demo data.sh ADMIN PUBLIC KEY ADMIN PRIVATE KEY
```

where ADMIN\_PUBLIC\_KEY and ADMIN\_PRIVATE\_KEY have been substituted by their respective values.

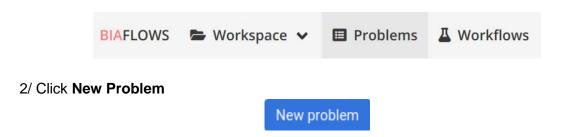
The script starts to download projects and import them in your local BIAFLOWS. The list of imported projects can be tweaked by editing the file Biaflows-bootstrap/configs/project\_migrator/projects.txt.

The whole data injection procedure can take several minutes, depending on your Internet connection and the number of projects being imported.

# Creating a new problem (project) in a local BIAFLOWS instance

To create a new problem, connect as regular user or admin.

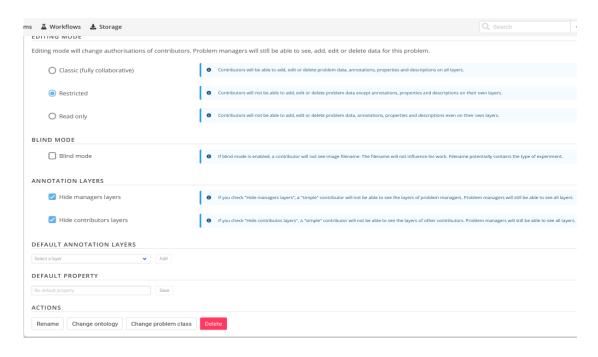
1/ Go to Problems tab



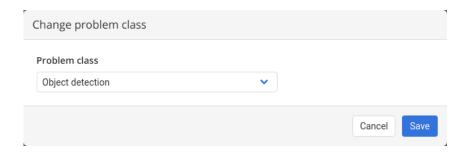
3/ Choose a meaningful problem name and save



4/ The problem is ready to be configured, the following configuration is recommended



5/ Map your problem to the corresponding problem class (see **Problem Class, Ground truth annotations formats and reported metrics**) by clicking on **Change problem class**. The problem class specifies the format of ground truth annotations (and workflow outputs), as well as the associated benchmark metrics to be computed.



6/ Configure project members. If you work alone, you can leave contributors and project managers to default user. This can be done from the "Members" tab in the problem configuration.

7/ The problem can be fully configured to display or hide panels / tabs / tools in the user interface. This is achieved from the **Custom UI** tab in the problem configuration.

8/ A description of the problem can optionally be added from the **Information** (left sidebar). The description is displayed in **Problems** list.

## Uploading images to a local BIAFLOWS instance

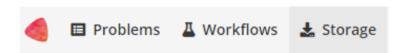
To upload new images, connect as regular user or admin.

## **Supported formats**

- **2D images**: 8-bit/16-bit TIFF (or OME-TIFF files)
- Multi-dimensional images (Z, C, T): single file 8-bit/16-bit OME-TIFF

Images name should not hold the text string **\_lbl** (reserved keyword for ground truth)

1/ Go to **Storage** section



2/ Select the **Problem** to which the images should be associated with (**Link with problem**)

Storage	admin storage
Link with problem	Select options
Files	No file
	Add files Start upload Cancel upload

**Note**: If a problem is not in the list, make sure you are a member for this problem

3/ Click on Add files... and select the files from the file browser

4/ Start upload with Start upload and wait until completion

The status can be:

- **DEPLOYED/CONVERTED**: The file is correctly imported to BIAFLOWS
- ERROR FORMAT: The file format is not supported
- ERROR EXTRACTION: Something went wrong during metadata extraction
- ERROR CONVERSION: Something went wrong during the conversion of the file into the BIAFLOWS internal image format
- ERROR DEPLOYMENT: Something went wrong during the communication with BIAFLOWS API. It can be due to access rights, or other unexpected error

<u>Note</u>: Images uploaded to storage can also be associated to a Problem after upload (Problem: **Add image**). This can be useful to associate the same image to several Problems.

#### Uploading ground truth annotations to an existing BIAFLOWS problem

If you plan to perform benchmarking, ground truth annotations should also be uploaded and associated to every image of a problem. The format of these annotations depends on the associated problem class (see **Problem Class**, ground truth annotations formats and reported metrics).

Image annotations (e.g. binary masks) should be uploaded as 16-bit TIFF (or OME-TIFF) for 2D images and single file 16-bit OME-TIFF for multidimensional (C,Z,T) images. They should be uploaded by following the procedure described in the previous section and should have the same name as their corresponding image + \_IbI suffix (e.g. AnImage.ome.tif is the original image and AnImage\_IbI.ome.tif is the ground-truth).

Other types of annotations (e.g. SWC, text file) should be added to the images as attached files. To do so, expand the image (blue arrow) in the list and click on **Add** next to **Attached files**.



# Adding existing workflows from trusted sources to a local BIAFLOWS instance

It is possible to integrate existing BIAFLOWS workflows to any BIAFLOWS instance. This operation requires configuring an external trusted source made of:

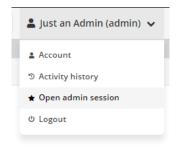
- 1. A source code registry (typically a Github user space)
- 2. An execution environment registry (typically a DockerHub user space)

If your workflow repositories are mixed with other repositories in your user space, you can specify a prefix to distinguish workflows repositories. For instance, all bioimage analysis workflows developed by NEUBIAS are prefixed by **W**\_ and available from this user space: <a href="https://github.com/Neubias-WG5">https://github.com/Neubias-WG5</a>.

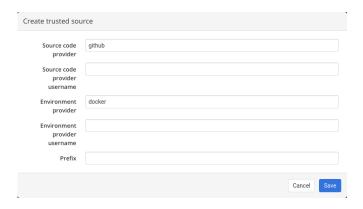
Some information regarding trusted sources is given below.

To manage trusted sources, you need to be administrator.

1/ Connect as administrator by clicking Open admin session:



2/ In administration page, go to Trusted sources tab and click Add trusted source



#### 3/ Fill the form and Save

For instance, to add NEUBIAS curated set of workflows, the trusted source has to be configured as follows:

Source code provider: github

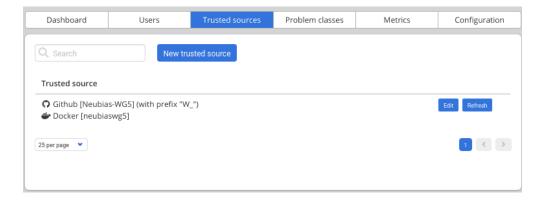
• Source code provider username: Neubias-WG5

• Environment provider: docker

Environment provider username: neubiaswg5

Prefix: W\_

4/ Trusted sources are periodically checked (about every 5/10 minutes) to automatically add new versions of existing workflows or new workflows, but you can also click on **Refresh** to trigger the detection.



5) Once a workflow is imported, it has to be linked to a given **Problem** so as to process all the images from this **Problem**. This can be performed in the Configuration panel of the **Problem** (**Workflows** tab) by toggling **Enable** for that workflow as illustrated below:

