### **Evaluate PQ README**

### General Panoptic Quality evaluation script

Find the panoptic quality match PQ between predicted and GT mask for vessels. The script can run as-is with the example folders supplied.

The script has a few modes class agnostic that will evaluate the PQ value ignoring classification errors.

## **Main Input Parameters**

**GTDir:** contains the path to the LabPics evaluation set that is used as ground truth

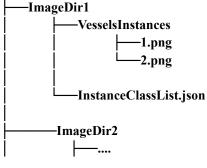
**PredDir:** Contain the link to the predicted instances of the vessels (See Subfolder **ExampleData/Predict/** for example in this folder).

#### **Predicition folder strucutre:**

See Subfolder ExampleData/Predict/ for example prediction folder.

The structure of the predicted dir should be as follows:

### PredDir |---ImageD



**ImageDir:** Should have the same name as the image folder in LabPics evaluation set for every image folder in the GT dir there should be a similar folder in the predicted folder.

ImageDir1\VesselsInstances: Mask of Vessels instances.

ImageDir1\Vessels\1.png: Binary mask for vessel instance 1. ImageDir1\Vessels\1.png: Binary mask for vessel instance 2.

**ImageDir1\InstanceClassList.json:** Contain classes for material instances in vessel 1. For example: {"1": ["Pippete", "Transparent"], "2": [ "Transparent", "Tube"]} means instance 1 is transparent tube and instance 2 is transparent pippete.

# **Additional parameters**

**ClassToUse:** classes that will be used for the evaluation. **LimitToVessel:** only this vessel types if "" check all.

IgnoreVesselsThatAreParts: ignore connectors condensers and stuff.

**MinPixelsInInstace:** Ignore smaller instances (In pixels). **MatchThresh:** IOU threshold for matching instances.

ClassAgnostic: Ignore misclassification errors only use IOU.