

# **HeLP Challenge**

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**Stroke onset time**

# File format

## ➤ Medical Image

- DCM(Digital Imaging and Communications in Medicine)
  - is the international standard for medical images and related information
  - raw data
  - 2D slice 를 여러 장으로 쌓은 형태
  - Included patient and image info
  - .DCM
- NIFTI(Neuroimaging Informatics Technology Initiative)
  - 3D Volume
  - .NII
- Analyze
  - hdr file for metadata, img file for image data

ref

<https://www.dicomstandard.org/>  
<https://nifti.nimh.nih.gov/nifti-1>

# 의료영상처리 라이브러리 (python)

## ➤ Medical Imaging

- SimpleITK: 전반적인 의료영상처리 관련 함수 제공 라이브러리
- opencv-python
- Pydicom
- medpy

## ➤ Neuroimaging (<http://nipy.org/index.html>)

- NiBabel: 의료영상 파일 읽고 쓰기 지원 라이브러리
- NiLearn: 영상 시각화, 머신러닝 등의 영상처리 라이브러리
- NiPype
- NiWidgets

## ➤ 기타

- Numpy
- Scipy
- matplotlib
- Pandas
- scikit-learn

# Radiographic Presentations

**Anterior**

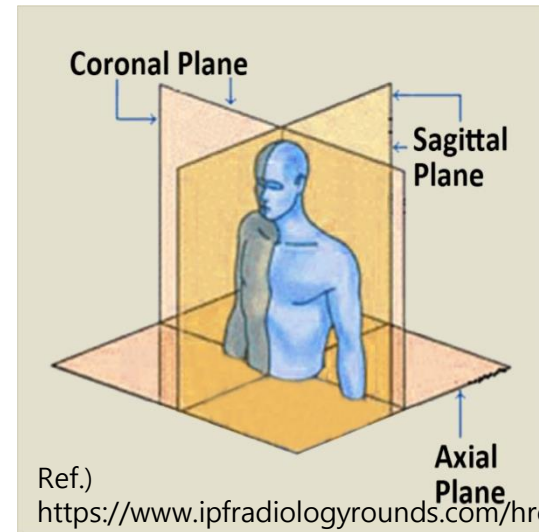


**Axial**

**Right**

**Left**

**Posterior**



Ref.)

<https://www.ipfradiologyrounds.com/hrct-primer/image-reconstruction/>

**Head**



**Coronal**

**Right**

**Left**

**Foot**

**Head**



**Sagittal**

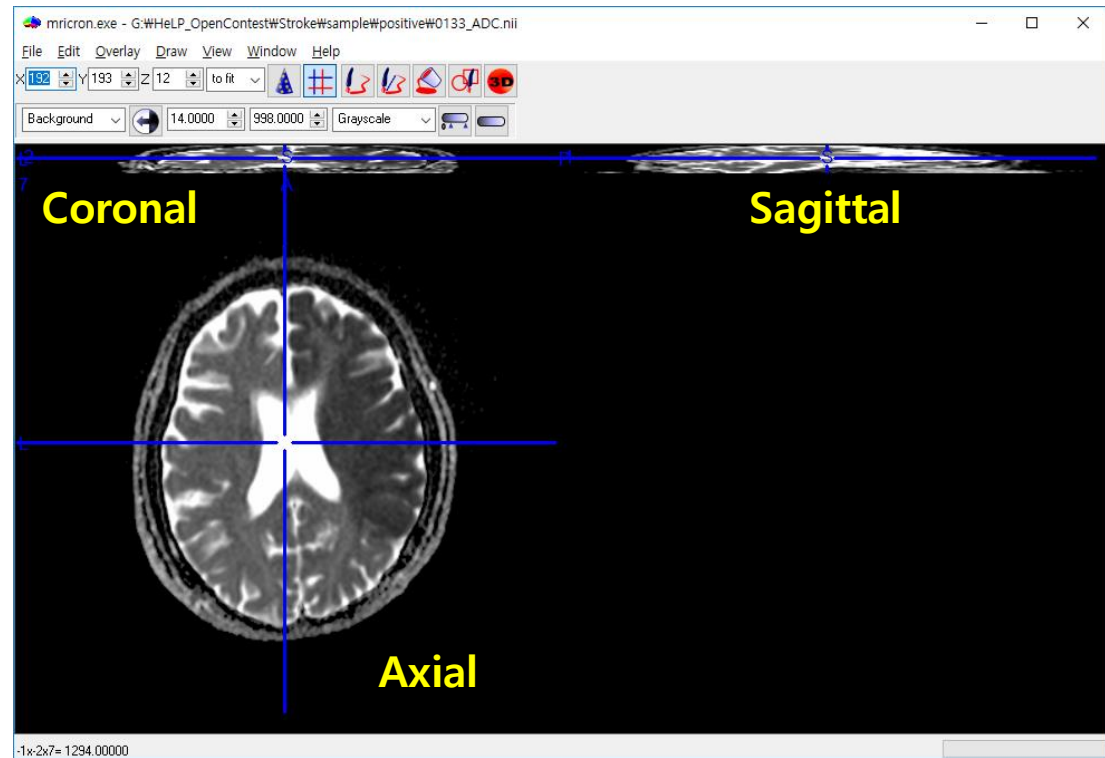
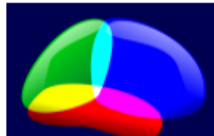
**Anterior**

**Posterior**

**Foot**

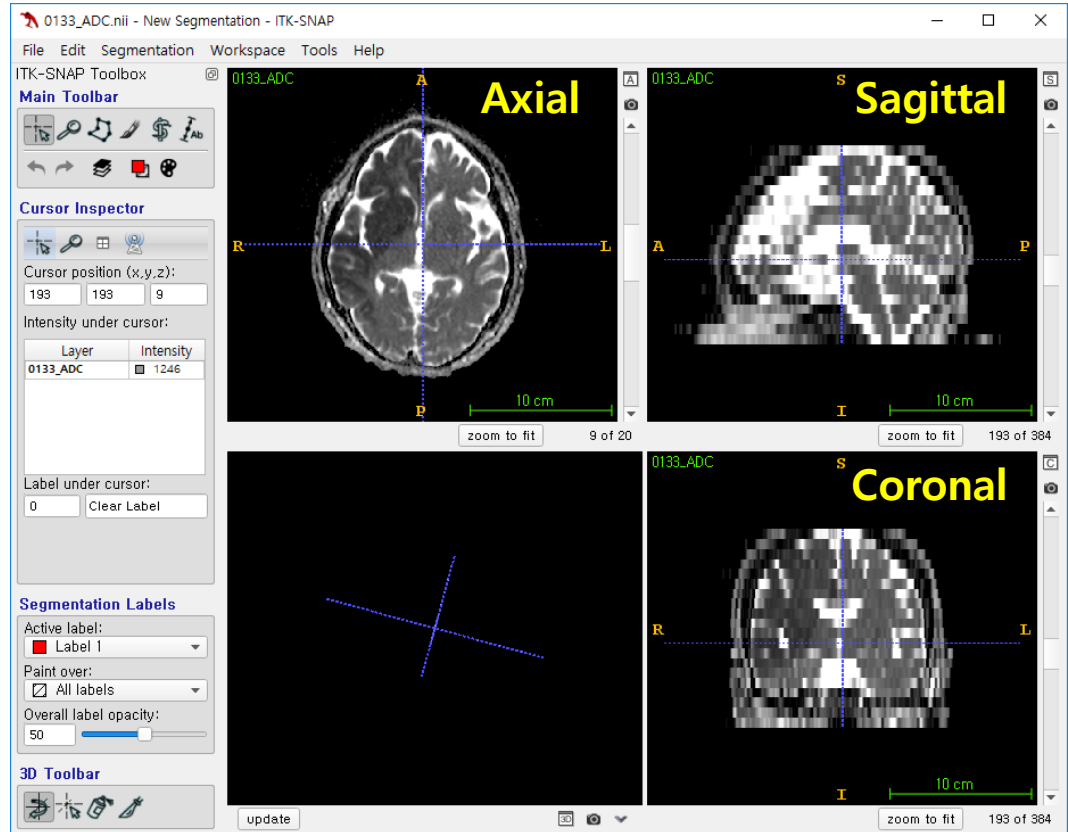
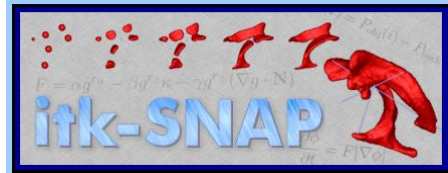
# Medical image viewer

## ➤ MRICron



# Medical image viewer

## ➤ ITK-SNAP



# NIfTI (.nii) file in python (using simpleITK)

## ➤ Read image

```
import SimpleITK as sitk  
imageInput = sitk.ReadImage(file_name_imageT1[n])
```

## ➤ Get size, origin, spacing, direction

```
imageInput = sitk.ReadImage(file_name_imageT1[0])  
  
origin3d = imageInput.GetOrigin()  
spacing3d = imageInput.GetSpacing()  
size3d = imageInput.GetSize()  
direction3d = imageInput.GetDirection()
```

## ➤ Convert itk image to array

```
imageT = sitk.GetArrayFromImage(image) # covert itk image to Array  
model.train_on_batch(imageT, LabelT1)
```

## ➤ Write image

```
model1.load_weights(path_model1)  
result11 = model1.predict(imageT)  
result1 = result11.reshape((img_rows, img_cols, img_dep))  
ResultImg11 = sitk.GetImageFromArray(result1)  
  
final_img1.SetSpacing(original_img.GetSpacing())  
final_img1.SetOrigin(original_img.GetOrigin())  
  
sitk.WriteImage(final_img1, output_filename)
```

# NIfTI (.nii) file in python (using simpleITK)

➤ <http://insightsoftwareconsortium.github.io/SimpleITK-Notebooks/>

➤ **IntensityWindowing**

```
isocubic_img_ww = sitk.Cast(sitk.IntensityWindowing(isocubic_img, LowT, HighT, 0, 255), sitk.sitkUInt8)
```

➤ **Extract**

`ExtractImageFilter()` - Crops an image to the selected region bounds using vectors; Collapses dimensions unless dimension is two

```
# Collapse along the x axis
extractSliceFilter = sitk.ExtractImageFilter()
size = list(mr_image.GetSize())
size[0] = 0
extractSliceFilter.SetSize( size )

index = (x, 0, 0)
extractSliceFilter.SetIndex(index)
sitk_xslic = extractSliceFilter.Execute(mr_image)
```

➤ **Compose**

`ComposeImageFilter()` - Combines several scalar images into a multicomponent vector image

```
# Recompose image (should be same as joined_image)
compose = sitk.ComposeImageFilter()
composed_image = compose.Execute(channel1_image, channel2_image, channel3_image)
```



# Stroke data

## ➤ MRI

- Multi-modal image
  - DWI (diffusion-weighted image) & ADC (apparent diffusion coefficient) map
    - “XXXX\_b1000.nii”
    - “XXXX\_ADC.nii”
  - FLAIR (fluid attenuated inversion recovery)
    - “XXXX\_FLAIR.nii”
    - Coregistered (rigid-transformed) into DWI
  - Brain & infarct masks
    - “XXXX\_brain.nii”
    - “XXXX\_infarct.nii”
    - Coregistered (rigid-transformed) into DWI

# Stroke data

## ➤ MRI

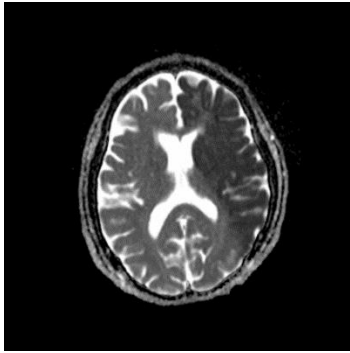
### ▪ Binary classification

- Positive (1): within 4.5 hours from symptom onset
- Negative (0): beyond 4.5 hours from symptom onset

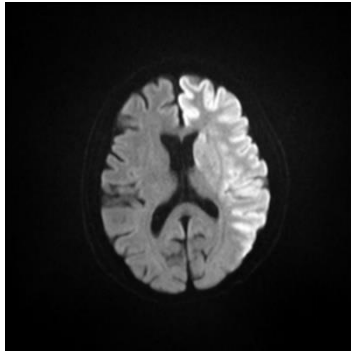
	Positive class	Negative class
Train	123	132
Validation	26	18
Test	33	23

# Stroke data

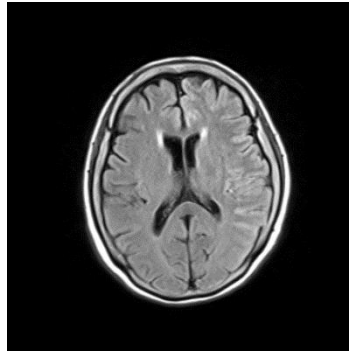
Positive case



ADC



b1000



FLAIR

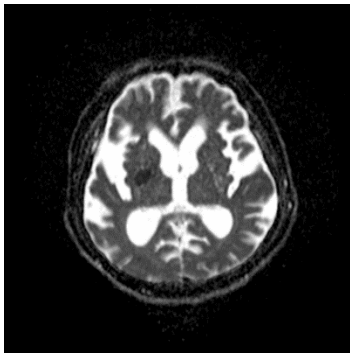


brain

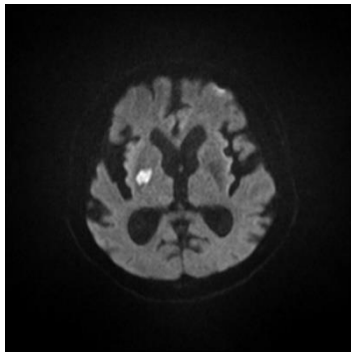


infarct

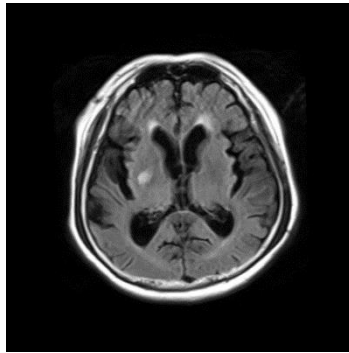
Negative case



ADC



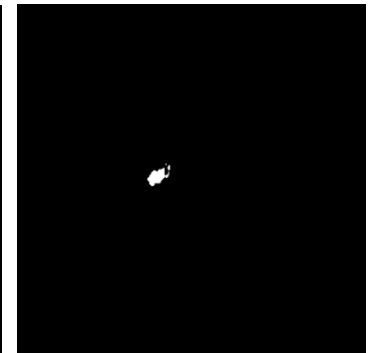
b1000



FLAIR



brain

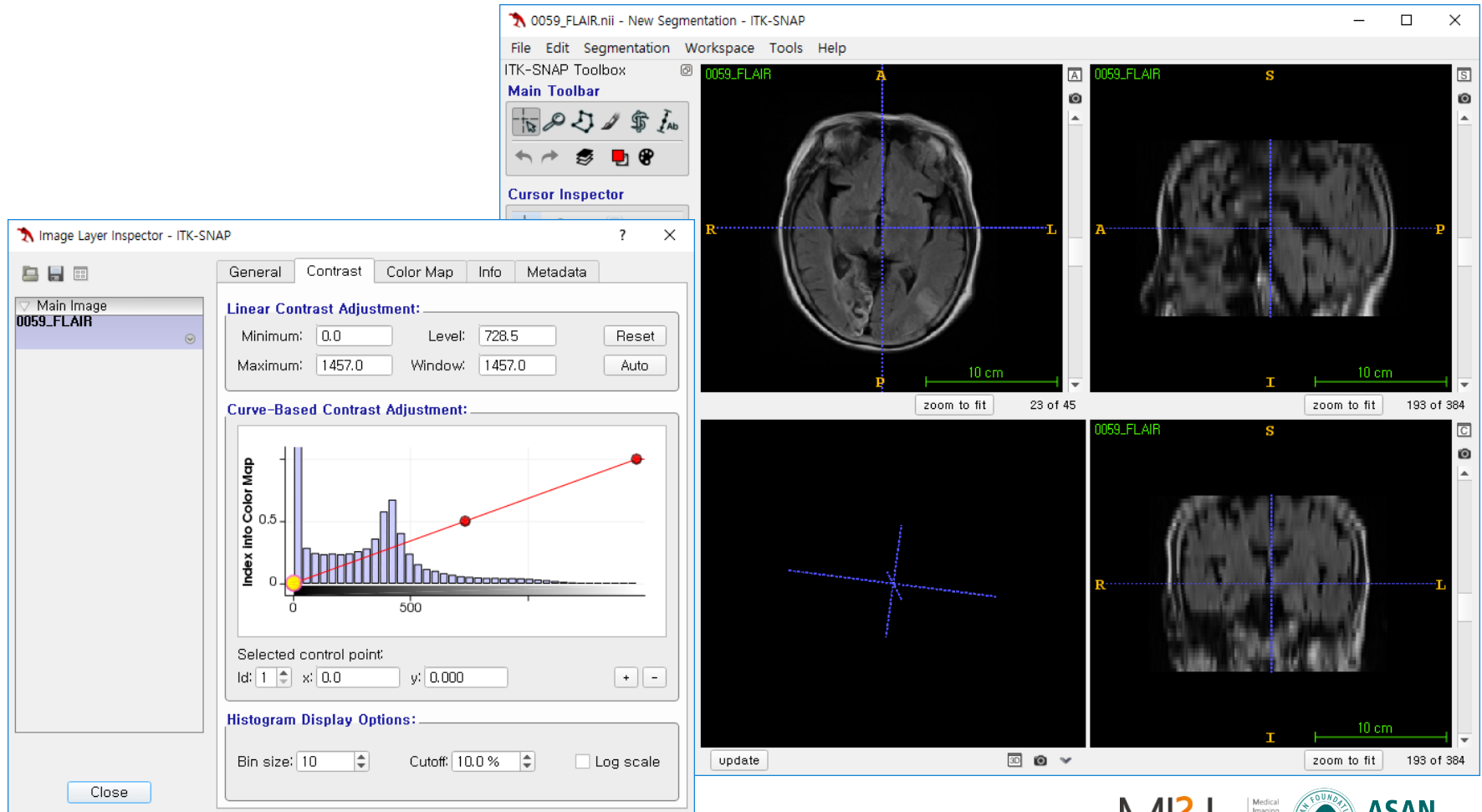


infarct

# Stroke data

## ➤ Preprocessing

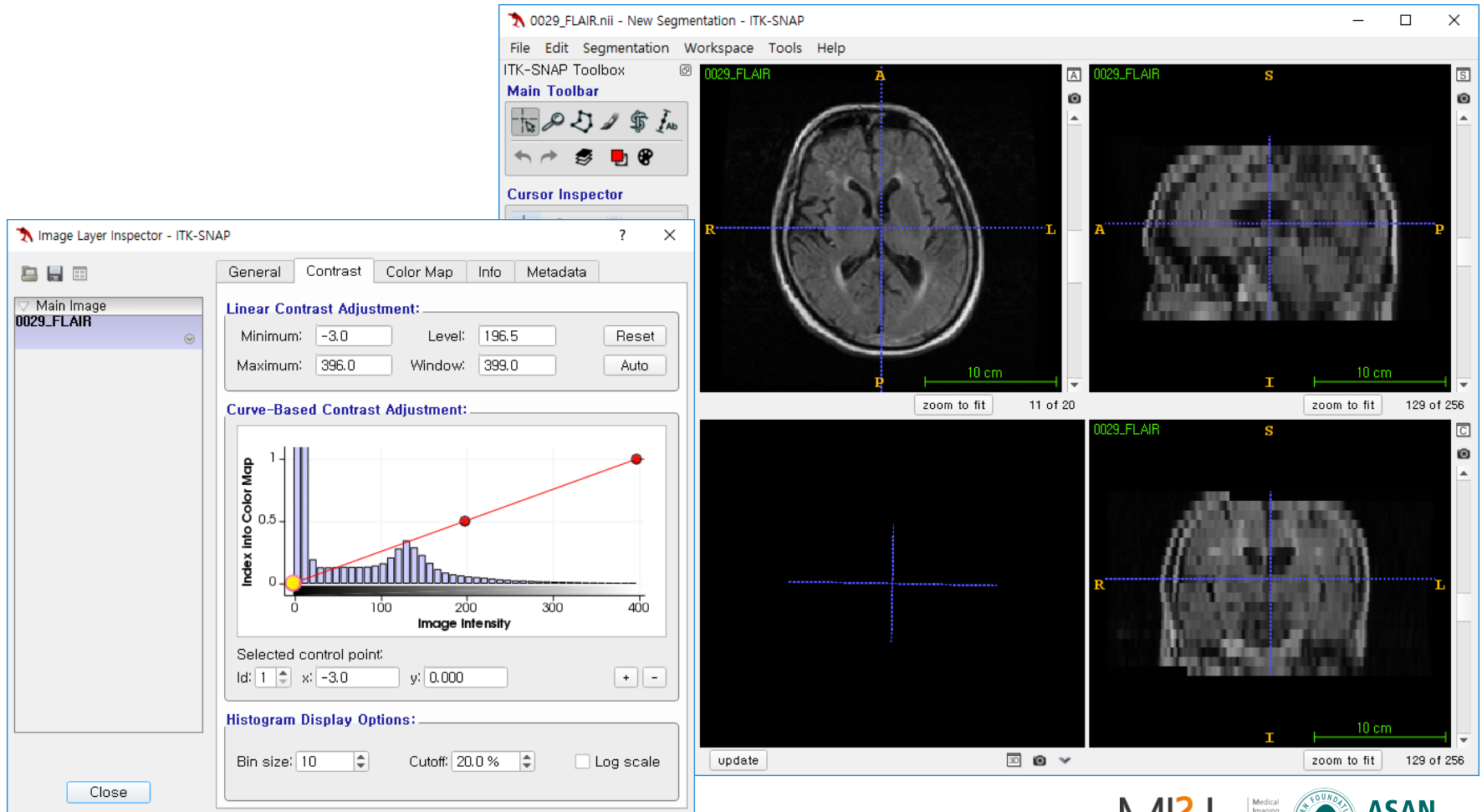
- Intensity normalization



# Stroke data

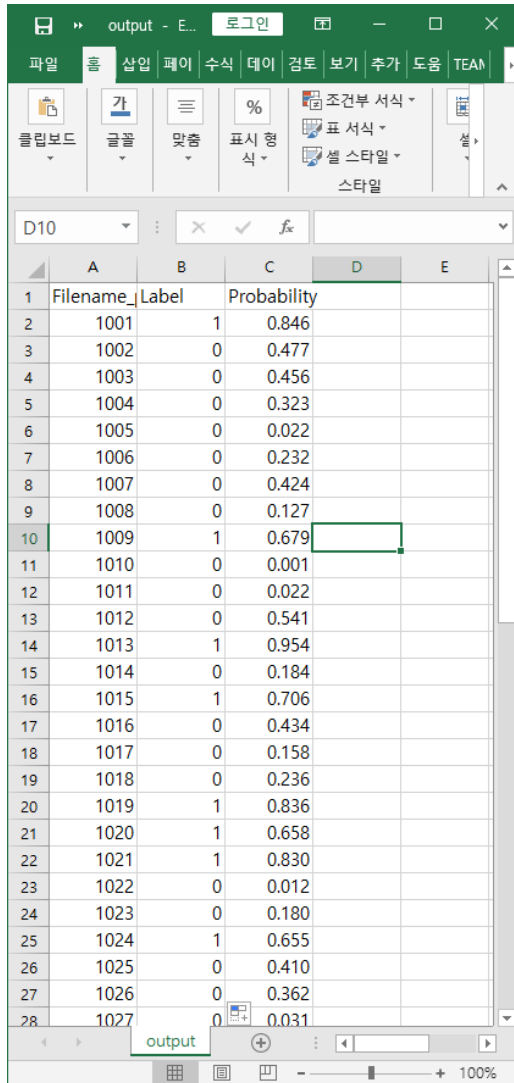
## ➤ Preprocessing

- Intensity normalization



# (Predicted) Output

## ➤ “output.csv”



	A	B	C	D	E
1	Filename_Label		Probability		
2	1001	1	0.846		
3	1002	0	0.477		
4	1003	0	0.456		
5	1004	0	0.323		
6	1005	0	0.022		
7	1006	0	0.232		
8	1007	0	0.424		
9	1008	0	0.127		
10	1009	1	0.679		
11	1010	0	0.001		
12	1011	0	0.022		
13	1012	0	0.541		
14	1013	1	0.954		
15	1014	0	0.184		
16	1015	1	0.706		
17	1016	0	0.434		
18	1017	0	0.158		
19	1018	0	0.236		
20	1019	1	0.836		
21	1020	1	0.658		
22	1021	1	0.830		
23	1022	0	0.012		
24	1023	0	0.180		
25	1024	1	0.655		
26	1025	0	0.410		
27	1026	0	0.362		
28	1027	0	0.031		

# Ranking scheme

## ➤ Evaluation

### ▪ $F_{0.5}$ score

- Weighs recall lower than precision (by attenuating the influence of false negatives)
- [https://en.wikipedia.org/wiki/F1\\_score](https://en.wikipedia.org/wiki/F1_score)

$$F_{\beta} = (1 + \beta^2) \cdot \frac{\text{precision} \cdot \text{recall}}{(\beta^2 \cdot \text{precision}) + \text{recall}}.$$

### ▪ Accuracy

$$\text{Accuracy} = (TP+TN)/(TP+TN+FP+FN)$$

### ▪ AUC (Area Under the Curve)

- [https://en.wikipedia.org/wiki/Receiver\\_operating\\_characteristic](https://en.wikipedia.org/wiki/Receiver_operating_characteristic)

## Clinical Collaborators @ Asan Medical Center

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