By AI

Intracranial Hemorrhage Detection

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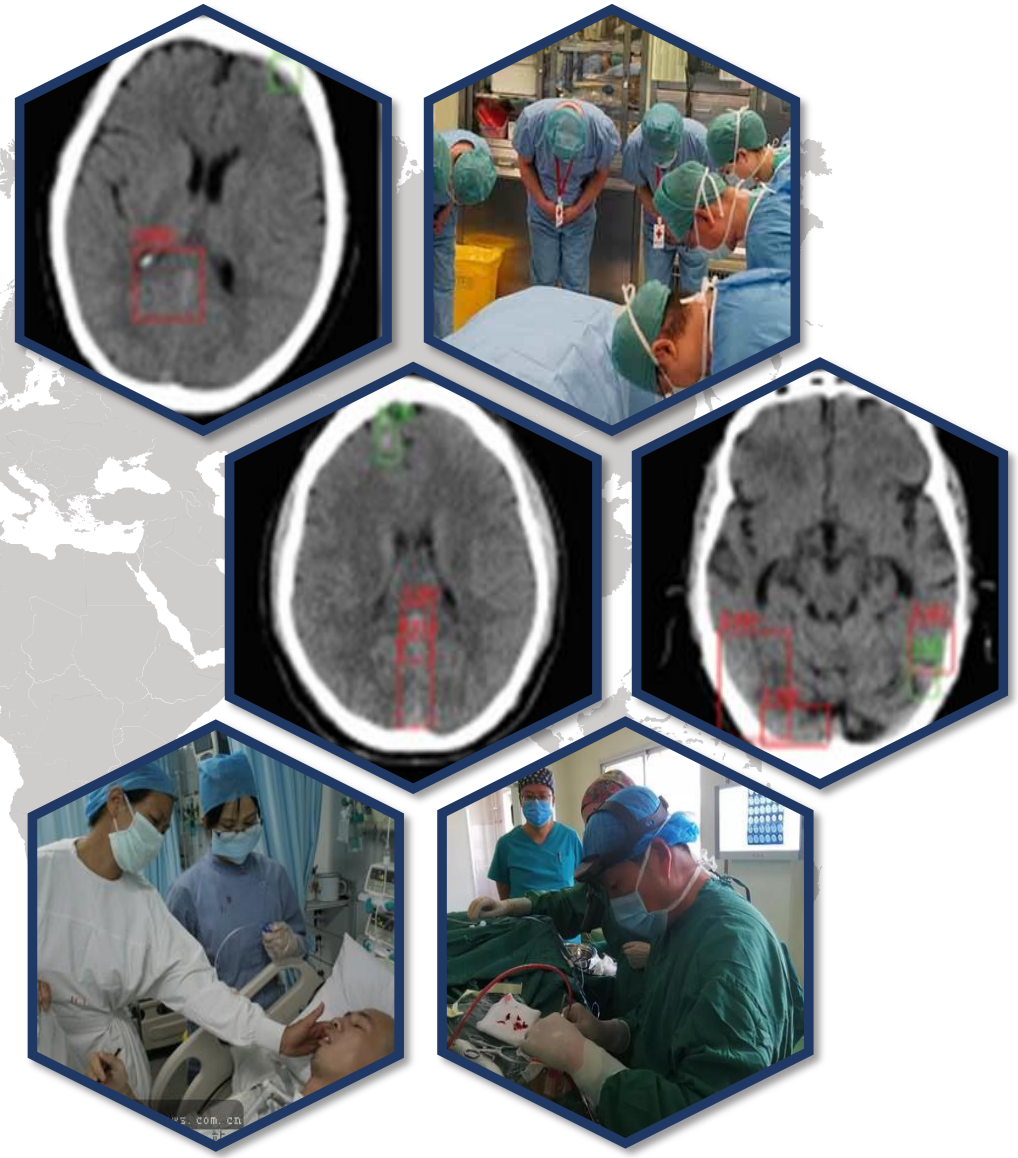
Project Overview

Overview

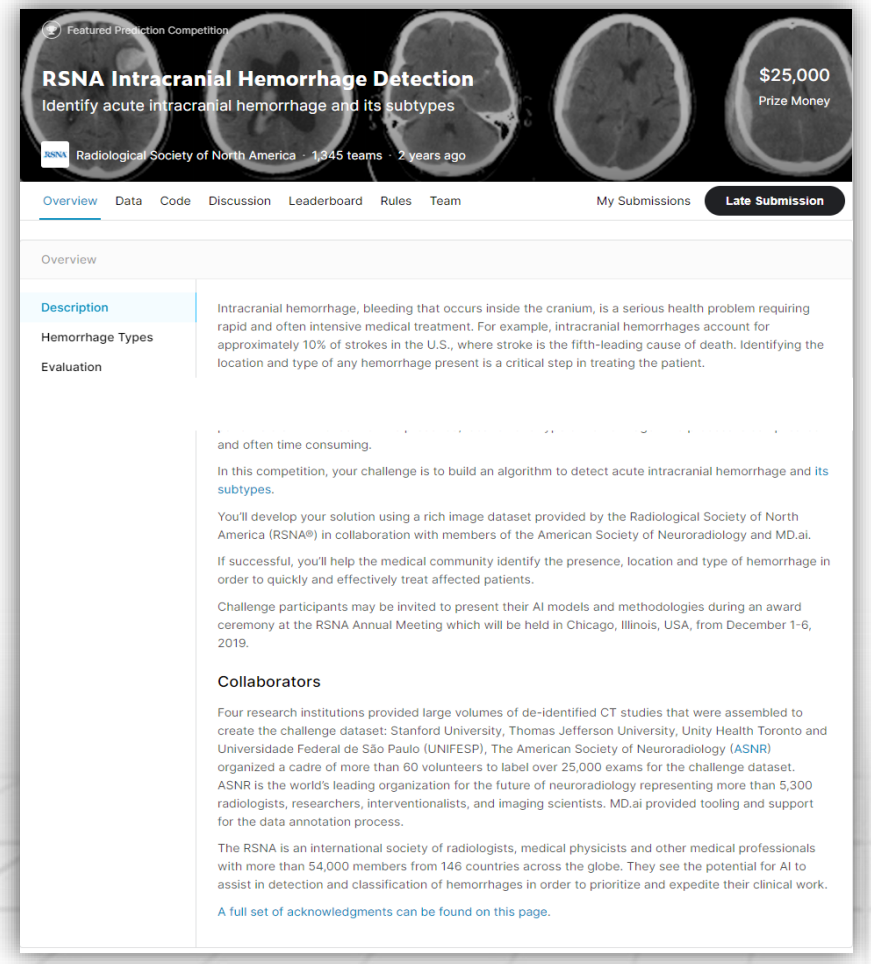
- **Serious** Health Problem
- **Rapid** & **Intensive** Treatment
- **Complicated** Process

- Detection Subtypes(5)
- **Multi**-class Classification
- InceptionNetV3 & Spark

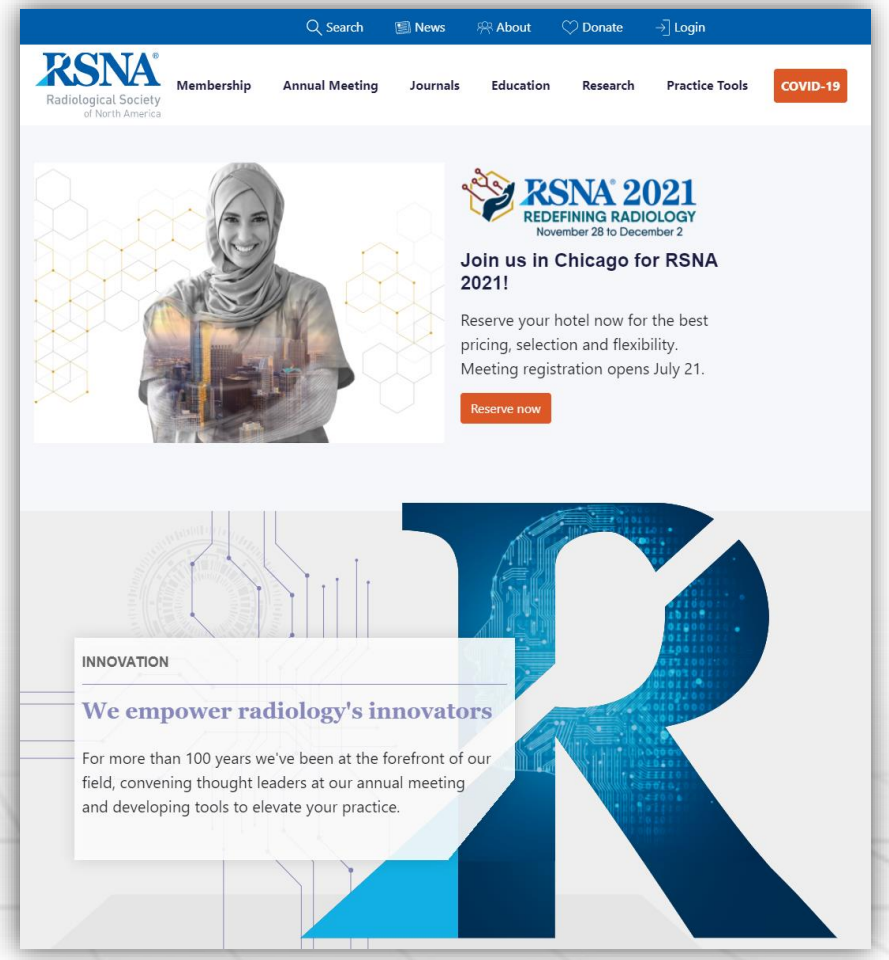


Data Information

Data Origin



kaggle www.kaggle.com



RSNA www.rsna.org

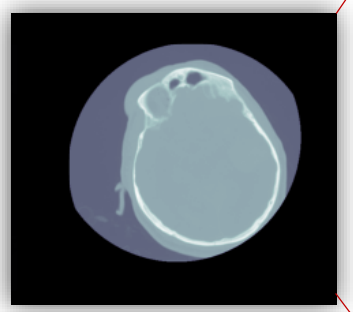
Data Format

CSV

19877

id	label
ID_52c205043	epidural
ID_15e6dc40a	intraventricular
ID_01a97add4	intraparenchymal
ID_9fed7e3db	subdural
ID_903d2ef48	subdural
ID_e0942808d	epidural
ID_d4d31f695	subdural
.....	
ID_c64f8db24	subarachnoid
ID_d202babf2	subarachnoid
ID_9ffd25faa	subdural

DICOM



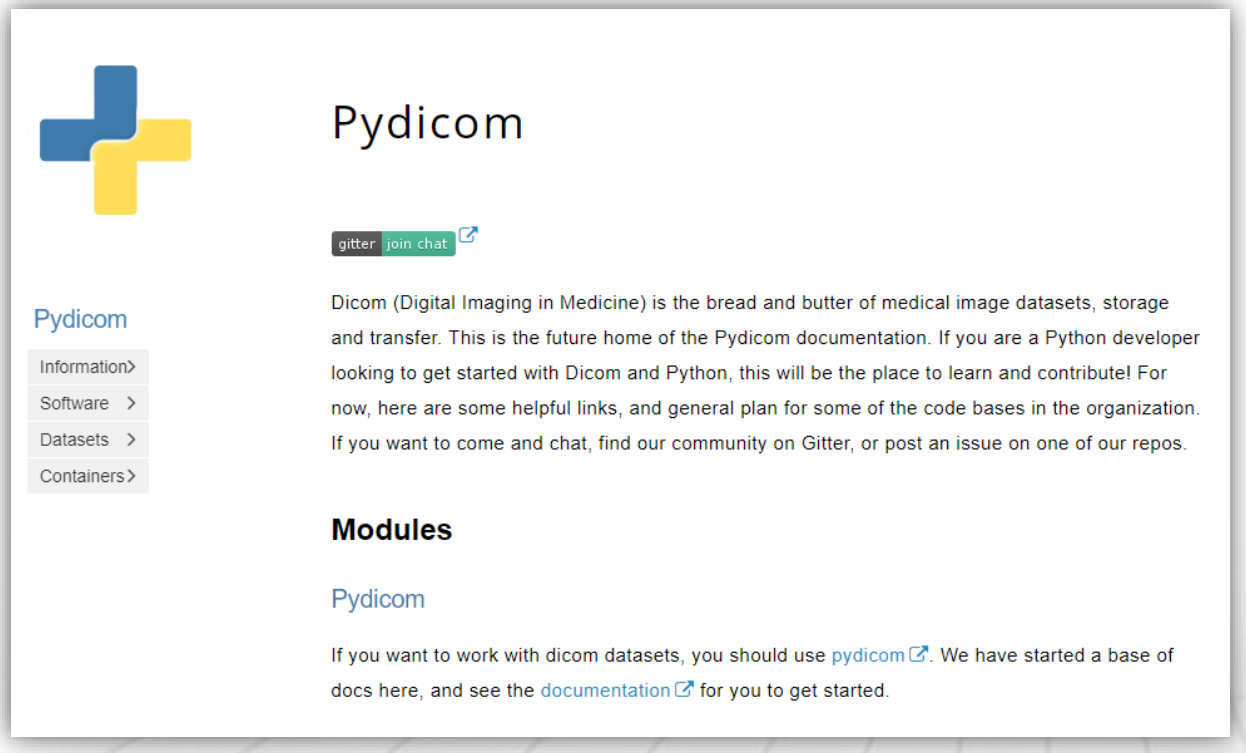
0.53MB

512X512

```
Dataset.file_meta ------------------------------------
(0002, 0000) File Meta Information Group Length  UL: 176
(0002, 0001) File Meta Information Version       OB: b'\x00\x01'
(0002, 0002) Media Storage SOP Class UID         UI: CT Image Storage
(0002, 0003) Media Storage SOP Instance UID      UI: 9999.134563857571520138645562911801058023921
(0002, 0010) Transfer Syntax UID                 UI: Explicit VR Little Endian
(0002, 0012) Implementation Class UID            UI: 1.2.40.0.13.1.1.1
(0002, 0013) Implementation Version Name         SH: 'dcm4che-1.4.38'
-------------------------------------------------
(0008, 0018) SOP Instance UID                    UI: ID_7ce5ae372
(0008, 0060) Modality                            CS: 'CT'
(0010, 0020) Patient ID                          LO: 'ID_314d3781'
(0020, 000d) Study Instance UID                  UI: ID_af0d670f5a
(0020, 000e) Series Instance UID                 UI: ID_4195a174b7
(0020, 0010) Study ID                            SH: ''
(0020, 0032) Image Position (Patient)            DS: [-126.408875, -126.408875, -244.165497]
(0020, 0037) Image Orientation (Patient)         DS: [1.000000, 0.000000, 0.000000, 0.000000, 1.000000, 0.000000]
(0028, 0002) Samples per Pixel                   US: 1
(0028, 0004) Photometric Interpretation          CS: 'MONOCHROME2'
(0028, 0010) Rows                                US: 512
(0028, 0011) Columns                             US: 512
(0028, 0030) Pixel Spacing                       DS: [0.494750976563, 0.494750976563]
(0028, 0100) Bits Allocated                      US: 16
(0028, 0101) Bits Stored                         US: 16
(0028, 0102) High Bit                            US: 15
(0028, 0103) Pixel Representation                US: 1
(0028, 1050) Window Center                       DS: "35.0"
(0028, 1051) Window Width                        DS: "135.0"
(0028, 1052) Rescale Intercept                   DS: "-1024.0"
(0028, 1053) Rescale Slope                       DS: "1.0"
(7fe0, 0010) Pixel Data                          OW: Array of 524288 elements
```

10434.73MB

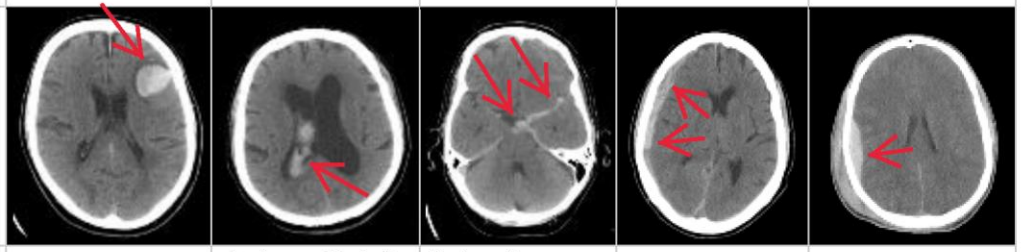
Data Format DICOM

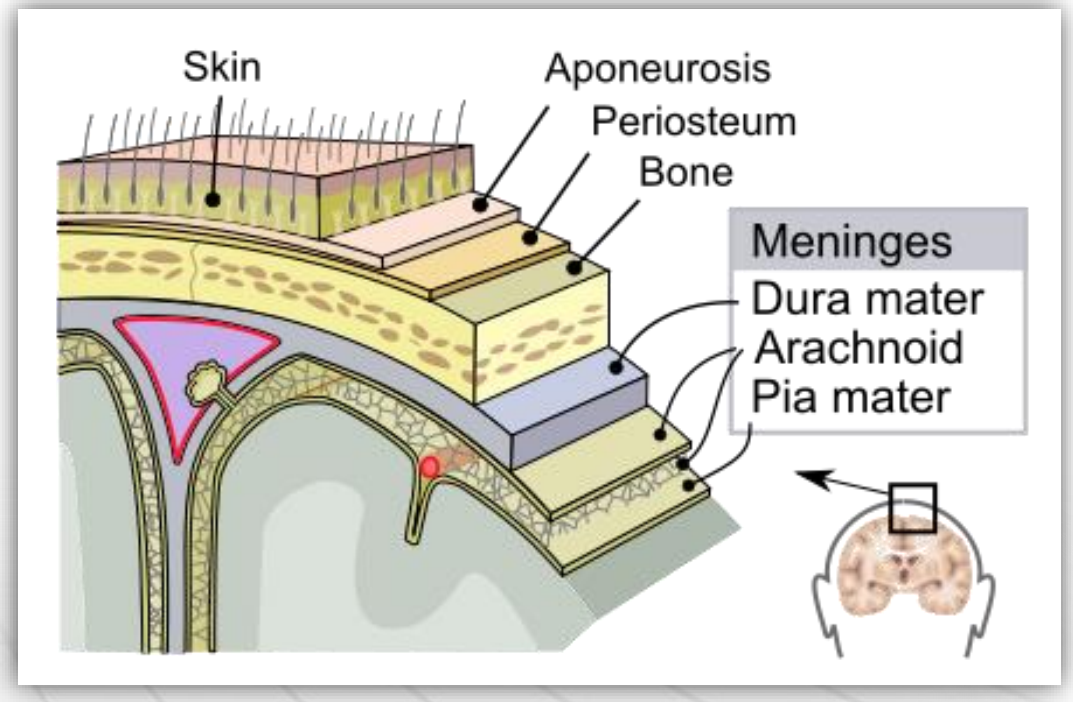


pydicom.github.io

DICOM Digital Imaging and Communications in Medicine www.dicomstandard.org

Classes

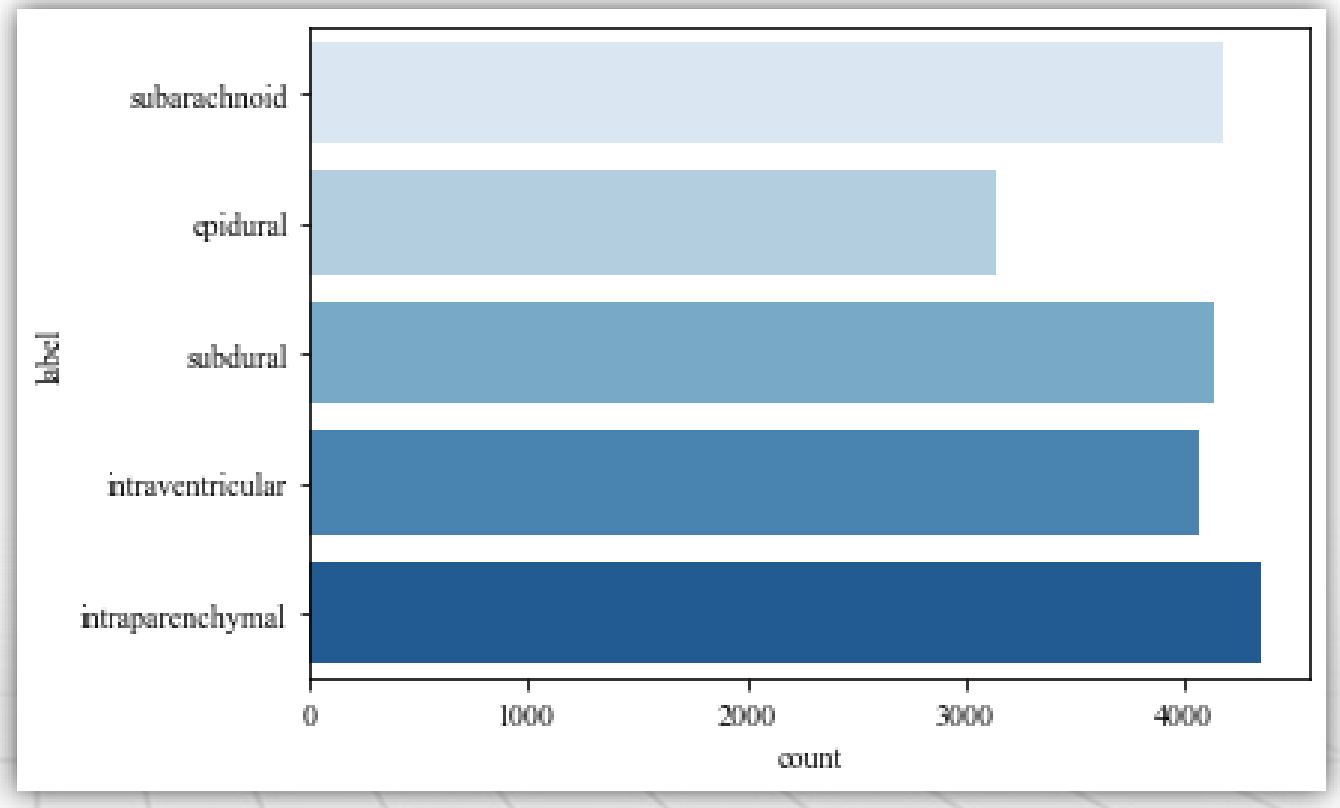
	Intraparenchymal	Intraventricular	Subarachnoid	Subdural	Epidural
Location	Inside of the brain	Inside of the ventricle	Between the arachnoid and the pia mater	Between the Dura and the arachnoid	Between the dura and the skull
Imaging					
Mechanism	High blood pressure, trauma, arteriovenous malformation, tumor, etc	Can be associated with both intraparenchymal and subarachnoid hemorrhages	Rupture of aneurysms or arteriovenous malformations or trauma	Trauma	Trauma or after surgery
Source	Arterial or venous	Arterial or venous	Predominantly arterial	Venous (bridging veins)	Arterial
Shape	Typically rounded	Conforms to ventricular shape	Tracks along the sulci and fissures	Crescent	Lentiform
Presentation	Acute (sudden onset of headache, nausea, vomiting)	Acute (sudden onset of headache, nausea, vomiting)	Acute (worst headache of life)	May be insidious (worsening headache)	Acute (skull fracture and altered mental status)



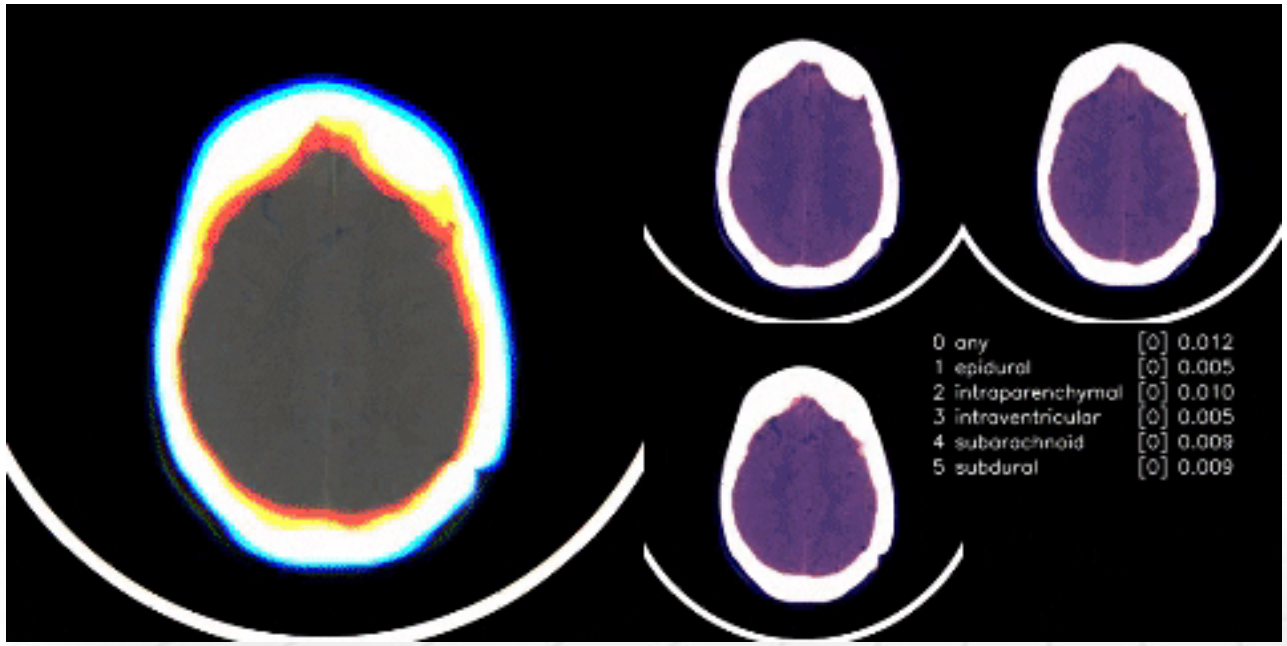
Exploratory Data Analysis

Classes

label	count
subarachnoid	4177
epidural	3145
subdural	4132
intraventricular	4072
intraparenchymal	4351



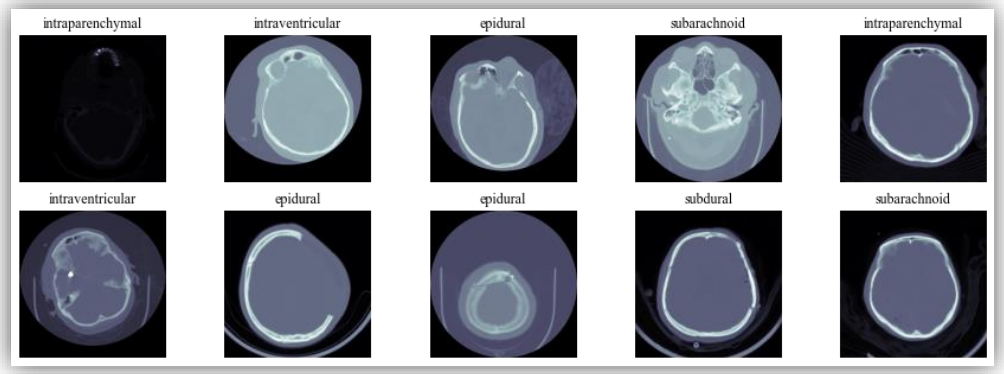
Visualization for Different Windows



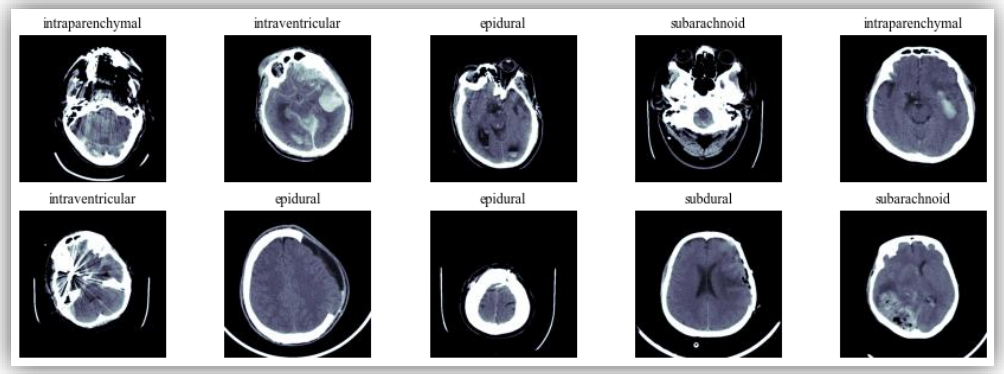
Window:also known as ~grey-level mapping，contrast stretching，histogram modification or contrast enhancement` is the process in which the CT image greyscale component of an image is manipulated via the CT numbers; doing this will change the appearance of the picture to highlight particular structures. The brightness of the image is, adjusted via the window level. The contrast is adjusted via the window width.

Visualization for Different Windows

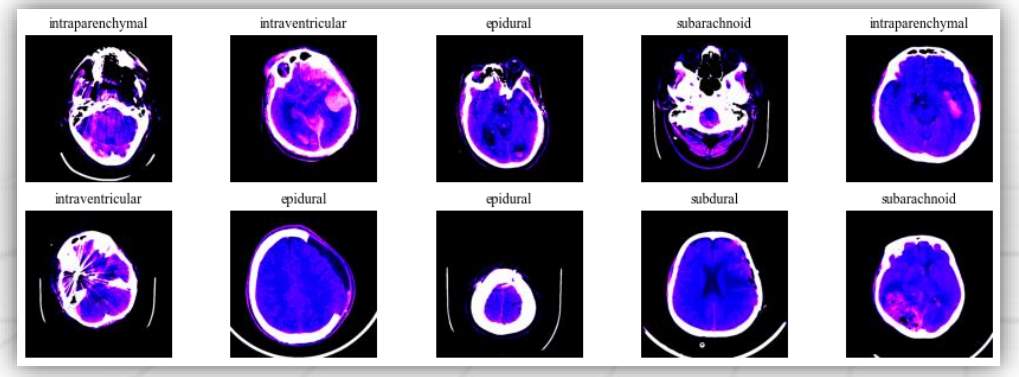
No Windows



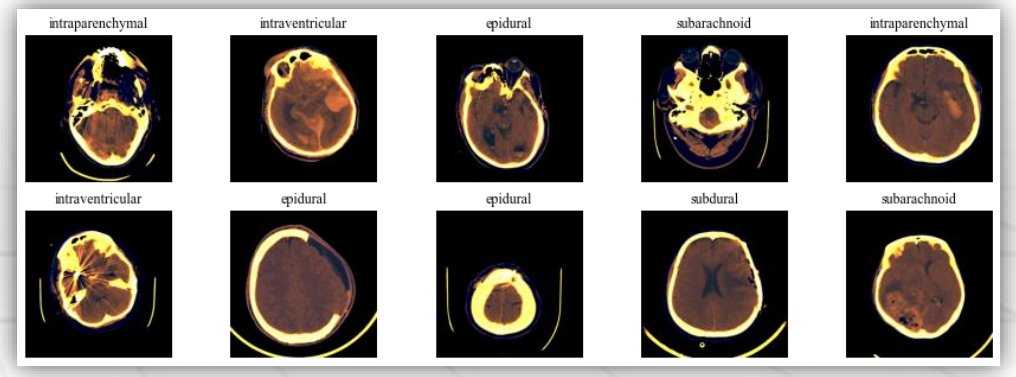
Brain Windows



Three Channels Windows

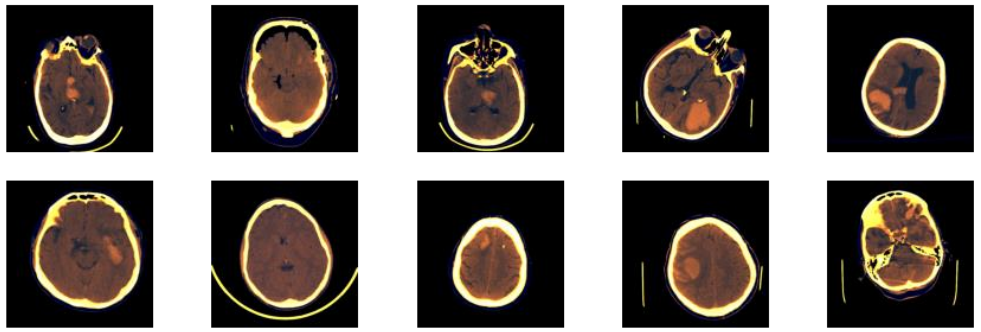


Brain + Subdural + Bone Windows

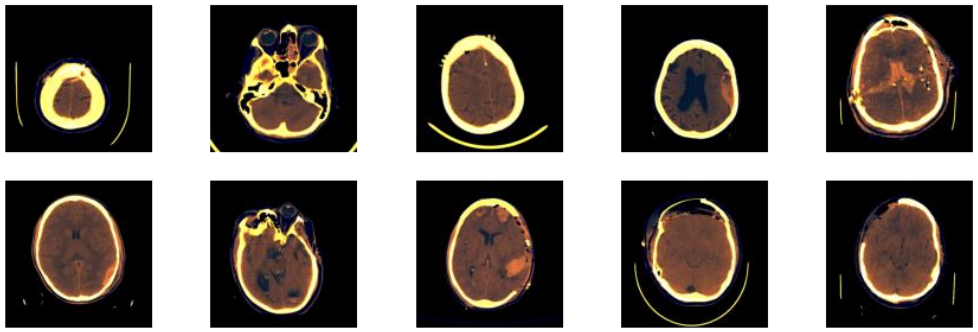


Visualization for Different Classes

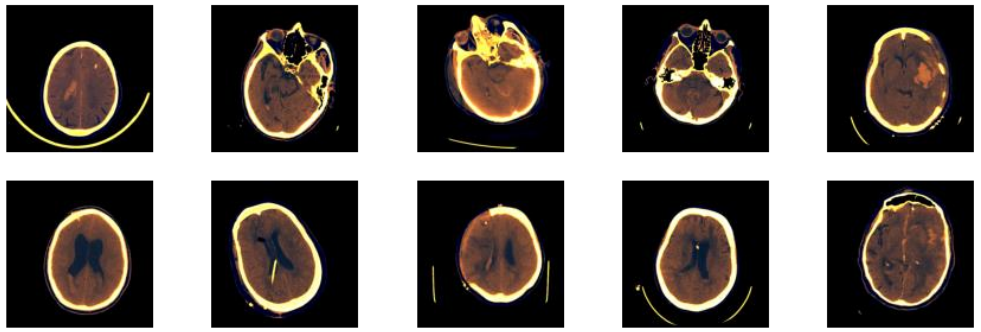
Instances with Intraparenchymal Hemorrhage



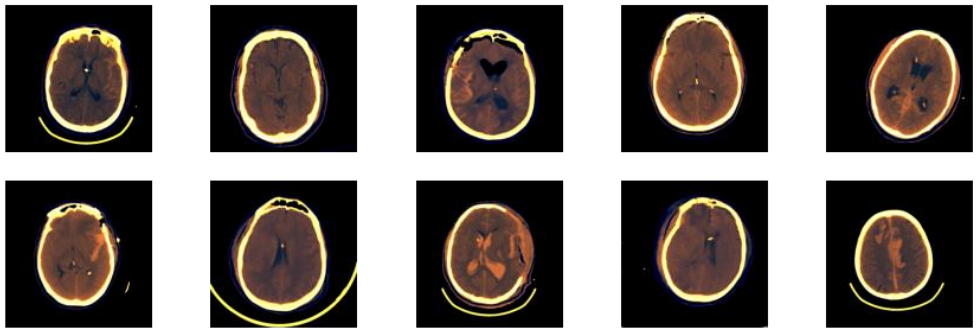
Instances with Epidural Hemorrhage



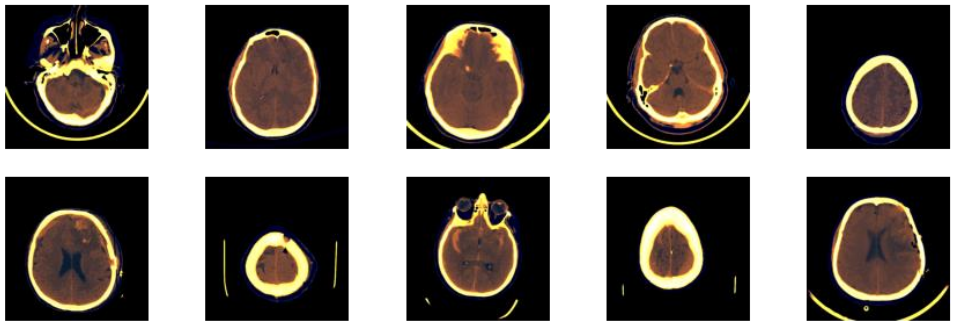
Instances with Intraventricular Hemorrhage



Subarachnoid



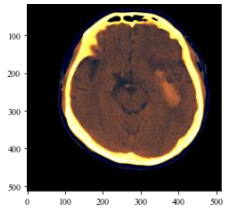
Subdural



Data Preprocess

Data Preprocess

Get windowed image



Rescale image into 299X299

Apply operation by RDD method

Flatten

```
+------------+--------------------+
|          id|               image|
+------------+--------------------+
|ID_7ce5ae372|[0.0, 0.0, 0.0, 0...|
|ID_ace564fe2|[0.0, 0.0, 0.0, 0...|
|ID_9416259b1|[0.0, 0.0, 0.0, 0...|
|ID_928806565|[0.0, 0.0, 0.0, 0...|
|ID_e97dd15eb|[0.0, 0.0, 0.0, 0...|
+------------+--------------------+
only showing top 5 rows
```

Add Label

```
+------------+--------------------+----------------+
|          id|               image|           label|
+------------+--------------------+----------------+
|ID_7ce5ae372|[0.0, 0.0, 0.0, 0...|intraparenchymal|
|ID_ace564fe2|[0.0, 0.0, 0.0, 0...|intraventricular|
|ID_9416259b1|[0.0, 0.0, 0.0, 0...|        epidural|
|ID_928806565|[0.0, 0.0, 0.0, 0...|     subarachnoid|
|ID_e97dd15eb|[0.0, 0.0, 0.0, 0...|intraparenchymal|
+------------+--------------------+----------------+
only showing top 5 rows
```

Map Label in to Numeric Value

```
+----------------+----------+
|           label|labelIndex|
+----------------+----------+
|     subarachnoid|       3.0|
|intraventricular|       2.0|
|        epidural|       0.0|
|intraparenchymal|       1.0|
|        subdural|       4.0|
+----------------+----------+
```

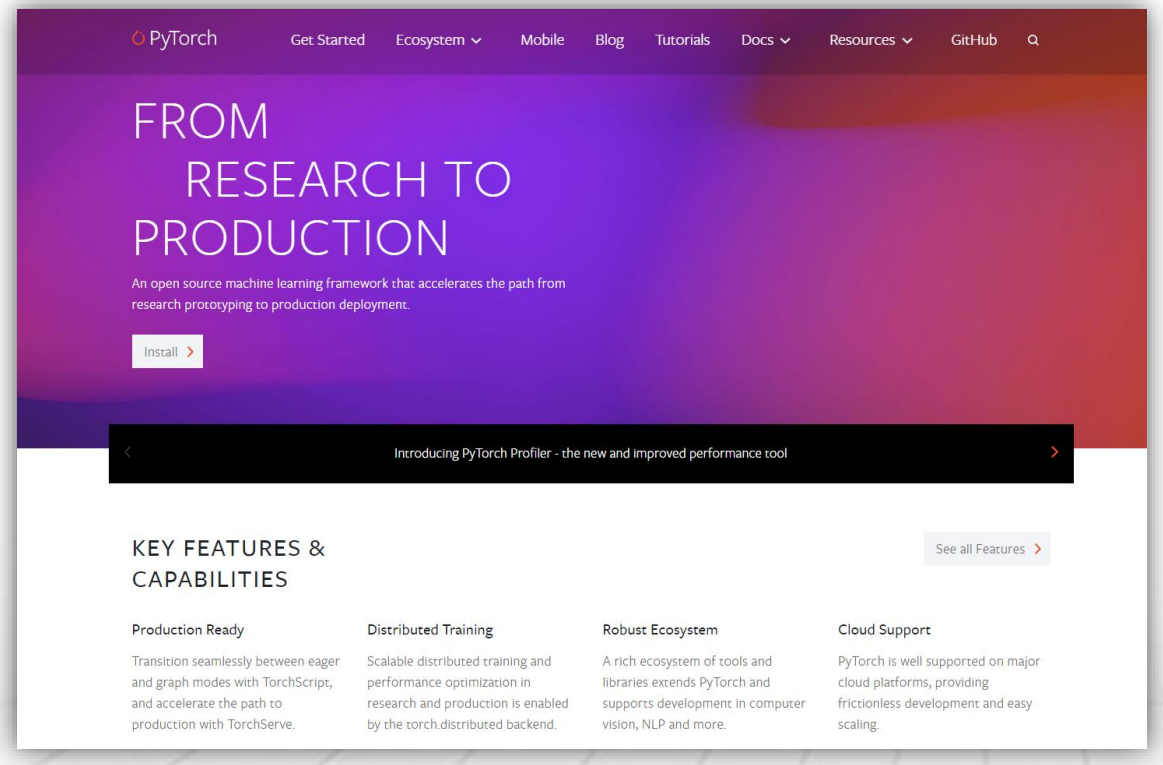
Split

```
Size of training dataset: 15951
Size of testing dataset: 3926
```

Save

Model

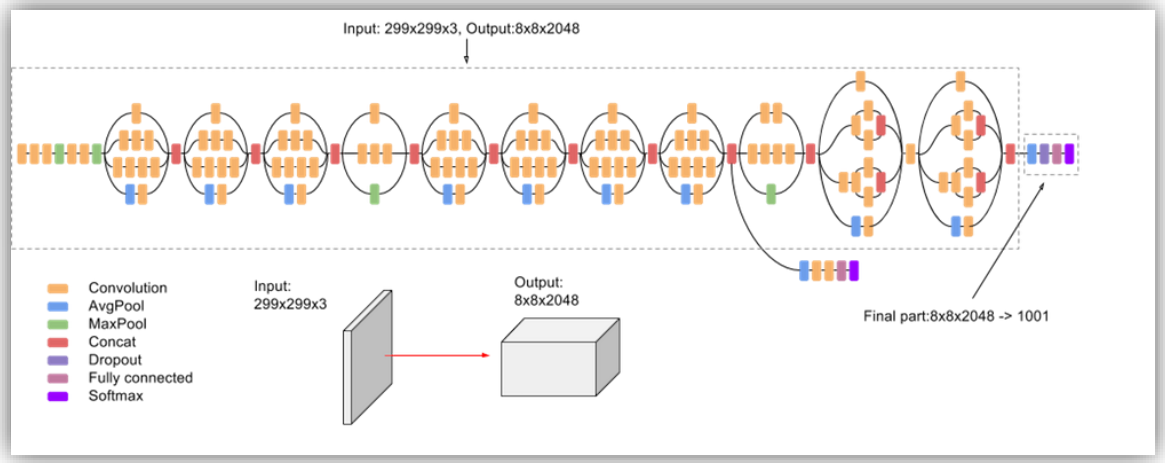
Tools

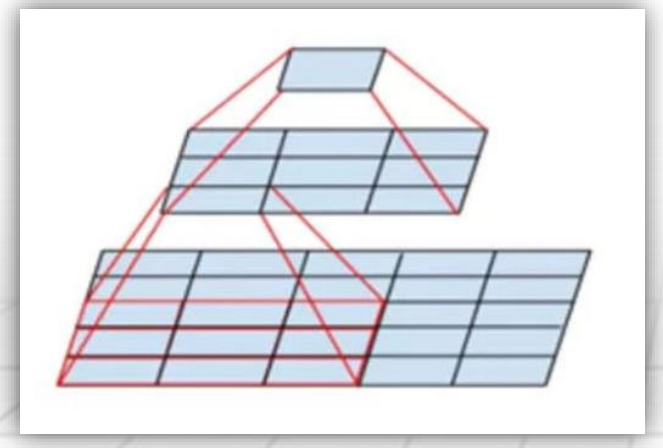


PyTorch **pytorch.org**

APACHE Spark PyTorch **github.com/dmmiller612/sparktorch**

InceptionNetV3





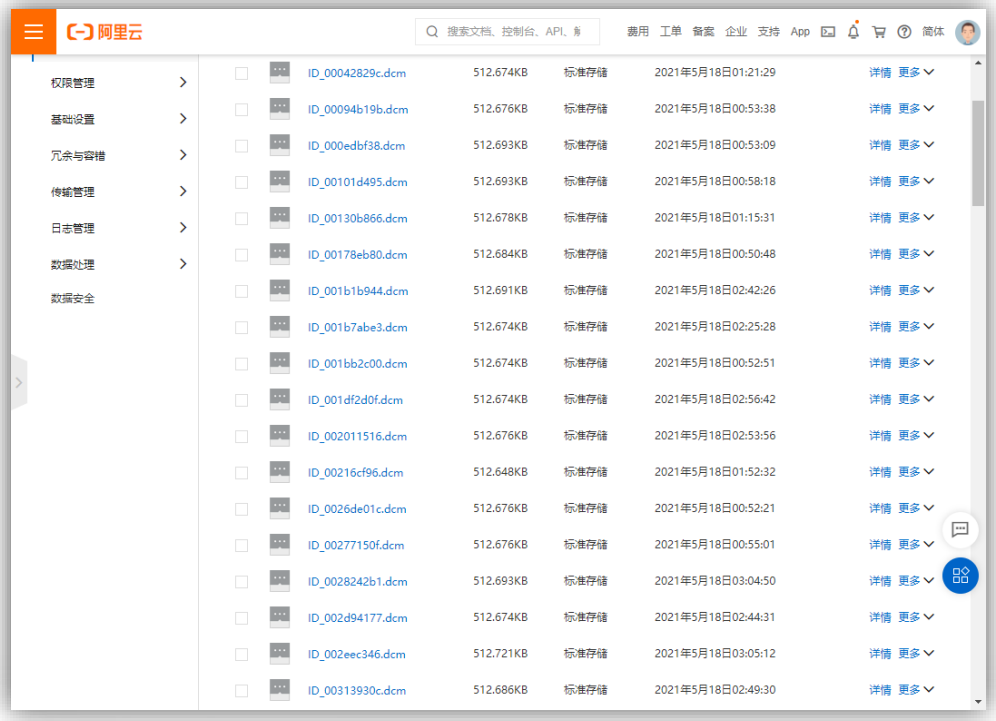
Model Structure

```python
class InceptionNet3(nn.Module):
    def __init__(self, num_classes=5):
        super(InceptionNet3, self).__init__()
        self.Conv2d_1a_3x3 = BasicConv2d(3, 32, kernel_size=
3, stride=2)
        self.Conv2d_2a_3x3 = BasicConv2d(32, 32, kernel_size
=3)
        self.Conv2d_2b_3x3 = BasicConv2d(32, 64, kernel_size
=3, padding=1)
        self.Conv2d_3b_1x1 = BasicConv2d(64, 80, kernel_size
=1)
        self.Conv2d_4a_3x3 = BasicConv2d(80, 192, kernel_siz
e=3)
        self.Mixed_5b = InceptionA(192, pool_features=32)
        self.Mixed_5c = InceptionA(256, pool_features=64)
        self.Mixed_5d = InceptionA(288, pool_features=64)
        self.Mixed_6a = InceptionB(288)
        self.Mixed_6b = InceptionC(768, channels_7x7=128)
        self.Mixed_6c = InceptionC(768, channels_7x7=160)
        self.Mixed_6d = InceptionC(768, channels_7x7=160)
        self.Mixed_6e = InceptionC(768, channels_7x7=192)
        self.Mixed_7a = InceptionD(768)
        self.Mixed_7b = InceptionE(1280)
        self.Mixed_7c = InceptionE(2048)
        self.fc = nn.Linear(2048, num_classes)
        for m in self.modules():
            if isinstance(m, nn.Conv2d) or isinstance(m, nn.
Linear):
                import scipy.stats as stats
                stddev = m.stddev if hasattr(m, 'stddev') e
lse 0.1 #if else
                X = stats.truncnorm(-2, 2, scale=stddev)
                values = torch.Tensor(X.rvs(m.weight.data.nu
mel()))
                values = values.view(m.weight.data.size())
                m.weight.data.copy_(values)
            elif isinstance(m, nn.BatchNorm2d):
                m.weight.data.fill_(1)
                m.bias.data.zero_()
```

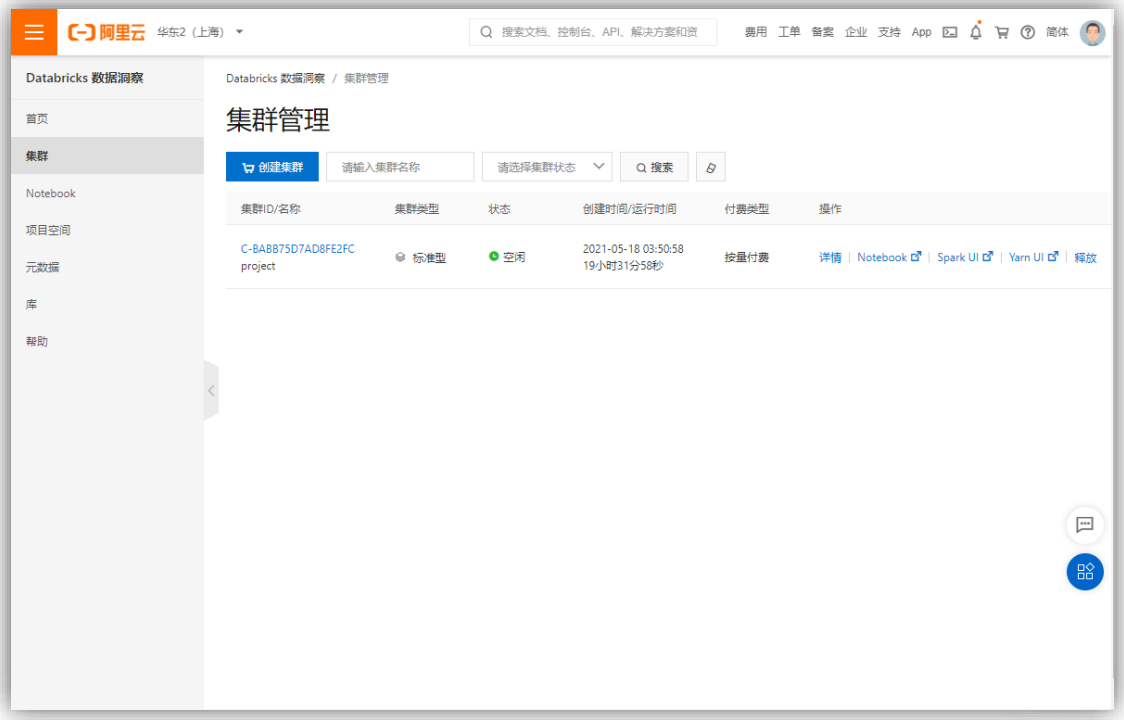
```python
def forward(self, x):
    x.view(-1,3,INPUT_SIZE[0],INPUT_SIZE[1])
    x = self.Conv2d_1a_3x3(x)
    x = self.Conv2d_2a_3x3(x)
    x = self.Conv2d_2b_3x3(x)
    x = F.max_pool2d(x, kernel_size=3, stride=2)
    x = self.Conv2d_3b_1x1(x)
    x = self.Conv2d_4a_3x3(x)
    x = F.max_pool2d(x, kernel_size=3, stride=2)
    x = self.Mixed_5b(x)
    x = self.Mixed_5c(x)
    x = self.Mixed_5d(x)
    x = self.Mixed_6a(x)
    x = self.Mixed_6b(x)
    x = self.Mixed_6c(x)
    x = self.Mixed_6d(x)
    x = self.Mixed_6e(x)
    x = self.Mixed_7a(x)
    x = self.Mixed_7b(x)
    x = self.Mixed_7c(x)
    x = F.avg_pool2d(x, kernel_size=8)
    x = F.dropout(x, training=self.training)
    x = x.view(x.size(0), -1)
    x = self.fc(x)
    return x
```

Model Code

Training



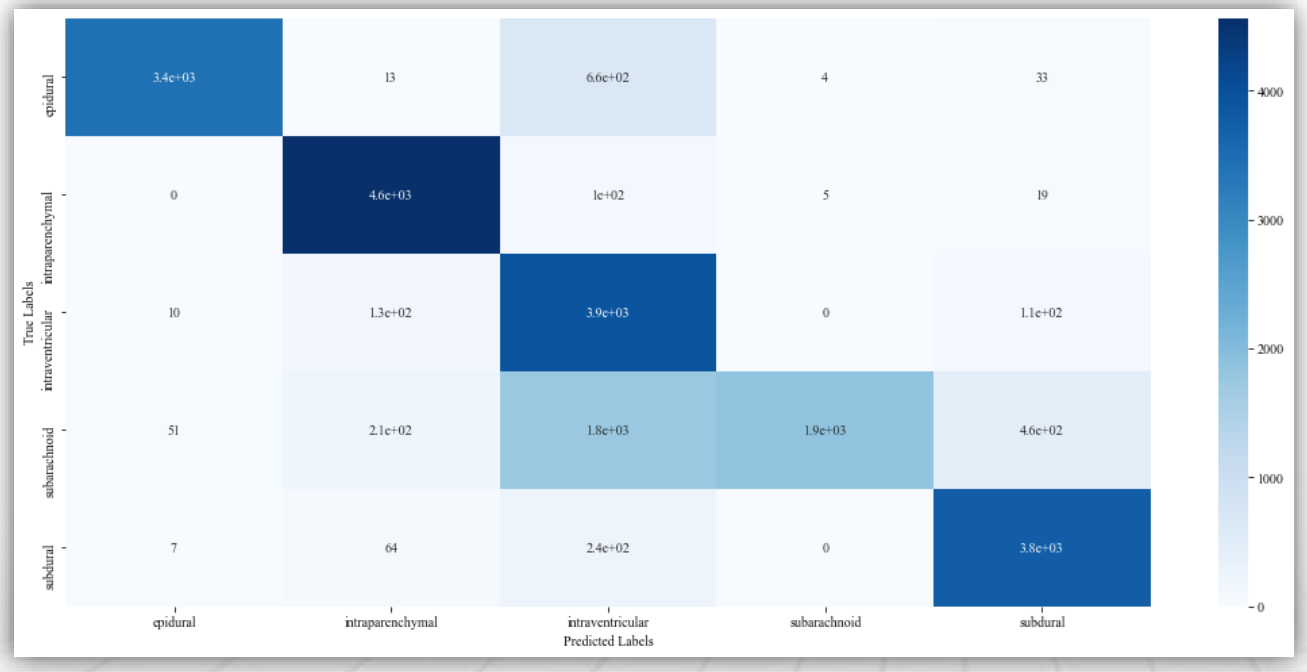
OSS



Databricks

Evaluation

Evaluation

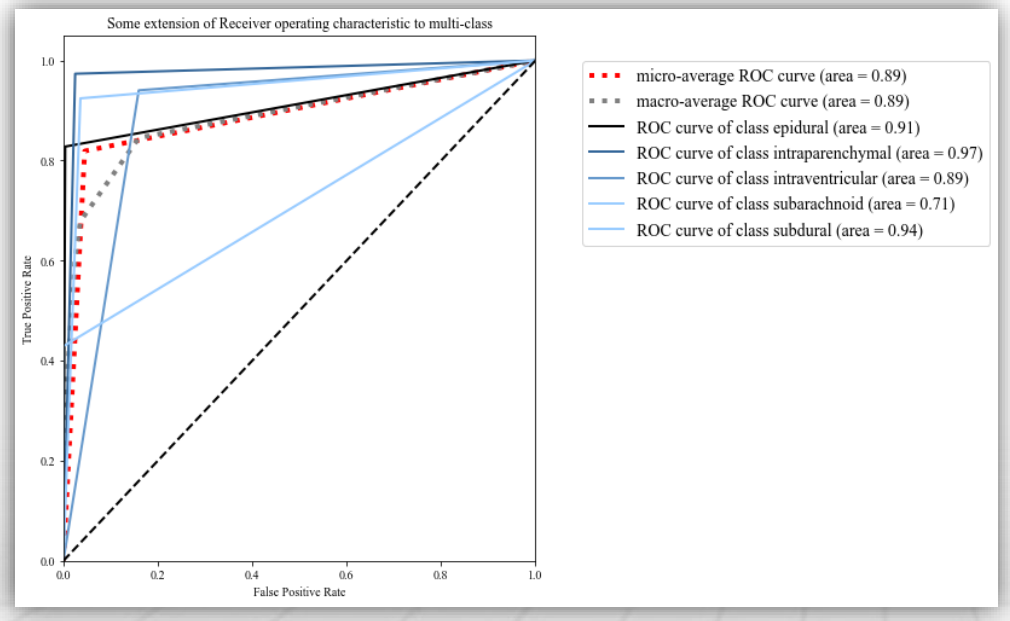


Confusion Matrix

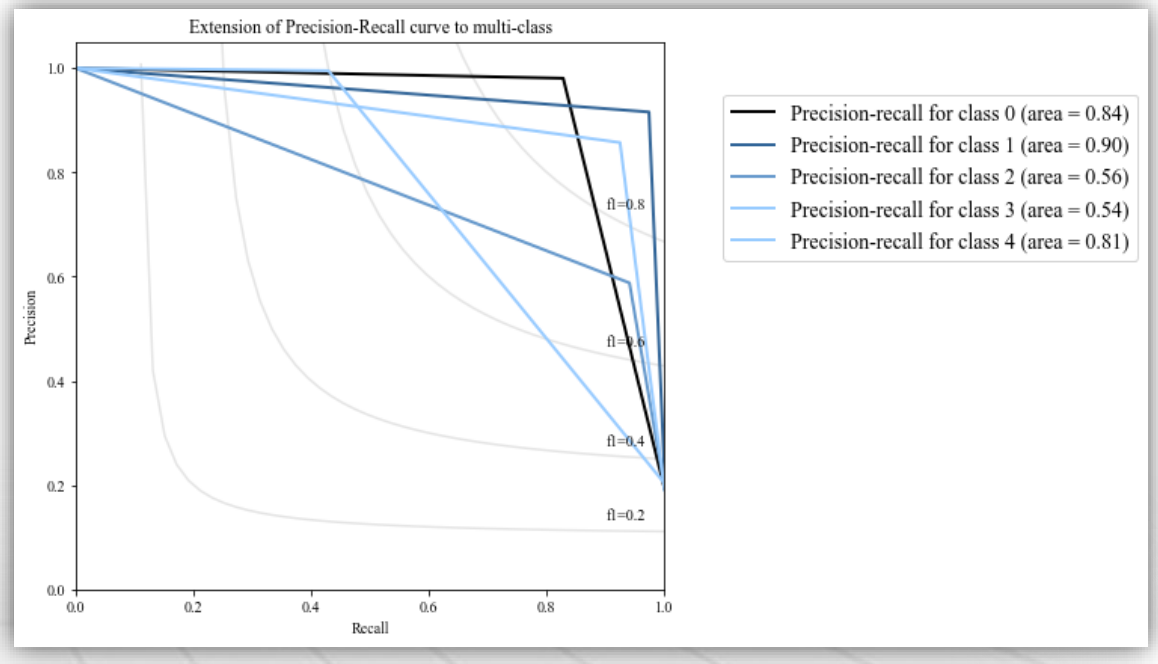
	Score
F1-Score	0.811718
Precision	0.86934
Recall	0.819014
Accuracy	0.819014

Metrics

Evaluation



ROC Curve



PR Curve

Thank for Listening!