!pip install livelossplot

Looking in indexes: https://us-python.pkg.dev/colab-wheels/public/simple/ Collecting livelossplot Downloading livelossplot-0.5.5-py3-none-any.whl (22 kB) Requirement already satisfied: bokeh in /usr/local/lib/python3.9/dist-packages (from livelossplot) (2.4 Requirement already satisfied: matplotlib in /usr/local/lib/python3.9/dist-packages (from livelossplot Requirement already satisfied: Jinja2>=2.9 in /usr/local/lib/python3.9/dist-packages (from bokeh->live) Requirement already satisfied: packaging>=16.8 in /usr/local/lib/python3.9/dist-packages (from bokeh-> Requirement already satisfied: typing-extensions>=3.10.0 in /usr/local/lib/python3.9/dist-packages (from Requirement already satisfied: numpy>=1.11.3 in /usr/local/lib/python3.9/dist-packages (from bokeh->liv Requirement already satisfied: PyYAML>=3.10 in /usr/local/lib/python3.9/dist-packages (from bokeh->live Requirement already satisfied: pillow>=7.1.0 in /usr/local/lib/python3.9/dist-packages (from bokeh->liv Requirement already satisfied: tornado>=5.1 in /usr/local/lib/python3.9/dist-packages (from bokeh->live Requirement already satisfied: pyparsing>=2.2.1 in /usr/local/lib/python3.9/dist-packages (from matplo Requirement already satisfied: fonttools>=4.22.0 in /usr/local/lib/python3.9/dist-packages (from matple Requirement already satisfied: cycler>=0.10 in /usr/local/lib/python3.9/dist-packages (from matplotlib-Requirement already satisfied: kiwisolver>=1.0.1 in /usr/local/lib/python3.9/dist-packages (from matple Requirement already satisfied: python-dateutil>=2.7 in /usr/local/lib/python3.9/dist-packages (from material) Requirement already satisfied: MarkupSafe>=2.0 in /usr/local/lib/python3.9/dist-packages (from Jinja2>: Requirement already satisfied: six>=1.5 in /usr/local/lib/python3.9/dist-packages (from python-dateuti Installing collected packages: livelossplot Successfully installed livelossplot-0.5.5

```
# Import basic required packages
import numpy as np
import pandas as pd
import cv2
import matplotlib.pyplot as plt
import time
import seaborn as sns
import os
from datetime import datetime as dt
# Import Tensor Flow and Scikit
import tensorflow as tf
from sklearn.metrics import classification_report
from sklearn.linear model import LogisticRegression
from sklearn.metrics import accuracy score
from sklearn.model_selection import train_test_split
from sklearn.metrics import classification_report
from sklearn.svm import SVC
from sklearn.model_selection import GridSearchCV
from sklearn.ensemble import RandomForestRegressor
from sklearn.neighbors import KNeighborsClassifier
from sklearn.ensemble import GradientBoostingClassifier
from sklearn.metrics import confusion matrix
from sklearn.model selection import KFold
from keras.models import Model
from keras.optimizers import Adam
from keras.applications import vgg16, vgg19, resnet, inception v3, MobileNetV3Small, mobilenet v3
from keras.preprocessing.image import ImageDataGenerator
from keras.callbacks import ModelCheckpoint, EarlyStopping
from keras.layers import Dense, Dropout, Flatten
from keras.utils import to_categorical
from livelossplot.inputs.keras import PlotLossesCallback
from tensorflow.keras import backend as K
import warnings
warnings.filterwarnings("ignore")
```

1. Load the datasets

For the project, we provide a training set with 50000 images in the directory ../data/images/ with:

- noisy labels for all images provided in .../data/noisy label.csv;
- clean labels for the first 10000 images provided in ../data/clean labels.csv.

```
from google.colab import drive
drive.mount('/content/drive')
```

Drive already mounted at /content/drive; to attempt to forcibly remount, call drive.mount("/content/drive.

```
# [DO NOT MODIFY THIS CELL]
# load the images
n_{img} = 50000
n \text{ noisy} = 40000
n_clean_noisy = n_img - n_noisy
\#imgs = np.empty((n_img, 32, 32, 3))
#for i in range(n_img):
    #img_fn = f'../data/images/{i+1:05d}.png'
    #imgs[i,:,:,:]=cv2.cvtColor(cv2.imread(img_fn),cv2.COLOR_BGR2RGB)
#converting above data to numpy file 'imgs.npy'
# load the labels
#clean_labels = np.genfromtxt('../data/clean_labels.csv', delimiter=',', dtype="int8")
#noisy_labels = np.genfromtxt('.../data/noisy_labels.csv', delimiter=',', dtype="int8")
# Converting the arrays from images above and reload
data path = '/content/drive/MyDrive/data/'
imgs = np.load(data_path + 'imgs.npy')
# load the labels
clean labels = np.genfromtxt(data path + 'clean labels.csv', delimiter=',', dtype="int8")
noisy labels = np.genfromtxt(data_path + 'noisy labels.csv', delimiter=',', dtype="int8")
```

For illustration, we present a small subset (of size 8) of the images with their clean and noisy labels in clean noisy trainset. You are encouraged to explore more characteristics of the label noises on the whole dataset.

```
# [DO NOT MODIFY THIS CELL]

fig = plt.figure()

ax1 = fig.add_subplot(2,4,1)
ax1.imshow(imgs[0]/255)
ax2 = fig.add_subplot(2,4,2)
ax2.imshow(imgs[1]/255)
ax3 = fig.add_subplot(2,4,3)
ax3.imshow(imgs[2]/255)
ax4 = fig.add_subplot(2,4,4)
ax4.imshow(imgs[3]/255)
ax1 = fig.add_subplot(2,4,5)
ax1.imshow(imgs[4]/255)
ax2 = fig.add_subplot(2,4,6)
ax2.imshow(imgs[5]/255)
ax3 = fig.add_subplot(2,4,7)
```

```
Inception_V3.ipynb - Colaboratory
uno - 119.000_200P100(2,1,1)
ax3.imshow(imgs[6]/255)
ax4 = fig.add_subplot(2,4,8)
ax4.imshow(imgs[7]/255)
# The class-label correspondence
classes = ('plane', 'car', 'bird', 'cat',
           'deer', 'dog', 'frog', 'horse', 'ship', 'truck')
# print clean labels
print('Clean labels:')
print(' '.join('%5s' % classes[clean_labels[j]] for j in range(8)))
# print noisy labels
print('Noisy labels:')
print(' '.join('%5s' % classes[noisy_labels[j]] for j in range(8)))
    Clean labels:
                                      car bird horse
     frog truck truck deer
                               car
    Noisy labels:
                                           bird deer
             dog truck
                        frog
                               dog
                                    ship
```

2. The predictive model

We consider a baseline model directly on the noisy dataset without any label corrections. RGB histogram features are extracted to fit a logistic regression model.

2.1. Baseline Model

Logistic regression is the classification algorithm that is usually used to predict probability of binary categorical outcome. In the image classification context, input features are the pixel values of images and the outcomes are class of label images.

```
# [DO NOT MODIFY THIS CELL]
# The dataset consists of two parts: a target vector of noisy labels (y), and a feature matrix of RGB histog
# RGB histogram dataset construction
no bins = 6
bins = np.linspace(0,255,no_bins) # the range of the rgb histogram
target vec = np.empty(n img)
feature mtx = np.empty((n img, 3*(len(bins)-1)))
i = 0
for i in range(n_img):
    # The target vector consists of noisy labels
    target_vec[i] = noisy_labels[i]
    # Use the numbers of pixels in each bin for all three channels as the features
    feature1 = np.histogram(imgs[i][:,:,0],bins=bins)[0]
    feature2 = np.histogram(imgs[i][:,:,1],bins=bins)[0]
    feature3 = np.histogram(imgs[i][:,:,2],bins=bins)[0]
```

```
# Concatenate three features
  feature_mtx[i,] = np.concatenate((feature1, feature2, feature3), axis=None)
  i += 1 # Each row in the feature matrix corresponds to a single image in the dataset, and the columns re

# [DO NOT MODIFY THIS CELL]
# Train a logistic regression model
clf = LogisticRegression(random_state=0).fit(feature_mtx, target_vec)
```

For the convenience of evaluation, we write the following function $predictive_model$ that does the label prediction. For your predictive model, feel free to modify the function, but make sure the function takes an RGB image of numpy.array format with dimension $32 \times 32 \times 3$ as input, and returns one single label as output.

```
# [DO NOT MODIFY THIS CELL]
def baseline_model(image):
    This is the baseline predictive model that takes in the image and returns a label prediction
    feature1 = np.histogram(image[:,:,0],bins=bins)[0]
    feature2 = np.histogram(image[:,:,1],bins=bins)[0]
    feature3 = np.histogram(image[:,:,2],bins=bins)[0]
    feature = np.concatenate((feature1, feature2, feature3), axis=None).reshape(1,-1)
    return clf.predict(feature)
fig = plt.figure()
ax1 = fig.add_subplot(2,4,1)
ax1.imshow(imgs[0]/255)
ax2 = fig.add_subplot(2,4,2)
ax2.imshow(imgs[1]/255)
ax3 = fig.add subplot(2,4,3)
ax3.imshow(imgs[2]/255)
ax4 = fig.add_subplot(2,4,4)
ax4.imshow(imgs[3]/255)
ax1 = fig.add_subplot(2,4,5)
ax1.imshow(imgs[4]/255)
ax2 = fig.add subplot(2,4,6)
ax2.imshow(imgs[5]/255)
ax3 = fig.add_subplot(2,4,7)
ax3.imshow(imgs[6]/255)
ax4 = fig.add subplot(2,4,8)
ax4.imshow(imgs[7]/255)
print('Clean labels:')
print(' '.join('%5s' % classes[clean_labels[j]] for j in range(8)))
print('Predicted baseline labels:')
print(' '.join('%5s' % classes[int(baseline model(imgs[j])[0])] for j in range(8)))
```

Clean labels:
frog truck truck deer car car bird horse
Predicted baseline labels:

2.2. Model I



In this model, the first thing to do is to implement data augmentation by resizing the images to 75 x 75 x 3 to capture
more details and features -- increasing the spatial resolution. This array of 'imgs_load' is then saved under 'img_load.py'
and splitted under 75:25 ratio.

```
#imgs_load = np.empty((n_img,75,75,3))
#for i in range(n_img):
    #img_fn = f'../data/images/{i+1:05d}.png'
    #imgs_load[i,:,:,:]=cv2.resize(cv2.imread(img_fn),(75,75),interpolation=cv2.INTER_CUBIC)

#Re-load the imgs_load.npy under imgs_load
imgs_load = np.load(data_path + 'imgs_load.npy')
```

• InceptionV3 requires its instances to be pre-processed using 'inception_v3.preprocess_input' built in function. The process in this function comprised of resizing images to 299 x 299 pixels, scalling the pixel values to the range [-1,1], and applying some normalization for the color channels RGB.

```
imgs_inception = inception_v3.preprocess_input(imgs_load)

# splitting the data into 75:25 ratio
imgs inception train, imgs inception test, noisy labels train, noisy labels test = train test split(imgs load)
```

• The next step is to load Inception V3 model with some modifications. First, it provides option to freze the pre-tained layers so that the features will follow those of ImageNet. Second, the fully-connected layers will be built based on new architecture with ReLU activation and drop out rate to prevent overfitting.

```
# loading the inception v3 model
def create_model_inception(input_shape, n_classes, optimizer, fine_tune):
   Deploy a compilation of a model integrated with inception v3 pretrained layers
   - input_shape: the shape of input images (width, height, color channels) in tuple
   - n classes: number of classes for the output layer in integer
   - optimizer: optimizer to use for training. Defaults to 'RMSProp', in this case this model uses Adam ir
   - fine_tune: The number of pre-trained layers to unfreeze in string.
              If set to 0, all pretrained layers will freeze during training & only top layers (dense/fc) v
              Freezing pre-trained layers means that the weights will not be updated and features from Imag
    .....
   # In pretrained convolutional layers, weights are loaded using the ImageNet weights.
   # Excluding the fully-connected layers to make customization for the model -- matching the classes that
   conv base = inception v3.InceptionV3(include top=False,
                           weights='imagenet',input shape=input shape)
   # Defines how many layers to freeze during training.
   # Layers in the convolutional base are switched from trainable to non-trainable
   # If the fine_tune is set to 0, all pre-trained layers will be frozen and left un-trainable, by default
   # Otherwise, the last n layers will be made available for training.
   if fine tune > 0:
       for layer in conv_base.layers[:-fine_tune]:
```

```
layer.trainable = False
   else:
       for layer in conv_base.layers:
           layer.trainable = False
   # Customize the architecture of the top_model(i.e. fully-connected layers or Dense layers)
   # The intuition is to use the 'conv base' outputs and flattening them.
   # The method uses 'bootstrapping' approach to a new top model onto the pretrained layers.
   top_model = conv_base.output
   top model = Flatten()(top model)
   top_model = Dense(n_classes*8, activation='relu')(top_model) #the more the nodes, the higher chance of t
   top_model = Dense(n_classes*4, activation='relu')(top_model) #reduce the dimensionality from previous la
   top_model = Dropout(0.2)(top_model) #20% of the input will be randomly set to 0 to prevent overfitting
   output_layer = Dense(n_classes, activation='softmax')(top_model)
   # Compiling the convolutional base and new fully-connected layers into a Model object.
   model = Model(inputs=conv_base.input, outputs=output_layer)
   # Compiles the model for training.
   model.compile(optimizer=optimizer,
                 loss='categorical crossentropy',
                 metrics=['accuracy'])
   return model
input\_shape = (75, 75,3)
n classes=10
optim = Adam(learning_rate=0.001)
inception_model = create_model_inception(input_shape, n_classes, optim, fine_tune = 0)
    Downloading data from https://storage.googleapis.com/tensorflow/keras-applications/inception v3/incept;
    87910968/87910968 [============] - 0s Ous/step
```

• In training the model, the data input is divided into training sets and text sets froom img_inception_training data. Some of the callback functions are used to prevent overfitting and endorse the model to learn more generalized patterns.

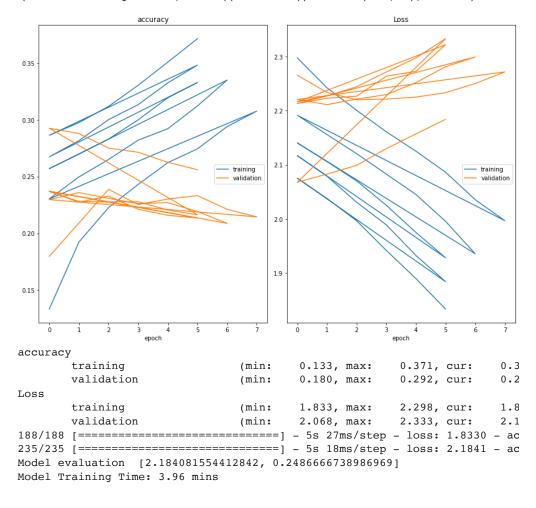
```
# Train the incep model
# ModelCheckpoint: callback function to save the best model weights based on validation loss so that the model
# EarlyStopping: another callback function to monitor validation loss during training and stop it earlier if
# PlotLossesCallBack: callback function, not too essential, but beneficial to plot the training and validati
start = time.time()
inception_checkpoint = ModelCheckpoint(filepath= data_path + 'inception.weights.best.hdf5', save_best_only=""]
early_stop = EarlyStopping(monitor='val_loss', patience=5, restore_best_weights=True, mode='min')
plot_loss = PlotLossesCallback()
# Splitting the data with K-Fold cross validation method with 5 fold
# Each fold is trained with 'inception model' for 20 epochs and batch size of 128
# tf.one_hot is to convert noisy labels into categorical format
# validation split is a hyperparameter to specify the fraction of input data to be used as validation set
n_split=5
for train index, test index in KFold(n split).split(imgs inception train):
  x_train,x_test=imgs_inception_train[train_index],imgs_inception_train[test_index]
 y_train,y_test=noisy_labels_train[train_index],noisy_labels_train[test_index]
  inception_fit = inception_model.fit(x_train,
                            tf.one_hot(y_train,10),
                            batch size=128, # Mini-batch gradient descent
                            epochs=20,# 20
```

```
validation_split=0.2,
callbacks=[inception_checkpoint, early_stop, plot_loss],
verbose=1)
```

K.clear_session()

Evaluates the trained inception model with x_text or test set that's not used within training mdoel
print('Model evaluation ',inception_model.evaluate(x_test,tf.one_hot(y_test,10)))

print('Model Training Time:', round((time.time() - start)/60, 2), 'mins')



• Deploy prediction and assess accuracy of the model using the best weights from previous model and test datasets.

```
# [BUILD A MORE SOPHISTICATED PREDICTIVE MODEL]
def model_I(image):
   Model 1 uses Inception V3 methods of CNN
    inception model.load weights(data path + 'inception.weights.best.hdf5')
    inception_preds = inception_model.predict([np.expand_dims(image, axis=0)], verbose=0); #adding batch size
    inception_pred_classes = np.argmax(inception_preds, axis=1) #return the highest probability
    return int(inception pred classes[0])
fig = plt.figure()
ax1 = fig.add_subplot(2,4,1)
ax1.imshow(imgs[0]/255)
ax2 = fig.add subplot(2,4,2)
ax2.imshow(imgs[1]/255)
ax3 = fig.add_subplot(2,4,3)
ax3.imshow(imgs[2]/255)
ax4 = fig.add_subplot(2,4,4)
ax4.imshow(imgs[3]/255)
ax1 = fig.add_subplot(2,4,5)
ax1.imshow(imgs[4]/255)
ax2 = fig.add_subplot(2,4,6)
ax2.imshow(imgs[5]/255)
ax3 = fig.add subplot(2,4,7)
ax3.imshow(imgs[6]/255)
ax4 = fig.add_subplot(2,4,8)
ax4.imshow(imgs[7]/255)
print('Clean labels:')
print(' '.join('%5s' % classes[clean_labels[j]] for j in range(8)))
print('Predicted baseline labels:')
print(' '.join('%5s' % classes[model_I(imgs_inception[j])] for j in range(8)))
    Clean labels:
     frog truck truck deer
                                     car bird horse
                               car
    Predicted baseline labels:
      cat ship plane deer
                                    deer horse horse
                               car
```

2.3. Model II

```
# [ADD WEAKLY SUPERVISED LEARNING FEATURE TO MODEL I]
# write your code here...
def model_II(image):
```

This function should takes in the image of dimension 32*32*3 as input and returns a label prediction

""

write your code here...

3. Evaluation

For assessment, we will evaluate your final model on a hidden test dataset with clean labels by the evaluation function defined as follows. Although you will not have the access to the test set, the function would be useful for the model developments. For example, you can split the small training set, using one portion for weakly supervised learning and the other for validation purpose.

```
# [DO NOT MODIFY THIS CELL]
def evaluation(model, test_labels, test_imgs):
   y true = test labels
   y_pred = []
   for image in test_imgs:
       y_pred.append(model(image))
   print(classification report(y true, y pred))
# [DO NOT MODIFY THIS CELL]
# This is the code for evaluating the prediction performance on a testset
# You will get an error if running this cell, as you do not have the testset
# Nonetheless, you can create your own validation set to run the evlauation
n test = 10000
test_labels = np.genfromtxt('../data/test_labels.csv', delimiter=',', dtype="int8")
test_imgs = np.empty((n_test, 32, 32, 3))
for i in range(n_test):
   img_fn = f'../data/test_images/test{i+1:05d}.png'
   test_imgs[i,:,:,:]=cv2.cvtColor(cv2.imread(img_fn),cv2.COLOR_BGR2RGB)
evaluation(baseline_model, test_labels, test_imgs)
!wget -nc https://raw.githubusercontent.com/brpy/colab-pdf/master/colab pdf.py
from colab pdf import colab pdf
colab pdf('Inception V3.ipynb')
```

```
--2023-03-08 21:06:10-- https://raw.githubusercontent.com/brpy/colab-pdf
Resolving raw.githubusercontent.com (raw.githubusercontent.com)... 185.19
Connecting to raw.githubusercontent.com (raw.githubusercontent.com) 185.1
HTTP request sent, awaiting response... 200 OK
Length: 1864 (1.8K) [text/plain]
Saving to: 'colab pdf.py'
colab_pdf.py
                   100%[========>]
                                              1.82K --.-KB/s
                                                                   in 0s
2023-03-08 21:06:11 (28.6 MB/s) - 'colab_pdf.py' saved [1864/1864]
WARNING: apt does not have a stable CLI interface. Use with caution in sc
WARNING: apt does not have a stable CLI interface. Use with caution in sc
E: Unable to locate package texlive-generic-recommended
[NbConvertApp] WARNING | pattern '$notebookpath$file name' matched no fil
This application is used to convert notebook files (*.ipynb)
       to various other formats.
        WARNING: THE COMMANDLINE INTERFACE MAY CHANGE IN FUTURE RELEASES.
Options
======
The options below are convenience aliases to configurable class-options,
as listed in the "Equivalent to" description-line of the aliases.
To see all configurable class-options for some <cmd>, use:
   <cmd> --help-all
```

The overall accuracy is 0.24, which is better than random guess (which should have a accuracy around 0.10). For the project, you should try to improve the performance by the following strategies:

- Consider a better choice of model architectures, hyperparameters, or training scheme for the predictive model;
- Use both clean_noisy_trainset and noisy_trainset for model training via weakly supervised learning methods.
 One possible solution is to train a "label-correction" model using the former, correct the labels in the latter, and train the final predictive model using the corrected dataset.
- Apply techniques such as k-fold cross validation to avoid overfitting;
- · Any other reasonable strategies.

```
Equivalent to: [--JupyterApp.answer_yes=True]
--execute
    Execute the notebook prior to export.
    Equivalent to: [--ExecutePreprocessor.enabled=True]
--allow-errors
    Continue notebook execution even if one of the cells throws an error
    Equivalent to: [--ExecutePreprocessor.allow_errors=True]
--stdin
    read a single notebook file from stdin. Write the resulting notebook
    Equivalent to: [--NbConvertApp.from_stdin=True]
--stdout
```

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