Image Classification

In this project, you'll classify images from the <u>CIFAR-10 dataset</u> (https://www.cs.toronto.edu/~kriz/cifar.html). The dataset consists of airplanes, dogs, cats, and other objects. You'll preprocess the images, then train a convolutional neural network on all the samples. The images need to be normalized and the labels need to be one-hot encoded. You'll get to apply what you learned and build a convolutional, max pooling, dropout, and fully connected layers. At the end, you'll get to see your neural network's predictions on the sample images.

Get the Data

Run the following cell to download the <u>CIFAR-10 dataset for python</u> (https://www.cs.toronto.edu/~kriz/cifar-10-python.tar.gz).

Data

CIFAR-10 is an established computer-vision dataset used for object recognition. It is a subset of the 80 million tiny images dataset and consists of 60,000 32x32 color images containing one of 10 object classes, with 6000 images per class. It was collected by Alex Krizhevsky, Vinod Nair, and Geoffrey Hinton.

Let's get the data by running the following function

```
In [2]: from urllib.request import urlretrieve
        from os.path import isfile, isdir
        from tqdm import tqdm
        import tarfile
        cifar10_dataset_folder_path = 'cifar-10-batches-py'
        class DLProgress(tqdm):
            last block = 0
            def hook(self, block num=1, block size=1, total size=None):
                 self.total = total size
                 self.update((block_num - self.last_block) * block_size)
                 self.last block = block num
        if not isfile('cifar-10-python.tar.gz'):
            with DLProgress(unit='B', unit scale=True, miniters=1, desc='CIFAR-10 Dataset
                 urlretrieve(
                     'https://www.cs.toronto.edu/~kriz/cifar-10-python.tar.gz',
                     'cifar-10-python.tar.gz',
                     pbar.hook)
        if not isdir(cifar10 dataset folder path):
            with tarfile.open('cifar-10-python.tar.gz') as tar:
                tar.extractall()
                tar.close()
```

Explore the Data

The dataset is broken into batches to prevent your machine from running out of memory. The CIFAR-10 dataset consists of 5 batches, named data_batch_1, data_batch_2, etc.. Each batch contains the labels and images that are one of the following:

- airplane
- · automobile
- bird
- cat
- deer
- dog
- frog
- horse
- ship
- truck

Understanding a dataset is part of making predictions on the data. Play around with the code cell below by changing the batch_id and sample_id. The batch_id is the id for a batch (1-5). The sample id is the id for a image and label pair in the batch.

Ask yourself "What are all possible labels?", "What is the range of values for the image data?", "Are the labels in order or random?". Answers to questions like these will help you preprocess the data and end up with better predictions.

The following are some helper functions students can use in their code

```
In [3]: import pickle
        import numpy as np
        import matplotlib.pyplot as plt
        from sklearn.preprocessing import LabelBinarizer
        def _load_label_names():
            Load the label names from file
            return ['airplane', 'automobile', 'bird', 'cat', 'deer', 'dog', 'frog', 'hors
        def load cfar10 batch(cifar10 dataset folder path, batch id):
            Load a batch of the dataset
            with open(cifar10_dataset_folder_path + '/data_batch_' + str(batch_id), mode=
                batch = pickle.load(file, encoding='latin1')
            features = batch['data'].reshape((len(batch['data']), 3, 32, 32)).transpose(0)
            labels = batch['labels']
            return features, labels
        def display_stats(cifar10_dataset_folder_path, batch_id, sample_id):
            Display Stats of the the dataset
            batch ids = list(range(1, 6))
            if batch id not in batch ids:
                 print('Batch Id out of Range. Possible Batch Ids: {}'.format(batch ids))
                 return None
            features, labels = load cfar10 batch(cifar10 dataset folder path, batch id)
            if not (0 <= sample id < len(features)):</pre>
                 print('{} samples in batch {}. {} is out of range.'.format(len(features)
                return None
            print('\nStats of batch {}:'.format(batch_id))
            print('Samples: {}'.format(len(features)))
            print('Label Counts: {}'.format(dict(zip(*np.unique(labels, return counts=Tru
            print('First 20 Labels: {}'.format(labels[:20]))
            sample image = features[sample id]
            sample label = labels[sample id]
            label names = load label names()
            print('\nExample of Image {}:'.format(sample_id))
            print('Image - Min Value: {} Max Value: {}'.format(sample_image.min(), sample
            print('Image - Shape: {}'.format(sample_image.shape))
            print('Label - Label Id: {} Name: {}'.format(sample_label, label_names[sample]
            plt.axis('off')
```

```
plt.imshow(sample image)
def _preprocess_and_save(normalize, one_hot_encode, features, labels, filename):
   Preprocess data and save it to file
   features = normalize(features)
   labels = one_hot_encode(labels)
   pickle.dump((features, labels), open(filename, 'wb'))
def preprocess_and_save_data(cifar10_dataset_folder_path, normalize, one_hot_enco
   Preprocess Training and Validation Data
   n batches = 5
   valid features = []
   valid labels = []
   for batch i in range(1, n batches + 1):
        features, labels = load cfar10 batch(cifar10 dataset folder path, batch i
        validation count = int(len(features) * 0.1)
       # Prprocess and save a batch of training data
        preprocess and save(
            normalize,
            one hot encode,
            features[:-validation count],
            labels[:-validation_count],
            'preprocess batch ' + str(batch i) + '.p')
       # Use a portion of training batch for validation
       valid features.extend(features[-validation count:])
        valid labels.extend(labels[-validation count:])
   # Preprocess and Save all validation data
    preprocess and save(
        normalize,
       one hot encode,
        np.array(valid features),
        np.array(valid_labels),
        'preprocess validation.p')
   with open(cifar10 dataset folder path + '/test batch', mode='rb') as file:
        batch = pickle.load(file, encoding='latin1')
   # load the training data
   test_features = batch['data'].reshape((len(batch['data']), 3, 32, 32)).transp
   test labels = batch['labels']
   # Preprocess and Save all training data
   preprocess and save(
        normalize,
        one_hot_encode,
        np.array(test features),
```

```
np.array(test labels),
        'preprocess_training.p')
def batch features labels(features, labels, batch size):
   Split features and labels into batches
   for start in range(0, len(features), batch_size):
        end = min(start + batch size, len(features))
       yield features[start:end], labels[start:end]
def load_preprocess_training_batch(batch_id, batch_size):
   Load the Preprocessed Training data and return them in batches of <batch size
   filename = 'preprocess_batch_' + str(batch_id) + '.p'
   features, labels = pickle.load(open(filename, mode='rb'))
   # Return the training data in batches of size <batch_size> or less
   return batch features labels(features, labels, batch size)
def display_image_predictions(features, labels, predictions):
   n classes = 10
   label names = load label names()
   label_binarizer = LabelBinarizer()
   label binarizer.fit(range(n classes))
   label_ids = label_binarizer.inverse_transform(np.array(labels))
   fig, axies = plt.subplots(nrows=4, ncols=2)
   fig.tight_layout()
   fig.suptitle('Softmax Predictions', fontsize=20, y=1.1)
   n predictions = 3
   margin = 0.05
   ind = np.arange(n_predictions)
   width = (1. - 2. * margin) / n predictions
   for image i, (feature, label id, pred indicies, pred values) in enumerate(zip
        pred names = [label names[pred i] for pred i in pred indicies]
        correct_name = label_names[label_id]
        axies[image i][0].imshow(feature*255)
        axies[image i][0].set title(correct name)
        axies[image_i][0].set_axis_off()
        axies[image_i][1].barh(ind + margin, pred_values[::-1], width)
        axies[image_i][1].set_yticks(ind + margin)
        axies[image i][1].set yticklabels(pred names[::-1])
        axies[image_i][1].set_xticks([0, 0.5, 1.0])
```

```
In [21]: %matplotlib inline
%config InlineBackend.figure_format = 'retina'

import numpy as np

# Explore the dataset
batch_id = 3
sample_id = 5
display_stats(cifar10_dataset_folder_path, batch_id, sample_id)
```

```
Stats of batch 3:
Samples: 10000
Label Counts: {0: 994, 1: 1042, 2: 965, 3: 997, 4: 990, 5: 1029, 6: 978, 7: 101
5, 8: 961, 9: 1029}
First 20 Labels: [8, 5, 0, 6, 9, 2, 8, 3, 6, 2, 7, 4, 6, 9, 0, 0, 7, 3, 7, 2]

Example of Image 5:
Image - Min Value: 9 Max Value: 255
Image - Shape: (32, 32, 3)
```

