Spring 2023 CS 4641\7641 A: Machine Learning Homework 4

Instructor: Dr. Mahdi Roozbahani

Deadline: Friday, April 21, 2022 11:59 pm EST

- No unapproved extension of the deadline is allowed. Submission past our 48-hour penalized acceptance period will lead to 0 credit.
- Discussion is encouraged on Ed as part of the Q/A. However, all assignments should be done individually.
- Plagiarism is a serious offense. You are responsible for completing your own work. You are not allowed to copy and paste, or paraphrase, or submit materials created or published by others, as if you created the materials. All materials submitted must be your own.
- All incidents of suspected dishonesty, plagiarism, or violations of the Georgia Tech Honor Code will be subject to the institute's
 Academic Integrity procedures. If we observe any (even small) similarities/plagiarisms detected by Gradescope or our TAs, WE
 WILL DIRECTLY REPORT ALL CASES TO OSI, which may, unfortunately, lead to a very harsh outcome. Consequences can be
 severe, e.g., academic probation or dismissal, grade penalties, a 0 grade for assignments concerned, and prohibition
 from withdrawing from the class.

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Instructions for the assignment

- This assignment consists of both programming and theory questions.
- Unless a theory question explicitly states that no work is required to be shown, you must provide an explanation, justification, or calculation for your answer.
- To switch between cell for code and for markdown, see the menu -> Cell -> Cell Type
- You can directly type Latex equations into markdown cells.
- If a question requires a picture, you could use this syntax to include them within your ipython notebook.
- Your write up must be submitted in PDF form. You may use either Latex, markdown, or any word processing software. We will **NOT** accept handwritten work. Make sure that your work is formatted correctly, for example submit $\sum_{i=0} x_i$ instead of \text{sum_{i=0} x_i}
- When submitting the non-programming part of your assignment, you must correctly map pages of your PDF to each question/subquestion to reflect where they appear. **Improperly mapped questions may not be graded correctly and/or will result in point deductions for the error.**
- All assignments should be done individually, and each student must write up and submit their own answers.
- Graduate Students: You are required to complete any sections marked as Bonus for Undergrads

Using the autograder

- You will find three assignments (for grads) on Gradescope that correspond to HW4: "Assignment 4 Programming", "Assignment 4 Non-programming" and "Assignment 4 Programming Bonus for all". Undergrads will have an additional assignment called "Assignment 4 Programming Bonus for Undergrads".
- You will submit your code for the autograder in the Assignment 4 Programming sections. Please refer to the Deliverables and Point Distribution section for what parts are considered required, bonus for undergrads, and bonus for all".
- We provided you different .py files and we added libraries in those files please DO NOT remove those lines and add your code after those lines. Note that these are the only allowed libraries that you can use for the homework
- You are allowed to make as many submissions until the deadline as you like. Additionally, note that the autograder tests each function separately, therefore it can serve as a useful tool to help you debug your code if you are not sure of what part of your implementation might have an issue
- For the "Assignment 4 Non-programming" part, you will need to submit to Gradescope a PDF copy of your Jupyter Notebook with the cells ran. See this EdStem Post for multiple ways on to convert your .ipynb into a .pdf file. Please refer to the Deliverables and Point Distribution section for an outline of the non-programming questions.
- When submitting to Gradescope, please make sure to mark the page(s) corresponding to each problem/sub-problem. The pages in the PDF should be of size 8.5" x 11", otherwise there may be a deduction in points for extra long sheets.
- You **MUST** pass the Autograder Test to gain points for the programming section. There will not be any partial credit or manual grading for this part.

Using the local tests

- For some of the programming questions we have included a local test using a small toy dataset to aid in debugging. The local test sample data and outputs are stored in localtests.py
- There are no points associated with passing or failing the local tests, you must still pass the autograder to get points.
- It is possible to fail the local test and pass the autograder since the autograder has a certain allowed error tolerance while the local test allowed error may be smaller. Likewise, passing the local tests does not guarantee passing the autograder.
- You do not need to pass both local and autograder tests to get points, passing the Gradescope autograder is sufficient for credit.
- It might be helpful to comment out the tests for functions that have not been completed yet.
- It is recommended to test the functions as it gets completed instead of completing the whole class and then testing. This may help in isolating errors. Do not solely rely on the local tests, continue to test on the autograder regularly as well.

Deliverables and Points Distribution

Q1: Two Layer NN [80 pts; 55pts + 25pts Undergrad Bonus]

Deliverables: NN.py and Notebook Graphs

- 1.1 NN Implementation [65pts; 50pts + 15pts Bonus for Undergrad] programming
 - Leaky_relu [5pts]
 - tanh [5pts]
 - loss [5pts]
 - dropout [5pts]
 - forward propagation and with and without dropout [5pts + 5pts]
 - compute gradients and update weights [2.5pts + 2.5pts]
 - backward without momentum [5pt]
 - Gradient Descent [10pts]
 - Batch Gradient Descent [10pts Bonus for Undergrad]
 - Momentum [5pts Bonus for Undergrad]
- 1.2 Loss plot and MSE for Gradient Descent [5pts] non-programming
- 1.3 Loss plot and MSE for Batch Gradient Descent [5pts Bonus for Undergrad] non-programming
- 1.4 Loss plot and MSE value for NN with Gradient Descent with Momentum [5pts Bonus for Undergrad] non-programming

Q2: CNN [25pts; 20pts Bonus for Undergrad + 5pts Bonus for All]

Deliverables: cnn.py and Written Report

- 2.1 Image Classification using Keras CNN [20pts Bonus for Undergrad]
 - 2.1.1 Loading the Model [5pts Bonus for Undergrad] programming
 - 2.1.3 Building the Model [5pts **Bonus for Undergrad**] non-programming
 - 2.1.4 Training the Model [8pts **Bonus for Undergrad**] non-programming
 - 2.1.5 Examining Accuracy and Loss [2pts **Bonus for Undergrad**] non-programming
- 2.2 Exploring Deep CNN Architectures [5pts Bonus for All] non-programming

Q3: Random Forest [45pts; 40pts + 5pts Bonus for All]

Deliverables: random_forest.py and Written Report

- 3.1 Random Forest Implementation [35pts] programming
- 3.2 Hyperparameter Tuning with a Random Forest [5pts] programming
- 3.3 Plotting Feature Importance [5pts **Bonus for All**] non-programming

Q4: SVM [30pts Bonus for all]

Deliverables: feature.py and Written Report

- 4.1: Fitting an SVM Classifier by hand [20pts] non programming
- 4.2: Feature Mapping [10pts] programming

Environment Setup

numpy: 1.23.5

```
In [ ]:
        import sys
        import matplotlib
        import numpy as np
        import matplotlib.pyplot as plt
        from sklearn.datasets import load diabetes, fetch california housing
        from sklearn.preprocessing import MinMaxScaler
        from sklearn.model_selection import train_test_split
        from sklearn.metrics import classification report
        from sklearn.metrics import mean squared error
        from collections import Counter
        from scipy import stats
        from math import log2, sqrt
        import pandas as pd
        import time
        from sklearn.model selection import train test split
        from sklearn.preprocessing import LabelEncoder
        from sklearn.tree import DecisionTreeClassifier
        from sklearn.datasets import make moons
        from sklearn.metrics import accuracy score
        from sklearn import svm
        print('Version information')
        print('python: {}'.format(sys.version))
        print('matplotlib: {}'.format(matplotlib.__version__))
        print('numpy: {}'.format(np. version ))
        %load_ext autoreload
        %autoreload 2
        %reload ext autoreload
        Version information
        python: 3.10.10 | packaged by Anaconda, Inc. | (main, Mar 21 2023, 18:39:17) [MSC v.1916 64 bit (AMD64)]
        matplotlib: 3.7.1
```

%reload_ext autoreload

The autoreload extension is already loaded. To reload it, use:

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1: Two Layer Neural Network [80 pts; 55pts + 25pts Undergrad Bonus] **[P]****[W]**

1.1 NN Implementation [65pts; 50pts + 15pts Bonus for Undergrad] **[P]**

In this section, you will implement a two layer fully connected neural network. You will also experiment with different activation functions and optimization techniques. We provide two activation functions here - Leaky Relu and Tanh. You will implement a neural network where the first hidden layer has a Leaky Relu activation and the second hidden layer has a Tanh layer.

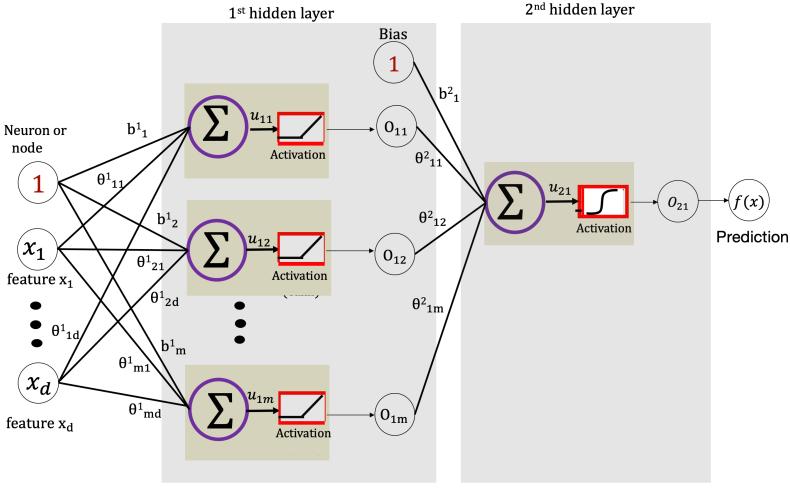
You'll also implement Gradient Descent (GD) and Batch Gradient Descent (BGD) algorithms for training these neural nets. **GD is** mandatory for all. **BGD** is bonus for undergraduate students but mandatory for graduate students.

In the **NN.py** file, complete the following functions:

- Leaky_Relu: Note Hint 1
- Tanh: Note Hint 1
- nloss
- _dropout
- forward
- compute_gradients
- update weights
- backward: Note Hint 2, if you still have issues passing the autograder make sure to address Hint 1 as well.
- gradient_descent
- batch_gradient_descent: **Mandatory for graduate students, bonus for undergraduate students.** Please batch your data in a wraparound manner. For example, given a dataset of 9 numbers, [1, 2, 3, 4, 5, 6, 7, 8, 9], and a batch size of 6, the first iteration batch will be [1, 2, 3, 4, 5, 6], the second iteration batch will be [7, 8, 9, 1, 2, 3], the third iteration batch will be [4, 5, 6, 7, 8, 9], etc...

We'll train this neural net on sklearn's California Housing dataset.

Perceptron



Notation clarification – superscript represents the layer number, subscripts represent the specific units in two adjacent layers being connected by theta

 θ^1_{21} - theta of the 1st layer connecting the 2nd hidden unit of the 1st layer and the 1st input unit θ^2_{12} - theta of the 2nd layer connecting the 1st hidden unit of the 2nd layer and the 2nd hidden unit of the 1st layer θ^1_{11} - bias of the 1st hidden unit of the 1st layer

A single layer perceptron can be thought of as a linear hyperplane as in logistic regression followed by a non-linear activation function.

$$u_i = \sum_{j=1}^d heta_{ij} x_j + b_i$$

$$o_i = \phi\left(\sum_{j=1}^d heta_{ij} x_j + b_i
ight) = \phi(heta_i^T x + b_i)$$

where x is a d-dimensional vector i.e. $x \in R^d$. It is one datapoint with d features. $\theta_i \in R^d$ is the weight vector for the i^{th} hidden unit, $b_i \in R$ is the bias element for the i^{th} hidden unit and $\phi(.)$ is a non-linear activation function that has been described below. u_i is a linear combination of the features in x_i weighted by θ_i whereas o_i is the i^{th} output unit from the activation layer.

Fully connected Layer

Typically, a modern neural network contains millions of perceptrons as the one shown in the previous image. Perceptrons interact in different configurations such as cascaded or parallel. In this part, we describe a fully connected layer configuration in a neural network which comprises multiple parallel perceptrons forming one layer.

We extend the previous notation to describe a fully connected layer. Each layer in a fully connected network has a number of input/hidden/output units cascaded in parallel. Let us a define a single layer of the neural net as follows: m denotes the number of hidden units in a single layer l whereas l denotes the number of units in the previous layer l-1.

$$u^{[l]} = \theta^{[l]} o^{[l-1]} + b^{[l]}$$

where $u^{[l]} \in R^m$ is a m-dimensional vector pertaining to the hidden units of the l^{th} layer of the neural network after applying linear operations. Similarly, $o^{[l-1]}$ is the n-dimensional output vector corresponding to the hidden units of the $(l-1)^{th}$ activation layer. $\theta^{[l]} \in R^{m \times n}$ is the weight matrix of the l^{th} layer where each row of $\theta^{[l]}$ is analogous to θ_i described in the previous section i.e. each row corresponds to one hidden unit of the l^{th} layer. $b^{[l]} \in R^m$ is the bias vector of the layer where each element of b pertains to one hidden unit of the l^{th} layer. This is followed by element wise non-linear activation function $o^{[l]} = \phi(u^{[l]})$. The whole operation can be summarized as,

$$o^{[l]} = \phi(heta^{[l]}o^{[l-1]} + b^{[l]})$$

. . . .

where $o^{\lfloor l-1 \rfloor}$ is the output of the previous layer.

Activation Function

There are many activation functions in the literature but for this question we are going to use Leaky Relu and Tanh only.

HINT 1: When calculating the tanh and leaky relu function, make sure you are not modifying the values in the original passed in matrix. You may find np.copy() helpful (`u` should not be modified in the method.).

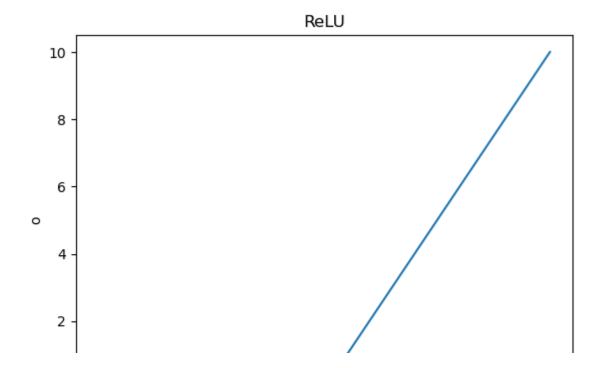
ReLU and Leaky ReLU

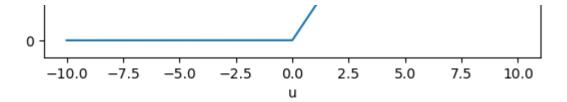
The rectified linear unit (ReLU) is one of the most commonly used activation functions in deep learning models. The mathematical form is

$$o = \phi(u) = max(0, u)$$

One of the advantages of Relu is that it is a fast nonlinear activation function.

The derivative of relu function is given as
$$o'=\phi'(u)=\left\{egin{array}{ll} 0 & u\leq 0 \\ 1 & u>0 \end{array}
ight.$$



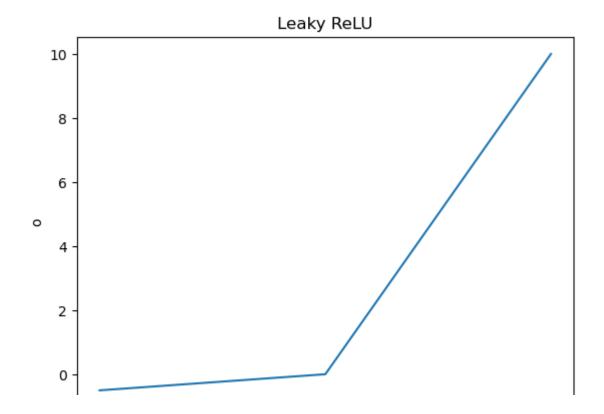


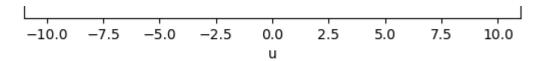
Leaky ReLU is a type of activation function based on a ReLU. The difference is that it has a small slope (such as $\alpha=0.05$) for negative values instead of a flat slope. The slope coefficient is determined before training. Leaky relu is a popular solution for sparse gradients, for example training generative adversarial networks.

It takes the form

$$o = \phi(u) = \left\{egin{array}{ll} lpha u & u \leq 0 \ u & u > 0 \end{array}
ight.$$

Here in our homework, we are going to implement Leaky ReLU.





In the **NN.py** file, complete the following functions:

• Leaky Relu: Recall Hint 1

```
In [ ]: from utilities.localtests import TestNN

TestNN('test_leaky_relu').test_leaky_relu()
TestNN('test_d_leaky_relu').test_d_leaky_relu()

test_leaky_relu passed!
test_d_leaky_relu passed!
```

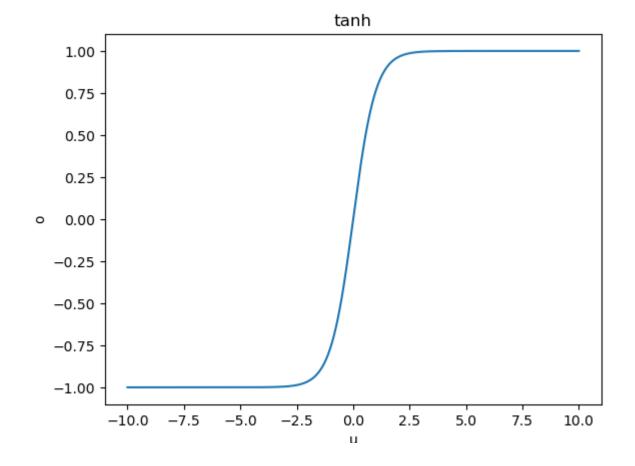
Tanh

Tanh also known as hyperbolic tangent is like a shifted version of sigmoid activation function with its range going from -1 to 1. Tanh almost always proves to be better than the sigmoid function since the mean of the activations are closer to zero. Tanh has an effect of centering data that makes learning for the next layer a bit easier. The mathematical form of tanh is given as

$$o=\phi(u)=tanh(u)=rac{e^u-e^{-u}}{e^u+e^{-u}}$$

The derivative of tanh is given as

$$o' = \phi'(u) = 1 - \left(rac{e^u - e^{-u}}{e^u + e^{-u}}
ight)^2 = 1 - o^2$$



-

In the **NN.py** file, complete the following functions:

• Tanh: Recall Hint 1

```
In [ ]: from utilities.localtests import TestNN

TestNN('test_tanh').test_tanh()
TestNN('test_d_tanh').test_d_tanh()

test_tanh passed!
test_d_tanh passed!
```

Sigmoid

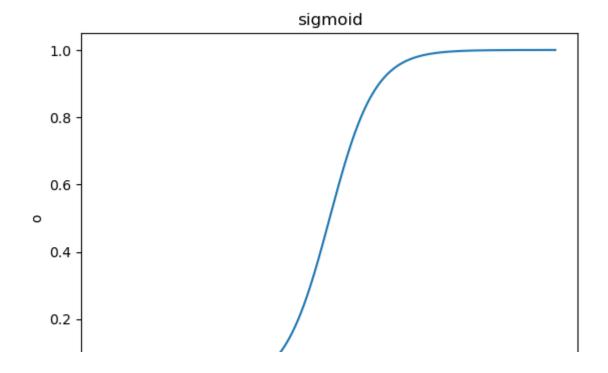
The sigmoid function is another non-linear function with S-shaped curve. This function is useful in the case of binary classification as its output is between 0 and 1. The mathematical form of the function is

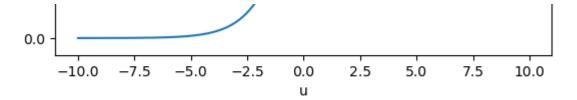
$$o=\phi(u)=rac{1}{1+e^{-u}}$$

The derivation of the sigmoid function has a nice form and is given as

$$o' = \phi'(u) = rac{1}{1 + e^{-u}} igg(1 - rac{1}{1 + e^{-u}} igg) = \phi(u) (1 - \phi(u))$$

Note: We will not be using sigmoid activation function for this assignment. This is included only for the sake of completeness.





Dropout

A dropout layer is a regularization technique used in neural networks to reduce overfitting. During training, a dropout layer looks at each input unit and randomly decide if it will drop (i.e., sets to zero) the input unit. The random decisions are made independently for each unit. Formally, given N x K input units (N is the number of data and K is the width of the layer), it samples N x K i.i.d. from Bernoulli(p). This forces the network to learn more robust and generalizable features, since it cannot rely too much on any one input unit. During inference, the dropout layer is turned off, and the full network is used to make predictions. Usually, p is passed to the neural network as a hyperparameter. Setting p=0 is equivalent to no dropout.

Note that the derivative of dropout(u) with respect to u has the same shape as u. The values of the derivative depend on the random mask.

Use this as a reference for your implementation.

Hint: Use np.random.choice() to determine which nodes to drop. Use this function to generate a mask consisting 0s (drop) and 1s (keep) to apply to the input. Apply the mask, and then scale the result by a factor of 1/(1-p). Why do we do this? The dropout mask may also be helpful for the backward method.

In the **NN.py** file, complete the following functions:

• _dropout: Use the hint above.

```
In [ ]: from utilities.localtests import TestNN
    TestNN('test_dropout').test_dropout()
    test dropout passed!
```

Mean Squared Error

Mean squared error (MSE) is an estimator that measures the average of the squares of the errors i.e. the average squared difference between the actual and the estimated values. MSE estimates the quality of the learnt hypothesis between the actual and the predicted values. Note that MSE is non-negative. If it is closer to zero, the better the learnt function is.

Implementation details

For regression problems as in this exercise, we compute the loss as follows:

$$MSE = rac{1}{2N} \sum_{i=1}^{N} \left(y_i - \hat{y}_i
ight)^2$$

where y_i is the true label and $\hat{y_i}$ is the estimated label. We use a factor of $\frac{1}{2N}$ instead of $\frac{1}{N}$ to simply the derivative of loss function.

In the **NN.py** file, complete the following functions:

nloss

```
In [ ]: from utilities.localtests import TestNN
    TestNN('test_loss').test_loss()
```

test_loss passed!

Initialization

We start by initializing the weights of the fully connected layer using Xavier initialization (At a high level, we are using a uniform distribution for weight initialization). This is already implemented for you.

Forward Propagation

During training, we pass all the data points through the network layer by layer using forward propagation. The main equations for forward prop have been described below.

$$egin{array}{lll} u^{[0]} = & x \ u^{[1]} = & heta^{[1]} u^{[0]} + b^{[1]} \ o^{[1]} = & ext{Dropout(LeakyRelu}(u^{[1]}))} \ u^{[2]} = & heta^{[2]} o^{[1]} + b^{[2]} \ \hat{y} = o^{[2]} = & ext{Tanh}(u^{[2]}) \end{array}$$

Then we get the output and compute the loss

$$l=rac{1}{2N}\sum_{i=1}^{N}\left(y_{i}-\hat{y_{i}}
ight)^{2}$$

In the **NN.py** file, complete the following functions:

• **forward**: Be sure to check the value of the *use_dropout* parameter and calculate the output accordingly.

```
In [ ]: from utilities.localtests import TestNN

    TestNN('test_forward').test_forward()
    TestNN('test_forward_without_dropout').test_forward_without_dropout()

    test_forward passed!
    test_forward_without_dropout passed!
```

Backward Propagation: Update Weights and Compute Gradients

After the forward pass, we do back propagation to update the weights and biases in the direction of the negative gradient of the loss function.

Update Weights

So, we update the weights and biases using the following formulas

$$egin{align} heta^{[2]} &:= heta^{[2]} - lr imes rac{\partial l}{\partial heta^{[2]}} \ b^{[2]} &:= b^{[2]} - lr imes rac{\partial l}{\partial b^{[2]}} \ heta^{[1]} &:= heta^{[1]} - lr imes rac{\partial l}{\partial heta^{[1]}} \ b^{[1]} &:= b^{[1]} - lr imes rac{\partial l}{\partial b^{[1]}} \end{split}$$

where lr is the learning rate. It decides the step size we want to take in the direction of the negative gradient.

In the NN.py file, complete the following functions:

• update_weights: the following local test will test when use_momentum is False

Update Weights with Momentum [Bonus for Undergrad]

Gradient descent does a generally good job of facilitating the convergence of the model's parameters to minimize the loss function. However, the process of doing so can be slow and/or noisy. **Momentum** is a technique used to stabilize this convergence.

As a reminder, vanilla gradient descent applies the following update function to the parameters:

$$\theta_{t+1} = \theta_t - \alpha \nabla f(\theta_t) \tag{1}$$

where θ_t represents the parameters at time t, α represents the learning rate, and f is the loss function.

Momentum proposes the following tweak to our parameter update function:

$$egin{aligned} z_{t+1} &= eta z_t +
abla f(heta_t) \ heta_{t+1} &= heta_t - lpha z_{t+1} \end{aligned}$$

where $\beta \in [0,1]$ is the momentum constant and z_t represents the momentum records at time t.

You can think of momentum as taking our previous changes into consideration. If we've been moving in a certain direction recently, it's likely we should keep moving in that direction. The recurrence relation given shows that we use an exponentially-weighted average of the previous updates for our current update.

A useful analogy about momentum from this great article on Distill:

Here's a popular story about momentum: gradient descent is a man walking down a hill. He follows the steepest path downwards; his progress is slow, but steady. Momentum is a heavy ball rolling down the same hill. The added inertia acts both as a smoother and an accelerator, dampening oscillations and causing us to barrel through narrow valleys, small humps and local minima.

In the **NN.py** file, complete the following functions:

• update_weights: Make to sure to update the weights using momentum

HINT: z is stored in self.change

Compute Gradients

To compute the terms $\frac{\partial l}{\partial \theta^{[i]}}$ and $\frac{\partial l}{\partial b^{[i]}}$ we use chain rule for differentiation as follows:

$$egin{array}{ll} rac{\partial l}{\partial heta^{[2]}} = & rac{\partial l}{\partial o^{[2]}} rac{\partial o^{[2]}}{\partial u^{[2]}} rac{\partial u^{[2]}}{\partial heta^{[2]}} \ rac{\partial l}{\partial b^{[2]}} = & rac{\partial l}{\partial o^{[2]}} rac{\partial o^{[2]}}{\partial u^{[2]}} rac{\partial u^{[2]}}{\partial b^{[2]}} \end{array}$$

So, $rac{\partial l}{\partial o^{[2]}}$ is the differentiation of the loss function at point $o^{[2]}$

 $rac{\partial o^{[2]}}{\partial u^{[2]}}$ is the differentiation of the Tanh function at point $u^{[2]}$

$$rac{\partial u^{[2]}}{\partial heta^{[2]}}$$
 is equal to $o^{[1]}$

$$\frac{\partial u^{[2]}}{\partial b^{[2]}}$$
 is equal to 1.

To compute $\frac{\partial l}{\partial \theta^{[2]}}$, we need $o^{[2]}, u^{[2]} \& o^{[1]}$ which are calculated during forward propagation. So we need to store these values in cache variables during forward propagation to be able to access them during backward propagation. Similarly for calculating other partial derivatives, we store the values we'll be needing for chain rule in cache. These values are obtained from the forward propagation and used in backward propagation. The cache is implemented as a dictionary here where the keys are the variable names and the values are the variables values.

Also, the functional form of the MSE differentiation and Leaky Relu differentiation are given by

$$egin{array}{ll} rac{\partial l}{\partial o^{[2]}} = & \left(o^{[2]} - y
ight) \ rac{\partial l}{\partial u^{[2]}} = & rac{\partial l}{\partial o^{[2]}} * (1 - (tanh(u^{[2]}))^2) \ rac{\partial u^{[2]}}{\partial heta^{[2]}} = & o^{[1]} \ rac{\partial u^{[2]}}{\partial u^{[2]}} = & 1 \end{array}$$

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 $\partial b^{[2]}$

On vectorization, the above equations become:

$$egin{array}{ll} rac{\partial l}{\partial heta^{[2]}} = & rac{1}{n} rac{\partial l}{\partial u^{[2]}} o^{[1]} \ rac{\partial l}{\partial b^{[2]}} = & rac{1}{n} \sum rac{\partial l}{\partial u^{[2]}} \end{array}$$

!!!!! IMPORTANT !!!!! HINT 2: Division by N only needs to occur ONCE for any derivative that requires a division by N. Make sure you avoid cascading divisions by N where you might accidentally divide your derivative by N^2 or greater.

This completes the differentiation of loss function w.r.t to parameters in the second layer. We now move on to the first layer, the equations for which are given as follows:

$$\frac{\partial l}{\partial \theta^{[1]}} = \qquad \frac{\partial l}{\partial o^{[2]}} \frac{\partial o^{[2]}}{\partial u^{[2]}} \frac{\partial u^{[2]}}{\partial o^{[1]}} \frac{\partial o^{[1]}}{\partial u^{[1]}} \frac{\partial u^{[1]}}{\partial \theta^{[1]}}$$

$$\frac{\partial l}{\partial b^{[1]}} = \qquad \frac{\partial l}{\partial o^{[2]}} \frac{\partial o^{[2]}}{\partial u^{[2]}} \frac{\partial u^{[2]}}{\partial o^{[1]}} \frac{\partial o^{[1]}}{\partial u^{[1]}} \frac{\partial u^{[1]}}{\partial b^{[1]}}$$

Where

$$\begin{split} \frac{\partial u^{[2]}}{\partial o^{[1]}} &= \theta^{[2]} \\ \frac{\partial o^{[1]}}{\partial u^{[1]}} &= \begin{cases} \text{DropoutMask} \cdot \text{ScalingFactor} \cdot [1*(o^{[1]} > 0) \text{ and } \alpha*(o^{[1]} <= 0)] & \text{dropout=true} \\ 1*(o^{[1]} > 0) \text{ and } \alpha*(o^{[1]} <= 0) & \text{dropout=false} \end{cases} \\ \frac{\partial u^{[1]}}{\partial \theta^{[1]}} &= x \\ \frac{\partial u^{[1]}}{\partial b^{[1]}} &= 1 \end{split}$$

Note that $\frac{\partial o^{[1]}}{\partial u^{[1]}}$ is the derivative of the compostion of leaky ReLU and dropout at $u^{[1]}$. The dropout mask you stored at forward may be helpful.

The above equations outline the forward and backward propagation process for a 2-layer fully connected neural net with leaky Relu

as the first activation layer and lann has the second one. The same process can be extended to different neural networks with different activation layers.

Code Implementation:

$$dLoss_o2 = \dfrac{\partial l}{\partial o^{[2]}} \Longrightarrow dim = (1,375)$$
 $dLoss_u2 = dLoss_o2 \dfrac{\partial o^{[2]}}{\partial u^{[2]}} \Longrightarrow dim = (1,375)$
 $dLoss_theta2 = dLoss_u2 \dfrac{\partial u^{[2]}}{\partial \theta^{[2]}} \Longrightarrow dim = (1,15)$
 $dLoss_b2 = dLoss_u2 \dfrac{\partial u^{[2]}}{\partial b^{[2]}} \Longrightarrow dim = (1,1)$
 $dLoss_o1 = dLoss_u2 \dfrac{\partial u^{[2]}}{\partial o^{[1]}} \Longrightarrow dim = (15,375)$
 $dLoss_u1 = dLoss_o1 \dfrac{\partial o^{[1]}}{\partial u^{[1]}} \Longrightarrow dim = (15,375)$
 $dLoss_theta1 = dLoss_u1 \dfrac{\partial u^{[1]}}{\partial \theta^{[1]}} \Longrightarrow dim = (15,8)$
 $dLoss_b1 = dLoss_u1 \dfrac{\partial u^{[1]}}{\partial \theta^{[1]}} \Longrightarrow dim = (15,1)$

One common confusion is when to use matrix multiplication versus element-wise multiplication. A good rule of thumb here is that whenever you use matrix multiplication in forward propagation, you'll likely have to use it for the backward step as well. The easiest way to make sure you're on the right track is to check the shapes of the arrays you're working with.

Note: Training set has 375 examples.

In the **NN.py** file, complete the following functions:

• compute_gradients: Be sure to account for the different values of use_dropout and calculate accordingly.

```
In [ ]: from utilities.localtests import TestNN

TestNN('test_compute_gradients_without_dropout').test_compute_gradients_without_dropout()
TestNN('test_compute_gradients').test_compute_gradients()
```

```
test_compute_gradients_withou_dropout passed!
test_compute_gradients passed!
```

1.1.1 Local Test: Helper Functions and Compute Gradient

You may test your implementation of the hepler functions and compute_gradient contained in **NN.py** in the cell below. See Using the Local Tests for more details.

```
In [ ]:
         #####################################
         ### DO NOT CHANGE THIS CELL ###
         ####################################
         from utilities.localtests import TestNN
         TestNN('test leaky relu').test leaky relu()
         TestNN('test d leaky relu').test d leaky relu()
         TestNN('test_tanh').test_tanh()
         TestNN('test_d_tanh').test_d_tanh()
         TestNN('test dropout').test dropout()
        TestNN('test_loss').test_loss()
        TestNN('test_forward').test_forward()
         TestNN('test_forward_without_dropout').test_forward_without_dropout()
        TestNN('test_update_weights').test_update_weights()
        TestNN('test update weights with momentum').test update weights with momentum()
         TestNN('test_compute_gradients_without_dropout').test_compute_gradients_without_dropout()
         TestNN('test compute gradients').test compute gradients()
         test leaky relu passed!
         test d leaky relu passed!
         test tanh passed!
        test d tanh passed!
         test_dropout passed!
         test loss passed!
         test forward passed!
         test_forward_without_dropout passed!
         test update weights passed!
         test update weights with momentum passed!
         test compute gradients withou dropout passed!
         test compute gradients passed!
```

1.1.2 Local Test: Gradient Descent

You may test your implementation of the GD function contained in **NN.py** in the cell below. See Using the Local Tests for more details.

1.1.3 Local Test: Batch Gradient Descent [No Points]

You may test your implementation of the BGD function contained in **NN.py** in the cell below. See Using the Local Tests for more details.

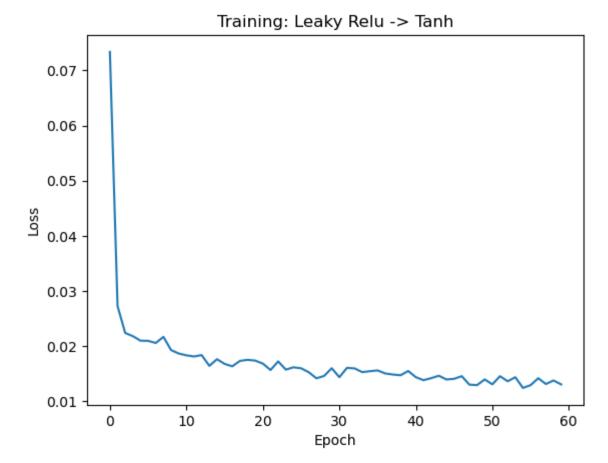
1.1.4 Local Test: Gradient Descent with Momentum

You may test your implementation of the GD function with momentum contained in **NN.py** in the cell below. See Using the Local Tests for more details.

1.2 Loss plot and MSE value for NN with Gradient Descent [5pts] **[W]**

Train your neural net implementation with gradient descent and print out the loss at every 1000th iteration (starting at iteration 0). The following cells will plot the loss vs epoch graph and calculate the final test MSE.

```
### DO NOT CHANGE THIS CELL ###
from NN import dlnet
dataset = fetch_california_housing() # Load the dataset
x, y = dataset.data, dataset.target
y = y.reshape(-1,1)
perm = np.random.RandomState(seed=3).permutation(x.shape[0])[:500]
y = y[perm]
x_train, x_test, y_train, y_test = train_test_split(x, y, random_state=1) #split data
x scale = MinMaxScaler()
x_train = x_scale.fit_transform(x_train) #normalize data/
x test = x scale.transform(x test)
y_scale = MinMaxScaler()
y_train = y_scale.fit_transform(y_train)
y test = y scale.transform(y test)
x_train, x_test, y_train, y_test = x_train.T, x_test.T, y_train.reshape(1,-1), y_test #condition data
nn = dlnet(y_train,lr=0.01,use_dropout=True, use_momentum=False) # initalize neural net class
nn.gradient descent(x train, y train, iter = 60000) #train
# create figure
fig = plt.plot(np.array(nn.loss).squeeze())
plt.title(f'Training: {nn.neural net type}')
plt.xlabel("Epoch")
plt.ylabel("Loss")
plt.show()
```



Mean Squared Error (MSE) 0.018511096794416558

1.3 Loss plot and MSE value for NN with BGD [5pts Bonus for Undergrad] **[W]**

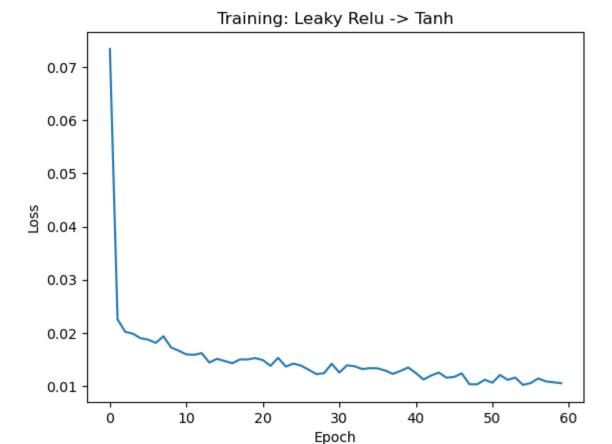
Train your neural net implementation with batch gradient descent and print out the loss at every 1000th iteration (starting at iteration 0). The following cells will plot the loss vs epoch graph and calculate the final test MSE.

```
### DO NOT CHANGE THIS CELL ###
from NN import dlnet
dataset = fetch_california_housing() # Load the dataset
x, y = dataset.data, dataset.target
y = y.reshape(-1,1)
perm = np.random.RandomState(seed=3).permutation(x.shape[0])[:500]
y = y[perm]
x train, x test, y train, y test = train test split(x, y, random state=1) #split data
x scale = MinMaxScaler()
x train = x scale.fit transform(x train) #normalize data
x_test = x_scale.transform(x_test)
y_scale = MinMaxScaler()
y train = y scale.fit transform(y train)
y_test = y_scale.transform(y_test)
x_train, x_test, y_train, y_test = x_train.T, x_test.T, y_train.reshape(1,-1), y_test #condition data
nn = dlnet(y_train,lr=0.01,use_dropout=True, use_momentum=False) # initalize neural net class
nn.batch_gradient_descent(x_train, y_train, iter = 60000, use_momentum=False) #train
# create figure
fig = plt.plot(np.array(nn.loss).squeeze())
plt.title(f'Training: {nn.neural_net_type}')
plt.xlabel("Epoch")
plt.ylabel("Loss")
plt.show()
```

1.4 Loss plot and MSE value for NN with Gradient Descent with Momentum [5pts Bonus for Undergrad] **[W]**

Train your neural net implementation with gradient descent with momentum and print out the loss at every 1000th iteration (starting at iteration 0). The following cells will plot the loss vs epoch graph and calculate the final test MSE.

```
### DO NOT CHANGE THIS CELL ###
from NN import dlnet
dataset = fetch_california_housing() # Load the dataset
x, y = dataset.data, dataset.target
y = y.reshape(-1,1)
perm = np.random.RandomState(seed=3).permutation(x.shape[0])[:500]
y = y[perm]
x_train, x_test, y_train, y_test = train_test_split(x, y, random_state=1) #split data
x scale = MinMaxScaler()
x_train = x_scale.fit_transform(x_train) #normalize data
x test = x scale.transform(x test)
y_scale = MinMaxScaler()
y_train = y_scale.fit_transform(y_train)
y test = y scale.transform(y test)
x_train, x_test, y_train, y_test = x_train.T, x_test.T, y_train.reshape(1,-1), y_test #condition data
nn = dlnet(y_train,lr=0.01,use_dropout=True, use_momentum=True) # initalize neural net class
nn.gradient descent(x train, y train, iter = 60000, use momentum=True) #train
# create figure
fig = plt.plot(np.array(nn.loss).squeeze())
plt.title(f'Training: {nn.neural net type}')
plt.xlabel("Epoch")
plt.ylabel("Loss")
plt.show()
```



Mean Squared Error (MSE) 0.017073819526876123

2: Image Classification based on Convolutional Neural Networks [25pts; 20pts Bonus for Undergrad + 5pts Bonus for all] **[P]****[W]**

2.1 Image Classification using Keras and CNN

- TransorFlow is a popular platform for machine learning.
- Keras is a deep learning API written in Python that runs on top of the machine learning platform TensorFlow.

TensorFlow and Keras

Tensorflow is a machine learning package developed by Google. In 2019, Google integrated Keras into Tensorflow. It creates a simple, layer-centric interface to Tensorflow.

Helpful Links

- Install Keras
- CS231n Tutorial(Layers used to build ConvNets)

Setup TensorFlow

You can see here to install TensorFlow. Make sure you have upgraded to the latest pip to install the TensorFlow 2 package. Alternatively, you can train your model by uploading HW4 notebook to Colab directly. Colab contains all packages you need for this section.

Please also see TensorFlow 2 quickstart for beginners to see how to load a data set, build a tf.keras.Sequential model, and train and evaluate the model.

For Mac M1 users, please see this EdStem Post

Environment Setup

```
In []: from __future__ import print_function
    import tensorflow as tf
    from keras.datasets import cifar10
    from sklearn.model_selection import train_test_split
    from tensorflow.keras.models import Sequential
    from tensorflow.keras.layers import Conv2D, MaxPooling2D, Flatten, Dense, Activation, Dropout
    from tensorflow.keras.layers import LeakyReLU
    from tensorflow.keras.layers import BatchNormalization
    from sklearn.utils import shuffle
    import numpy as np
    import matplotlib.pyplot as plt
```

2.1.1 Load CIFAR-10 Dataset and Data Augmentation [5pts - Bonus for Undergrad]**[P]**

HW4 Spring23 Student

We use CIFAR-10 dataset to train our model. This is a dataset of 60,000 32x32 colour images in 10 classes, with 6,000 images per class. There are 50,000 training images and 10,000 test images. We provide code for you to download CIFAR-10 dataset below.

Data Augmentation [5pts]

Data augmentation is a technique to increase the diversity of your training set by applying random (but realistic) transformations, such as image rotation. If the dataset in a machine learning model is rich and sufficient, the model performs better and more accurately.

In the **image_preprocessing.py** file, complete the following functions to understand the common practices used for preprocessing the image data:

• data_preprocessing

• data_augmentation

 HORIZONTAL AND VERTICAL FLIP: We flip the images horizontally or vertically based on the mode attribute which can be horizontal, vertical or both.

■ RANDOM ROTATION: We apply random rotations to each image, filling empty space according to fill_mode. The fill mode can be nearest (using nearest pixels to determine the fill value), reflect, constant or wrap.

■ RANDOM ZOOM: We apply zooming on images by a factor which represents lower and upper bound for zooming. It could be applied along either height or width or both, which positive factor values mean zooming out, and negetive values mean zooming in.

Please note that the Gradescope only checks if expected preprocessing layers are existent.

You can go through the Tensorflow documentation to see the different image processing techniques we can use for our computer vision projects.

References

tf.data Module

tf.data.Dataset

tf.data: Build TensorFlow input pipelines

Andrew Control of the property

Better performance with the tf.data API

Random Rotation

Random Flip

Random Zoom

```
#####################################
### DO NOT CHANGE THIS CELL ###
####################################
from image_preprocessing import data_augmentation,data_preprocessing
# split data between train and test sets
(x train, y train),(x test, y test)=cifar10.load data()
# input image dimensions
img rows, img cols = 32, 32
number channels = 3
#set num of classes
num classes = 10
#create augmented data
augmented x=np.zeros((10000,32,32,3))
augmented_y=np.zeros((10000,1),dtype=y_train.dtype)
y=y_train.squeeze()
i=0
for c in range(10):
    augmented_x[i:i+1000,:,:,:]=x_train[y==c][:1000]
    augmented_y[i:i+1000,:]=c
    i=i+1000
# Create a tf.data pipeline of train images (and their labels)
train_dataset = tf.data.Dataset.from_tensor_slices((x_train, y_train))
train dataset = train dataset.map(lambda x, y: (data preprocessing()(x), y))
aug dataset=tf.data.Dataset.from tensor slices((augmented x, augmented y))
aug_dataset=aug_dataset.shuffle(5000)
aug_dataset = aug_dataset.map(lambda x, y: (data_preprocessing()(x), y))
aug dataset = aug dataset.map(lambda x, y: (data augmentation()(x), y))
train_dataset=train_dataset.concatenate(aug_dataset)
# Create a tf.data pipeline of test images (and their labels)
test dataset = tf data Dataset from tensor slices((v test v test))
```

```
test dataset = test dataset.map(lambda x, y: (data preprocessing()(x), y))
        x_train, y_train = zip(*train_dataset)
        x train = np.array(x train)
        y train = np.array(y train)
        x_test, y_test = zip(*test_dataset)
        x_test = np.array(x_test)
        y test = np.array(y test)
        if tf.keras.backend.image_data_format() == 'channels_first':
           x_train = x_train.reshape(x_train.shape[0], number_channels, img_rows, img_cols)
           x_test = x_test.reshape(x_test.shape[0], number_channels, img_rows, img_cols)
           input_shape = (number_channels, img_rows, img_cols)
        else:
           x_train = x_train.reshape(x_train.shape[0], img_rows, img_cols, number_channels)
           x_test = x_test.reshape(x_test.shape[0], img_rows, img_cols, number_channels)
           input_shape = (img_rows, img_cols, number_channels)
        print('x_train shape:', x_train.shape)
        print('x_test shape:', x_test.shape)
        print(x_train.shape[0], 'train samples')
### DO NOT CHANGE THIS CELL ###
        #####################################
        # Show some images from CIFAR-10
        fig = plt.figure(figsize=(10, 10))
        for i in range(9):
           random_index = np.random.randint(0, len(y_train))
           ax = fig.add_subplot(3, 3, i+1)
           ax.imshow(x train[random index, :])
        plt.show()
```

As you can see from above, the CIFAR-10 dataset contains different types of objects. The images have been size-normalized and objects remain centered in fixed-size images.

2.1.3 Build convolutional neural network model [5pts] **[W]**

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In this part, you need to build a convolutional neural network as described below. The architecture of the model is outlined.

In the **cnn.py** file, complete the following functions:

- _init_: See Defining Variables section
- **create_net**: See Defining Model section
- compile_net: See Compiling Model section

[INPUT - CONV - CONV - MAXPOOL - DROPOUT - CONV - CONV - MAXPOOL - DROPOUT - FC1 - DROPOUT - FC2 - DROPOUT - FC3]

INPUT: $[32 \times 32 \times 3]$ will hold the raw pixel values of the image, in this case, an image of width 32, height 32. This layer should give 8 filters and have appropriate padding to maintain shape.

CONV: Conv. layer will compute the output of neurons that are connected to local regions in the input, each computing a dot product between their weights and a small region they are connected to the input volume. We decide to set the kernel_size 3×3 for both Conv. layers. For example, the output of the Conv. layer may look like $[32 \times 32 \times 32]$ if we use 32 filters. Again, we use padding to maintain shape.

MAXPOOL: MAXPOOL layer will perform a downsampling operation along the spatial dimensions (width, height). With pool size of 2×2 , resulting shape takes form 16×16 .

DROPOUT: DROPOUT layer with the dropout rate of 0.30 to prevent overfitting.

CONV: Additional Conv. layer takes outputs from above layers and applies more filters. The Conv. layer may look like $[16 \times 16 \times 32]$. We set the kernel_size 3×3 and use padding to maintain shape for both Conv. layers.

CONV: Additional Conv. layer takes outputs from above layers and applies more filters. The Conv. layer may look like $[16 \times 16 \times 64]$.

MAXPOOL: MAXPOOL layer will perform a downsampling operation along the spatial dimensions (width, height).

DROPOUT: Dropout layer with the dropout rate of 0.30 to prevent overfitting.

FC1: Dense layer which takes output from above layers, and has 256 neurons. Flatten() operations may be useful.

DROPOLIT: Dropout laver with the dropout rate of 0.5 to prevent overfitting

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FC2: Dense layer which takes output from above layers, and has 128 neurons.

DROPOUT: Dropout layer with the dropout rate of 0.5 to prevent overfitting.

FC3: Dense layer with 10 neurons, and Softmax activation, is the final layer. The dimension of the output space is the number of classes.

Activation function: Use LeakyReLU with negative_slope 0.1 as the activation function for Conv. layers and Dense layers unless otherwise indicated to build you model architecture

Please specify the loss, the optimizer, and metrics. Note that while this is a suggested model design, you may use other architectures and experiment with different layers for better results.

Use the following keras links to reference crucial layers of the model in keras API:

- Conv2d
- Dense
- Flatten
- MaxPool
- Dropout
- LeakyReLU

Lastly, if you would like to experiment with additional layers, explore the keras layers API.

Defining Variables [No Points]

You now need to set training variables in the __init__() function in **cnn.py**. Once you have defined variables you may use the cell below to see them.

- Recommended Batch Sizes fall in the range 32-512 (use powers of 2)
- Recommended Epoch Counts fall in the range 5-20
- Recommended Learning Rates fall in the range .0001-.01

Defining model [5pts Bonus for Undergrad]**[W]**

You now need to complete the <code>create_net()</code> function in **cnn.py** to define your model structure. Once you have defined a model structure you may use the cell below to examine your architecture.

Your model is required to have at least 2 convolutional layers and at least 2 dense layers. Ensuring that these requirements are met will earn you 5pts.

Compiling model [No Points]

Next prepare the model for training by completing <code>compile_model()</code> in **cnn.py**. Remember we are performing 10-way clasification when selecting a loss function. Loss function can be categorical crossentropy.

Note that the compile method configures the model for training. You can get more insights into this method here.

2.1.4 Train the network [8pts total (3pts, 3pts, 2pts) Bonus for Undergrad] **[W]**

Tuning: Training the network is the next thing to try. You can set your parameter at the **Defining Variable** section. If your parameters are set properly, you should see the loss of the validation set decreased and the value of accuracy increased. **It may take more than 15 minutes to train your model.**

Expected Result: You should be able to achieve more than 79% accuracy on the test set to get full points. If you achieve accuracy between 60% to 69%, you will only get 3 points. An accuracy between 69% to 79% will earn an additional 3pts.

Note: If you would like to automate the tuning process, you can use a nested for loop to search for the hyperparameter that achieves the accuracy.

- 60% to 69% earns 3pts
- \bullet 69% to 79% earns 3pts more (6pts total)
- 79%+ earns 2pts more (8pts total)

Train your own CNN model

```
###################################
### DO NOT CHANGE THIS CELL ###
######################################
from cnn import CNN
net = CNN()
batch_size, epochs, init_lr = net.get_vars()
def lr scheduler(epoch):
    new_lr = init_lr * 0.9 ** epoch
    print("Learning rate:", new_lr)
    return new lr
history = model.fit(
    x train, y train,
    batch_size=batch_size,
    epochs=epochs,
    callbacks=[tf.keras.callbacks.LearningRateScheduler(lr_scheduler)],
    shuffle=True,
    verbose=1,
    initial_epoch=0,
    validation_data=(x_test, y_test)
score = model.evaluate(x_test, y_test, verbose=0)
print('Test loss:', score[0])
print('Test accuracy:', score[1])
```

2.1.5 Examine accuracy and loss [2pts Bonus for Undergrad] **[W]**

You should expect to see gradually decreasing loss and gradually increasing accuracy. Examine loss and accuracy by running the cell below, no editing is necessary. Having appropriate looking loss and accuracy plots will earn you the last 2pts for your convolutional neural net.

```
In [ ]:
        ### DO NOT CHANGE THIS CELL ###
        # list all data in history
        print(history.history.keys())
        # summarize history for accuracy and loss
        plt.plot(history.history['accuracy'])
        plt.plot(history.history['val accuracy'])
        plt.title('model accuracy')
        plt.ylabel('accuracy')
        plt.xlabel('epoch')
        plt.legend(['train', 'test'], loc='upper left')
        plt.show()
        plt.plot(history.history['loss'])
        plt.plot(history.history['val loss'])
        plt.title('model loss')
        plt.ylabel('loss')
        plt.xlabel('epoch')
        plt.legend(['train', 'test'], loc='upper left')
        plt.show()
        #####################################
        ### DO NOT CHANGE THIS CELL ###
        # make predictions
        y_pred = model.predict(x_test)
        y pred classes = np.argmax(y pred, axis=1)
        y pred prob = np.max(y pred, axis=1)
        y_gt_classes = np.argmax(y_test, axis=1)
        from sklearn.metrics import confusion matrix, accuracy score
        plt.figure(figsize=(8, 7))
        plt.imshow(confusion_matrix(y_gt_classes, y_pred_classes))
        plt.title('Confusion matrix', fontsize=16)
        plt.xticks(np.arange(10), class names, rotation=90, fontsize=12)
        plt.yticks(np.arange(10), class names, fontsize=12)
        plt.colorbar()
        plt.show()
```

2.2 Exploring Deep CNN Architectures [5pts Bonus for All] **[W]**

The network you have produced is rather simple relative to many of those used in industry and research. Researchers have worked to make CNN models deeper and deeper over the past years in an effort to gain higher accuracy in predictions. While your model is only a handful of layers deep, some state of the art deep architectures may include up to 150 layers. However, this process has not been without challenges.

One such problem is the problem of the vanishing gradient. The weights of a neural network are updated using the backpropagation algorithm. The backpropagation algorithm makes a small change to each weight in such a way that the loss of the model decreases. Using the chain rule, we can find this gradient for each weight. But, as this gradient keeps flowing backwards to the initial layers, this value keeps getting multiplied by each local gradient. Hence, the gradient becomes smaller and smaller, making the updates to the initial layers very small, increasing the training time considerably.

Many tactics have been used in an effort to solve this problem. One architecture, named ResNet, solves the vanishing gradient problem in a unique way. ResNet was developed at Microsoft Research to find better ways to train deep networks. Take a moment to explore how ResNet tackles the vanishing gradient problem by reading the original research paper here: https://arxiv.org/pdf/1512.03385.pdf (also included as PDF in papers directory).

Question: In your own words, explain how ResNet addresses the vanishing gradient problem in 1-2 sentences below: (Please type answers directly in the cell below.)

Answer:

•••

3: Random Forests [45pts; 40pts + 5pts Bonus for All] **[P]** **[W]**

NOTE: Please use sklearn's ExtraTreeClassifier in your Random Forest implementation. You can find more details about this classifier here.

For context, the general difference between an extra tree and decision tree classifier is that the decision tree optimizes which feature to reduce entropy on and at what value to split, while an extra tree randomly splits on the features given.

3.1 Random Forest Implementation [35pts] **[P]**

The decision boundaries drawn by decision or extra trees are very sharp, and fitting a tree of unbounded depth to a list of examples almost inevitably leads to **overfitting**. In an attempt to decrease the variance of an extra tree, we're going to use a technique called 'Bootstrap Aggregating' (often abbreviated 'bagging'). This stems from the idea that a collection of weak learners can learn decision boundaries as well as a strong learner. This is commonly called a Random Forest.

We can build a Random Forest as a collection of extra trees, as follows:

- 1. For every tree in the random forest, we're going to
 - a) Subsample the examples with replacement. Note that in this question, the size of the subsample data is equal to the original dataset.
 - b) From the subsamples in part a, choose attributes at random without replacement to learn on in accordance with a provided attribute subsampling rate. Based on what it was mentioned in the class, we randomly pick features in each split. We use a more general approach here to make the programming part easier. Let's randomly pick some features (65% percent of features) and grow the tree based on the pre-determined randomly selected features. Therefore, there is no need to find random features in each split.
 - c) Fit an extra tree to the subsample of data we've chosen to a certain depth.

Classification for a random forest is then done by taking a majority vote of the classifications yielded by each tree in the forest after it classifies an example.

In the random_forest.py file, complete the following functions:

- _bootstrapping: this function will be used in bootstrapping()
- **fit**: Fit the extra trees initialized in __init__ with the datasets created in bootstrapping() . You will need to call bootstrapping() .

NOTES:

1. In the Random Forest Class, X is assumed to be a matrix with num_training rows and num_features columns where num_training is the number of total records and num_features is the number of features of each record. y is assumed to be a

vector of labels of length num_training.

- 2. Look out for TODO's for the parts that need to be implemented
- 3. If you receive any SettingWithCopyWarning warnings from the Pandas library, you can safely ignore them.
- 4. Hint: when bootstrapping, set replace = False while creating col_idx

3.2 Hyperparameter Tuning with a Random Forest [5pts] **[P]**

In machine learning, hyperparameters are parameters that are set before the learning process begins. The max_depth, num_estimators, or max_features variables from 3.1 are examples of different hyperparameters for a random forest model. Let's first review the dataset in a bit more detail.

Dataset Objective

Imagine that we are a team of researchers in the US Census Bureau working to track and document various information pertaining to income for a machine learning model that predicts whether or not a person makes over 50K a year. We know that there are multiple things to track such as hours worked per week in order to determine how much a person makes in the year. We will use the information we track and document in order to publish it for the general public, which will be helpful for people interested in diversifying their incomes and making more money.

After much deliberation amongst the team of researchers, you come to a conclusion that we can use the past observations of how much people have earned in the past to create a model.

We will use our random forest algorithm from Q3.1 to predict if person earns above 50K or not a year.

You can find more information on the dataset here.

Loading the dataset

The dataset that the company has collected has the following features:

There were 13 features used in this dataset.

Inputs:

- 1. age: continuous.
- 2. workclass: Private, Self-emp-not-inc, Self-emp-inc, Federal-gov, Local-gov, State-gov, Without-pay, Never-worked.
- 3. fnlwgt: continuous.
- 4. education: Bachelors, Some-college, 11th, HS-grad, Prof-school, Assoc-acdm, Assoc-voc, 9th, 7th-8th, 12th, Masters, 1st-4th, 10th, Doctorate, 5th-6th, Preschool.
- 5. education-num: continuous.
- 6. marital-status: Married-civ-spouse, Divorced, Never-married, Separated, Widowed, Married-spouse-absent, Married-AF-spouse.
- 7. occupation: Tech-support, Craft-repair, Other-service, Sales, Exec-managerial, Prof-specialty, Handlers-cleaners, Machine-opinspect, Adm-clerical, Farming-fishing, Transport-moving, Priv-house-serv, Protective-serv, Armed-Forces.
- 8. relationship: Wife, Own-child, Husband, Not-in-family, Other-relative, Unmarried.
- 9. sex at birth: Female, Male.
- 10. capital-gain: continuous.
- 11. capital-loss: continuous.
- 12. hours-per-week: continuous.
- 13. native-country: United-States, Cambodia, England, Puerto-Rico, Canada, Germany, Outlying-US(Guam-USVI-etc), India, Japan, Greece, South, China, Cuba, Iran, Honduras, Philippines, Italy, Poland, Jamaica, Vietnam, Mexico, Portugal, Ireland, France, Dominican-Republic, Laos, Ecuador, Taiwan, Haiti, Columbia, Hungary, Guatemala, Nicaragua, Scotland, Thailand, Yugoslavia, El-Salvador, Trinadad&Tobago, Peru, Hong, Holand-Netherlands.

Output:

- 1. (high income) target value:
 - h (aka > 50K) means the someone's income is high
 - I (aka <=50K) means the someone's income is low

Your random forest model will try to predict this variable.

```
### DO NOT CHANGE THIS CELL ###
        from sklearn import preprocessing
        import pandas as pd
        import numpy as np
        preprocessor = preprocessing.OrdinalEncoder()
        data train = pd.read csv("./data/income train clean.csv")
        data test = pd.read csv("./data/income test clean.csv")
       numerical_features = ["age", "fnlwgt", "education-num", "capital-gain", "capital-loss", "hours-per-week"]
        data_train_numerical = data_train[numerical_features]
        data test numerical = data test[numerical features]
        data train categorical = data train.drop(numerical features, axis=1)
        data_test_categorical = data_test.drop(numerical_features, axis=1)
        preprocessor.fit(data train categorical)
        data train categorical = pd.DataFrame(preprocessor.transform(data train categorical), columns=data train categorical
       data_test_categorical = pd.DataFrame(preprocessor.transform(data_test_categorical), columns=data_test_categorical.col
        data train = pd.concat([data train categorical, data train numerical], axis=1)
        data test = pd.concat([data test categorical, data test numerical], axis=1)
        X train = data train.drop(columns = 'income')
        y_train = data_train['income']
        y train = y train.to numpy()
        X test = data test.drop(columns = 'income')
        X_test = np.array(X_test)
       y_test = data_test['income']
       y test = y test.to numpy()
        X_train, y_train, X_test, y_test = np.array(X_train), np.array(y_train), np.array(X_test), np.array(y_test)
       In [ ]:
        ### DO NOT CHANGE THIS CELL ###
        print(X_train.shape, y_train.shape, X_test.shape, y_test.shape)
        assert X train.shape == (30162, 13)
        assert y_train.shape == (30162,)
        assert X test.shape == (12966, 13)
        assert y test.shape == (12966,)
        (30162, 13) (30162,) (12966, 13) (12966,)
```

In the following codeblock, train your random forest model with different values for max_depth, n_estimators, or max_features and evaluate each model on the held-out test set. Try to choose a combination of hyperparameters that maximizes your prediction accuracy on the test set (aim for 80%+).

In random_forest.py, once you are satisfied with your chosen parameters, update the following function:

• select_hyperparameters: change the values for <code>max_depth</code> , <code>n_estimators</code> , and <code>max_features</code> to your chosen values

Submit this file to Gradescope. You must achieve at least a **80% accuracy** against the test set in Gradescope to receive full credit for this section.

test_bootstrapping passed!

```
0.000
In [ ]:
       TODO:
       n_estimators defines how many Extra trees are fitted for the random forest.
       max_depth defines a stop condition when the tree reaches to a certain depth.
       max features controls the percentage of features that are used to fit each extra tree.
       Tune these three parameters to achieve a better accuracy. n_estimators and max_depth must both
       be at least 3 in value for moderately reliable answers. While you can use the provided test set
       to evaluate your implementation, you will need to obtain 80% on the test set to receive full
       credit for this section.
       from random_forest import RandomForest
       from sklearn import preprocessing
       import sklearn.ensemble
       ############# DO NOT CHANGE THIS RANDOM SEED ####################
       student random seed = 4641 + 7641
       n estimators = 4 #Hint: Consider values between 3-15.
       max depth = 15 # Hint: Consider values betweeen 3-15.
       max features = 1.0 # Hint: Consider values betweeen 0.3-1.0.
       random forest = RandomForest(n estimators, max depth, max features, random seed=student random seed)
       random_forest.fit(X_train, y_train)
       accuracy=random_forest.00B_score(X_test, y_test)
       print("accuracy: %.4f" % accuracy)
```

accuracy: 0.8178

DON'T FORGET: Once you are satisfied with your chosen parameters, change the values for <code>max_depth</code>, <code>n_estimators</code>, and <code>max_features</code> in the <code>select_hyperparameters()</code> function of your RandomForest class in <code>random_forest.py</code> to your chosen values, and then submit this file to Gradescope. You must achieve at least a **80% accuracy** against the test set in Gradescope to receive full credit for this section.

3.3 Plotting Feature Importance [5pts Bonus for All] **[W]**

While building tree-based models, it's common to quantify how well splitting on a particular feature in an extra tree helps with predicting the target label in a dataset. Machine learning practicioners typically use "Gini importance", or the (normalized) total reduction in entropy brought by that feature to evaluate how important that feature is for predicting the target variable.

Gini importance is typically calculated as the reduction in entropy from reaching a split in an extra tree weighted by the probability of reaching that split in the extra tree. Sklearn internally computes the probability for reaching a split by finding the total number of samples that reaches it during the training phase divided by the total number of samples in the dataset. This weighted value is our feature importance.

Let's think about what this metric means with an example. A high probability of reaching a split on feature A in an extra tree trained on a dataset (many samples will reach this split for a decision) and a large reduction in entropy from splitting on feature A will result in a high feature importance value for feature A. This could mean feature A is a very important feature for predicting the probability of the target label. On the other hand, a low probability of reaching a split on feature B in an extra tree and a low reduction in entropy from splitting on feature B will result in a low feature importance value. This could mean feature B is not a very informative feature for predicting the target label. Thus, the higher the feature importance value, the more important the feature is to predicting the target label.

Fortunately for us, fitting a sklearn. Extra Tree Classifier to a dataset automatically computes the Gini importance for every feature in the extra tree and stores these values in a **feature_importances_** variable. Review the docs for more details on how to access this variable

In the **random_forest.py** file, complete the following function:

• plot_feature_importance: Make sure to sort the bars in descending order and remove any features with feature importance of 0

In the cell below, call your implementation of plot_feature_importance() and display a bar plot that shows the feature importance values for at least one extra tree in your tuned random forest from Q3.2.

Note that there isn't one "correct" answer here. We simply want you to investigate how different features in your random forest contribute to predicting the target variable.

Also note that: the number of features can be different if you change max_features value since it ends up changing the number of features considered in bootstrapped datasets.

4: (Bonus for All) SVM [30 pts] **[W]** **[P]**

4.1 Fitting an SVM classifier by hand [20 pts] **[W]**

Consider a dataset with the following points in 2-dimensional space:

x_1	x_2	y
0	0	-1
0	2	-1
2	0	-1
3	3	1
3	4	1
2	4	1

Here, x_1 and x_2 are features and y is the label.

The max margin classifier has the formulation,

$$egin{aligned} \min \left| \left| heta
ight|
ight|^2 \ s.\,t.\,\,y_i(\mathbf{x_i} heta+b) \geq 1 \quad orall\,\,i \end{aligned}$$

Hint: $\mathbf{x_i}$ are the support vectors. Margin is equal to $\frac{1}{||\theta||}$ and full margin is equal to $\frac{2}{||\theta||}$. You might find it useful to plot the points in a 2D plane. When calculating the θ you don't need to consider the bias term.

- (1) Are the points linearly separable? Does adding the point $\mathbf{x}=(3,2)$, y=1 change the separability? (2 pts)
- (2) According to the max-margin formulation, find the separating hyperplane. Do not consider the new point from part 1 in your calculations for this current question or subsequent parts. (You should give some kind of explanation or calculation on how you found the hyperplane, you may solve this question graphically.) (4 pts)
- (3) Find a vector parallel to the optimal vector θ . (Hint: Recall whether the optimal vector is parallel or perpendicular to the separating hyperplane.) (4 pts)

- (4) Calculate the value of the margin (single-sided) achieved by this θ ? (4 pts)
- (5) Solve for θ , given that the margin is equal to $1/||\theta||$. (4 pts)
- (6) If we remove one of the points from the original data the SVM solution might change. Find all such points which change the solution. (2 pts)
- (7) Consider the optimization formulation stated above. Why do we want to optimzie $||\theta||^2$ instead of $|\theta||^2$ (2 pts)

4.2 Feature Mapping [10 pts] **[P]**

Let's look at a dataset where the datapoint can't be classified with a good accuracy using a linear classifier. Run the below cell to generate the dataset.

We will also see what happens when we try to fit a linear classifier to the dataset.

```
### DO NOT CHANGE THIS CELL ###
# Generate dataset
random_state = 1
np.random.seed(0)
theta = np.linspace(0, 2*np.pi, 1000)
r = np.random.uniform(0.8, 1.2, 1000)
X = np.column stack([r * np.cos(theta), r * np.sin(theta)])
y = np.logical_or(theta < np.pi, theta >= 2 * np.pi)
X[y == 0, 0] += 1
X[y == 0, 1] += 0.5
X_train, X_test, y_train, y_test = train_test_split(X, y,
                                               test size=0.20,
                                               random_state=random_state)
f, ax = plt.subplots(nrows=1, ncols=1,figsize=(5,5))
plt.scatter(X[:, 0], X[:, 1], c = y, marker = 'o', s=12)
plt.xlabel('X_1')
plt.ylabel('X_2')
plt.show()
```

```
###################################
### DO NOT CHANGE THIS CELL ###
######################################
def visualize decision boundary(X, y, feature new=None, h=0.02):
    You don't have to modify this function
    Function to vizualize decision boundary
    feature new is a function to get X with additional features
    x1 \text{ min}, x1 \text{ max} = X[:, 0].min() - 1, X[:, 0].max() + 1
    x2 min, x2 max = X[:, 1].min() - 1, X[:, 1].max() + 1
    xx_1, xx_2 = np.meshgrid(np.arange(x1_min, x1_max, h),
                          np.arange(x2 min, x2 max, h))
    if X.shape[1] == 2:
        Z = svm_cls.predict(np.c_[xx_1.ravel(), xx_2.ravel()])
    else:
        X_{conc} = np.c_[xx_1.ravel(), xx_2.ravel()]
        X new = feature new(X conc)
        Z = svm_cls.predict(X_new)
    Z = Z.reshape(xx_1.shape)
    f, ax = plt.subplots(nrows=1, ncols=1, figsize=(5,5))
    plt.contourf(xx_1, xx_2, Z, cmap=plt.cm.coolwarm, alpha=0.8)
    plt.scatter(X[:, 0], X[:, 1], c=y, cmap=plt.cm.coolwarm)
    plt.xlabel('X 1')
    plt.ylabel('X 2')
    plt.xlim(xx_1.min(), xx_1.max())
    plt.ylim(xx_2.min(), xx_2.max())
    plt.xticks(())
    plt.yticks(())
    plt.show()
```

We can see that we need a non-linear boundary to be able to successfully classify data in this dataset. By mapping the current feature x to a higher space with more features, linear SVM could be performed on the features in the higher space to learn a non-linear decision boundary. In feature.py, modify create_nl_feature() to add additional features which can help classify in the above dataset. After creating the additional features use code in the further cells to see how well the features perform on the test set.

Note: You should get a test accuracy above 90%

Hint: Think of the shape of the decision boundary that would best separate the above points. What additional features could help map the linear boundary to the non-linear one? Look at this for a detailed analysis of doing the same for points separable with a circular boundary