```
In [43]: # This mounts your Google Drive to the Colab VM.
         from google.colab import drive
         drive.mount('/content/drive')
         # TODO: Enter the foldername in your Drive where you have saved the unzipped
         # assignment folder, e.g. 'cs231n/assignments/assignment1/'
         FOLDERNAME = 'Coursework/ENPM703/assignment2'
         assert FOLDERNAME is not None, "[!] Enter the foldername."
         # Now that we've mounted your Drive, this ensures that
         # the Python interpreter of the Colab VM can load
         # python files from within it.
         import sys
         sys.path.append('/content/drive/My Drive/{}'.format(FOLDERNAME))
         # This downloads the CIFAR-10 dataset to your Drive
         # if it doesn't already exist.
         %cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/datasets/
         !bash get datasets.sh
         %cd /content/drive/My\ Drive/$FOLDERNAME
```

Drive already mounted at /content/drive; to attempt to forcibly remount, cal l drive.mount("/content/drive", force_remount=True). /content/drive/My Drive/Coursework/ENPM703/assignment2/cs231n/datasets /content/drive/My Drive/Coursework/ENPM703/assignment2

Multi-Layer Fully Connected Network

In this exercise, you will implement a fully connected network with an arbitrary number of hidden layers.

```
Read through the FullyConnectedNet class in the file cs231n/classifiers/fc_net.py.
```

Implement the network initialization, forward pass, and backward pass. Throughout this assignment, you will be implementing layers in cs231n/layers.py. You can re-use your implementations for affine_forward, affine_backward, relu_forward, relu_backward, and softmax_loss from Assignment 1. For right now, don't worry about implementing dropout or batch/layer normalization yet, as you will add those features later.

```
In [44]: # Setup cell.
import time
import numpy as np
import matplotlib.pyplot as plt
from cs231n.classifiers.fc_net import *
from cs231n.data_utils import get_CIFAR10_data
from cs231n.gradient_check import eval_numerical_gradient, eval_numerical_gr
```

```
from cs231n.solver import Solver

%matplotlib inline
plt.rcParams["figure.figsize"] = (10.0, 8.0) # Set default size of plots.
plt.rcParams["image.interpolation"] = "nearest"
plt.rcParams["image.cmap"] = "gray"

%load_ext autoreload
%autoreload 2

def rel_error(x, y):
    """Returns relative error."""
    return np.max(np.abs(x - y) / (np.maximum(1e-8, np.abs(x) + np.abs(y))))
```

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

```
In [45]: # Load the (preprocessed) CIFAR-10 data.
data = get_CIFAR10_data()
    for k, v in list(data.items()):
        print(f"{k}: {v.shape}")

X_train: (49000, 3, 32, 32)
    y_train: (49000,)
    X_val: (1000, 3, 32, 32)
    y_val: (1000,)
    X_test: (1000, 3, 32, 32)
    y_test: (1000,)
```

Initial Loss and Gradient Check

As a sanity check, run the following to check the initial loss and to gradient check the network both with and without regularization. This is a good way to see if the initial losses seem reasonable.

For gradient checking, you should expect to see errors around 1e-7 or less.

```
In [46]: np.random.seed(231)
N, D, H1, H2, C = 2, 15, 20, 30, 10
X = np.random.randn(N, D)
y = np.random.randint(C, size=(N,))

for reg in [0, 3.14]:
    print("Running check with reg = ", reg)
    model = FullyConnectedNet(
        [H1, H2],
        input_dim=D,
        num_classes=C,
        reg=reg,
        weight_scale=5e-2,
        dtype=np.float64
)

loss, grads = model.loss(X, y)
```

```
print("Initial loss: ", loss)
     # Most of the errors should be on the order of e-7 or smaller.
     # NOTE: It is fine however to see an error for W2 on the order of e-5
     # for the check when reg = 0.0
     for name in sorted(grads):
         f = lambda : model.loss(X, y)[0]
         grad num = eval numerical gradient(f, model.params[name], verbose=Fa
         print(f"{name} relative error: {rel error(grad num, grads[name])}")
Running check with reg = 0
Initial loss: 2.3004790897684924
W1 relative error: 7.696803870986541e-08
W2 relative error: 1.7087519140575808e-05
W3 relative error: 2.9508423118300657e-07
b1 relative error: 4.660094650186831e-09
b2 relative error: 2.085654124402131e-09
b3 relative error: 6.598642296022133e-11
Running check with reg = 3.14
Initial loss: 7.052114776533016
W1 relative error: 3.904542008453064e-09
W2 relative error: 6.86942277940646e-08
```

As another sanity check, make sure your network can overfit on a small dataset of 50 images. First, we will try a three-layer network with 100 units in each hidden layer. In the following cell, tweak the **learning rate** and **weight initialization scale** to overfit and achieve 100% training accuracy within 20 epochs.

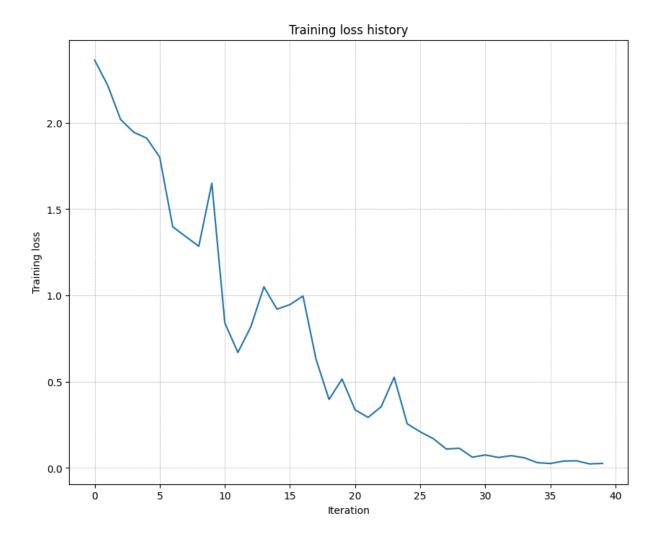
W3 relative error: 2.1311298702113723e-08 b1 relative error: 1.1683196894962977e-08 b2 relative error: 1.7223751746766738e-09 b3 relative error: 1.3200479211447775e-10

```
In [47]: # TODO: Use a three-layer Net to overfit 50 training examples by
         # tweaking just the learning rate and initialization scale.
         num train = 50
         small data = {
           "X train": data["X train"][:num train],
           "y_train": data["y_train"][:num_train],
           "X val": data["X val"],
           "y val": data["y val"],
         weight scale = 1e-2 # Experiment with this!
         learning rate = 1e-2 # Experiment with this!
         model = FullyConnectedNet(
             [100, 100],
             weight scale=weight scale,
             dtype=np.float64
         solver = Solver(
             model,
             small data,
             print every=10,
```

```
num epochs=20,
     batch size=25,
     update rule="sqd",
     optim config={"learning rate": learning rate},
 solver.train()
 plt.plot(solver.loss history)
 plt.title("Training loss history")
 plt.xlabel("Iteration")
 plt.ylabel("Training loss")
 plt.grid(linestyle='--', linewidth=0.5)
 plt.show()
(Iteration 1 / 40) loss: 2.363364
(Epoch 0 / 20) train acc: 0.180000; val acc: 0.108000
(Epoch 1 / 20) train acc: 0.320000; val acc: 0.127000
(Epoch 2 / 20) train acc: 0.440000; val acc: 0.172000
(Epoch 3 / 20) train acc: 0.500000; val acc: 0.184000
(Epoch 4 / 20) train acc: 0.540000; val acc: 0.181000
(Epoch 5 / 20) train acc: 0.740000; val acc: 0.190000
(Iteration 11 / 40) loss: 0.839976
(Epoch 6 / 20) train acc: 0.740000; val acc: 0.187000
(Epoch 7 / 20) train acc: 0.740000; val acc: 0.183000
(Epoch 8 / 20) train acc: 0.820000; val acc: 0.177000
(Epoch 9 / 20) train acc: 0.860000; val acc: 0.200000
(Epoch 10 / 20) train acc: 0.920000; val acc: 0.191000
(Iteration 21 / 40) loss: 0.337174
(Epoch 11 / 20) train acc: 0.960000; val acc: 0.189000
(Epoch 12 / 20) train acc: 0.940000; val acc: 0.180000
(Epoch 13 / 20) train acc: 1.000000; val acc: 0.199000
(Epoch 14 / 20) train acc: 1.000000; val acc: 0.199000
(Epoch 15 / 20) train acc: 1.000000; val acc: 0.195000
```

(Iteration 31 / 40) loss: 0.075911

(Epoch 16 / 20) train acc: 1.000000; val_acc: 0.182000 (Epoch 17 / 20) train acc: 1.000000; val_acc: 0.201000 (Epoch 18 / 20) train acc: 1.000000; val_acc: 0.207000 (Epoch 19 / 20) train acc: 1.000000; val_acc: 0.185000 (Epoch 20 / 20) train acc: 1.000000; val acc: 0.192000



Now, try to use a five-layer network with 100 units on each layer to overfit on 50 training examples. Again, you will have to adjust the learning rate and weight initialization scale, but you should be able to achieve 100% training accuracy within 20 epochs.

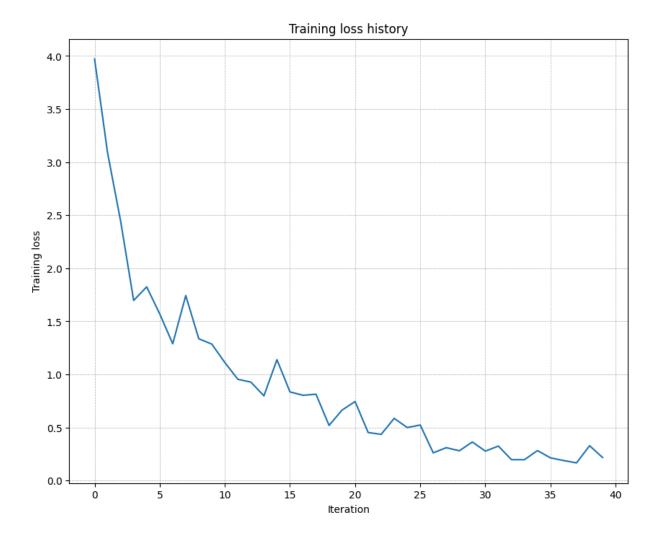
```
In [55]: # TODO: Use a five-layer Net to overfit 50 training examples by
    # tweaking just the learning rate and initialization scale.

num_train = 50
small_data = {
    'X_train': data['X_train'][:num_train],
    'y_train': data['y_train'][:num_train],
    'X_val': data['X_val'],
    'y_val': data['y_val'],
}

learning_rate = 2e-3  # Experiment with this!
weight_scale = 5e-2  # Experiment with this!
model = FullyConnectedNet(
    [100, 100, 100, 100],
    weight_scale=weight_scale,
    dtype=np.float64
)
```

```
solver = Solver(
    model,
     small data,
     print every=10,
     num epochs=20,
     batch size=25,
     update rule='sqd',
     optim config={'learning rate': learning rate},
 solver.train()
 plt.plot(solver.loss history)
 plt.title('Training loss history')
 plt.xlabel('Iteration')
 plt.ylabel('Training loss')
 plt.grid(linestyle='--', linewidth=0.5)
 plt.show()
(Iteration 1 / 40) loss: 3.971181
(Epoch 0 / 20) train acc: 0.180000; val acc: 0.139000
(Epoch 1 / 20) train acc: 0.260000; val acc: 0.127000
(Epoch 2 / 20) train acc: 0.480000; val acc: 0.120000
(Epoch 3 / 20) train acc: 0.560000; val acc: 0.152000
(Epoch 4 / 20) train acc: 0.760000; val acc: 0.158000
(Epoch 5 / 20) train acc: 0.780000; val acc: 0.148000
(Iteration 11 / 40) loss: 1.111498
(Epoch 6 / 20) train acc: 0.800000; val acc: 0.150000
(Epoch 7 / 20) train acc: 0.820000; val acc: 0.144000
(Epoch 8 / 20) train acc: 0.840000; val acc: 0.149000
(Epoch 9 / 20) train acc: 0.860000; val acc: 0.151000
(Epoch 10 / 20) train acc: 0.880000; val acc: 0.138000
(Iteration 21 / 40) loss: 0.743830
(Epoch 11 / 20) train acc: 0.940000; val acc: 0.155000
(Epoch 12 / 20) train acc: 0.960000; val acc: 0.159000
(Epoch 13 / 20) train acc: 0.960000; val acc: 0.159000
(Epoch 14 / 20) train acc: 0.960000; val acc: 0.147000
(Epoch 15 / 20) train acc: 1.000000; val acc: 0.159000
(Iteration 31 / 40) loss: 0.276849
(Epoch 16 / 20) train acc: 0.980000; val acc: 0.166000
(Epoch 17 / 20) train acc: 0.980000; val acc: 0.167000
```

(Epoch 18 / 20) train acc: 0.980000; val_acc: 0.168000 (Epoch 19 / 20) train acc: 0.980000; val_acc: 0.168000 (Epoch 20 / 20) train acc: 1.000000; val acc: 0.158000



Inline Question 1:

Did you notice anything about the comparative difficulty of training the three-layer network vs. training the five-layer network? In particular, based on your experience, which network seemed more sensitive to the initialization scale? Why do you think that is the case?

Answer:

When the three-layer and five-layer networks were trained side by side, it was found that the five-layer network was considerably more sensitive to the weight initialization scale. Its deeper focus increases the sensitivity since it increases the chance of seeing disappearing or bursting gradients.

Training Notes

Network in Three Layers:

• Scale of Weight: 1e-2

• **Training Accuracy**: By Epoch 15, **100**% was attained.

• Loss: Showing successful learning, it decreased from **2.31** to around **0.08**.

Gradient Behavior: Stabler gradients that make optimization simpler.

Network of Five Layers:

Scale of Weight: 5e-2

• **Training Accuracy**: Starting at just **22%**, it reached **100%** by Epoch 19.

• **Loss**: Commenced at **3.08**, indicating initial training difficulties.

Gradient Behavior: Because of the possibility of disappearing or bursting gradients, increased sensitivity to weight scale resulted in more intricate tuning.

Conclusion

Both networks attained 100% training accuracy; however, because of its depth, the five-layer network needed more cautious weight initialization. The threelayer network, on the other hand, provided a more seamless training process and a more noticeable ideal weight scale.

Update rules

So far we have used vanilla stochastic gradient descent (SGD) as our update rule. More sophisticated update rules can make it easier to train deep networks. We will implement a few of the most commonly used update rules and compare them to vanilla SGD.

SGD+Momentum

Stochastic gradient descent with momentum is a widely used update rule that tends to make deep networks converge faster than vanilla stochastic gradient descent. See the Momentum Update section at http://cs231n.github.io/neuralnetworks-3/#sgd for more information.

Open the file cs231n/optim.py and read the documentation at the top of the file to make sure you understand the API. Implement the SGD+momentum update rule in the function sgd momentum and run the following to check your implementation. You should see errors less than e-8.

```
N, D = 4, 5
w = np.linspace(-0.4, 0.6, num=N*D).reshape(N, D)
dw = np.linspace(-0.6, 0.4, num=N*D).reshape(N, D)
v = np.linspace(0.6, 0.9, num=N*D).reshape(N, D)
config = {"learning rate": 1e-3, "velocity": v}
next w, = sgd momentum(w, dw, config=config)
expected next w = np.asarray([
 [ 0.1406,
            0.20738947, 0.27417895, 0.34096842, 0.40775789],
 [0.47454737, 0.54133684, 0.60812632, 0.67491579, 0.74170526],
 [ 0.80849474, 0.87528421, 0.94207368, 1.00886316, 1.07565263],
 [ 1.14244211, 1.20923158, 1.27602105, 1.34281053, 1.4096
                                                              11)
expected velocity = np.asarray([
 [0.5406, 0.55475789, 0.56891579, 0.58307368, 0.59723158],
 [ 0.61138947, 0.62554737, 0.63970526, 0.65386316, 0.66802105],
 [0.68217895, 0.69633684, 0.71049474, 0.72465263, 0.73881053],
 [ 0.75296842, 0.76712632, 0.78128421, 0.79544211, 0.8096
                                                              ]])
# Should see relative errors around e-8 or less
print("next w error: ", rel error(next w, expected next w))
print("velocity error: ", rel_error(expected_velocity, config["velocity"]))
```

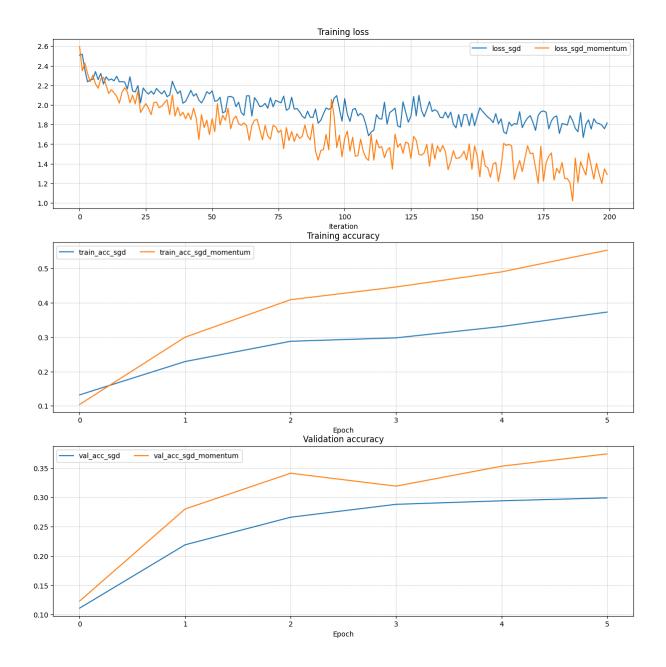
next_w error: 8.882347033505819e-09 velocity error: 4.269287743278663e-09

Once you have done so, run the following to train a six-layer network with both SGD and SGD+momentum. You should see the SGD+momentum update rule converge faster.

```
In [57]: num train = 4000
         small data = {
           'X train': data['X train'][:num train],
           'y_train': data['y_train'][:num_train],
           'X val': data['X val'],
           'y_val': data['y val'],
         solvers = {}
         for update_rule in ['sgd', 'sgd_momentum']:
             print('Running with ', update rule)
             model = FullyConnectedNet(
                  [100, 100, 100, 100, 100],
                 weight scale=5e-2
             solver = Solver(
                 model,
                  small data,
                  num epochs=5,
                  batch size=100,
                  update rule=update rule,
                  optim config={'learning rate': 5e-3},
                 verbose=True,
```

```
solvers[update_rule] = solver
    solver.train()
fig, axes = plt.subplots(3, 1, figsize=(15, 15))
axes[0].set title('Training loss')
axes[0].set_xlabel('Iteration')
axes[1].set_title('Training accuracy')
axes[1].set xlabel('Epoch')
axes[2].set_title('Validation accuracy')
axes[2].set xlabel('Epoch')
for update rule, solver in solvers.items():
    axes[0].plot(solver.loss history, label=f"loss {update rule}")
    axes[1].plot(solver.train acc history, label=f"train acc {update rule}")
    axes[2].plot(solver.val_acc_history, label=f"val_acc_{update_rule}")
for ax in axes:
   ax.legend(loc="best", ncol=4)
   ax.grid(linestyle='--', linewidth=0.5)
plt.show()
```

```
Running with sqd
(Iteration 1 / 200) loss: 2.508084
(Epoch 0 / 5) train acc: 0.132000; val acc: 0.111000
(Iteration 11 / 200) loss: 2.287128
(Iteration 21 / 200) loss: 2.143093
(Iteration 31 / 200) loss: 2.137349
(Epoch 1 / 5) train acc: 0.229000; val acc: 0.219000
(Iteration 41 / 200) loss: 2.035900
(Iteration 51 / 200) loss: 2.144458
(Iteration 61 / 200) loss: 2.027986
(Iteration 71 / 200) loss: 2.016312
(Epoch 2 / 5) train acc: 0.288000; val acc: 0.266000
(Iteration 81 / 200) loss: 2.079021
(Iteration 91 / 200) loss: 1.813389
(Iteration 101 / 200) loss: 2.064546
(Iteration 111 / 200) loss: 1.723232
(Epoch 3 / 5) train acc: 0.298000; val acc: 0.288000
(Iteration 121 / 200) loss: 1.787481
(Iteration 131 / 200) loss: 1.880817
(Iteration 141 / 200) loss: 1.927259
(Iteration 151 / 200) loss: 1.883699
(Epoch 4 / 5) train acc: 0.331000; val acc: 0.294000
(Iteration 161 / 200) loss: 1.725596
(Iteration 171 / 200) loss: 1.891697
(Iteration 181 / 200) loss: 1.887813
(Iteration 191 / 200) loss: 1.667553
(Epoch 5 / 5) train acc: 0.373000; val acc: 0.299000
Running with sgd momentum
(Iteration 1 / 200) loss: 2.595847
(Epoch 0 / 5) train acc: 0.104000; val acc: 0.123000
(Iteration 11 / 200) loss: 2.202265
(Iteration 21 / 200) loss: 2.100635
(Iteration 31 / 200) loss: 1.971155
(Epoch 1 / 5) train acc: 0.300000; val acc: 0.280000
(Iteration 41 / 200) loss: 1.858637
(Iteration 51 / 200) loss: 1.859528
(Iteration 61 / 200) loss: 1.809259
(Iteration 71 / 200) loss: 1.791472
(Epoch 2 / 5) train acc: 0.409000; val acc: 0.341000
(Iteration 81 / 200) loss: 1.728877
(Iteration 91 / 200) loss: 1.436205
(Iteration 101 / 200) loss: 1.659388
(Iteration 111 / 200) loss: 1.699924
(Epoch 3 / 5) train acc: 0.446000; val acc: 0.319000
(Iteration 121 / 200) loss: 1.568407
(Iteration 131 / 200) loss: 1.512991
(Iteration 141 / 200) loss: 1.420275
(Iteration 151 / 200) loss: 1.478698
(Epoch 4 / 5) train acc: 0.490000; val acc: 0.353000
(Iteration 161 / 200) loss: 1.607737
(Iteration 171 / 200) loss: 1.503822
(Iteration 181 / 200) loss: 1.353460
(Iteration 191 / 200) loss: 1.349060
(Epoch 5 / 5) train acc: 0.553000; val acc: 0.374000
```



RMSProp and Adam

RMSProp [1] and Adam [2] are update rules that set per-parameter learning rates by using a running average of the second moments of gradients.

In the file cs231n/optim.py , implement the RMSProp update rule in the rmsprop function and implement the Adam update rule in the adam function, and check your implementations using the tests below.

NOTE: Please implement the *complete* Adam update rule (with the bias correction mechanism), not the first simplified version mentioned in the course notes.

[1] Tijmen Tieleman and Geoffrey Hinton. "Lecture 6.5-rmsprop: Divide the gradient by a running average of its recent magnitude." COURSERA: Neural

Networks for Machine Learning 4 (2012).

[2] Diederik Kingma and Jimmy Ba, "Adam: A Method for Stochastic Optimization", ICLR 2015.

```
In [58]: # Test RMSProp implementation
         from cs231n.optim import rmsprop
         N, D = 4, 5
         w = np.linspace(-0.4, 0.6, num=N*D).reshape(N, D)
         dw = np.linspace(-0.6, 0.4, num=N*D).reshape(N, D)
         cache = np.linspace(0.6, 0.9, num=N*D).reshape(N, D)
         config = {'learning rate': 1e-2, 'cache': cache}
         next w, = rmsprop(w, dw, config=config)
         expected next w = np.asarray([
          [-0.39223849, -0.34037513, -0.28849239, -0.23659121, -0.18467247],
           [-0.132737, -0.08078555, -0.02881884, 0.02316247, 0.07515774],
           [0.12716641, 0.17918792, 0.23122175, 0.28326742, 0.33532447],
           [0.38739248, 0.43947102, 0.49155973, 0.54365823, 0.59576619]])
         expected cache = np.asarray([
          [0.5976, 0.6126277, 0.6277108, 0.64284931, 0.65804321],
           [0.67329252, 0.68859723, 0.70395734, 0.71937285, 0.73484377],
           [0.75037008, 0.7659518, 0.78158892, 0.79728144, 0.81302936],
          [ 0.82883269, 0.84469141, 0.86060554, 0.87657507, 0.8926 ]])
         # You should see relative errors around e-7 or less
         print('next w error: ', rel error(expected next w, next w))
         print('cache error: ', rel error(expected cache, config['cache']))
        next w error: 9.524687511038133e-08
```

next_w error: 9.524687511038133e-08 cache error: 2.6477955807156126e-09

```
In [59]: # Test Adam implementation
         from cs231n.optim import adam
         N, D = 4, 5
         w = np.linspace(-0.4, 0.6, num=N*D).reshape(N, D)
         dw = np.linspace(-0.6, 0.4, num=N*D).reshape(N, D)
         m = np.linspace(0.6, 0.9, num=N*D).reshape(N, D)
         v = np.linspace(0.7, 0.5, num=N*D).reshape(N, D)
         config = {'learning rate': 1e-2, 'm': m, 'v': v, 't': 5}
         next w, = adam(w, dw, config=config)
         expected next w = np.asarray([
          [-0.40094747, -0.34836187, -0.29577703, -0.24319299, -0.19060977],
           [-0.1380274, -0.08544591, -0.03286534, 0.01971428, 0.0722929],
                       0.17744702, 0.23002243, 0.28259667, 0.33516969],
           [ 0.1248705,
           [0.38774145, 0.44031188, 0.49288093, 0.54544852, 0.59801459]])
         expected v = np.asarray([
          [0.69966, 0.68908382, 0.67851319, 0.66794809, 0.65738853,],
           [0.64683452, 0.63628604, 0.6257431, 0.61520571, 0.60467385,],
           [0.59414753, 0.58362676, 0.57311152, 0.56260183, 0.55209767,],
```

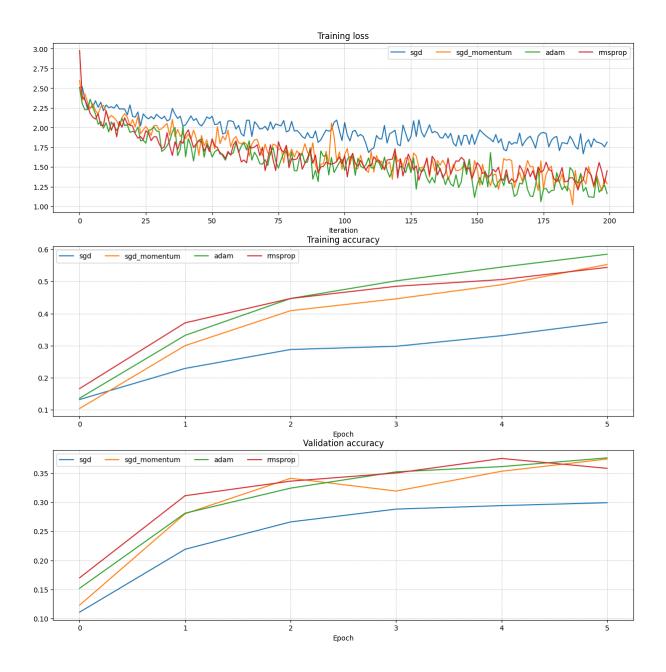
next_w error: 1.1395691798535431e-07 v error: 4.208314038113071e-09 m error: 4.214963193114416e-09

Once you have debugged your RMSProp and Adam implementations, run the following to train a pair of deep networks using these new update rules:

```
In [60]: learning rates = {'rmsprop': le-4, 'adam': le-3}
         for update_rule in ['adam', 'rmsprop']:
             print('Running with ', update rule)
             model = FullyConnectedNet(
                 [100, 100, 100, 100, 100],
                 weight scale=5e-2
             solver = Solver(
                 model,
                 small data,
                 num epochs=5,
                 batch size=100,
                 update rule=update rule,
                 optim config={'learning rate': learning rates[update rule]},
                 verbose=True
             solvers[update rule] = solver
             solver.train()
             print()
         fig, axes = plt.subplots(3, 1, figsize=(15, 15))
         axes[0].set title('Training loss')
         axes[0].set xlabel('Iteration')
         axes[1].set title('Training accuracy')
         axes[1].set xlabel('Epoch')
         axes[2].set title('Validation accuracy')
         axes[2].set xlabel('Epoch')
         for update rule, solver in solvers.items():
             axes[0].plot(solver.loss history, label=f"{update rule}")
             axes[1].plot(solver.train acc history, label=f"{update rule}")
             axes[2].plot(solver.val acc history, label=f"{update rule}")
         for ax in axes:
             ax.legend(loc='best', ncol=4)
             ax.grid(linestyle='--', linewidth=0.5)
```

plt.show()

```
Running with adam
(Iteration 1 / 200) loss: 2.503933
(Epoch 0 / 5) train acc: 0.136000; val acc: 0.152000
(Iteration 11 / 200) loss: 2.061754
(Iteration 21 / 200) loss: 1.952930
(Iteration 31 / 200) loss: 1.949460
(Epoch 1 / 5) train acc: 0.332000; val acc: 0.281000
(Iteration 41 / 200) loss: 1.629484
(Iteration 51 / 200) loss: 1.627666
(Iteration 61 / 200) loss: 1.683181
(Iteration 71 / 200) loss: 1.593157
(Epoch 2 / 5) train acc: 0.447000; val acc: 0.324000
(Iteration 81 / 200) loss: 1.691935
(Iteration 91 / 200) loss: 1.316360
(Iteration 101 / 200) loss: 1.457656
(Iteration 111 / 200) loss: 1.490421
(Epoch 3 / 5) train acc: 0.502000; val acc: 0.352000
(Iteration 121 / 200) loss: 1.437246
(Iteration 131 / 200) loss: 1.472117
(Iteration 141 / 200) loss: 1.284863
(Iteration 151 / 200) loss: 1.276522
(Epoch 4 / 5) train acc: 0.545000; val acc: 0.361000
(Iteration 161 / 200) loss: 1.265843
(Iteration 171 / 200) loss: 1.134311
(Iteration 181 / 200) loss: 1.429556
(Iteration 191 / 200) loss: 1.193171
(Epoch 5 / 5) train acc: 0.585000; val acc: 0.376000
Running with rmsprop
(Iteration 1 / 200) loss: 2.976445
(Epoch 0 / 5) train acc: 0.166000; val acc: 0.170000
(Iteration 11 / 200) loss: 2.215317
(Iteration 21 / 200) loss: 2.019541
(Iteration 31 / 200) loss: 1.796836
(Epoch 1 / 5) train acc: 0.371000; val acc: 0.311000
(Iteration 41 / 200) loss: 1.914885
(Iteration 51 / 200) loss: 1.854153
(Iteration 61 / 200) loss: 1.698718
(Iteration 71 / 200) loss: 1.455506
(Epoch 2 / 5) train acc: 0.447000; val acc: 0.336000
(Iteration 81 / 200) loss: 1.607075
(Iteration 91 / 200) loss: 1.421859
(Iteration 101 / 200) loss: 1.644586
(Iteration 111 / 200) loss: 1.517447
(Epoch 3 / 5) train acc: 0.485000; val acc: 0.350000
(Iteration 121 / 200) loss: 1.361570
(Iteration 131 / 200) loss: 1.493687
(Iteration 141 / 200) loss: 1.408904
(Iteration 151 / 200) loss: 1.555976
(Epoch 4 / 5) train acc: 0.506000; val acc: 0.375000
(Iteration 161 / 200) loss: 1.332950
(Iteration 171 / 200) loss: 1.378604
(Iteration 181 / 200) loss: 1.314109
(Iteration 191 / 200) loss: 1.334453
(Epoch 5 / 5) train acc: 0.544000; val acc: 0.358000
```



Inline Question 2:

AdaGrad, like Adam, is a per-parameter optimization method that uses the following update rule:

```
cache += dw**2
w += - learning rate * dw / (np.sqrt(cache) + eps)
```

John notices that when he was training a network with AdaGrad that the updates became very small, and that his network was learning slowly. Using your knowledge of the AdaGrad update rule, why do you think the updates would become very small? Would Adam have the same issue?

Answer:

Smaller updates result from a decreasing step size over time due to the monotonic growth in the cache's squared gradients' cumulative value. Slower updates at the global minimum might be beneficial if the optimization function is convex. On the other hand, this may result in the algorithm becoming trapped in local minima for non-convex functions.

The Adam algorithm uses bias correction to lessen the effects of the growing time, therefore avoiding this problem.

 $b_1 = m/(1-\beta_1^t)$ and $b_2 = v/(1-\beta_2^t)$ are the update rules for Adam. The numerator of the updates will also rise when (t) increases, as both (b_1) and (b_2) increase as well. This keeps the updates from being too tiny.

Train a Good Model!

Train the best fully connected model that you can on CIFAR-10, storing your best model in the best_model variable. We require you to get at least 50% accuracy on the validation set using a fully connected network.

If you are careful it should be possible to get accuracies above 55%, but we don't require it for this part and won't assign extra credit for doing so. Later in the assignment we will ask you to train the best convolutional network that you can on CIFAR-10, and we would prefer that you spend your effort working on convolutional networks rather than fully connected networks.

Note: You might find it useful to complete the BatchNormalization.ipynb and Dropout.ipynb notebooks before completing this part, since those techniques can help you train powerful models.

```
reg=reg,
    weight scale=weight scales,
    dtype=np.float64
 # Set up solver
 solver = Solver(
    model,
    data,
    num epochs=10,
    batch size=100,
    update rule="adam",
    optim config={"learning rate": learning rates},
    verbose=True,
    print every=1000
 # Train model
 solver.train()
# Save best model
best model = model
 # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
END OF YOUR CODE
(Iteration 1 / 4900) loss: 2.309740
(Epoch 0 / 10) train acc: 0.103000; val acc: 0.098000
(Epoch 1 / 10) train acc: 0.374000; val acc: 0.355000
(Epoch 2 / 10) train acc: 0.418000; val acc: 0.419000
(Iteration 1001 / 4900) loss: 1.801674
(Epoch 3 / 10) train acc: 0.484000; val acc: 0.442000
(Epoch 4 / 10) train acc: 0.479000; val acc: 0.443000
(Iteration 2001 / 4900) loss: 1.545944
(Epoch 5 / 10) train acc: 0.495000; val acc: 0.472000
(Epoch 6 / 10) train acc: 0.532000; val acc: 0.494000
(Iteration 3001 / 4900) loss: 1.353906
(Epoch 7 / 10) train acc: 0.547000; val acc: 0.510000
(Epoch 8 / 10) train acc: 0.512000; val acc: 0.504000
(Iteration 4001 / 4900) loss: 1.293499
(Epoch 9 / 10) train acc: 0.538000; val acc: 0.516000
(Epoch 10 / 10) train acc: 0.544000; val acc: 0.500000
```

Test Your Model!

Run your best model on the validation and test sets. You should achieve at least 50% accuracy on the validation set.

```
In [62]: y_test_pred = np.argmax(best_model.loss(data['X_test']), axis=1)
    y_val_pred = np.argmax(best_model.loss(data['X_val']), axis=1)
```

```
print('Validation set accuracy: ', (y_val_pred == data['y_val']).mean())
print('Test set accuracy: ', (y_test_pred == data['y_test']).mean())
```

Validation set accuracy: 0.516

Test set accuracy: 0.525

This notebook was converted to PDF with convert.ploomber.io

```
In [ ]: # This mounts your Google Drive to the Colab VM.
        from google.colab import drive
        drive.mount('/content/drive')
        # TODO: Enter the foldername in your Drive where you have saved the unzipped
        # assignment folder, e.g. 'cs231n/assignments/assignment1/'
        FOLDERNAME = None
        assert FOLDERNAME is not None, "[!] Enter the foldername."
        # Now that we've mounted your Drive, this ensures that
        # the Python interpreter of the Colab VM can load
        # python files from within it.
        import sys
        sys.path.append('/content/drive/My Drive/{}'.format(FOLDERNAME))
        # This downloads the CIFAR-10 dataset to your Drive
        # if it doesn't already exist.
        %cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/datasets/
        !bash get datasets.sh
        %cd /content/drive/My\ Drive/$FOLDERNAME
```

Batch Normalization

One way to make deep networks easier to train is to use more sophisticated optimization procedures such as SGD+momentum, RMSProp, or Adam. Another strategy is to change the architecture of the network to make it easier to train. One idea along these lines is batch normalization, proposed by [1] in 2015.

To understand the goal of batch normalization, it is important to first recognize that machine learning methods tend to perform better with input data consisting of uncorrelated features with zero mean and unit variance. When training a neural network, we can preprocess the data before feeding it to the network to explicitly decorrelate its features. This will ensure that the first layer of the network sees data that follows a nice distribution. However, even if we preprocess the input data, the activations at deeper layers of the network will likely no longer be decorrelated and will no longer have zero mean or unit variance, since they are output from earlier layers in the network. Even worse, during the training process the distribution of features at each layer of the network will shift as the weights of each layer are updated.

The authors of [1] hypothesize that the shifting distribution of features inside deep neural networks may make training deep networks more difficult. To overcome this problem, they propose to insert into the network layers that normalize batches. At training time, such a layer uses a minibatch of data to estimate the mean and standard deviation of each feature. These estimated means and standard deviations are then used to center and normalize the

features of the minibatch. A running average of these means and standard deviations is kept during training, and at test time these running averages are used to center and normalize features.

It is possible that this normalization strategy could reduce the representational power of the network, since it may sometimes be optimal for certain layers to have features that are not zero-mean or unit variance. To this end, the batch normalization layer includes learnable shift and scale parameters for each feature dimension.

[1] Sergey Ioffe and Christian Szegedy, "Batch Normalization: Accelerating Deep Network Training by Reducing Internal Covariate Shift", ICML 2015.

```
In [ ]: # Setup cell.
        import time
        import numpy as np
        import matplotlib.pyplot as plt
        from cs231n.classifiers.fc net import *
        from cs231n.data utils import get CIFAR10 data
        from cs231n.gradient check import eval numerical gradient, eval numerical gr
        from cs231n.solver import Solver
        %matplotlib inline
        plt.rcParams["figure.figsize"] = (10.0, 8.0) # Set default size of plots.
        plt.rcParams["image.interpolation"] = "nearest"
        plt.rcParams["image.cmap"] = "gray"
        %load ext autoreload
        %autoreload 2
        def rel error(x, y):
            """Returns relative error."""
            return np.max(np.abs(x - y) / (np.maximum(1e-8, np.abs(x) + np.abs(y))))
        def print mean std(x,axis=0):
            print(f" means: {x.mean(axis=axis)}")
            print(f" stds: {x.std(axis=axis)}\n")
In [ ]: # Load the (preprocessed) CIFAR-10 data.
        data = get CIFAR10 data()
        for k, v in list(data.items()):
            print(f"{k}: {v.shape}")
```

Batch Normalization: Forward Pass

In the file cs231n/layers.py , implement the batch normalization forward pass in the function batchnorm_forward . Once you have done so, run the following to test your implementation.

```
In [ ]: # Check the training-time forward pass by checking means and variances
        # of features both before and after batch normalization
        # Simulate the forward pass for a two-layer network.
        np.random.seed(231)
        N, D1, D2, D3 = 200, 50, 60, 3
        X = np.random.randn(N, D1)
        W1 = np.random.randn(D1, D2)
        W2 = np.random.randn(D2, D3)
        a = np.maximum(0, X.dot(W1)).dot(W2)
        print('Before batch normalization:')
        print mean std(a,axis=0)
        gamma = np.ones((D3,))
        beta = np.zeros((D3,))
        # Means should be close to zero and stds close to one.
        print('After batch normalization (gamma=1, beta=0)')
        a norm, = batchnorm forward(a, gamma, beta, {'mode': 'train'})
        print mean std(a norm,axis=0)
        gamma = np.asarray([1.0, 2.0, 3.0])
        beta = np.asarray([11.0, 12.0, 13.0])
        # Now means should be close to beta and stds close to gamma.
        print('After batch normalization (gamma=', gamma, ', beta=', beta, ')')
        a norm, = batchnorm forward(a, gamma, beta, {'mode': 'train'})
        print mean std(a norm,axis=0)
In [ ]: # Check the test-time forward pass by running the training-time
        # forward pass many times to warm up the running averages, and then
        # checking the means and variances of activations after a test-time
        # forward pass.
        np.random.seed(231)
        N, D1, D2, D3 = 200, 50, 60, 3
        W1 = np.random.randn(D1, D2)
        W2 = np.random.randn(D2, D3)
        bn param = {'mode': 'train'}
        gamma = np.ones(D3)
        beta = np.zeros(D3)
        for t in range(50):
          X = np.random.randn(N, D1)
          a = np.maximum(0, X.dot(W1)).dot(W2)
          batchnorm forward(a, gamma, beta, bn param)
        bn param['mode'] = 'test'
        X = np.random.randn(N, D1)
        a = np.maximum(0, X.dot(W1)).dot(W2)
        a norm, = batchnorm forward(a, gamma, beta, bn param)
```

```
# Means should be close to zero and stds close to one, but will be
# noisier than training-time forward passes.
print('After batch normalization (test-time):')
print_mean_std(a_norm,axis=0)
```

Batch Normalization: Backward Pass

Now implement the backward pass for batch normalization in the function batchnorm_backward .

To derive the backward pass you should write out the computation graph for batch normalization and backprop through each of the intermediate nodes. Some intermediates may have multiple outgoing branches; make sure to sum gradients across these branches in the backward pass.

Once you have finished, run the following to numerically check your backward pass.

```
In [ ]: # Gradient check batchnorm backward pass.
        np.random.seed(231)
        N, D = 4, 5
        x = 5 * np.random.randn(N, D) + 12
        gamma = np.random.randn(D)
        beta = np.random.randn(D)
        dout = np.random.randn(N, D)
        bn param = {'mode': 'train'}
        fx = lambda x: batchnorm forward(x, gamma, beta, bn param)[0]
        fg = lambda a: batchnorm forward(x, a, beta, bn param)[0]
        fb = lambda b: batchnorm forward(x, gamma, b, bn param)[0]
        dx num = eval numerical gradient array(fx, x, dout)
        da num = eval numerical gradient array(fg, gamma.copy(), dout)
        db num = eval numerical gradient array(fb, beta.copy(), dout)
        _, cache = batchnorm_forward(x, gamma, beta, bn_param)
        dx, dgamma, dbeta = batchnorm backward(dout, cache)
        # You should expect to see relative errors between 1e-13 and 1e-8.
        print('dx error: ', rel error(dx num, dx))
        print('dgamma error: ', rel error(da num, dgamma))
        print('dbeta error: ', rel_error(db_num, dbeta))
```

Batch Normalization: Alternative Backward Pass

In class we talked about two different implementations for the sigmoid backward pass. One strategy is to write out a computation graph composed of simple operations and backprop through all intermediate values. Another strategy is to work out the derivatives on paper. For example, you can derive a very simple formula for the sigmoid function's backward pass by simplifying gradients on paper.

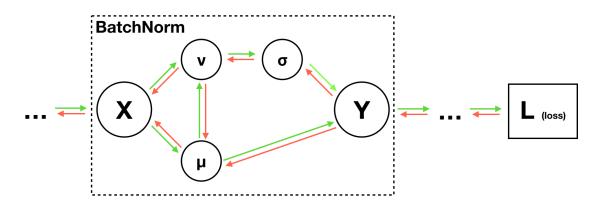
Surprisingly, it turns out that you can do a similar simplification for the batch normalization backward pass too!

In the forward pass, given a set of inputs
$$X = \begin{bmatrix} x_1 \\ x_2 \\ \dots \\ x_N \end{bmatrix}$$
 ,

we first calculate the mean μ and variance v. With μ and v calculated, we can calculate the standard deviation σ and normalized data Y. The equations and graph illustration below describe the computation (y_i is the i-th element of the vector Y).

$$\mu = \frac{1}{N} \sum_{k=1}^{N} x_k \qquad v = \frac{1}{N} \sum_{k=1}^{N} (x_k - \mu)^2$$
 (1)

$$\sigma = \sqrt{v + \epsilon}$$
 $y_i = \frac{x_i - \mu}{\sigma}$ (2)



The meat of our problem during backpropagation is to compute $\frac{\partial L}{\partial X}$, given the upstream gradient we receive, $\frac{\partial L}{\partial Y}$. To do this, recall the chain rule in calculus gives us $\frac{\partial L}{\partial X} = \frac{\partial L}{\partial Y} \cdot \frac{\partial Y}{\partial X}$.

The unknown/hard part is $\frac{\partial Y}{\partial X}$. We can find this by first deriving step-by-step our local gradients at $\frac{\partial v}{\partial X}$, $\frac{\partial \mu}{\partial X}$, $\frac{\partial \sigma}{\partial v}$, $\frac{\partial Y}{\partial \sigma}$, and then use the chain rule to

compose these gradients (which appear in the form of vectors!) appropriately to compute $\frac{\partial Y}{\partial X}$.

If it's challenging to directly reason about the gradients over X and Y which require matrix multiplication, try reasoning about the gradients in terms of individual elements x_i and y_i first: in that case, you will need to come up with the derivations for $\frac{\partial L}{\partial x_i}$, by relying on the Chain Rule to first calculate the intermediate $\frac{\partial \mu}{\partial x_i}$, $\frac{\partial v}{\partial x_i}$, $\frac{\partial \sigma}{\partial x_i}$, then assemble these pieces to calculate $\frac{\partial y_i}{\partial x_i}$.

You should make sure each of the intermediary gradient derivations are all as simplified as possible, for ease of implementation.

After doing so, implement the simplified batch normalization backward pass in the function batchnorm_backward_alt and compare the two implementations by running the following. Your two implementations should compute nearly identical results, but the alternative implementation should be a bit faster.

```
In [ ]: np.random.seed(231)
        N, D = 100, 500
        x = 5 * np.random.randn(N, D) + 12
        gamma = np.random.randn(D)
        beta = np.random.randn(D)
        dout = np.random.randn(N, D)
        bn param = {'mode': 'train'}
        out, cache = batchnorm_forward(x, gamma, beta, bn_param)
        t1 = time.time()
        dx1, dgamma1, dbeta1 = batchnorm_backward(dout, cache)
        t2 = time.time()
        dx2, dgamma2, dbeta2 = batchnorm backward alt(dout, cache)
        t3 = time.time()
        print('dx difference: ', rel error(dx1, dx2))
        print('dgamma difference: ', rel_error(dgamma1, dgamma2))
        print('dbeta difference: ', rel error(dbeta1, dbeta2))
        print('speedup: %.2fx' % ((t2 - t1) / (t3 - t2)))
```

Fully Connected Networks with Batch Normalization

Now that you have a working implementation for batch normalization, go back to your FullyConnectedNet in the file cs231n/classifiers/fc_net.py . Modify your implementation to add batch normalization.

Concretely, when the normalization flag is set to "batchnorm" in the constructor, you should insert a batch normalization layer before each ReLU

nonlinearity. The outputs from the last layer of the network should not be normalized. Once you are done, run the following to gradient-check your implementation.

Hint: You might find it useful to define an additional helper layer similar to those in the file cs231n/layer_utils.py .

```
In [ ]: np.random.seed(231)
        N, D, H1, H2, C = 2, 15, 20, 30, 10
        X = np.random.randn(N, D)
        y = np.random.randint(C, size=(N,))
        # You should expect losses between 1e-4~1e-10 for W,
        # losses between 1e-08~1e-10 for b,
        # and losses between 1e-08~1e-09 for beta and gammas.
        for reg in [0, 3.14]:
          print('Running check with reg = ', reg)
          model = FullyConnectedNet([H1, H2], input dim=D, num classes=C,
                                    reg=reg, weight scale=5e-2, dtype=np.float64,
                                    normalization='batchnorm')
          loss, grads = model.loss(X, y)
          print('Initial loss: ', loss)
          for name in sorted(grads):
            f = lambda : model.loss(X, y)[0]
            grad num = eval numerical gradient(f, model.params[name], verbose=False,
            print('%s relative error: %.2e' % (name, rel error(grad num, grads[name]
          if reg == 0: print()
```

Batch Normalization for Deep Networks

Run the following to train a six-layer network on a subset of 1000 training examples both with and without batch normalization.

```
In []: np.random.seed(231)

# Try training a very deep net with batchnorm.
hidden_dims = [100, 100, 100, 100]

num_train = 1000
small_data = {
    'X_train': data['X_train'][:num_train],
    'y_train': data['y_train'][:num_train],
    'X_val': data['X_val'],
    'y_val': data['y_val'],
}

weight_scale = 2e-2
```

```
bn model = FullyConnectedNet(hidden dims, weight scale=weight scale, normali
model = FullyConnectedNet(hidden dims, weight scale=weight scale, normalizat
print('Solver with batch norm:')
bn solver = Solver(bn model, small data,
                num epochs=10, batch size=50,
                update rule='adam',
                optim config={
                  'learning rate': 1e-3,
                verbose=True,print every=20)
bn solver.train()
print('\nSolver without batch norm:')
solver = Solver(model, small data,
                num epochs=10, batch size=50,
                update rule='adam',
                optim config={
                  'learning rate': 1e-3,
                verbose=True, print every=20)
solver.train()
```

Run the following to visualize the results from two networks trained above. You should find that using batch normalization helps the network to converge much faster.

```
In [ ]: def plot training history(title, label, baseline, bn solvers, plot fn, bl ma
            """utility function for plotting training history"""
            plt.title(title)
            plt.xlabel(label)
            bn plots = [plot fn(bn solver) for bn solver in bn solvers]
            bl plot = plot fn(baseline)
            num bn = len(bn plots)
            for i in range(num bn):
                label='with norm'
                if labels is not None:
                    label += str(labels[i])
                plt.plot(bn plots[i], bn marker, label=label)
            label='baseline'
            if labels is not None:
                label += str(labels[0])
            plt.plot(bl plot, bl marker, label=label)
            plt.legend(loc='lower center', ncol=num bn+1)
        plt.subplot(3, 1, 1)
        plot training history('Training loss','Iteration', solver, [bn solver], \
                              lambda x: x.loss history, bl marker='o', bn marker='o'
        plt.subplot(3, 1, 2)
        plot training history('Training accuracy','Epoch', solver, [bn solver], \
                              lambda x: x.train acc history, bl marker='-o', bn mark
        plt.subplot(3, 1, 3)
        plot training history('Validation accuracy', 'Epoch', solver, [bn solver], \
                              lambda x: x.val acc history, bl marker='-o', bn marker
```

```
plt.gcf().set_size_inches(15, 15)
plt.show()
```

Batch Normalization and Initialization

We will now run a small experiment to study the interaction of batch normalization and weight initialization.

The first cell will train eight-layer networks both with and without batch normalization using different scales for weight initialization. The second layer will plot training accuracy, validation set accuracy, and training loss as a function of the weight initialization scale.

```
In [ ]: np.random.seed(231)
        # Try training a very deep net with batchnorm.
        hidden dims = [50, 50, 50, 50, 50, 50, 50]
        num train = 1000
        small data = {
          'X train': data['X train'][:num train],
          'y train': data['y train'][:num train],
          'X val': data['X val'],
          'y_val': data['y_val'],
        bn solvers ws = {}
        solvers ws = {}
        weight scales = np.logspace(-4, 0, num=20)
        for i, weight scale in enumerate(weight scales):
            print('Running weight scale %d / %d' % (i + 1, len(weight scales)))
            bn model = FullyConnectedNet(hidden dims, weight scale=weight scale, nor
            model = FullyConnectedNet(hidden dims, weight scale=weight scale, normal
            bn solver = Solver(bn model, small data,
                           num epochs=10, batch size=50,
                           update rule='adam',
                          optim config={
                             'learning rate': 1e-3,
                           },
                          verbose=False, print every=200)
            bn solver.train()
            bn solvers ws[weight scale] = bn solver
            solver = Solver(model, small data,
                           num epochs=10, batch size=50,
                           update rule='adam',
                          optim config={
                             'learning rate': 1e-3,
                           },
                          verbose=False, print every=200)
```

```
solver.train()
solvers_ws[weight_scale] = solver
```

```
In [ ]: # Plot results of weight scale experiment.
        best train accs, bn best train accs = [], []
        best val accs, bn best val accs = [], []
        final train loss, bn final train loss = [], []
        for ws in weight scales:
          best train accs.append(max(solvers ws[ws].train acc history))
          bn best train accs.append(max(bn solvers ws[ws].train acc history))
          best val accs.append(max(solvers ws[ws].val acc history))
          bn best val accs.append(max(bn solvers ws[ws].val acc history))
          final train loss.append(np.mean(solvers ws[ws].loss history[-100:]))
          bn final train loss.append(np.mean(bn solvers ws[ws].loss history[-100:]))
        plt.subplot(3, 1, 1)
        plt.title('Best val accuracy vs. weight initialization scale')
        plt.xlabel('Weight initialization scale')
        plt.ylabel('Best val accuracy')
        plt.semilogx(weight scales, best val accs, '-o', label='baseline')
        plt.semilogx(weight scales, bn best val accs, '-o', label='batchnorm')
        plt.legend(ncol=2, loc='lower right')
        plt.subplot(3, 1, 2)
        plt.title('Best train accuracy vs. weight initialization scale')
        plt.xlabel('Weight initialization scale')
        plt.ylabel('Best training accuracy')
        plt.semilogx(weight scales, best train accs, '-o', label='baseline')
        plt.semilogx(weight scales, bn best train accs, '-o', label='batchnorm')
        plt.legend()
        plt.subplot(3, 1, 3)
        plt.title('Final training loss vs. weight initialization scale')
        plt.xlabel('Weight initialization scale')
        plt.ylabel('Final training loss')
        plt.semilogx(weight scales, final train loss, '-o', label='baseline')
        plt.semilogx(weight scales, bn_final_train_loss, '-o', label='batchnorm')
        plt.legend()
        plt.gca().set ylim(1.0, 3.5)
        plt.gcf().set size inches(15, 15)
        plt.show()
```

Inline Question 1:

Describe the results of this experiment. How does the weight initialization scale affect models with/without batch normalization differently, and why?

Answer:

Batch Normalization and Batch Size

We will now run a small experiment to study the interaction of batch normalization and batch size.

The first cell will train 6-layer networks both with and without batch normalization using different batch sizes. The second layer will plot training accuracy and validation set accuracy over time.

```
In [ ]: def run batchsize experiments(normalization mode):
            np.random.seed(231)
            # Try training a very deep net with batchnorm.
            hidden dims = [100, 100, 100, 100, 100]
            num train = 1000
            small data = {
              'X train': data['X train'][:num train],
              'y train': data['y train'][:num train],
              'X val': data['X val'],
              'y val': data['y val'],
            n epochs=10
            weight scale = 2e-2
            batch sizes = [5,10,50]
            lr = 10**(-3.5)
            solver_bsize = batch sizes[0]
            print('No normalization: batch size = ',solver bsize)
            model = FullyConnectedNet(hidden dims, weight scale=weight scale, normal
            solver = Solver(model, small data,
                            num epochs=n epochs, batch size=solver bsize,
                             update rule='adam',
                             optim config={
                               'learning rate': lr,
                            verbose=False)
            solver.train()
            bn solvers = []
            for i in range(len(batch sizes)):
                b size=batch sizes[i]
                print('Normalization: batch size = ',b size)
                bn_model = FullyConnectedNet(hidden_dims, weight scale=weight scale,
                bn solver = Solver(bn model, small data,
                                 num epochs=n epochs, batch size=b size,
                                 update rule='adam',
                                 optim config={
                                   'learning rate': lr,
                                 },
                                 verbose=False)
```

Inline Question 2:

Describe the results of this experiment. What does this imply about the relationship between batch normalization and batch size? Why is this relationship observed?

Answer:

[FILL THIS IN]

Layer Normalization

Batch normalization has proved to be effective in making networks easier to train, but the dependency on batch size makes it less useful in complex networks which have a cap on the input batch size due to hardware limitations.

Several alternatives to batch normalization have been proposed to mitigate this problem; one such technique is Layer Normalization [2]. Instead of normalizing over the batch, we normalize over the features. In other words, when using Layer Normalization, each feature vector corresponding to a single datapoint is normalized based on the sum of all terms within that feature vector.

[2] Ba, Jimmy Lei, Jamie Ryan Kiros, and Geoffrey E. Hinton. "Layer Normalization." stat 1050 (2016): 21.

Inline Question 3:

Which of these data preprocessing steps is analogous to batch normalization, and which is analogous to layer normalization?

- 1. Scaling each image in the dataset, so that the RGB channels for each row of pixels within an image sums up to 1.
- 2. Scaling each image in the dataset, so that the RGB channels for all pixels within an image sums up to 1.
- 3. Subtracting the mean image of the dataset from each image in the dataset.
- 4. Setting all RGB values to either 0 or 1 depending on a given threshold.

Answer:

[FILL THIS IN]

Layer Normalization: Implementation

Now you'll implement layer normalization. This step should be relatively straightforward, as conceptually the implementation is almost identical to that of batch normalization. One significant difference though is that for layer normalization, we do not keep track of the moving moments, and the testing phase is identical to the training phase, where the mean and variance are directly calculated per datapoint.

Here's what you need to do:

• In cs231n/layers.py , implement the forward pass for layer normalization in the function layernorm forward.

Run the cell below to check your results.

 In cs231n/layers.py , implement the backward pass for layer normalization in the function layernorm backward.

Run the second cell below to check your results.

• Modify cs231n/classifiers/fc net.py to add layer normalization to the FullyConnectedNet . When the normalization flag is set to "layernorm" in the constructor, you should insert a layer normalization layer before each ReLU nonlinearity.

Run the third cell below to run the batch size experiment on layer normalization.

```
# Simulate the forward pass for a two-layer network.
        np.random.seed(231)
        N, D1, D2, D3 = 4, 50, 60, 3
        X = np.random.randn(N, D1)
        W1 = np.random.randn(D1, D2)
        W2 = np.random.randn(D2, D3)
        a = np.maximum(0, X.dot(W1)).dot(W2)
        print('Before layer normalization:')
        print mean std(a,axis=1)
        qamma = np.ones(D3)
        beta = np.zeros(D3)
        # Means should be close to zero and stds close to one.
        print('After layer normalization (gamma=1, beta=0)')
        a norm, = layernorm forward(a, gamma, beta, {'mode': 'train'})
        print mean std(a norm,axis=1)
        gamma = np.asarray([3.0,3.0,3.0])
        beta = np.asarray([5.0,5.0,5.0])
        # Now means should be close to beta and stds close to gamma.
        print('After layer normalization (gamma=', gamma, ', beta=', beta, ')')
        a norm, = layernorm forward(a, gamma, beta, {'mode': 'train'})
        print mean std(a norm,axis=1)
In [ ]: # Gradient check batchnorm backward pass.
        np.random.seed(231)
        N, D = 4, 5
        x = 5 * np.random.randn(N, D) + 12
        gamma = np.random.randn(D)
        beta = np.random.randn(D)
        dout = np.random.randn(N, D)
        ln param = {}
        fx = lambda x: layernorm forward(x, gamma, beta, ln param)[0]
        fg = lambda a: layernorm forward(x, a, beta, ln param)[0]
        fb = lambda b: layernorm forward(x, gamma, b, ln param)[0]
        dx num = eval numerical gradient array(fx, x, dout)
        da num = eval numerical gradient array(fg, gamma.copy(), dout)
        db num = eval numerical gradient array(fb, beta.copy(), dout)
        , cache = layernorm forward(x, gamma, beta, ln param)
        dx, dgamma, dbeta = layernorm backward(dout, cache)
        # You should expect to see relative errors between 1e-12 and 1e-8.
        print('dx error: ', rel_error(dx_num, dx))
        print('dgamma error: ', rel_error(da_num, dgamma))
        print('dbeta error: ', rel error(db num, dbeta))
```

Layer Normalization and Batch Size

We will now run the previous batch size experiment with layer normalization instead of batch normalization. Compared to the previous experiment, you should see a markedly smaller influence of batch size on the training history!

Inline Question 4:

When is layer normalization likely to not work well, and why?

- 1. Using it in a very deep network
- 2. Having a very small dimension of features
- 3. Having a high regularization term

Answer:

[FILL THIS IN]

This notebook was converted to PDF with convert.ploomber.io

```
In [ ]: # This mounts your Google Drive to the Colab VM.
        from google.colab import drive
        drive.mount('/content/drive')
        # TODO: Enter the foldername in your Drive where you have saved the unzipped
        # assignment folder, e.g. 'cs231n/assignments/assignment1/'
        FOLDERNAME = None
        assert FOLDERNAME is not None, "[!] Enter the foldername."
        # Now that we've mounted your Drive, this ensures that
        # the Python interpreter of the Colab VM can load
        # python files from within it.
        import sys
        sys.path.append('/content/drive/My Drive/{}'.format(FOLDERNAME))
        # This downloads the CIFAR-10 dataset to your Drive
        # if it doesn't already exist.
        %cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/datasets/
        !bash get datasets.sh
        %cd /content/drive/My\ Drive/$F0LDERNAME
```

Dropout

Dropout [1] is a technique for regularizing neural networks by randomly setting some output activations to zero during the forward pass. In this exercise, you will implement a dropout layer and modify your fully connected network to optionally use dropout.

[1] Geoffrey E. Hinton et al, "Improving neural networks by preventing coadaptation of feature detectors", arXiv 2012

```
In []: # Setup cell.
import time
import numpy as np
import matplotlib.pyplot as plt
from cs231n.classifiers.fc_net import *
from cs231n.data_utils import get_CIFAR10_data
from cs231n.gradient_check import eval_numerical_gradient, eval_numerical_gr
from cs231n.solver import Solver

%matplotlib inline
plt.rcParams["figure.figsize"] = (10.0, 8.0) # Set default size of plots.
plt.rcParams["image.interpolation"] = "nearest"
plt.rcParams["image.cmap"] = "gray"

%load_ext autoreload
%autoreload 2

def rel_error(x, y):
```

```
"""Returns relative error."""
    return np.max(np.abs(x - y) / (np.maximum(le-8, np.abs(x) + np.abs(y))))
In []: # Load the (preprocessed) CIFAR-10 data.
data = get_CIFAR10_data()
for k, v in list(data.items()):
    print(f"{k}: {v.shape}")
```

Dropout: Forward Pass

In the file cs231n/layers.py , implement the forward pass for dropout. Since dropout behaves differently during training and testing, make sure to implement the operation for both modes.

Once you have done so, run the cell below to test your implementation.

Dropout: Backward Pass

In the file cs231n/layers.py , implement the backward pass for dropout. After doing so, run the following cell to numerically gradient-check your implementation.

```
In []: np.random.seed(231)
    x = np.random.randn(10, 10) + 10
    dout = np.random.randn(*x.shape)

dropout_param = {'mode': 'train', 'p': 0.2, 'seed': 123}
    out, cache = dropout_forward(x, dropout_param)
    dx = dropout_backward(dout, cache)
    dx_num = eval_numerical_gradient_array(lambda xx: dropout_forward(xx, dropout_start))

# Error should be around e-10 or less.
    print('dx relative error: ', rel_error(dx, dx_num))
```

Inline Question 1:

What happens if we do not divide the values being passed through inverse dropout by p in the dropout layer? Why does that happen?

Answer:

[FILL THIS IN]

Fully Connected Networks with Dropout

In the file cs231n/classifiers/fc_net.py , modify your implementation to use dropout. Specifically, if the constructor of the network receives a value that is not 1 for the dropout_keep_ratio parameter, then the net should add a dropout layer immediately after every ReLU nonlinearity. After doing so, run the following to numerically gradient-check your implementation.

```
In []: np.random.seed(231)
        N, D, H1, H2, C = 2, 15, 20, 30, 10
        X = np.random.randn(N, D)
        y = np.random.randint(C, size=(N,))
        for dropout keep ratio in [1, 0.75, 0.5]:
            print('Running check with dropout = ', dropout keep ratio)
            model = FullyConnectedNet(
                [H1, H2],
                input dim=D,
                num classes=C,
                weight scale=5e-2,
                dtype=np.float64,
                dropout keep ratio=dropout keep ratio,
                seed=123
            )
            loss, grads = model.loss(X, y)
            print('Initial loss: ', loss)
            # Relative errors should be around e-6 or less.
            # Note that it's fine if for dropout keep ratio=1 you have W2 error be d
            for name in sorted(grads):
                f = lambda : model.loss(X, y)[0]
                grad num = eval numerical gradient(f, model.params[name], verbose=Fa
                print('%s relative error: %.2e' % (name, rel error(grad num, grads[r
            print()
```

Regularization Experiment

As an experiment, we will train a pair of two-layer networks on 500 training examples: one will use no dropout, and one will use a keep probability of 0.25. We will then visualize the training and validation accuracies of the two networks over time.

```
In [ ]: # Train two identical nets, one with dropout and one without.
        np.random.seed(231)
        num train = 500
        small data = {
            'X train': data['X train'][:num train],
            'y train': data['y train'][:num train],
            'X val': data['X val'],
            'y val': data['y val'],
        solvers = {}
        dropout choices = [1, 0.25]
        for dropout keep ratio in dropout choices:
            model = FullyConnectedNet(
                [500],
                dropout keep ratio=dropout keep ratio
            print(dropout keep ratio)
            solver = Solver(
                model,
                small data,
                num epochs=25,
                batch size=100,
                update rule='adam',
                optim config={'learning rate': 5e-4,},
                verbose=True,
                print every=100
            solver.train()
            solvers[dropout keep ratio] = solver
            print()
```

```
In []: # Plot train and validation accuracies of the two models.
    train_accs = []
    val_accs = []
    for dropout_keep_ratio in dropout_choices:
        solver = solvers[dropout_keep_ratio]
        train_accs.append(solver.train_acc_history[-1])
        val_accs.append(solver.val_acc_history[-1])

plt.subplot(3, 1, 1)
    for dropout_keep_ratio in dropout_choices:
        plt.plot(
            solvers[dropout_keep_ratio].train_acc_history, 'o', label='%.2f dropout_keep_ratio].train_acc_history, 'o', label='%.2f dropout
```

```
plt.title('Train accuracy')
plt.xlabel('Epoch')
plt.ylabel('Accuracy')
plt.legend(ncol=2, loc='lower right')

plt.subplot(3, 1, 2)
for dropout_keep_ratio in dropout_choices:
    plt.plot(
        solvers[dropout_keep_ratio].val_acc_history, 'o', label='%.2f dropout_title('Val accuracy')
plt.xlabel('Epoch')
plt.ylabel('Accuracy')
plt.legend(ncol=2, loc='lower right')

plt.gcf().set_size_inches(15, 15)
plt.show()
```

Inline Question 2:

Compare the validation and training accuracies with and without dropout -- what do your results suggest about dropout as a regularizer?

Answer:

[FILL THIS IN]

Inline Question 3:

Suppose we are training a deep fully connected network for image classification, with dropout after hidden layers (parameterized by keep probability p). If we are concerned about overfitting, how should we modify p (if at all) when we decide to decrease the size of the hidden layers (that is, the number of nodes in each layer)?

Answer:

[FILL THIS IN]

In []:

```
In [ ]: # This mounts your Google Drive to the Colab VM.
        from google.colab import drive
        drive.mount('/content/drive')
        # TODO: Enter the foldername in your Drive where you have saved the unzipped
        # assignment folder, e.g. 'cs231n/assignments/assignment1/'
        FOLDERNAME = None
        assert FOLDERNAME is not None, "[!] Enter the foldername."
        # Now that we've mounted your Drive, this ensures that
        # the Python interpreter of the Colab VM can load
        # python files from within it.
        import sys
        sys.path.append('/content/drive/My Drive/{}'.format(FOLDERNAME))
        # This downloads the CIFAR-10 dataset to your Drive
        # if it doesn't already exist.
        %cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/datasets/
        !bash get datasets.sh
        %cd /content/drive/My\ Drive/$FOLDERNAME
```

Convolutional Networks

So far we have worked with deep fully connected networks, using them to explore different optimization strategies and network architectures. Fully connected networks are a good testbed for experimentation because they are very computationally efficient, but in practice all state-of-the-art results use convolutional networks instead.

First you will implement several layer types that are used in convolutional networks. You will then use these layers to train a convolutional network on the CIFAR-10 dataset.

```
In []: # Setup cell.
    import numpy as np
    import matplotlib.pyplot as plt
    from cs23ln.classifiers.cnn import *
    from cs23ln.data_utils import get_CIFAR10_data
    from cs23ln.gradient_check import eval_numerical_gradient_array, eval_numeri
    from cs23ln.layers import *
    from cs23ln.fast_layers import *
    from cs23ln.solver import Solver

%matplotlib inline
    plt.rcParams['figure.figsize'] = (10.0, 8.0) # set default size of plots
    plt.rcParams['image.interpolation'] = 'nearest'
    plt.rcParams['image.cmap'] = 'gray'

# for auto-reloading external modules
# see http://stackoverflow.com/questions/1907993/autoreload-of-modules-in-ip
```

```
%load_ext autoreload
%autoreload 2

def rel_error(x, y):
    """ returns relative error """
    return np.max(np.abs(x - y) / (np.maximum(1e-8, np.abs(x) + np.abs(y))))

In []: # Load the (preprocessed) CIFAR-10 data.
data = get_CIFAR10_data()
for k, v in list(data.items()):
    print(f"{k}: {v.shape}")
```

Convolution: Naive Forward Pass

The core of a convolutional network is the convolution operation. In the file cs231n/layers.py , implement the forward pass for the convolution layer in the function conv forward naive .

You don't have to worry too much about efficiency at this point; just write the code in whatever way you find most clear.

You can test your implementation by running the following:

```
In []: x \text{ shape} = (2, 3, 4, 4)
        w \text{ shape} = (3, 3, 4, 4)
        x = np.linspace(-0.1, 0.5, num=np.prod(x shape)).reshape(x shape)
        w = np.linspace(-0.2, 0.3, num=np.prod(w shape)).reshape(w shape)
        b = np.linspace(-0.1, 0.2, num=3)
        conv param = {'stride': 2, 'pad': 1}
        out, = conv forward naive(x, w, b, conv param)
        correct out = np.array([[[-0.08759809, -0.10987781],
                                   [-0.18387192, -0.2109216]],
                                  [[ 0.21027089, 0.21661097],
                                   [ 0.22847626, 0.23004637]],
                                   [[ 0.50813986, 0.54309974],
                                   [ 0.64082444, 0.67101435]]],
                                  [[[-0.98053589, -1.03143541],
                                   [-1.19128892, -1.24695841]],
                                  [[ 0.69108355, 0.66880383],
                                   [ 0.59480972, 0.56776003]],
                                   [[ 2.36270298, 2.36904306],
                                   [ 2.38090835, 2.38247847]]])
        # Compare your output to ours; difference should be around e-8
        print('Testing conv forward naive')
        print('difference: ', rel error(out, correct out))
```

Aside: Image Processing via Convolutions

As fun way to both check your implementation and gain a better understanding of the type of operation that convolutional layers can perform, we will set up an input containing two images and manually set up filters that perform common image processing operations (grayscale conversion and edge detection). The convolution forward pass will apply these operations to each of the input images. We can then visualize the results as a sanity check.

```
In [ ]: from imageio import imread
        from PIL import Image
        kitten = imread('cs231n/notebook images/kitten.jpg')
        puppy = imread('cs231n/notebook images/puppy.jpg')
        # kitten is wide, and puppy is already square
        d = kitten.shape[1] - kitten.shape[0]
        kitten cropped = kitten[:, d//2:-d//2, :]
        img size = 200  # Make this smaller if it runs too slow
        resized puppy = np.array(Image.fromarray(puppy).resize((img size, img size))
        resized kitten = np.array(Image.fromarray(kitten cropped).resize((img size,
        x = np.zeros((2, 3, img size, img size))
        x[0, :, :, :] = resized puppy.transpose((2, 0, 1))
        x[1, :, :, :] = resized kitten.transpose((2, 0, 1))
        # Set up a convolutional weights holding 2 filters, each 3x3
        w = np.zeros((2, 3, 3, 3))
        # The first filter converts the image to grayscale.
        # Set up the red, green, and blue channels of the filter.
        w[0, 0, :, :] = [[0, 0, 0], [0, 0.3, 0], [0, 0, 0]]
        w[0, 1, :, :] = [[0, 0, 0], [0, 0.6, 0], [0, 0, 0]]
        w[0, 2, :, :] = [[0, 0, 0], [0, 0.1, 0], [0, 0, 0]]
        # Second filter detects horizontal edges in the blue channel.
        w[1, 2, :, :] = [[1, 2, 1], [0, 0, 0], [-1, -2, -1]]
        # Vector of biases. We don't need any bias for the grayscale
        # filter, but for the edge detection filter we want to add 128
        # to each output so that nothing is negative.
        b = np.array([0, 128])
        # Compute the result of convolving each input in x with each filter in w,
        # offsetting by b, and storing the results in out.
        out, = conv forward naive(x, w, b, {'stride': 1, 'pad': 1})
        def imshow no ax(img, normalize=True):
            """ Tiny helper to show images as uint8 and remove axis labels """
            if normalize:
                img max, img min = np.max(img), np.min(img)
                img = 255.0 * (img - img min) / (img max - img min)
            plt.imshow(img.astype('uint8'))
            plt.gca().axis('off')
        # Show the original images and the results of the conv operation
        plt.subplot(2, 3, 1)
```

```
imshow_no_ax(puppy, normalize=False)
plt.title('Original image')
plt.subplot(2, 3, 2)
imshow_no_ax(out[0, 0])
plt.title('Grayscale')
plt.subplot(2, 3, 3)
imshow_no_ax(out[0, 1])
plt.title('Edges')
plt.subplot(2, 3, 4)
imshow_no_ax(kitten_cropped, normalize=False)
plt.subplot(2, 3, 5)
imshow_no_ax(out[1, 0])
plt.subplot(2, 3, 6)
imshow_no_ax(out[1, 1])
plt.show()
```

Convolution: Naive Backward Pass

Implement the backward pass for the convolution operation in the function conv_backward_naive in the file cs231n/layers.py . Again, you don't need to worry too much about computational efficiency.

When you are done, run the following to check your backward pass with a numeric gradient check.

```
In [ ]: np.random.seed(231)
        x = np.random.randn(4, 3, 5, 5)
        w = np.random.randn(2, 3, 3, 3)
        b = np.random.randn(2,)
        dout = np.random.randn(4, 2, 5, 5)
        conv param = {'stride': 1, 'pad': 1}
        dx num = eval numerical gradient array(lambda x: conv forward naive(x, w, b,
        dw num = eval numerical gradient array(lambda w: conv forward naive(x, w, b,
        db_num = eval_numerical_gradient_array(lambda b: conv_forward_naive(x, w, b,
        out, cache = conv forward naive(x, w, b, conv param)
        dx, dw, db = conv backward naive(dout, cache)
        # Your errors should be around e-8 or less.
        print('Testing conv backward naive function')
        print('dx error: ', rel_error(dx, dx_num))
        print('dw error: ', rel_error(dw, dw_num))
        print('db error: ', rel error(db, db num))
```

Max-Pooling: Naive Forward Pass

Implement the forward pass for the max-pooling operation in the function max_pool_forward_naive in the file cs231n/layers.py . Again, don't worry

too much about computational efficiency.

Check your implementation by running the following:

```
In []: x \text{ shape} = (2, 3, 4, 4)
        x = np.linspace(-0.3, 0.4, num=np.prod(x shape)).reshape(x shape)
        pool param = {'pool width': 2, 'pool height': 2, 'stride': 2}
        out, = max pool forward naive(x, pool param)
        correct out = np.array([[[[-0.26315789, -0.24842105],
                                  [-0.20421053, -0.18947368]],
                                 [[-0.14526316, -0.13052632],
                                 [-0.08631579, -0.07157895]],
                                 [[-0.02736842, -0.01263158],
                                 [ 0.03157895, 0.04631579]]],
                                [[[ 0.09052632, 0.10526316],
                                 [ 0.14947368, 0.16421053]],
                                 [[ 0.20842105, 0.22315789],
                                 [ 0.26736842, 0.28210526]],
                                 [[ 0.32631579, 0.34105263],
                                  [ 0.38526316, 0.4
                                                           ]]]])
        # Compare your output with ours. Difference should be on the order of e-8.
        print('Testing max pool forward naive function:')
        print('difference: ', rel_error(out, correct_out))
```

Max-Pooling: Naive Backward

Implement the backward pass for the max-pooling operation in the function max_pool_backward_naive in the file cs231n/layers.py . You don't need to worry about computational efficiency.

Check your implementation with numeric gradient checking by running the following:

```
In []: np.random.seed(231)
    x = np.random.randn(3, 2, 8, 8)
    dout = np.random.randn(3, 2, 4, 4)
    pool_param = {'pool_height': 2, 'pool_width': 2, 'stride': 2}

    dx_num = eval_numerical_gradient_array(lambda x: max_pool_forward_naive(x, pout, cache = max_pool_forward_naive(x, pool_param)
    dx = max_pool_backward_naive(dout, cache)

# Your error should be on the order of e-12
    print('Testing max_pool_backward_naive function:')
    print('dx error: ', rel_error(dx, dx_num))
```

Fast Layers

Making convolution and pooling layers fast can be challenging. To spare you the pain, we've provided fast implementations of the forward and backward passes for convolution and pooling layers in the file cs231n/fast layers.py.

Execute the below cell, save the notebook, and restart the runtime

The fast convolution implementation depends on a Cython extension; to compile it, run the cell below. Next, save the Colab notebook (File > Save) and restart the runtime (Runtime > Restart runtime). You can then re-execute the preceding cells from top to bottom and skip the cell below as you only need to run it once for the compilation step.

```
In [ ]: # Remember to restart the runtime after executing this cell!
%cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/
!python setup.py build_ext --inplace
%cd /content/drive/My\ Drive/$FOLDERNAME/
```

The API for the fast versions of the convolution and pooling layers is exactly the same as the naive versions that you implemented above: the forward pass receives data, weights, and parameters and produces outputs and a cache object; the backward pass recieves upstream derivatives and the cache object and produces gradients with respect to the data and weights.

Note: The fast implementation for pooling will only perform optimally if the pooling regions are non-overlapping and tile the input. If these conditions are not met then the fast pooling implementation will not be much faster than the naive implementation.

You can compare the performance of the naive and fast versions of these layers by running the following:

```
In []: # Rel errors should be around e-9 or less.
    from cs231n.fast_layers import conv_forward_fast, conv_backward_fast
    from time import time
    np.random.seed(231)
    x = np.random.randn(100, 3, 31, 31)
    w = np.random.randn(25, 3, 3, 3)
    b = np.random.randn(25,)
    dout = np.random.randn(100, 25, 16, 16)
    conv_param = {'stride': 2, 'pad': 1}

    t0 = time()
    out_naive, cache_naive = conv_forward_naive(x, w, b, conv_param)
    t1 = time()
```

```
t2 = time()
        print('Testing conv forward fast:')
        print('Naive: %fs' % (t1 - t0))
        print('Fast: %fs' % (t2 - t1))
        print('Speedup: %fx' % ((t1 - t0) / (t2 - t1)))
        print('Difference: ', rel_error(out_naive, out_fast))
        t0 = time()
        dx naive, dw naive, db naive = conv backward naive(dout, cache naive)
        t1 = time()
        dx fast, dw fast, db fast = conv backward fast(dout, cache fast)
        t2 = time()
        print('\nTesting conv backward fast:')
        print('Naive: %fs' % (t1 - t0))
        print('Fast: %fs' % (t2 - t1))
        print('Speedup: %fx' % ((t1 - t0) / (t2 - t1)))
        print('dx difference: ', rel_error(dx_naive, dx_fast))
        print('dw difference: ', rel_error(dw_naive, dw_fast))
        print('db difference: ', rel_error(db_naive, db_fast))
In []: # Relative errors should be close to 0.0.
        from cs231n.fast layers import max pool forward fast, max pool backward fast
        np.random.seed(231)
        x = np.random.randn(100, 3, 32, 32)
        dout = np.random.randn(100, 3, 16, 16)
        pool param = {'pool height': 2, 'pool width': 2, 'stride': 2}
        t0 = time()
        out naive, cache naive = max pool forward naive(x, pool param)
        t1 = time()
        out fast, cache fast = \max pool forward fast(x, pool param)
        t2 = time()
        print('Testing pool forward fast:')
        print('Naive: %fs' % (t1 - t0))
        print('fast: %fs' % (t2 - t1))
        print('speedup: %fx' % ((t1 - t0) / (t2 - t1)))
        print('difference: ', rel error(out naive, out fast))
        t0 = time()
        dx naive = max pool backward naive(dout, cache naive)
        t1 = time()
        dx fast = max pool backward fast(dout, cache fast)
        t2 = time()
        print('\nTesting pool backward fast:')
        print('Naive: %fs' % (t1 - t0))
        print('fast: %fs' % (t2 - t1))
        print('speedup: %fx' % ((t1 - t0) / (t2 - t1)))
        print('dx difference: ', rel error(dx naive, dx fast))
```

out fast, cache fast = conv forward fast(x, w, b, conv param)

Convolutional "Sandwich" Layers

In the previous assignment, we introduced the concept of "sandwich" layers that combine multiple operations into commonly used patterns. In the file cs231n/layer_utils.py you will find sandwich layers that implement a few commonly used patterns for convolutional networks. Run the cells below to sanity check their usage.

```
In [ ]: from cs231n.layer utils import conv relu pool forward, conv relu pool backwa
        np.random.seed(231)
        x = np.random.randn(2, 3, 16, 16)
        w = np.random.randn(3, 3, 3, 3)
         b = np.random.randn(3,)
         dout = np.random.randn(2, 3, 8, 8)
         conv param = {'stride': 1, 'pad': 1}
         pool param = {'pool height': 2, 'pool width': 2, 'stride': 2}
         out, cache = conv relu pool forward(x, w, b, conv param, pool param)
         dx, dw, db = conv relu pool backward(dout, cache)
         dx num = eval numerical gradient array(lambda x: conv relu pool forward(x, w
         dw num = eval numerical gradient array(lambda w: conv relu pool forward(x, w
         db num = eval numerical gradient array(lambda b: conv relu pool forward(x, w
         # Relative errors should be around e-8 or less
         print('Testing conv relu pool')
        print('dx error: ', rel_error(dx_num, dx))
        print('dw error: ', rel_error(dw_num, dw))
print('db error: ', rel_error(db_num, db))
In [ ]: from cs231n.layer utils import conv relu forward, conv relu backward
        np.random.seed(231)
        x = np.random.randn(2, 3, 8, 8)
        w = np.random.randn(3, 3, 3, 3)
         b = np.random.randn(3,)
         dout = np.random.randn(2, 3, 8, 8)
         conv param = {'stride': 1, 'pad': 1}
         out, cache = conv relu forward(x, w, b, conv param)
         dx, dw, db = conv relu backward(dout, cache)
         dx num = eval numerical gradient array(lambda x: conv relu forward(x, w, b,
         dw num = eval numerical gradient array(lambda w: conv relu forward(x, w, b,
         db num = eval numerical gradient array(lambda b: conv relu forward(x, w, b,
         # Relative errors should be around e-8 or less
         print('Testing conv relu:')
         print('dx error: ', rel_error(dx_num, dx))
        print('dw error: ', rel_error(dw_num, dw))
print('db error: ', rel_error(db_num, db))
```

Three-Layer Convolutional Network

Now that you have implemented all the necessary layers, we can put them together into a simple convolutional network.

Open the file cs231n/classifiers/cnn.py and complete the implementation of the ThreeLayerConvNet class. Remember you can use the fast/sandwich layers (already imported for you) in your implementation. Run the following cells to help you debug:

Sanity Check Loss

After you build a new network, one of the first things you should do is sanity check the loss. When we use the softmax loss, we expect the loss for random weights (and no regularization) to be about log(C) for C classes. When we add regularization the loss should go up slightly.

```
In []: model = ThreeLayerConvNet()

N = 50
X = np.random.randn(N, 3, 32, 32)
y = np.random.randint(10, size=N)

loss, grads = model.loss(X, y)
print('Initial loss (no regularization): ', loss)

model.reg = 0.5
loss, grads = model.loss(X, y)
print('Initial loss (with regularization): ', loss)
```

Gradient Check

After the loss looks reasonable, use numeric gradient checking to make sure that your backward pass is correct. When you use numeric gradient checking you should use a small amount of artifical data and a small number of neurons at each layer. Note: correct implementations may still have relative errors up to the order of e-2.

```
In []: num_inputs = 2
   input_dim = (3, 16, 16)
   reg = 0.0
   num_classes = 10
   np.random.seed(231)
   X = np.random.randn(num_inputs, *input_dim)
   y = np.random.randint(num_classes, size=num_inputs)
```

```
model = ThreeLayerConvNet(
    num_filters=3,
    filter_size=3,
    input_dim=input_dim,
    hidden_dim=7,
    dtype=np.float64
)
loss, grads = model.loss(X, y)
# Errors should be small, but correct implementations may have
# relative errors up to the order of e-2
for param_name in sorted(grads):
    f = lambda _: model.loss(X, y)[0]
    param_grad_num = eval_numerical_gradient(f, model.params[param_name], vece = rel_error(param_grad_num, grads[param_name])
    print('%s max relative error: %e' % (param_name, rel_error(param_grad_num))
```

Overfit Small Data

A nice trick is to train your model with just a few training samples. You should be able to overfit small datasets, which will result in very high training accuracy and comparatively low validation accuracy.

```
In [ ]: np.random.seed(231)
        num train = 100
        small data = {
          'X train': data['X train'][:num train],
          'y train': data['y train'][:num train],
          'X val': data['X val'],
          'y_val': data['y val'],
        model = ThreeLayerConvNet(weight scale=1e-2)
        solver = Solver(
            model,
            small data,
            num epochs=15,
            batch size=50,
            update_rule='adam',
            optim config={'learning rate': 1e-3,},
            verbose=True,
            print every=1
        solver.train()
```

Plotting the loss, training accuracy, and validation accuracy should show clear overfitting:

```
In []: plt.subplot(2, 1, 1)
    plt.plot(solver.loss_history, 'o')
    plt.xlabel('iteration')
    plt.ylabel('loss')

plt.subplot(2, 1, 2)
    plt.plot(solver.train_acc_history, '-o')
    plt.plot(solver.val_acc_history, '-o')
    plt.legend(['train', 'val'], loc='upper left')
    plt.xlabel('epoch')
    plt.ylabel('accuracy')
    plt.show()
```

Train the Network

By training the three-layer convolutional network for one epoch, you should achieve greater than 40% accuracy on the training set:

```
In [ ]: model = ThreeLayerConvNet(weight scale=0.001, hidden dim=500, reg=0.001)
        solver = Solver(
            model.
            data,
            num epochs=1,
            batch size=50,
            update rule='adam',
            optim_config={'learning_rate': 1e-3,},
            verbose=True,
            print every=20
        solver.train()
In [ ]: # Print final training accuracy.
        print(
            "Full data training accuracy:",
            solver.check_accuracy(data['X_train'], data['y_train'])
In [ ]: # Print final validation accuracy.
        print(
            "Full data validation accuracy:",
```

```
solver.check_accuracy(data['X_val'], data['y_val'])
)
```

Visualize Filters

You can visualize the first-layer convolutional filters from the trained network by running the following:

```
In []: from cs231n.vis_utils import visualize_grid

    grid = visualize_grid(model.params['W1'].transpose(0, 2, 3, 1))
    plt.imshow(grid.astype('uint8'))
    plt.axis('off')
    plt.gcf().set_size_inches(5, 5)
    plt.show()
```

Spatial Batch Normalization

We already saw that batch normalization is a very useful technique for training deep fully connected networks. As proposed in the original paper (link in BatchNormalization.ipynb), batch normalization can also be used for convolutional networks, but we need to tweak it a bit; the modification will be called "spatial batch normalization."

Normally, batch-normalization accepts inputs of shape (N, D) and produces outputs of shape (N, D), where we normalize across the minibatch dimension N. For data coming from convolutional layers, batch normalization needs to accept inputs of shape (N, C, H, W) and produce outputs of shape (N, C, H, W) where the N dimension gives the minibatch size and the (H, W) dimensions give the spatial size of the feature map.

If the feature map was produced using convolutions, then we expect every feature channel's statistics e.g. mean, variance to be relatively consistent both between different images, and different locations within the same image -- after all, every feature channel is produced by the same convolutional filter! Therefore, spatial batch normalization computes a mean and variance for each of the ${\tt C}$ feature channels by computing statistics over the minibatch dimension ${\tt N}$ as well the spatial dimensions ${\tt H}$ and ${\tt W}$.

[1] Sergey Ioffe and Christian Szegedy, "Batch Normalization: Accelerating Deep Network Training by Reducing Internal Covariate Shift", ICML 2015.

Spatial Batch Normalization: Forward Pass

In the file cs231n/layers.py , implement the forward pass for spatial batch normalization in the function spatial_batchnorm_forward . Check your implementation by running the following:

```
In [ ]: np.random.seed(231)
        # Check the training-time forward pass by checking means and variances
        # of features both before and after spatial batch normalization.
        N, C, H, W = 2, 3, 4, 5
        x = 4 * np.random.randn(N, C, H, W) + 10
        print('Before spatial batch normalization:')
        print(' shape: ', x.shape)
        print(' means: ', x.mean(axis=(0, 2, 3)))
        print(' stds: ', x.std(axis=(0, 2, 3)))
        # Means should be close to zero and stds close to one
        gamma, beta = np.ones(C), np.zeros(C)
        bn param = {'mode': 'train'}
        out, = spatial batchnorm forward(x, gamma, beta, bn param)
        print('After spatial batch normalization:')
        print(' shape: ', out.shape)
        print(' means: ', out.mean(axis=(0, 2, 3)))
        print(' stds: ', out.std(axis=(0, 2, 3)))
        # Means should be close to beta and stds close to gamma
        gamma, beta = np.asarray([3, 4, 5]), np.asarray([6, 7, 8])
        out, = spatial batchnorm forward(x, gamma, beta, bn param)
        print('After spatial batch normalization (nontrivial gamma, beta):')
        print(' shape: ', out.shape)
        print(' means: ', out.mean(axis=(0, 2, 3)))
        print(' stds: ', out.std(axis=(0, 2, 3)))
In [ ]: np.random.seed(231)
        # Check the test-time forward pass by running the training-time
        # forward pass many times to warm up the running averages, and then
        # checking the means and variances of activations after a test-time
        # forward pass.
        N, C, H, W = 10, 4, 11, 12
        bn param = {'mode': 'train'}
        gamma = np.ones(C)
        beta = np.zeros(C)
        for t in range (50):
          x = 2.3 * np.random.randn(N, C, H, W) + 13
          spatial batchnorm forward(x, gamma, beta, bn param)
        bn param['mode'] = 'test'
```

```
x = 2.3 * np.random.randn(N, C, H, W) + 13
a_norm, _ = spatial_batchnorm_forward(x, gamma, beta, bn_param)

# Means should be close to zero and stds close to one, but will be
# noisier than training-time forward passes.
print('After spatial batch normalization (test-time):')
print(' means: ', a_norm.mean(axis=(0, 2, 3)))
print(' stds: ', a_norm.std(axis=(0, 2, 3)))
```

Spatial Batch Normalization: Backward Pass

In the file cs231n/layers.py , implement the backward pass for spatial batch normalization in the function spatial_batchnorm_backward . Run the following to check your implementation using a numeric gradient check:

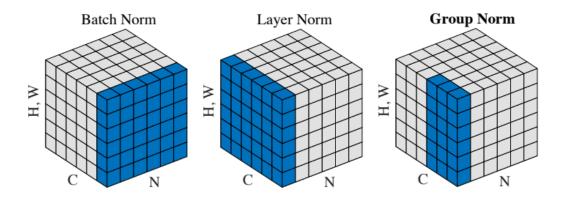
```
In [ ]: np.random.seed(231)
        N, C, H, W = 2, 3, 4, 5
        x = 5 * np.random.randn(N, C, H, W) + 12
        gamma = np.random.randn(C)
        beta = np.random.randn(C)
        dout = np.random.randn(N, C, H, W)
        bn param = {'mode': 'train'}
        fx = lambda x: spatial batchnorm forward(x, gamma, beta, bn param)[0]
        fg = lambda a: spatial batchnorm forward(x, gamma, beta, bn param)[0]
        fb = lambda b: spatial batchnorm forward(x, gamma, beta, bn param)[0]
        dx num = eval numerical gradient array(fx, x, dout)
        da num = eval numerical gradient array(fg, gamma, dout)
        db num = eval numerical gradient array(fb, beta, dout)
        #You should expect errors of magnitudes between 1e-12~1e-06
        , cache = spatial batchnorm forward(x, gamma, beta, bn param)
        dx, dgamma, dbeta = spatial_batchnorm_backward(dout, cache)
        print('dx error: ', rel error(dx num, dx))
        print('dgamma error: ', rel_error(da_num, dgamma))
        print('dbeta error: ', rel error(db num, dbeta))
```

Spatial Group Normalization

In the previous notebook, we mentioned that Layer Normalization is an alternative normalization technique that mitigates the batch size limitations of Batch Normalization. However, as the authors of [2] observed, Layer Normalization does not perform as well as Batch Normalization when used with Convolutional Layers:

With fully connected layers, all the hidden units in a layer tend to make similar contributions to the final prediction, and re-centering and rescaling the summed inputs to a layer works well. However, the assumption of similar contributions is no longer true for convolutional neural networks. The large number of the hidden units whose receptive fields lie near the boundary of the image are rarely turned on and thus have very different statistics from the rest of the hidden units within the same layer.

The authors of [3] propose an intermediary technique. In contrast to Layer Normalization, where you normalize over the entire feature per-datapoint, they suggest a consistent splitting of each per-datapoint feature into G groups and a per-group per-datapoint normalization instead.



Visual comparison of the normalization techniques discussed so far (image edited from [3])

Even though an assumption of equal contribution is still being made within each group, the authors hypothesize that this is not as problematic, as innate grouping arises within features for visual recognition. One example they use to illustrate this is that many high-performance handcrafted features in traditional computer vision have terms that are explicitly grouped together. Take for example Histogram of Oriented Gradients [4] -- after computing histograms per spatially local block, each per-block histogram is normalized before being concatenated together to form the final feature vector.

You will now implement Group Normalization.

[2] Ba, Jimmy Lei, Jamie Ryan Kiros, and Geoffrey E. Hinton. "Layer Normalization." stat 1050 (2016): 21.

[3] Wu, Yuxin, and Kaiming He. "Group Normalization." arXiv preprint arXiv:1803.08494 (2018).

Spatial Group Normalization: Forward Pass

In the file cs231n/layers.py , implement the forward pass for group normalization in the function spatial_groupnorm_forward . Check your implementation by running the following:

```
In [ ]: np.random.seed(231)
        # Check the training-time forward pass by checking means and variances
        # of features both before and after spatial batch normalization.
        N, C, H, W = 2, 6, 4, 5
        G = 2
        x = 4 * np.random.randn(N, C, H, W) + 10
        x g = x.reshape((N*G, -1))
        print('Before spatial group normalization:')
        print(' shape: ', x.shape)
        print(' means: ', x_g.mean(axis=1))
        print(' stds: ', x g.std(axis=1))
        # Means should be close to zero and stds close to one
        gamma, beta = np.ones((1,C,1,1)), np.zeros((1,C,1,1))
        bn param = {'mode': 'train'}
        out, = spatial groupnorm forward(x, gamma, beta, G, bn param)
        out g = out.reshape((N*G, -1))
        print('After spatial group normalization:')
        print(' shape: ', out.shape)
        print(' means: ', out_g.mean(axis=1))
        print(' stds: ', out_g.std(axis=1))
```

Spatial Group Normalization: Backward Pass

In the file cs231n/layers.py , implement the backward pass for spatial batch normalization in the function spatial_groupnorm_backward . Run the following to check your implementation using a numeric gradient check:

```
In []: np.random.seed(231)
    N, C, H, W = 2, 6, 4, 5
    G = 2
    x = 5 * np.random.randn(N, C, H, W) + 12
    gamma = np.random.randn(1,C,1,1)
    beta = np.random.randn(1,C,1,1)
```

```
dout = np.random.randn(N, C, H, W)

gn_param = {}
fx = lambda x: spatial_groupnorm_forward(x, gamma, beta, G, gn_param)[0]
fg = lambda a: spatial_groupnorm_forward(x, gamma, beta, G, gn_param)[0]
fb = lambda b: spatial_groupnorm_forward(x, gamma, beta, G, gn_param)[0]

dx_num = eval_numerical_gradient_array(fx, x, dout)
da_num = eval_numerical_gradient_array(fg, gamma, dout)
db_num = eval_numerical_gradient_array(fb, beta, dout)

_, cache = spatial_groupnorm_forward(x, gamma, beta, G, gn_param)
dx, dgamma, dbeta = spatial_groupnorm_backward(dout, cache)

# You should expect errors of magnitudes between 1e-12 and 1e-07.
print('dx error: ', rel_error(dx_num, dx))
print('dgamma error: ', rel_error(da_num, dgamma))
print('dbeta error: ', rel_error(db_num, dbeta))
```

This notebook was converted to PDF with convert.ploomber.io

```
In [ ]: # This mounts your Google Drive to the Colab VM.
        from google.colab import drive
        drive.mount('/content/drive')
        # TODO: Enter the foldername in your Drive where you have saved the unzipped
        # assignment folder, e.g. 'cs231n/assignments/assignment1/'
        FOLDERNAME = None
        assert FOLDERNAME is not None, "[!] Enter the foldername."
        # Now that we've mounted your Drive, this ensures that
        # the Python interpreter of the Colab VM can load
        # python files from within it.
        import sys
        sys.path.append('/content/drive/My Drive/{}'.format(FOLDERNAME))
        # This downloads the CIFAR-10 dataset to your Drive
        # if it doesn't already exist.
        %cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/datasets/
        !bash get datasets.sh
        %cd /content/drive/My\ Drive/$F0LDERNAME
```

Introduction to PyTorch

You've written a lot of code in this assignment to provide a whole host of neural network functionality. Dropout, Batch Norm, and 2D convolutions are some of the workhorses of deep learning in computer vision. You've also worked hard to make your code efficient and vectorized.

For the last part of this assignment, though, we're going to leave behind your beautiful codebase and instead migrate to one of two popular deep learning frameworks: in this instance, PyTorch (or TensorFlow, if you choose to work with that notebook).

Why do we use deep learning frameworks?

- Our code will now run on GPUs! This will allow our models to train much faster. When using a framework like PyTorch or TensorFlow you can harness the power of the GPU for your own custom neural network architectures without having to write CUDA code directly (which is beyond the scope of this class).
- In this class, we want you to be ready to use one of these frameworks for your project so you can experiment more efficiently than if you were writing every feature you want to use by hand.
- We want you to stand on the shoulders of giants! TensorFlow and PyTorch are both excellent frameworks that will make your lives a lot easier, and now

- that you understand their guts, you are free to use them :)
- Finally, we want you to be exposed to the sort of deep learning code you might run into in academia or industry.

What is PyTorch?

PyTorch is a system for executing dynamic computational graphs over Tensor objects that behave similarly as numpy ndarray. It comes with a powerful automatic differentiation engine that removes the need for manual backpropagation.

How do I learn PyTorch?

One of our former instructors, Justin Johnson, made an excellent tutorial for PyTorch.

You can also find the detailed API doc here. If you have other questions that are not addressed by the API docs, the PyTorch forum is a much better place to ask than StackOverflow.

Table of Contents

This assignment has 5 parts. You will learn PyTorch on **three different levels of abstraction**, which will help you understand it better and prepare you for the final project.

- 1. Part I, Preparation: we will use CIFAR-10 dataset.
- 2. Part II, Barebones PyTorch: **Abstraction level 1**, we will work directly with the lowest-level PyTorch Tensors.
- 3. Part III, PyTorch Module API: **Abstraction level 2**, we will use nn.Module to define arbitrary neural network architecture.
- 4. Part IV, PyTorch Sequential API: **Abstraction level 3**, we will use nn.Sequential to define a linear feed-forward network very conveniently.
- 5. Part V, CIFAR-10 open-ended challenge: please implement your own network to get as high accuracy as possible on CIFAR-10. You can experiment with any layer, optimizer, hyperparameters or other advanced features.

Here is a table of comparison:

API	Flexibility	Convenience
Barebone	High	Low
nn.Module	High	Medium

API	Flexibility	Convenience
nn.Sequential	Low	High

GPU

You can manually switch to a GPU device on Colab by clicking Runtime -> Change runtime type and selecting GPU under Hardware Accelerator. You should do this before running the following cells to import packages, since the kernel gets restarted upon switching runtimes.

```
In [ ]: import torch
        import torch.nn as nn
        import torch.optim as optim
        from torch.utils.data import DataLoader
        from torch.utils.data import sampler
        import torchvision.datasets as dset
        import torchvision.transforms as T
        import numpy as np
        USE GPU = True
        dtype = torch.float32 # We will be using float throughout this tutorial.
        if USE GPU and torch.cuda.is available():
            device = torch.device('cuda')
        else:
            device = torch.device('cpu')
        # Constant to control how frequently we print train loss.
        print every = 100
        print('using device:', device)
```

Part I. Preparation

Now, let's load the CIFAR-10 dataset. This might take a couple minutes the first time you do it, but the files should stay cached after that.

In previous parts of the assignment we had to write our own code to download the CIFAR-10 dataset, preprocess it, and iterate through it in minibatches; PyTorch provides convenient tools to automate this process for us.

```
In [ ]: NUM_TRAIN = 49000

# The torchvision.transforms package provides tools for preprocessing data
# and for performing data augmentation; here we set up a transform to
# preprocess the data by subtracting the mean RGB value and dividing by the
```

```
# standard deviation of each RGB value; we've hardcoded the mean and std.
transform = T.Compose([
                T.ToTensor().
                T.Normalize((0.4914, 0.4822, 0.4465), (0.2023, 0.1994, 0.201
            ])
# We set up a Dataset object for each split (train / val / test); Datasets l
# training examples one at a time, so we wrap each Dataset in a DataLoader w
\# iterates through the Dataset and forms minibatches. We divide the CIFAR-10
# training set into train and val sets by passing a Sampler object to the
# DataLoader telling how it should sample from the underlying Dataset.
cifar10 train = dset.CIFAR10('./cs231n/datasets', train=True, download=True,
                             transform=transform)
loader train = DataLoader(cifar10 train, batch size=64,
                          sampler=sampler.SubsetRandomSampler(range(NUM TRAI
cifar10 val = dset.CIFAR10('./cs231n/datasets', train=True, download=True,
                           transform=transform)
loader val = DataLoader(cifar10 val, batch size=64,
                        sampler=sampler.SubsetRandomSampler(range(NUM TRAIN,
cifar10 test = dset.CIFAR10('./cs231n/datasets', train=False, download=True,
                            transform=transform)
loader test = DataLoader(cifar10 test, batch size=64)
```

Part II. Barebones PyTorch

PyTorch ships with high-level APIs to help us define model architectures conveniently, which we will cover in Part II of this tutorial. In this section, we will start with the barebone PyTorch elements to understand the autograd engine better. After this exercise, you will come to appreciate the high-level model API more.

We will start with a simple fully-connected ReLU network with two hidden layers and no biases for CIFAR classification. This implementation computes the forward pass using operations on PyTorch Tensors, and uses PyTorch autograd to compute gradients. It is important that you understand every line, because you will write a harder version after the example.

When we create a PyTorch Tensor with requires_grad=True, then operations involving that Tensor will not just compute values; they will also build up a computational graph in the background, allowing us to easily backpropagate through the graph to compute gradients of some Tensors with respect to a downstream loss. Concretely if x is a Tensor with x.requires_grad == True then after backpropagation x.grad will be another Tensor holding the gradient of x with respect to the scalar loss at the end.

PyTorch Tensors: Flatten Function

A PyTorch Tensor is conceptionally similar to a numpy array: it is an n-dimensional grid of numbers, and like numpy PyTorch provides many functions to efficiently operate on Tensors. As a simple example, we provide a flatten function below which reshapes image data for use in a fully-connected neural network.

Recall that image data is typically stored in a Tensor of shape $N \times C \times H \times W$, where:

- N is the number of datapoints
- C is the number of channels
- H is the height of the intermediate feature map in pixels
- W is the height of the intermediate feature map in pixels

This is the right way to represent the data when we are doing something like a 2D convolution, that needs spatial understanding of where the intermediate features are relative to each other. When we use fully connected affine layers to process the image, however, we want each datapoint to be represented by a single vector -- it's no longer useful to segregate the different channels, rows, and columns of the data. So, we use a "flatten" operation to collapse the C x H x W values per representation into a single long vector. The flatten function below first reads in the N, C, H, and W values from a given batch of data, and then returns a "view" of that data. "View" is analogous to numpy's "reshape" method: it reshapes x's dimensions to be N x ??, where ?? is allowed to be anything (in this case, it will be C x H x W, but we don't need to specify that explicitly).

```
In []: def flatten(x):
    N = x.shape[0] # read in N, C, H, W
    return x.view(N, -1) # "flatten" the C * H * W values into a single vec

def test_flatten():
    x = torch.arange(12).view(2, 1, 3, 2)
    print('Before flattening: ', x)
    print('After flattening: ', flatten(x))

test_flatten()
```

Barebones PyTorch: Two-Layer Network

Here we define a function two_layer_fc which performs the forward pass of a two-layer fully-connected ReLU network on a batch of image data. After defining the forward pass we check that it doesn't crash and that it produces outputs of the right shape by running zeros through the network.

You don't have to write any code here, but it's important that you read and understand the implementation.

```
In [ ]: import torch.nn.functional as F # useful stateless functions
        def two layer fc(x, params):
            A fully-connected neural networks; the architecture is:
            NN is fully connected -> ReLU -> fully connected layer.
            Note that this function only defines the forward pass;
            PyTorch will take care of the backward pass for us.
            The input to the network will be a minibatch of data, of shape
            (N, d1, ..., dM) where d1 * ... * dM = D. The hidden layer will have H \iota
            and the output layer will produce scores for C classes.
            Inputs:
            - x: A PyTorch Tensor of shape (N, d1, ..., dM) giving a minibatch of
              input data.
            - params: A list [w1, w2] of PyTorch Tensors giving weights for the netw
              w1 has shape (D, H) and w2 has shape (H, C).
            - scores: A PyTorch Tensor of shape (N, C) giving classification scores
             the input data x.
            # first we flatten the image
            x = flatten(x) # shape: [batch size, C x H x W]
            w1, w2 = params
            # Forward pass: compute predicted y using operations on Tensors. Since w
            # w2 have requires grad=True, operations involving these Tensors will ca
            # PyTorch to build a computational graph, allowing automatic computation
            # gradients. Since we are no longer implementing the backward pass by ha
            # don't need to keep references to intermediate values.
            # you can also use `.clamp(min=0)`, equivalent to F.relu()
            x = F.relu(x.mm(w1))
            x = x.mm(w2)
            return x
        def two layer fc test():
            hidden layer size = 42
            x = torch.zeros((64, 50), dtype=dtype) # minibatch size 64, feature dim
            w1 = torch.zeros((50, hidden layer size), dtype=dtype)
            w2 = torch.zeros((hidden layer size, 10), dtype=dtype)
            scores = two_layer_fc(x, [w1, w2])
            print(scores.size()) # you should see [64, 10]
        two layer fc test()
```

Here you will complete the implementation of the function three_layer_convnet, which will perform the forward pass of a three-layer convolutional network. Like above, we can immediately test our implementation by passing zeros through the network. The network should have the following architecture:

- 1. A convolutional layer (with bias) with channel_1 filters, each with shape KW1 x KH1, and zero-padding of two
- 2. ReLU nonlinearity
- 3. A convolutional layer (with bias) with channel_2 filters, each with shape KW2 x KH2, and zero-padding of one
- 4. ReLU nonlinearity
- 5. Fully-connected layer with bias, producing scores for C classes.

Note that we have **no softmax activation** here after our fully-connected layer: this is because PyTorch's cross entropy loss performs a softmax activation for you, and by bundling that step in makes computation more efficient.

HINT: For convolutions:

http://pytorch.org/docs/stable/nn.html#torch.nn.functional.conv2d; pay attention to the shapes of convolutional filters!

```
In [ ]: def three layer convnet(x, params):
            Performs the forward pass of a three-layer convolutional network with the
            architecture defined above.
            Inputs:
            - x: A PyTorch Tensor of shape (N, 3, H, W) giving a minibatch of images
            - params: A list of PyTorch Tensors giving the weights and biases for th
              network; should contain the following:
              - conv w1: PyTorch Tensor of shape (channel 1, 3, KH1, KW1) giving wei
               for the first convolutional layer
              - conv b1: PyTorch Tensor of shape (channel 1,) giving biases for the
                convolutional layer
              - conv w2: PyTorch Tensor of shape (channel 2, channel 1, KH2, KW2) qi
                weights for the second convolutional layer
              - conv b2: PyTorch Tensor of shape (channel 2,) giving biases for the
                convolutional layer
              - fc w: PyTorch Tensor giving weights for the fully-connected layer. (
                figure out what the shape should be?
              - fc b: PyTorch Tensor giving biases for the fully-connected layer. Ca
                figure out what the shape should be?
            Returns:
            - scores: PyTorch Tensor of shape (N, C) giving classification scores fd
            conv w1, conv b1, conv w2, conv b2, fc w, fc b = params
            scores = None
```

After defining the forward pass of the ConvNet above, run the following cell to test your implementation.

When you run this function, scores should have shape (64, 10).

```
In []: def three_layer_convnet_test():
    x = torch.zeros((64, 3, 32, 32), dtype=dtype) # minibatch size 64, imag
    conv_wl = torch.zeros((6, 3, 5, 5), dtype=dtype) # [out_channel, in_chateonv_bl = torch.zeros((6,)) # out_channel
    conv_w2 = torch.zeros((9, 6, 3, 3), dtype=dtype) # [out_channel, in_chateonv_b2 = torch.zeros((9,)) # out_channel

# you must calculate the shape of the tensor after two conv layers, before fc_w = torch.zeros((9 * 32 * 32, 10))
    fc_b = torch.zeros(10)

scores = three_layer_convnet(x, [conv_wl, conv_bl, conv_w2, conv_b2, fc_print(scores.size()) # you should see [64, 10]
three_layer_convnet_test()
```

Barebones PyTorch: Initialization

Let's write a couple utility methods to initialize the weight matrices for our models.

- random_weight(shape) initializes a weight tensor with the Kaiming normalization method.
- zero_weight(shape) initializes a weight tensor with all zeros. Useful for instantiating bias parameters.

The random_weight function uses the Kaiming normal initialization method, described in:

He et al, Delving Deep into Rectifiers: Surpassing Human-Level Performance on ImageNet Classification, ICCV 2015, https://arxiv.org/abs/1502.01852

```
In [ ]: def random_weight(shape):
```

```
Create random Tensors for weights; setting requires grad=True means that
   want to compute gradients for these Tensors during the backward pass.
   We use Kaiming normalization: sgrt(2 / fan in)
   if len(shape) == 2: # FC weight
       fan in = shape[0]
        fan in = np.prod(shape[1:]) # conv weight [out channel, in channel,
   # randn is standard normal distribution generator.
   w = torch.randn(shape, device=device, dtype=dtype) * np.sqrt(2. / fan ir
   w.requires grad = True
   return w
def zero weight(shape):
   return torch zeros(shape, device=device, dtype=dtype, requires grad=True
# create a weight of shape [3 \times 5]
# you should see the type `torch.cuda.FloatTensor` if you use GPU.
# Otherwise it should be `torch.FloatTensor`
random weight((3, 5))
```

Barebones PyTorch: Check Accuracy

When training the model we will use the following function to check the accuracy of our model on the training or validation sets.

When checking accuracy we don't need to compute any gradients; as a result we don't need PyTorch to build a computational graph for us when we compute scores. To prevent a graph from being built we scope our computation under a torch.no_grad() context manager.

```
In [ ]: def check accuracy part2(loader, model fn, params):
            Check the accuracy of a classification model.
            Inputs:
            - loader: A DataLoader for the data split we want to check
            - model fn: A function that performs the forward pass of the model,
             with the signature scores = model fn(x, params)
            - params: List of PyTorch Tensors giving parameters of the model
            Returns: Nothing, but prints the accuracy of the model
            split = 'val' if loader.dataset.train else 'test'
            print('Checking accuracy on the %s set' % split)
            num correct, num samples = 0, 0
            with torch.no grad():
                for x, y in loader:
                    x = x.to(device=device, dtype=dtype) # move to device, e.g. GPU
                    y = y.to(device=device, dtype=torch.int64)
                    scores = model fn(x, params)
                    , preds = scores.max(1)
                    num correct += (preds == y).sum()
```

```
num_samples += preds.size(0)
acc = float(num_correct) / num_samples
print('Got %d / %d correct (%.2f%%)' % (num_correct, num_samples, 16
```

BareBones PyTorch: Training Loop

We can now set up a basic training loop to train our network. We will train the model using stochastic gradient descent without momentum. We will use torch.functional.cross_entropy to compute the loss; you can read about it here.

The training loop takes as input the neural network function, a list of initialized parameters ([w1, w2] in our example), and learning rate.

```
In [ ]: def train part2(model fn, params, learning rate):
            Train a model on CIFAR-10.
            Inputs:
            - model fn: A Python function that performs the forward pass of the mode
              It should have the signature scores = model fn(x, params) where x is a
              PyTorch Tensor of image data, params is a list of PyTorch Tensors givi
              model weights, and scores is a PyTorch Tensor of shape (N, C) giving
              scores for the elements in x.
            - params: List of PyTorch Tensors giving weights for the model
            - learning rate: Python scalar giving the learning rate to use for SGD
            Returns: Nothing
            for t, (x, y) in enumerate(loader train):
                # Move the data to the proper device (GPU or CPU)
                x = x.to(device=device, dtype=dtype)
                y = y.to(device=device, dtype=torch.long)
                # Forward pass: compute scores and loss
                scores = model fn(x, params)
                loss = F.cross entropy(scores, y)
                # Backward pass: PyTorch figures out which Tensors in the computation
                \# graph has requires grad=True and uses backpropagation to compute t
                # gradient of the loss with respect to these Tensors, and stores the
                # gradients in the .grad attribute of each Tensor.
                loss.backward()
                # Update parameters. We don't want to backpropagate through the
                # parameter updates, so we scope the updates under a torch.no grad()
                # context manager to prevent a computational graph from being built.
                with torch.no grad():
                    for w in params:
                        w -= learning rate * w.grad
                        # Manually zero the gradients after running the backward pas
                        w.grad.zero ()
```

```
if t % print_every == 0:
    print('Iteration %d, loss = %.4f' % (t, loss.item()))
    check_accuracy_part2(loader_val, model_fn, params)
    print()
```

BareBones PyTorch: Train a Two-Layer Network

Now we are ready to run the training loop. We need to explicitly allocate tensors for the fully connected weights, w1 and w2.

Each minibatch of CIFAR has 64 examples, so the tensor shape is [64, 3, 32, 32].

After flattening, x shape should be [64, 3 * 32 * 32]. This will be the size of the first dimension of w1. The second dimension of w1 is the hidden layer size, which will also be the first dimension of w2.

Finally, the output of the network is a 10-dimensional vector that represents the probability distribution over 10 classes.

You don't need to tune any hyperparameters but you should see accuracies above 40% after training for one epoch.

```
In []: hidden_layer_size = 4000
learning_rate = 1e-2

w1 = random_weight((3 * 32 * 32, hidden_layer_size))
w2 = random_weight((hidden_layer_size, 10))

train_part2(two_layer_fc, [w1, w2], learning_rate)
```

BareBones PyTorch: Training a ConvNet

In the below you should use the functions defined above to train a three-layer convolutional network on CIFAR. The network should have the following architecture:

- 1. Convolutional layer (with bias) with 32 5x5 filters, with zero-padding of 2
- 2. ReLU
- 3. Convolutional layer (with bias) with 16 3x3 filters, with zero-padding of 1
- 4. ReLU
- 5. Fully-connected layer (with bias) to compute scores for 10 classes

You should initialize your weight matrices using the random_weight function defined above, and you should initialize your bias vectors using the zero weight function above.

You don't need to tune any hyperparameters, but if everything works correctly you should achieve an accuracy above 42% after one epoch.

```
In [ ]: learning rate = 3e-3
     channel 1 = 32
     channel 2 = 16
     conv w1 = None
     conv b1 = None
     conv w2 = None
     conv b2 = None
     fc w = None
     fc b = None
     # TODO: Initialize the parameters of a three-layer ConvNet.
     # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
     pass
     # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
     END OF YOUR CODE
     params = [conv w1, conv b1, conv w2, conv b2, fc w, fc b]
     train_part2(three_layer_convnet, params, learning_rate)
```

Part III. PyTorch Module API

Barebone PyTorch requires that we track all the parameter tensors by hand. This is fine for small networks with a few tensors, but it would be extremely inconvenient and error-prone to track tens or hundreds of tensors in larger networks.

PyTorch provides the nn.Module API for you to define arbitrary network architectures, while tracking every learnable parameters for you. In Part II, we implemented SGD ourselves. PyTorch also provides the torch.optim package that implements all the common optimizers, such as RMSProp, Adagrad, and Adam. It even supports approximate second-order methods like L-BFGS! You can refer to the doc for the exact specifications of each optimizer.

To use the Module API, follow the steps below:

1. Subclass nn.Module . Give your network class an intuitive name like TwoLayerFC .

- 2. In the constructor __init__() , define all the layers you need as class attributes. Layer objects like nn.Linear and nn.Conv2d are themselves nn.Module subclasses and contain learnable parameters, so that you don't have to instantiate the raw tensors yourself. nn.Module will track these internal parameters for you. Refer to the doc to learn more about the dozens of builtin layers. Warning: don't forget to call the super().__init__() first!
- 3. In the forward() method, define the *connectivity* of your network. You should use the attributes defined in __init__ as function calls that take tensor as input and output the "transformed" tensor. Do *not* create any new layers with learnable parameters in forward()! All of them must be declared upfront in __init__.

After you define your Module subclass, you can instantiate it as an object and call it just like the NN forward function in part II.

Module API: Two-Layer Network

Here is a concrete example of a 2-layer fully connected network:

```
In [ ]: class TwoLayerFC(nn.Module):
            def __init__(self, input_size, hidden_size, num_classes):
                super().__init ()
                # assign layer objects to class attributes
                self.fc1 = nn.Linear(input size, hidden size)
                # nn.init package contains convenient initialization methods
                # http://pytorch.org/docs/master/nn.html#torch-nn-init
                nn.init.kaiming normal (self.fc1.weight)
                self.fc2 = nn.Linear(hidden size, num classes)
                nn.init.kaiming normal (self.fc2.weight)
            def forward(self, x):
                # forward always defines connectivity
                x = flatten(x)
                scores = self.fc2(F.relu(self.fc1(x)))
                return scores
        def test TwoLayerFC():
            input size = 50
            x = torch.zeros((64, input size), dtype=dtype) # minibatch size 64, fee
            model = TwoLayerFC(input size, 42, 10)
            scores = model(x)
            print(scores.size()) # you should see [64, 10]
        test TwoLayerFC()
```

Module API: Three-Layer ConvNet

It's your turn to implement a 3-layer ConvNet followed by a fully connected layer. The network architecture should be the same as in Part II:

- 1. Convolutional layer with channel 1 5x5 filters with zero-padding of 2
- 2. ReLU
- 3. Convolutional layer with channel 2 3x3 filters with zero-padding of 1
- 4 Rell
- 5. Fully-connected layer to num classes classes

You should initialize the weight matrices of the model using the Kaiming normal initialization method.

HINT: http://pytorch.org/docs/stable/nn.html#conv2d

After you implement the three-layer ConvNet, the test_ThreeLayerConvNet function will run your implementation; it should print (64, 10) for the shape of the output scores.

```
In [ ]: class ThreeLayerConvNet(nn.Module):
       def init (self, in channel, channel 1, channel 2, num classes):
         super(). init ()
         # TODO: Set up the layers you need for a three-layer ConvNet with th
         # architecture defined above.
         # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
         pass
         # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
         END OF YOUR CODE
         def forward(self, x):
         scores = None
         # TODO: Implement the forward function for a 3-layer ConvNet. you
         # should use the layers you defined in init and specify the
         # connectivity of those layers in forward()
         # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
         pass
         # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
         END OF YOUR CODE
         return scores
```

```
def test_ThreeLayerConvNet():
    x = torch.zeros((64, 3, 32, 32), dtype=dtype) # minibatch size 64, imag
    model = ThreeLayerConvNet(in_channel=3, channel_1=12, channel_2=8, num_c
    scores = model(x)
    print(scores.size()) # you should see [64, 10]
test_ThreeLayerConvNet()
```

Module API: Check Accuracy

Given the validation or test set, we can check the classification accuracy of a neural network.

This version is slightly different from the one in part II. You don't manually pass in the parameters anymore.

```
In [ ]: def check accuracy part34(loader, model):
            if loader.dataset.train:
                print('Checking accuracy on validation set')
            else:
                print('Checking accuracy on test set')
            num correct = 0
            num samples = 0
            model.eval() # set model to evaluation mode
            with torch.no grad():
                for x, y in loader:
                    x = x.to(device=device, dtype=dtype) # move to device, e.g. GPU
                    y = y.to(device=device, dtype=torch.long)
                    scores = model(x)
                    _, preds = scores.max(1)
                    num correct += (preds == y).sum()
                    num samples += preds.size(0)
                acc = float(num_correct) / num samples
                print('Got %d / %d correct (%.2f)' % (num correct, num samples, 100
```

Module API: Training Loop

We also use a slightly different training loop. Rather than updating the values of the weights ourselves, we use an Optimizer object from the torch.optim package, which abstract the notion of an optimization algorithm and provides implementations of most of the algorithms commonly used to optimize neural networks.

```
In []: def train_part34(model, optimizer, epochs=1):
    """
    Train a model on CIFAR-10 using the PyTorch Module API.

Inputs:
    - model: A PyTorch Module giving the model to train.
    - optimizer: An Optimizer object we will use to train the model
    - epochs: (Optional) A Python integer giving the number of epochs to train.
```

```
Returns: Nothing, but prints model accuracies during training.
model = model.to(device=device) # move the model parameters to CPU/GPU
for e in range(epochs):
    for t, (x, y) in enumerate(loader train):
        model.train() # put model to training mode
        x = x.to(device=device, dtype=dtype) # move to device, e.g. GPU
        y = y.to(device=device, dtype=torch.long)
        scores = model(x)
        loss = F.cross entropy(scores, y)
        # Zero out all of the gradients for the variables which the opti
        # will update.
        optimizer.zero grad()
        # This is the backwards pass: compute the gradient of the loss w
        # respect to each parameter of the model.
        loss.backward()
        # Actually update the parameters of the model using the gradient
        # computed by the backwards pass.
        optimizer.step()
        if t % print every == 0:
            print('Iteration %d, loss = %.4f' % (t, loss.item()))
            check accuracy part34(loader val, model)
            print()
```

Module API: Train a Two-Layer Network

Now we are ready to run the training loop. In contrast to part II, we don't explicitly allocate parameter tensors anymore.

Simply pass the input size, hidden layer size, and number of classes (i.e. output size) to the constructor of TwoLayerFC.

You also need to define an optimizer that tracks all the learnable parameters inside TwoLayerFC .

You don't need to tune any hyperparameters, but you should see model accuracies above 40% after training for one epoch.

```
In [ ]: hidden_layer_size = 4000
learning_rate = 1e-2
model = TwoLayerFC(3 * 32 * 32, hidden_layer_size, 10)
optimizer = optim.SGD(model.parameters(), lr=learning_rate)
train_part34(model, optimizer)
```

Module API: Train a Three-Layer ConvNet

You should now use the Module API to train a three-layer ConvNet on CIFAR. This should look very similar to training the two-layer network! You don't need to tune any hyperparameters, but you should achieve above 45% after training for one epoch.

You should train the model using stochastic gradient descent without momentum.

Part IV. PyTorch Sequential API

Part III introduced the PyTorch Module API, which allows you to define arbitrary learnable layers and their connectivity.

For simple models like a stack of feed forward layers, you still need to go through 3 steps: subclass nn.Module, assign layers to class attributes in __init__, and call each layer one by one in forward(). Is there a more convenient way?

Fortunately, PyTorch provides a container Module called nn.Sequential, which merges the above steps into one. It is not as flexible as nn.Module, because you cannot specify more complex topology than a feed-forward stack, but it's good enough for many use cases.

Sequential API: Two-Layer Network

Let's see how to rewrite our two-layer fully connected network example with nn.Sequential, and train it using the training loop defined above.

Again, you don't need to tune any hyperparameters here, but you should achieve above 40% accuracy after one epoch of training.

```
In [ ]: # We need to wrap `flatten` function in a module in order to stack it
        # in nn.Sequential
        class Flatten(nn.Module):
            def forward(self, x):
                return flatten(x)
        hidden layer size = 4000
        learning rate = 1e-2
        model = nn.Sequential(
            Flatten(),
            nn.Linear(3 * 32 * 32, hidden layer size),
            nn.ReLU(),
            nn.Linear(hidden layer size, 10),
        # you can use Nesterov momentum in optim.SGD
        optimizer = optim.SGD(model.parameters(), lr=learning rate,
                             momentum=0.9, nesterov=True)
        train part34(model, optimizer)
```

Sequential API: Three-Layer ConvNet

Here you should use nn.Sequential to define and train a three-layer ConvNet with the same architecture we used in Part III:

- 1. Convolutional layer (with bias) with 32 5x5 filters, with zero-padding of 2
- 2. ReLU
- 3. Convolutional layer (with bias) with 16 3x3 filters, with zero-padding of 1
- 4. ReLU
- 5. Fully-connected layer (with bias) to compute scores for 10 classes

You can use the default PyTorch weight initialization.

You should optimize your model using stochastic gradient descent with Nesterov momentum 0.9.

Again, you don't need to tune any hyperparameters but you should see accuracy above 55% after one epoch of training.

```
In []: channel_1 = 32
    channel_2 = 16
    learning_rate = 1e-2

model = None
    optimizer = None
```

Part V. CIFAR-10 open-ended challenge

In this section, you can experiment with whatever ConvNet architecture you'd like on CIFAR-10.

Now it's your job to experiment with architectures, hyperparameters, loss functions, and optimizers to train a model that achieves **at least 70**% accuracy on the CIFAR-10 **validation** set within 10 epochs. You can use the check_accuracy and train functions from above. You can use either nn.Module or nn.Sequential API.

Describe what you did at the end of this notebook.

Here are the official API documentation for each component. One note: what we call in the class "spatial batch norm" is called "BatchNorm2D" in PyTorch.

- Layers in torch.nn package: http://pytorch.org/docs/stable/nn.html
- Activations: http://pytorch.org/docs/stable/nn.html#non-linear-activations
- Loss functions: http://pytorch.org/docs/stable/nn.html#loss-functions
- Optimizers: http://pytorch.org/docs/stable/optim.html

Things you might try:

- Filter size: Above we used 5x5: would smaller filters be more efficient?
- Number of filters: Above we used 32 filters. Do more or fewer do better?
- Pooling vs Strided Convolution: Do you use max pooling or just stride convolutions?
- **Batch normalization**: Try adding spatial batch normalization after convolution layers and vanilla batch normalization after affine layers. Do your networks train faster?
- **Network architecture**: The network above has two layers of trainable parameters. Can you do better with a deep network? Good architectures to

try include:

- [conv-relu-pool]xN -> [affine]xM -> [softmax or SVM]
- [conv-relu-conv-relu-pool]xN -> [affine]xM -> [softmax or SVM]
- [batchnorm-relu-conv]xN -> [affine]xM -> [softmax or SVM]
- Global Average Pooling: Instead of flattening and then having multiple affine layers, perform convolutions until your image gets small (7x7 or so) and then perform an average pooling operation to get to a 1x1 image picture (1, 1, Filter#), which is then reshaped into a (Filter#) vector. This is used in Google's Inception Network (See Table 1 for their architecture).
- **Regularization**: Add I2 weight regularization, or perhaps use Dropout.

Tips for training

For each network architecture that you try, you should tune the learning rate and other hyperparameters. When doing this there are a couple important things to keep in mind:

- If the parameters are working well, you should see improvement within a few hundred iterations
- Remember the coarse-to-fine approach for hyperparameter tuning: start by testing a large range of hyperparameters for just a few training iterations to find the combinations of parameters that are working at all.
- Once you have found some sets of parameters that seem to work, search more finely around these parameters. You may need to train for more epochs.
- You should use the validation set for hyperparameter search, and save your test set for evaluating your architecture on the best parameters as selected by the validation set.

Going above and beyond

If you are feeling adventurous there are many other features you can implement to try and improve your performance. You are **not required** to implement any of these, but don't miss the fun if you have time!

- Alternative optimizers: you can try Adam, Adagrad, RMSprop, etc.
- Alternative activation functions such as leaky ReLU, parametric ReLU, ELU, or MaxOut.
- Model ensembles
- Data augmentation
- New Architectures
 - ResNets where the input from the previous layer is added to the output.
 - DenseNets where inputs into previous layers are concatenated together.
 - This blog has an in-depth overview

Have fun and happy training!

```
# Experiment with any architectures, optimizers, and hyperparameters.
     # Achieve AT LEAST 70% accuracy on the *validation set* within 10 epochs.
     # Note that you can use the check accuracy function to evaluate on either
     # the test set or the validation set, by passing either loader test or
      # loader val as the second argument to check accuracy. You should not touch
     # the test set until you have finished your architecture and hyperparameter
      # tuning, and only run the test set once at the end to report a final value.
      model = None
      optimizer = None
     # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
      pass
      # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
      END OF YOUR CODE
      # You should get at least 70% accuracy
      train part34(model, optimizer, epochs=10)
```

Describe what you did

In the cell below you should write an explanation of what you did, any additional features that you implemented, and/or any graphs that you made in the process of training and evaluating your network.

Answer:

Test set -- run this only once

Now that we've gotten a result we're happy with, we test our final model on the test set (which you should store in best_model). Think about how this compares to your validation set accuracy.

```
In [ ]: best_model = model
    check_accuracy_part34(loader_test, best_model)
```

```
In [ ]: # This mounts your Google Drive to the Colab VM.
        from google.colab import drive
        drive.mount('/content/drive')
        # TODO: Enter the foldername in your Drive where you have saved the unzipped
        # assignment folder, e.g. 'cs231n/assignments/assignment1/'
        FOLDERNAME = None
        assert FOLDERNAME is not None, "[!] Enter the foldername."
        # Now that we've mounted your Drive, this ensures that
        # the Python interpreter of the Colab VM can load
        # python files from within it.
        import sys
        sys.path.append('/content/drive/My Drive/{}'.format(FOLDERNAME))
        # This downloads the CIFAR-10 dataset to your Drive
        # if it doesn't already exist.
        %cd /content/drive/My\ Drive/$FOLDERNAME/cs231n/datasets/
        !bash get datasets.sh
        %cd /content/drive/My\ Drive/$F0LDERNAME
```

Introduction to TensorFlow

You've written a lot of code in this assignment to provide a whole host of neural network functionality. Dropout, Batch Norm, and 2D convolutions are some of the workhorses of deep learning in computer vision. You've also worked hard to make your code efficient and vectorized.

For the last part of this assignment, though, we're going to leave behind your beautiful codebase and instead migrate to one of two popular deep learning frameworks: in this instance, TensorFlow (or PyTorch, if you choose to work with that notebook).

Why do we use deep learning frameworks?

- Our code will now run on GPUs! This will allow our models to train much faster. When using a framework like PyTorch or TensorFlow you can harness the power of the GPU for your own custom neural network architectures without having to write CUDA code directly (which is beyond the scope of this class).
- In this class, we want you to be ready to use one of these frameworks for your project so you can experiment more efficiently than if you were writing every feature you want to use by hand.
- We want you to stand on the shoulders of giants! TensorFlow and PyTorch are both excellent frameworks that will make your lives a lot easier, and now

- that you understand their guts, you are free to use them :)
- Finally, we want you to be exposed to the sort of deep learning code you might run into in academia or industry.

What is TensorFlow?

TensorFlow is a system for executing computational graphs over Tensor objects, with native support for performing backpropogation for its Variables. In it, we work with Tensors which are n-dimensional arrays analogous to the numpy ndarray.

How do I learn TensorFlow?

TensorFlow has many excellent tutorials available, including those from Google themselves.

Otherwise, this notebook will walk you through much of what you need to do to train models in TensorFlow. See the end of the notebook for some links to helpful tutorials if you want to learn more or need further clarification on topics that aren't fully explained here.

Note: This notebook is meant to teach you Tensorflow 2.x. Most examples on the web today are still in 1.x, so be careful not to confuse the two when looking up documentation.

Table of Contents

This notebook has 5 parts. We will walk through TensorFlow at **three different levels of abstraction**, which should help you better understand it and prepare you for working on your project.

- 1. Part I, Preparation: load the CIFAR-10 dataset.
- 2. Part II, Barebone TensorFlow: **Abstraction Level 1**, we will work directly with low-level TensorFlow graphs.
- 3. Part III, Keras Model API: **Abstraction Level 2**, we will use tf.keras.Model to define arbitrary neural network architecture.
- 4. Part IV, Keras Sequential + Functional API: **Abstraction Level 3**, we will use tf.keras.Sequential to define a linear feed-forward network very conveniently, and then explore the functional libraries for building unique and uncommon models that require more flexibility.
- 5. Part V, CIFAR-10 open-ended challenge: please implement your own network to get as high accuracy as possible on CIFAR-10. You can experiment with

any layer, optimizer, hyperparameters or other advanced features.

We will discuss Keras in more detail later in the notebook.

Here is a table of comparison:

API	Flexibility	Convenience
Barebone	High	Low
tf.keras.Model	High	Medium
tf.keras.Sequential	Low	High

GPU

You can manually switch to a GPU device on Colab by clicking Runtime -> Change runtime type and selecting GPU under Hardware Accelerator . You should do this before running the following cells to import packages, since the kernel gets restarted upon switching runtimes.

```
In []:
    import os
    import tensorflow as tf
    import numpy as np
    import timeit
    import matplotlib.pyplot as plt

%matplotlib inline

USE_GPU = True

if USE_GPU:
    device = '/device:GPU:0'
else:
    device = '/cpu:0'

# Constant to control how often we print when training models.
print_every = 100
print('Using device: ', device)
```

Part I: Preparation

First, we load the CIFAR-10 dataset. This might take a few minutes to download the first time you run it, but after that the files should be cached on disk and loading should be faster.

In previous parts of the assignment we used CS231N-specific code to download and read the CIFAR-10 dataset; however the tf.keras.datasets package in

TensorFlow provides prebuilt utility functions for loading many common datasets.

For the purposes of this assignment we will still write our own code to preprocess the data and iterate through it in minibatches. The tf.data package in TensorFlow provides tools for automating this process, but working with this package adds extra complication and is beyond the scope of this notebook. However using tf.data can be much more efficient than the simple approach used in this notebook, so you should consider using it for your project.

```
In [ ]: def load cifar10(num training=49000, num validation=1000, num test=10000):
            Fetch the CIFAR-10 dataset from the web and perform preprocessing to pre
            it for the two-layer neural net classifier. These are the same steps as
            we used for the SVM, but condensed to a single function.
            # Load the raw CIFAR-10 dataset and use appropriate data types and shape
            cifar10 = tf.keras.datasets.cifar10.load data()
            (X train, y train), (X test, y test) = cifar10
            X train = np.asarray(X train, dtype=np.float32)
            y train = np.asarray(y train, dtype=np.int32).flatten()
            X test = np.asarray(X test, dtype=np.float32)
            y test = np.asarray(y test, dtype=np.int32).flatten()
            # Subsample the data
            mask = range(num training, num training + num validation)
            X val = X train[mask]
            y val = y train[mask]
            mask = range(num training)
            X train = X train[mask]
            y train = y train[mask]
            mask = range(num test)
            X \text{ test} = X \text{ test[mask]}
            y test = y test[mask]
            # Normalize the data: subtract the mean pixel and divide by std
            mean pixel = X train.mean(axis=(0, 1, 2), keepdims=True)
            std pixel = X train.std(axis=(0, 1, 2), keepdims=True)
            X train = (X train - mean pixel) / std pixel
            X val = (X val - mean pixel) / std pixel
            X test = (X test - mean pixel) / std pixel
            return X train, y train, X val, y val, X test, y test
        # If there are errors with SSL downloading involving self-signed certificate
        # it may be that your Python version was recently installed on the current n
        # See: https://github.com/tensorflow/tensorflow/issues/10779
        # To fix, run the command: /Applications/Python\ 3.7/Install\ Certificates.c
        # ...replacing paths as necessary.
        # Invoke the above function to get our data.
        NHW = (0, 1, 2)
        X_train, y_train, X_val, y_val, X_test, y_test = load_cifar10()
        print('Train data shape: ', X_train.shape)
        print('Train labels shape: ', y_train.shape, y_train.dtype)
```

```
print('Validation data shape: ', X_val.shape)
        print('Validation labels shape: ', y val.shape)
        print('Test data shape: ', X_test.shape)
        print('Test labels shape: ', y test.shape)
In [ ]: class Dataset(object):
            def __init__(self, X, y, batch_size, shuffle=False):
                Construct a Dataset object to iterate over data X and labels y
                Inputs:
                - X: Numpy array of data, of any shape
                - y: Numpy array of labels, of any shape but with y.shape[0] == X.sh
                - batch size: Integer giving number of elements per minibatch
                - shuffle: (optional) Boolean, whether to shuffle the data on each \epsilon
                assert X.shape[0] == y.shape[0], 'Got different numbers of data and
                self.X, self.y = X, y
                self.batch size, self.shuffle = batch size, shuffle
            def __iter__(self):
                N, B = self.X.shape[0], self.batch size
                idxs = np.arange(N)
                if self.shuffle:
                    np.random.shuffle(idxs)
                return iter((self.X[i:i+B], self.y[i:i+B]) for i in range(0, N, B))
        train dset = Dataset(X train, y train, batch size=64, shuffle=True)
        val dset = Dataset(X val, y val, batch size=64, shuffle=False)
        test_dset = Dataset(X_test, y_test, batch_size=64)
In [ ]: # We can iterate through a dataset like this:
        for t, (x, y) in enumerate(train dset):
            print(t, x.shape, y.shape)
            if t > 5: break
```

Part II: Barebones TensorFlow

TensorFlow ships with various high-level APIs which make it very convenient to define and train neural networks; we will cover some of these constructs in Part III and Part IV of this notebook. In this section we will start by building a model with basic TensorFlow constructs to help you better understand what's going on under the hood of the higher-level APIs.

"Barebones Tensorflow" is important to understanding the building blocks of TensorFlow, but much of it involves concepts from TensorFlow 1.x. We will be working with legacy modules such as tf.Variable.

Therefore, please read and understand the differences between legacy (1.x) TF and the new (2.0) TF.

Historical background on TensorFlow 1.x

TensorFlow 1.x is primarily a framework for working with **static computational graphs**. Nodes in the computational graph are Tensors which will hold n-dimensional arrays when the graph is run; edges in the graph represent functions that will operate on Tensors when the graph is run to actually perform useful computation.

Before Tensorflow 2.0, we had to configure the graph into two phases. There are plenty of tutorials online that explain this two-step process. The process generally looks like the following for TF 1.x:

- 1. Build a computational graph that describes the computation that you want to perform. This stage doesn't actually perform any computation; it just builds up a symbolic representation of your computation. This stage will typically define one or more placeholder objects that represent inputs to the computational graph.
- 2. **Run the computational graph many times.** Each time the graph is run (e.g. for one gradient descent step) you will specify which parts of the graph you want to compute, and pass a feed_dict dictionary that will give concrete values to any placeholder s in the graph.

The new paradigm in Tensorflow 2.0

Now, with Tensorflow 2.0, we can simply adopt a functional form that is more Pythonic and similar in spirit to PyTorch and direct Numpy operation. Instead of the 2-step paradigm with computation graphs, making it (among other things) easier to debug TF code. You can read more details at https://www.tensorflow.org/guide/eager.

The main difference between the TF 1.x and 2.0 approach is that the 2.0 approach doesn't make use of tf.Session, tf.run, placeholder, feed_dict. To get more details of what's different between the two version and how to convert between the two, check out the official migration guide: https://www.tensorflow.org/alpha/guide/migration guide

Later, in the rest of this notebook we'll focus on this new, simpler approach.

TensorFlow warmup: Flatten Function

We can see this in action by defining a simple flatten function that will reshape image data for use in a fully-connected network.

In TensorFlow, data for convolutional feature maps is typically stored in a Tensor of shape $N \times H \times W \times C$ where:

- N is the number of datapoints (minibatch size)
- H is the height of the feature map
- W is the width of the feature map
- C is the number of channels in the feature map

This is the right way to represent the data when we are doing something like a 2D convolution, that needs spatial understanding of where the intermediate features are relative to each other. When we use fully connected affine layers to process the image, however, we want each datapoint to be represented by a single vector -- it's no longer useful to segregate the different channels, rows, and columns of the data. So, we use a "flatten" operation to collapse the $H \times W$ \times C values per representation into a single long vector.

Notice the <code>tf.reshape</code> call has the target shape as (N, -1), meaning it will reshape/keep the first dimension to be N, and then infer as necessary what the second dimension is in the output, so we can collapse the remaining dimensions from the input properly.

NOTE: TensorFlow and PyTorch differ on the default Tensor layout; TensorFlow uses $N \times H \times W \times C$ but PyTorch uses $N \times C \times H \times W$.

```
In []: def flatten(x):
    """
    Input:
        - TensorFlow Tensor of shape (N, D1, ..., DM)

    Output:
        - TensorFlow Tensor of shape (N, D1 * ... * DM)
        """
        N = tf.shape(x)[0]
        return tf.reshape(x, (N, -1))

In []: def test_flatten():
    # Construct concrete values of the input data x using numpy
        x_np = np.arange(24).reshape((2, 3, 4))
        print('x_np:\n', x_np, '\n')
        # Compute a concrete output value.
        x_flat_np = flatten(x_np)
        print('x_flat_np:\n', x_flat_np, '\n')

    test_flatten()
```

Barebones TensorFlow: Define a Two-Layer Network

We will now implement our first neural network with TensorFlow: a fullyconnected ReLU network with two hidden layers and no biases on the CIFAR10 dataset. For now we will use only low-level TensorFlow operators to define the network; later we will see how to use the higher-level abstractions provided by tf.keras to simplify the process.

We will define the forward pass of the network in the function two_layer_fc; this will accept TensorFlow Tensors for the inputs and weights of the network, and return a TensorFlow Tensor for the scores.

After defining the network architecture in the two_layer_fc function, we will test the implementation by checking the shape of the output.

It's important that you read and understand this implementation.

```
In [ ]: def two layer fc(x, params):
            A fully-connected neural network; the architecture is:
            fully-connected layer -> ReLU -> fully connected layer.
            Note that we only need to define the forward pass here; TensorFlow will
            care of computing the gradients for us.
            The input to the network will be a minibatch of data, of shape
            (N, d1, ..., dM) where d1 * ... * dM = D. The hidden layer will have H \iota
            and the output layer will produce scores for C classes.
            Inputs:
            - x: A TensorFlow Tensor of shape (N, d1, ..., dM) giving a minibatch of
             input data.
            - params: A list [w1, w2] of TensorFlow Tensors giving weights for the
              network, where w1 has shape (D, H) and w2 has shape (H, C).
            Returns:
            - scores: A TensorFlow Tensor of shape (N, C) giving classification score
             for the input data x.
            w1, w2 = params
                                            # Unpack the parameters
                                            # Flatten the input; now x has shape (
            x = flatten(x)
            h = tf.nn.relu(tf.matmul(x, w1))  # Hidden layer: h has shape (N, H)
            scores = tf.matmul(h, w2) # Compute scores of shape (N, C)
            return scores
In [ ]: def two layer fc test():
            hidden layer size = 42
            # Scoping our TF operations under a tf.device context manager
            # lets us tell TensorFlow where we want these Tensors to be
            # multiplied and/or operated on, e.g. on a CPU or a GPU.
            with tf.device(device):
                x = tf.zeros((64, 32, 32, 3))
                w1 = tf.zeros((32 * 32 * 3, hidden layer size))
                w2 = tf.zeros((hidden layer size, 10))
                # Call our two layer fc function for the forward pass of the network
                scores = two layer fc(x, [w1, w2])
```

```
print(scores.shape)

two_layer_fc_test()
```

Barebones TensorFlow: Three-Layer ConvNet

Here you will complete the implementation of the function

three_layer_convnet which will perform the forward pass of a three-layer
convolutional network. The network should have the following architecture:

- 1. A convolutional layer (with bias) with channel_1 filters, each with shape KW1 x KH1, and zero-padding of two
- 2. ReLU nonlinearity
- 3. A convolutional layer (with bias) with channel_2 filters, each with shape KW2 x KH2, and zero-padding of one
- 4. ReLU nonlinearity
- 5. Fully-connected layer with bias, producing scores for C classes.

HINT: For convolutions:

https://www.tensorflow.org/versions/r2.0/api_docs/python/tf/nn/conv2d; be careful with padding!

HINT: For biases: https://www.tensorflow.org/performance/xla/broadcasting

```
In [ ]: def three layer convnet(x, params):
          A three-layer convolutional network with the architecture described above
          Inputs:
           - x: A TensorFlow Tensor of shape (N, H, W, 3) giving a minibatch of ima
          - params: A list of TensorFlow Tensors giving the weights and biases for
            network; should contain the following:
            - conv w1: TensorFlow Tensor of shape (KH1, KW1, 3, channel 1) giving
              weights for the first convolutional layer.
            - conv bl: TensorFlow Tensor of shape (channel 1,) giving biases for t
              first convolutional layer.
            - conv w2: TensorFlow Tensor of shape (KH2, KW2, channel 1, channel 2)
              giving weights for the second convolutional layer
            - conv b2: TensorFlow Tensor of shape (channel 2,) giving biases for t
              second convolutional layer.
            - fc w: TensorFlow Tensor giving weights for the fully-connected layer
              Can you figure out what the shape should be?
            - fc b: TensorFlow Tensor giving biases for the fully-connected layer.
              Can you figure out what the shape should be?
          conv w1, conv b1, conv w2, conv b2, fc w, fc b = params
          scores = None
          # TODO: Implement the forward pass for the three-layer ConvNet.
```

After defing the forward pass of the three-layer ConvNet above, run the following cell to test your implementation. Like the two-layer network, we run the graph on a batch of zeros just to make sure the function doesn't crash, and produces outputs of the correct shape.

When you run this function, scores np should have shape (64, 10).

```
In []:
    with tf.device(device):
        x = tf.zeros((64, 32, 32, 3))
        conv_wl = tf.zeros((5, 5, 3, 6))
        conv_bl = tf.zeros((6,))
        conv_b2 = tf.zeros((3, 3, 6, 9))
        conv_b2 = tf.zeros((32 * 32 * 9, 10))
        fc_b = tf.zeros((10,))
        params = [conv_wl, conv_b1, conv_w2, conv_b2, fc_w, fc_b]
        scores = three_layer_convnet(x, params)

# Inputs to convolutional layers are 4-dimensional arrays with shape
    # [batch_size, height, width, channels]
        print('scores_np has shape: ', scores.shape)

three_layer_convnet_test()
```

Barebones TensorFlow: Training Step

We now define the training_step function performs a single training step. This will take three basic steps:

- 1. Compute the loss
- 2. Compute the gradient of the loss with respect to all network weights
- 3. Make a weight update step using (stochastic) gradient descent.

We need to use a few new TensorFlow functions to do all of this:

For computing the cross-entropy loss we'll use
 tf.nn.sparse_softmax_cross_entropy_with_logits:
 https://www.tensorflow.org/versions/r2.0/api docs/python/tf/nn/sparse softmax

For averaging the loss across a minibatch of data we'll use
 tf.reduce_mean:
 https://www.tensorflow.org/versions/r2.0/api docs/python/tf/reduce mean

- For computing gradients of the loss with respect to the weights we'll use tf.GradientTape (useful for Eager execution): https://www.tensorflow.org/versions/r2.0/api_docs/python/tf/GradientTape
- We'll mutate the weight values stored in a TensorFlow Tensor using tf.assign_sub ("sub" is for subtraction): https://www.tensorflow.org/api docs/python/tf/assign_sub

```
In [ ]: def training step(model fn, x, y, params, learning rate):
            with tf.GradientTape() as tape:
                scores = model fn(x, params) # Forward pass of the model
                loss = tf.nn.sparse softmax cross entropy with logits(labels=y, logi
                total loss = tf.reduce mean(loss)
                grad params = tape.gradient(total loss, params)
                # Make a vanilla gradient descent step on all of the model parameter
                # Manually update the weights using assign sub()
                for w, grad w in zip(params, grad params):
                    w.assign sub(learning rate * grad w)
                return total loss
In [ ]: def train part2(model fn, init fn, learning rate):
            Train a model on CIFAR-10.
            Inputs:
            - model fn: A Python function that performs the forward pass of the mode
              using TensorFlow; it should have the following signature:
              scores = model fn(x, params) where x is a TensorFlow Tensor giving a
              minibatch of image data, params is a list of TensorFlow Tensors holding
              the model weights, and scores is a TensorFlow Tensor of shape (N, C)
              giving scores for all elements of x.
            - init fn: A Python function that initializes the parameters of the mode
              It should have the signature params = init fn() where params is a list
              of TensorFlow Tensors holding the (randomly initialized) weights of the
              model.
            - learning rate: Python float giving the learning rate to use for SGD.
            params = init_fn() # Initialize the model parameters
            for t, (x np, y np) in enumerate(train dset):
                # Run the graph on a batch of training data.
                loss = training step(model fn, x np, y np, params, learning rate)
                # Periodically print the loss and check accuracy on the val set.
```

if t % print every == 0:

```
print('Iteration %d, loss = %.4f' % (t, loss))
                    check accuracy(val dset, x np, model fn, params)
In [ ]: def check accuracy(dset, x, model fn, params):
            Check accuracy on a classification model, e.g. for validation.
            Inputs:
            - dset: A Dataset object against which to check accuracy
            - x: A TensorFlow placeholder Tensor where input images should be fed
            - model fn: the Model we will be calling to make predictions on x
            - params: parameters for the model fn to work with
            Returns: Nothing, but prints the accuracy of the model
            num correct, num samples = 0, 0
            for x batch, y batch in dset:
                scores np = model fn(x batch, params).numpy()
                y pred = scores np.argmax(axis=1)
                num samples += x batch.shape[0]
                num correct += (y pred == y batch).sum()
            acc = float(num correct) / num samples
            print('Got %d / %d correct (%.2f%%)' % (num correct, num samples, 100 *
```

Barebones TensorFlow: Initialization

We'll use the following utility method to initialize the weight matrices for our models using Kaiming's normalization method.

[1] He et al, *Delving Deep into Rectifiers: Surpassing Human-Level Performance on ImageNet Classification *, ICCV 2015, https://arxiv.org/abs/1502.01852

```
In [ ]: def create_matrix_with_kaiming_normal(shape):
    if len(shape) == 2:
        fan_in, fan_out = shape[0], shape[1]
    elif len(shape) == 4:
        fan_in, fan_out = np.prod(shape[:3]), shape[3]
    return tf.keras.backend.random_normal(shape) * np.sqrt(2.0 / fan_in)
```

Barebones TensorFlow: Train a Two-Layer Network

We are finally ready to use all of the pieces defined above to train a two-layer fully-connected network on CIFAR-10.

We just need to define a function to initialize the weights of the model, and call train_part2.

Defining the weights of the network introduces another important piece of TensorFlow API: tf.Variable. A TensorFlow Variable is a Tensor whose value is stored in the graph and persists across runs of the computational graph;

however unlike constants defined with tf.zeros or tf.random_normal, the values of a Variable can be mutated as the graph runs; these mutations will persist across graph runs. Learnable parameters of the network are usually stored in Variables.

You don't need to tune any hyperparameters, but you should achieve validation accuracies above 40% after one epoch of training.

Barebones TensorFlow: Train a three-layer ConvNet

We will now use TensorFlow to train a three-layer ConvNet on CIFAR-10.

You need to implement the three_layer_convnet_init function. Recall that the architecture of the network is:

- 1. Convolutional layer (with bias) with 32 5x5 filters, with zero-padding 2
- 2. ReLU
- 3. Convolutional layer (with bias) with 16 3x3 filters, with zero-padding 1
- 4. ReLU
- 5. Fully-connected layer (with bias) to compute scores for 10 classes

You don't need to do any hyperparameter tuning, but you should see validation accuracies above 43% after one epoch of training.

```
Inputs: None
  Returns a list containing:
  - conv w1: TensorFlow tf. Variable giving weights for the first conv laye
  - conv b1: TensorFlow tf. Variable giving biases for the first conv layer
  - conv w2: TensorFlow tf. Variable giving weights for the second conv lay
  - conv b2: TensorFlow tf. Variable giving biases for the second conv layer

    fc w: TensorFlow tf.Variable giving weights for the fully-connected la

   - fc b: TensorFlow tf. Variable giving biases for the fully-connected lay
  params = None
  # TODO: Initialize the parameters of the three-layer network.
  # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
  pass
  # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
  END OF YOUR CODE
  return params
learning rate = 3e-3
train part2(three layer convnet, three layer convnet init, learning rate)
```

Part III: Keras Model Subclassing API

Implementing a neural network using the low-level TensorFlow API is a good way to understand how TensorFlow works, but it's a little inconvenient - we had to manually keep track of all Tensors holding learnable parameters. This was fine for a small network, but could quickly become unweildy for a large complex model.

Fortunately TensorFlow 2.0 provides higher-level APIs such as tf.keras which make it easy to build models out of modular, object-oriented layers. Further, TensorFlow 2.0 uses eager execution that evaluates operations immediately, without explicitly constructing any computational graphs. This makes it easy to write and debug models, and reduces the boilerplate code.

In this part of the notebook we will define neural network models using the tf.keras.Model API. To implement your own model, you need to do the following:

- 1. Define a new class which subclasses tf.keras.Model . Give your class an intuitive name that describes it, like TwoLayerFC or ThreeLayerConvNet .
- 2. In the initializer __init__() for your new class, define all the layers you need as class attributes. The tf.keras.layers package provides many

common neural-network layers, like tf.keras.layers.Dense for fullyconnected layers and tf.keras.layers.Conv2D for convolutional layers.

Under the hood, these layers will construct Variable Tensors for any
learnable parameters. Warning: Don't forget to call super(YourModelName,
self).__init__() as the first line in your initializer!

3. Implement the call() method for your class; this implements the forward pass of your model, and defines the *connectivity* of your network. Layers defined in __init__() implement __call__() so they can be used as function objects that transform input Tensors into output Tensors. Don't define any new layers in call(); any layers you want to use in the forward pass should be defined in __init__().

After you define your tf.keras.Model subclass, you can instantiate it and use it like the model functions from Part II.

Keras Model Subclassing API: Two-Layer Network

Here is a concrete example of using the tf.keras.Model API to define a two-layer network. There are a few new bits of API to be aware of here:

We use an Initializer object to set up the initial values of the learnable parameters of the layers; in particular tf.initializers.VarianceScaling gives behavior similar to the Kaiming initialization method we used in Part II. You can read more about it here:

https://www.tensorflow.org/versions/r2.0/api docs/python/tf/initializers/VarianceScali

We construct tf.keras.layers.Dense objects to represent the two fully-connected layers of the model. In addition to multiplying their input by a weight matrix and adding a bias vector, these layer can also apply a nonlinearity for you. For the first layer we specify a ReLU activation function by passing activation='relu' to the constructor; the second layer uses softmax activation function. Finally, we use tf.keras.layers.Flatten to flatten the output from the previous fully-connected layer.

```
x = self.fc2(x)
return x

def test_TwoLayerFC():
    """ A small unit test to exercise the TwoLayerFC model above. """
    input_size, hidden_size, num_classes = 50, 42, 10
    x = tf.zeros((64, input_size))
    model = TwoLayerFC(hidden_size, num_classes)
    with tf.device(device):
        scores = model(x)
        print(scores.shape)

test_TwoLayerFC()
```

Keras Model Subclassing API: Three-Layer ConvNet

Now it's your turn to implement a three-layer ConvNet using the tf.keras.Model API. Your model should have the same architecture used in Part II:

- 1. Convolutional layer with 5 x 5 kernels, with zero-padding of 2
- 2. ReLU nonlinearity
- 3. Convolutional layer with 3 x 3 kernels, with zero-padding of 1
- 4. ReLU nonlinearity
- 5. Fully-connected layer to give class scores
- 6. Softmax nonlinearity

You should initialize the weights of your network using the same initialization method as was used in the two-layer network above.

Hint: Refer to the documentation for tf.keras.layers.Conv2D and tf.keras.layers.Dense:

https://www.tensorflow.org/versions/r2.0/api_docs/python/tf/keras/layers/Conv2D

https://www.tensorflow.org/versions/r2.0/api_docs/python/tf/keras/layers/Dense

Once you complete the implementation of the ThreeLayerConvNet above you can run the following to ensure that your implementation does not crash and produces outputs of the expected shape.

```
In [ ]:
    def test_ThreeLayerConvNet():
        channel_1, channel_2, num_classes = 12, 8, 10
        model = ThreeLayerConvNet(channel_1, channel_2, num_classes)
        with tf.device(device):
            x = tf.zeros((64, 3, 32, 32))
            scores = model(x)
            print(scores.shape)

test_ThreeLayerConvNet()
```

Keras Model Subclassing API: Eager Training

While keras models have a builtin training loop (using the model.fit), sometimes you need more customization. Here's an example, of a training loop implemented with eager execution.

In particular, notice tf.GradientTape . Automatic differentiation is used in the backend for implementing backpropagation in frameworks like TensorFlow.

During eager execution, tf.GradientTape is used to trace operations for computing gradients later. A particular tf.GradientTape can only compute one gradient; subsequent calls to tape will throw a runtime error.

TensorFlow 2.0 ships with easy-to-use built-in metrics under tf.keras.metrics module. Each metric is an object, and we can use update_state() to add observations and reset_state() to clear all observations. We can get the current result of a metric by calling result() on the metric object.

```
In [ ]: def train part34(model init fn, optimizer init fn, num epochs=1, is training
            Simple training loop for use with models defined using tf.keras. It trai
            a model for one epoch on the CIFAR-10 training set and periodically check
            accuracy on the CIFAR-10 validation set.
            Inputs:
            - model init fn: A function that takes no parameters; when called it
              constructs the model we want to train: model = model init fn()
            - optimizer init fn: A function which takes no parameters; when called i
              constructs the Optimizer object we will use to optimize the model:
              optimizer = optimizer init fn()
            - num epochs: The number of epochs to train for
            Returns: Nothing, but prints progress during trainingn
            with tf.device(device):
                # Compute the loss like we did in Part II
                loss fn = tf.keras.losses.SparseCategoricalCrossentropy()
                model = model init fn()
                optimizer = optimizer init fn()
                train loss = tf.keras.metrics.Mean(name='train loss')
                train accuracy = tf.keras.metrics.SparseCategoricalAccuracy(name='tr
                val loss = tf.keras.metrics.Mean(name='val loss')
                val accuracy = tf.keras.metrics.SparseCategoricalAccuracy(name='val
                t = 0
                for epoch in range(num epochs):
                    # Reset the metrics - https://www.tensorflow.org/alpha/guide/mig
                    train loss.reset states()
                    train accuracy.reset states()
                    for x_np, y_np in train_dset:
                        with tf.GradientTape() as tape:
                            # Use the model function to build the forward pass.
                            scores = model(x np, training=is training)
                            loss = loss fn(y np, scores)
                            gradients = tape.gradient(loss, model.trainable variable
                            optimizer.apply gradients(zip(gradients, model.trainable
                            # Update the metrics
                            train loss.update state(loss)
                            train accuracy.update state(y np, scores)
                            if t % print every == 0:
                                val loss.reset states()
                                val accuracy.reset states()
                                for test_x, test_y in val_dset:
```

Keras Model Subclassing API: Train a Two-Layer Network

We can now use the tools defined above to train a two-layer network on CIFAR-10. We define the <code>model_init_fn</code> and <code>optimizer_init_fn</code> that construct the model and optimizer respectively when called. Here we want to train the model using stochastic gradient descent with no momentum, so we construct a <code>tf.keras.optimizers.SGD</code> function; you can read about it here.

You don't need to tune any hyperparameters here, but you should achieve validation accuracies above 40% after one epoch of training.

```
In []: hidden_size, num_classes = 4000, 10
learning_rate = 1e-2

def model_init_fn():
    return TwoLayerFC(hidden_size, num_classes)

def optimizer_init_fn():
    return tf.keras.optimizers.SGD(learning_rate=learning_rate)

train_part34(model_init_fn, optimizer_init_fn)
```

Keras Model Subclassing API: Train a Three-Layer ConvNet

Here you should use the tools we've defined above to train a three-layer ConvNet on CIFAR-10. Your ConvNet should use 32 filters in the first convolutional layer and 16 filters in the second layer.

To train the model you should use gradient descent with Nesterov momentum 0.9.

HINT:

https://www.tensorflow.org/versions/r2.0/api docs/python/tf/optimizers/SGD

You don't need to perform any hyperparameter tuning, but you should achieve validation accuracies above 50% after training for one epoch.

```
In [ ]: learning rate = 3e-3
    channel 1, channel 2, num classes = 32, 16, 10
    def model init fn():
      model = None
      # TODO: Complete the implementation of model fn.
      # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
      pass
      # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
      END OF YOUR CODE
      return model
    def optimizer init fn():
      optimizer = None
      # TODO: Complete the implementation of model fn.
      # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
      pass
      # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
      END OF YOUR CODE
      return optimizer
    train part34(model init fn, optimizer init fn)
```

Part IV: Keras Sequential API

In Part III we introduced the tf.keras.Model API, which allows you to define models with any number of learnable layers and with arbitrary connectivity between layers.

However for many models you don't need such flexibility - a lot of models can be expressed as a sequential stack of layers, with the output of each layer fed to the next layer as input. If your model fits this pattern, then there is an even easier way to define your model: using tf.keras.Sequential. You don't need to write any custom classes; you simply call the tf.keras.Sequential constructor with a list containing a sequence of layer objects.

One complication with tf.keras.Sequential is that you must define the shape of the input to the model by passing a value to the input_shape of the first layer in your model.

Keras Sequential API: Two-Layer Network

In this subsection, we will rewrite the two-layer fully-connected network using the training loop defined above.

You don't need to perform any hyperparameter tuning here, but you should see validation accuracies above 40% after training for one epoch.

```
In [ ]: learning rate = 1e-2
        def model init fn():
            input shape = (32, 32, 3)
            hidden layer size, num classes = 4000, 10
            initializer = tf.initializers.VarianceScaling(scale=2.0)
            layers = [
                tf.keras.layers.Flatten(input shape=input shape),
                tf.keras.layers.Dense(hidden layer size, activation='relu',
                                       kernel initializer=initializer),
                tf.keras.layers.Dense(num_classes, activation='softmax',
                                      kernel initializer=initializer),
            ]
            model = tf.keras.Sequential(layers)
            return model
        def optimizer init fn():
            return tf.keras.optimizers.SGD(learning rate=learning rate)
        train part34(model init fn, optimizer init fn)
```

Abstracting Away the Training Loop

In the previous examples, we used a customised training loop to train models (e.g. train_part34). Writing your own training loop is only required if you need more flexibility and control during training your model. Alternately, you can also use built-in APIs like tf.keras.Model.fit() and tf.keras.Model.evaluate to train and evaluate a model. Also remember to configure your model for training by calling `tf.keras.Model.compile.

You don't need to perform any hyperparameter tuning here, but you should see validation and test accuracies above 42% after training for one epoch.

```
model.fit(X_train, y_train, batch_size=64, epochs=1, validation_data=(X_val, model.evaluate(X_test, y_test)
```

Keras Sequential API: Three-Layer ConvNet

Here you should use tf.keras.Sequential to reimplement the same three-layer ConvNet architecture used in Part II and Part III. As a reminder, your model should have the following architecture:

- 1. Convolutional layer with 32 5x5 kernels, using zero padding of 2
- 2. ReLU nonlinearity
- 3. Convolutional layer with 16 3x3 kernels, using zero padding of 1
- 4. ReLU nonlinearity
- 5. Fully-connected layer giving class scores
- 6. Softmax nonlinearity

You should initialize the weights of the model using a tf.initializers.VarianceScaling as above.

You should train the model using Nesterov momentum 0.9.

You don't need to perform any hyperparameter search, but you should achieve accuracy above 45% after training for one epoch.

```
In [ ]: def model init fn():
      model = None
      # TODO: Construct a three-layer ConvNet using tf.keras.Seguential.
      # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
      pass
      # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
      END OF YOUR CODE
      return model
    learning rate = 5e-4
    def optimizer init fn():
      optimizer = None
      # TODO: Complete the implementation of model fn.
      # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
      pass
      # ****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
```

We will also train this model with the built-in training loop APIs provided by TensorFlow.

Part IV: Functional API

Demonstration with a Two-Layer Network

In the previous section, we saw how we can use tf.keras.Sequential to stack layers to quickly build simple models. But this comes at the cost of losing flexibility.

Often we will have to write complex models that have non-sequential data flows: a layer can have **multiple inputs and/or outputs**, such as stacking the output of 2 previous layers together to feed as input to a third! (Some examples are residual connections and dense blocks.)

In such cases, we can use Keras functional API to write models with complex topologies such as:

- 1. Multi-input models
- 2. Multi-output models
- 3. Models with shared layers (the same layer called several times)
- 4. Models with non-sequential data flows (e.g. residual connections)

Writing a model with Functional API requires us to create a tf.keras.Model instance and explicitly write input tensors and output tensors for this model.

```
# Instantiate the model given inputs and outputs.
model = tf.keras.Model(inputs=inputs, outputs=scores)
return model

def test_two_layer_fc_functional():
    """ A small unit test to exercise the TwoLayerFC model above. """
    input_size, hidden_size, num_classes = 50, 42, 10
    input_shape = (50,)

x = tf.zeros((64, input_size))
model = two_layer_fc_functional(input_shape, hidden_size, num_classes)

with tf.device(device):
    scores = model(x)
    print(scores.shape)

test_two_layer_fc_functional()
```

Keras Functional API: Train a Two-Layer Network

You can now train this two-layer network constructed using the functional API.

You don't need to perform any hyperparameter tuning here, but you should see validation accuracies above 40% after training for one epoch.

```
In []: input_shape = (32, 32, 3)
    hidden_size, num_classes = 4000, 10
    learning_rate = le-2

def model_init_fn():
    return two_layer_fc_functional(input_shape, hidden_size, num_classes)

def optimizer_init_fn():
    return tf.keras.optimizers.SGD(learning_rate=learning_rate)

train_part34(model_init_fn, optimizer_init_fn)
```

Part V: CIFAR-10 open-ended challenge

In this section you can experiment with whatever ConvNet architecture you'd like on CIFAR-10.

You should experiment with architectures, hyperparameters, loss functions, regularization, or anything else you can think of to train a model that achieves **at least 70%** accuracy on the **validation** set within 10 epochs. You can use the built-in train function, the train_part34 function from above, or implement your own training loop.

Describe what you did at the end of the notebook.

Some things you can try:

- **Filter size**: Above we used 5x5 and 3x3; is this optimal?
- **Number of filters**: Above we used 16 and 32 filters. Would more or fewer do better?
- **Pooling**: We didn't use any pooling above. Would this improve the model?
- Normalization: Would your model be improved with batch normalization, layer normalization, group normalization, or some other normalization strategy?
- **Network architecture**: The ConvNet above has only three layers of trainable parameters. Would a deeper model do better?
- **Global average pooling**: Instead of flattening after the final convolutional layer, would global average pooling do better? This strategy is used for example in Google's Inception network and in Residual Networks.
- **Regularization**: Would some kind of regularization improve performance? Maybe weight decay or dropout?

NOTE: Batch Normalization / Dropout

If you are using Batch Normalization and Dropout, remember to pass is_training=True if you use the train_part34() function. BatchNorm and Dropout layers have different behaviors at training and inference time.

training is a specific keyword argument reserved for this purpose in any tf.keras.Model 's call() function. Read more about this here:

https://www.tensorflow.org/versions/r2.0/api_docs/python/tf/keras/layers/BatchNormahttps://www.tensorflow.org/versions/r2.0/api_docs/python/tf/keras/layers/Dropout#m

Tips for training

For each network architecture that you try, you should tune the learning rate and other hyperparameters. When doing this there are a couple important things to keep in mind:

- If the parameters are working well, you should see improvement within a few hundred iterations
- Remember the coarse-to-fine approach for hyperparameter tuning: start by testing a large range of hyperparameters for just a few training iterations to find the combinations of parameters that are working at all.
- Once you have found some sets of parameters that seem to work, search more finely around these parameters. You may need to train for more epochs.
- You should use the validation set for hyperparameter search, and save your test set for evaluating your architecture on the best parameters as selected by the validation set.

Going above and beyond

If you are feeling adventurous there are many other features you can implement to try and improve your performance. You are **not required** to implement any of these, but don't miss the fun if you have time!

- Alternative optimizers: you can try Adam, Adagrad, RMSprop, etc.
- Alternative activation functions such as leaky ReLU, parametric ReLU, ELU, or MaxOut.
- Model ensembles
- Data augmentation
- New Architectures
 - ResNets where the input from the previous layer is added to the output.
 - DenseNets where inputs into previous layers are concatenated together.
 - This blog has an in-depth overview

Have fun and happy training!

```
In [ ]: class CustomConvNet(tf.keras.Model):
      def init (self):
        super(CustomConvNet, self). init ()
        # TODO: Construct a model that performs well on CIFAR-10
        # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
        pass
        # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)****
        END OF YOUR CODE
        def call(self, input tensor, training=False):
        # TODO: Construct a model that performs well on CIFAR-10
        # *****START OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
        pass
        # *****END OF YOUR CODE (DO NOT DELETE/MODIFY THIS LINE)*****
        END OF YOUR CODE
        return x
    print every = 700
    num epochs = 10
```

```
model = CustomConvNet()

def model_init_fn():
    return CustomConvNet()

def optimizer_init_fn():
    learning_rate = 1e-3
    return tf.keras.optimizers.Adam(learning_rate)

train_part34(model_init_fn, optimizer_init_fn, num_epochs=num_epochs, is_train_part34(model_init_fn, optimizer_init_fn, num_epochs=num_epochs, is_train_epochs.)
```

Describe what you did

In the cell below you should write an explanation of what you did, any additional features that you implemented, and/or any graphs that you made in the process of training and evaluating your network.

Answer:

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