# **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 which was proposed by [1] in 2015.

The idea is relatively straightforward. Machine learning methods tend to work better when their input data consists 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, [1] proposes to insert batch normalization layers into the network. At training time, a batch normalization 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] <u>Sergey Ioffe and Christian Szegedy, "Batch Normalization: Accelerating Deep Network Training by Reducing Internal Covariate Shift", ICML 2015.</u>
(<a href="https://arxiv.org/abs/1502.03167">https://arxiv.org/abs/1502.03167</a>)

```
In [1]:
```

```
# As usual, a bit of setup
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, eva
l numerical gradient array
from cs231n.solver import Solver
%matplotlib inline
plt.rcParams['figure.figsize'] = (10.0, 8.0) # set default siz
e 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-ipython
%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(' means: ', x.mean(axis=axis))
    print(' stds: ', x.std(axis=axis))
    print()
```

```
In [2]:
```

```
# Load the (preprocessed) CIFAR10 data.
data = get_CIFAR10_data()
for k, v in data.items():
   print('%s: ' % 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,)
```

### **Batch normalization: forward**

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.

Referencing the paper linked to above in [1] may be helpful!

In [3]:

```
# Check the training-time forward pass by checking means and v
ariances
# 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=', b
eta, ')')
a norm, = batchnorm forward(a, gamma, beta, {'mode': 'train'
})
print mean std(a norm,axis=0)
```

```
In [4]:
```

stds:

```
# Check the test-time forward pass by running the training-tim
e
# forward pass many times to warm up the running averages, and
# 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 wil
1 be
# noisier than training-time forward passes.
print('After batch normalization (test-time):')
print mean std(a norm,axis=0)
After batch normalization (test-time):
```

[-0.03927354 -0.04349152 -0.10452688]

[1.01531428 1.01238373 0.97819988]

### **Batch normalization: backward**

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 [5]:

```
# 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]
fq = 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))
```

```
dx error: 1.7029258328157158e-09
dgamma error: 7.420414216247087e-13
dbeta error: 2.8795057655839487e-12
```

# **Batch normalization: alternative backward**

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

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

$$\sigma = \sqrt{v + \epsilon} \qquad y_i = \frac{x_i - \mu}{\sigma}$$

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/hart 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  $\frac{\partial Y}{\partial \mu}$ , 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 [6]:

```
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)))
```

dx difference: 9.20004371222927e-13

dgamma difference: 0.0 dbeta difference: 0.0

speedup: 1.56x

# Fully Connected Nets 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 . If you decide to do so, do it in the file cs231n/classifiers/fc\_net.py .

#### In [7]:

```
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, h=1e-5)
    print('%s relative error: %.2e' % (name, rel error(grad nu
m, grads[name])))
  if reg == 0: print()
```

```
Running check with reg = 0
Initial loss: 2.2611955101340957
W1 relative error: 1.10e-04
W2 relative error: 5.65e-06
W3 relative error: 4.14e-10
b1 relative error: 2.22e-08
b2 relative error: 5.55e-09
b3 relative error: 1.02e-10
beta1 relative error: 7.33e-09
beta2 relative error: 1.17e-09
gamma1 relative error: 7.47e-09
gamma2 relative error: 3.35e-09
Running check with reg = 3.14
Initial loss: 6.996533220108303
W1 relative error: 1.98e-06
W2 relative error: 2.28e-06
W3 relative error: 1.11e-08
b1 relative error: 5.55e-09
b2 relative error: 2.22e-08
b3 relative error: 2.10e-10
betal relative error: 6.32e-09
beta2 relative error: 3.48e-09
gamma1 relative error: 6.27e-09
gamma2 relative error: 4.14e-09
```

# Batchnorm 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 [8]:
```

```
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'],
}
weight scale = 2e-2
bn model = FullyConnectedNet(hidden dims, weight scale=weight
scale, normalization='batchnorm')
model = FullyConnectedNet(hidden dims, weight scale=weight sca
le, normalization=None)
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()
Solver with batch norm:
```

```
(Iteration 1 / 200) loss: 2.340975

(Epoch 0 / 10) train acc: 0.107000; val_acc: 0.115

000

(Epoch 1 / 10) train acc: 0.313000; val_acc: 0.266

000

(Iteration 21 / 200) loss: 2.039365
```

```
(Epoch 2 / 10) train acc: 0.390000; val acc: 0.278
000
(Iteration 41 / 200) loss: 2.036710
(Epoch 3 / 10) train acc: 0.497000; val acc: 0.315
000
(Iteration 61 / 200) loss: 1.769537
(Epoch 4 / 10) train acc: 0.529000; val acc: 0.321
000
(Iteration 81 / 200) loss: 1.265761
(Epoch 5 / 10) train acc: 0.592000; val acc: 0.318
000
(Iteration 101 / 200) loss: 1.261353
(Epoch 6 / 10) train acc: 0.657000; val acc: 0.321
000
(Iteration 121 / 200) loss: 1.101407
(Epoch 7 / 10) train acc: 0.689000; val_acc: 0.318
000
(Iteration 141 / 200) loss: 1.193775
(Epoch 8 / 10) train acc: 0.686000; val acc: 0.298
000
(Iteration 161 / 200) loss: 0.796336
(Epoch 9 / 10) train acc: 0.779000; val acc: 0.320
000
(Iteration 181 / 200) loss: 0.873413
(Epoch 10 / 10) train acc: 0.774000; val acc: 0.32
5000
Solver without batch norm:
(Iteration 1 / 200) loss: 2.302332
(Epoch 0 / 10) train acc: 0.129000; val acc: 0.131
000
(Epoch 1 / 10) train acc: 0.283000; val acc: 0.250
000
(Iteration 21 / 200) loss: 2.041970
(Epoch 2 / 10) train acc: 0.316000; val acc: 0.277
000
(Iteration 41 / 200) loss: 1.900473
(Epoch 3 / 10) train acc: 0.373000; val acc: 0.282
000
(Iteration 61 / 200) loss: 1.713156
(Epoch 4 / 10) train acc: 0.390000; val acc: 0.310
000
(Iteration 81 / 200) loss: 1.662209
(Epoch 5 / 10) train acc: 0.434000; val acc: 0.300
000
(Iteration 101 / 200) loss: 1.696059
(Epoch 6 / 10) train acc: 0.535000; val acc: 0.345
000
```

```
(Iteration 121 / 200) loss: 1.557986

(Epoch 7 / 10) train acc: 0.530000; val_acc: 0.304

000

(Iteration 141 / 200) loss: 1.432189

(Epoch 8 / 10) train acc: 0.628000; val_acc: 0.339

000

(Iteration 161 / 200) loss: 1.033932

(Epoch 9 / 10) train acc: 0.661000; val_acc: 0.340

000

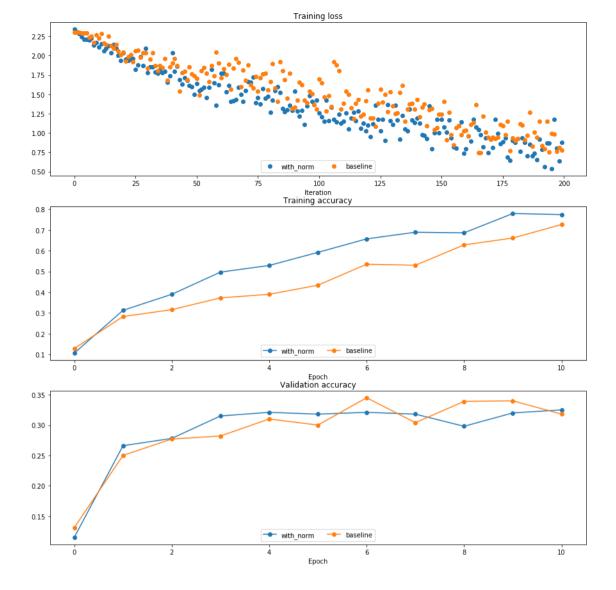
(Iteration 181 / 200) loss: 0.901035

(Epoch 10 / 10) train acc: 0.726000; val_acc: 0.31
```

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 [9]:
```

```
def plot training history(title, label, baseline, bn solvers,
plot fn, bl marker='.', bn marker='.', labels=None):
    """utility function for plotting training history"""
    plt.title(title)
    plt.xlabel(label)
    bn plots = [plot fn(bn solver) for bn solver in bn solvers
1
    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 marker='-o')
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='-o')
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.

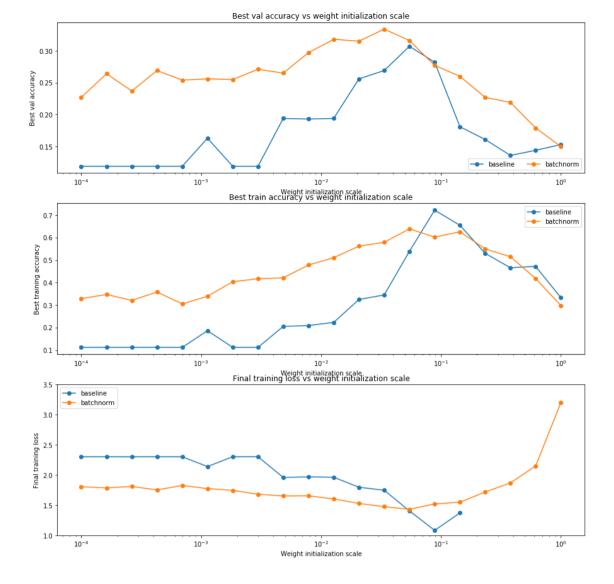
The first cell will train 8-layer networks both with and without batch normalization using different scales for weight initialization. The second layer will plot training accuracy, validation set accuracy, and training loss as a function of the weight initialization scale.

```
In [10]:
```

```
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 sc
ales)))
 bn model = FullyConnectedNet(hidden dims, weight scale=weigh
t scale, normalization='batchnorm')
 model = FullyConnectedNet(hidden dims, weight scale=weight s
cale, normalization=None)
  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
```

```
Running weight scale 1 / 20
Running weight scale 2 / 20
Running weight scale 3 / 20
Running weight scale 4 / 20
Running weight scale 5 / 20
Running weight scale 6 / 20
Running weight scale 7 / 20
Running weight scale 8 / 20
Running weight scale 9 / 20
Running weight scale 10 / 20
Running weight scale 11 / 20
Running weight scale 12 / 20
Running weight scale 13 / 20
Running weight scale 14 / 20
Running weight scale 15 / 20
Running weight scale 16 / 20
/home/ec2-user/workspace/Clean task/assignment2/cs
231n/layers.py:849: RuntimeWarning: divide by zero
encountered in log
  loss = -np.sum(np.log(probs[np.arange(N), y])) /
Ν
Running weight scale 17 / 20
Running weight scale 18 / 20
Running weight scale 19 / 20
Running weight scale 20 / 20
In [11]:
# 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_hi
story))
  best val accs.append(max(solvers ws[ws].val acc history))
  bn best val accs.append(max(bn solvers ws[ws].val acc histor
y))
  final train loss.append(np.mean(solvers ws[ws].loss history[
-100:1)
  bn final train loss.append(np.mean(bn solvers ws[ws].loss hi
```

```
story[-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='baseli
ne')
plt.semilogx(weight scales, bn best val accs, '-o', label='bat
chnorm')
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='base
line')
plt.semilogx(weight scales, bn best train accs, '-o', label='b
atchnorm')
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='bas
eline')
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 scale of weight initialization affect models with/without batch normalization differently, and why?

### **Answer:**

The second plot shows the problem of vanishing gradients (small initial weights). The baseline model is very sensitive to this problem (the accuracy is very low), therefore finding the correct weight scale is difficult. For this example, the baseline obtains the best result with a weight scale equal to 1e-1. On the other hand, we can see that the batchnorm model is less sensitive to weight initialization because its accuracy is around 30% for all the different weight scales.

The behaviour of the first plot is very similar to that of the second plot. The main difference is that the first plot shows that we are overfitting our model, besides that we can see that with the batchnorm model we obtained better results than the baseline model and that occurs because batch normalization has regularization properties.

The third plot depicts the problem of exploding gradients and it is very evident in the baseline model for weight scale values greater than 1e-1. However, the batchnorm model does not suffer from this problem.

# 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 [12]:
```

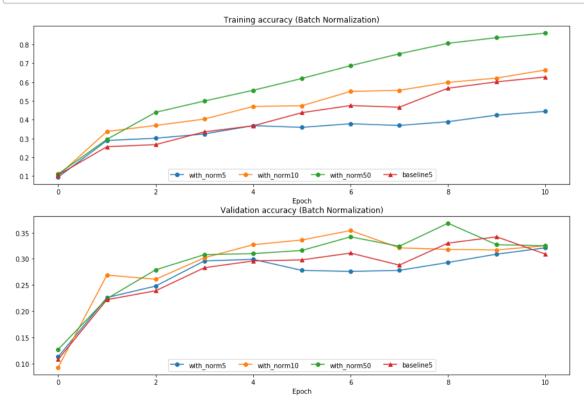
```
def run_batchsize_experiments(normalization_mode):
    np.random.seed(231)
    # Try training a very deep net with batchnorm
    hidden_dims = [100, 100, 100, 100]
```

```
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'],
    }
    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, normalization=None)
    solver = Solver(model, small data,
                    num epochs=n epochs, batch size=solver bsi
ze,
                    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, normalization=normalization mode)
        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)
        bn solver.train()
        bn solvers.append(bn solver)
    return bn solvers, solver, batch sizes
batch sizes = [5,10,50]
bn solvers bsize, solver bsize, batch sizes = run batchsize ex
```

periments('batchnorm')

No normalization: batch size = 5 Normalization: batch size = 5 Normalization: batch size = 10 Normalization: batch size = 50

#### In [13]:



# **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:**

According to the results, we can see that the batch size affects directly the performance of batch normalization (the smaller the batch size the worse). Even the baseline model outperforms the batchnorm model when using a very small batch size. This problem occurs because when we calculate the statistics of a batch, i.e., mean and variance, we try to find an approximation of the statistics of the entire dataset. Therefore with a small batch size, these statistics can be very noisy. On the other hand, with a large batch size we can obtain a better approximation.

# **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] <u>Ba, Jimmy Lei, Jamie Ryan Kiros, and Geoffrey E. Hinton. "Layer Normalization."</u> stat 1050 (2016): 21. (https://arxiv.org/pdf/1607.06450.pdf)

# **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:**

Number 2 is analogous to layer normalization when we consider: mean = 0, beta parameter = 0 (at this point we have gamma\*x/std where std=sqrt(sum( $x^2$ ))) and gamma=x/std. Thus the result of layer normalization will be  $x^2/sum(x^2)$ .

Number 3 is analogous to batch normalization when we consider: batch size = size of the dataset, gamma parameter = standard deviation and beta parameter = 0. Thus the result of batch normalization will be  $std^*(x-mean)/std + 0 = x-mean$ .

# **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 backward.

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.

In [14]:

```
# Check the training-time forward pass by checking means and v
ariances
# of features both before and after 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)
gamma = 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=', b
eta, ')')
a norm, = layernorm forward(a, gamma, beta, {'mode': 'train'
})
print_mean_std(a norm,axis=1)
```

```
Before layer normalization:
 means: [-59.06673243 -47.60782686 -43.31137368
-26.40991744]
  stds: [10.07429373 28.39478981 35.28360729 4.
01831507]
After layer normalization (gamma=1, beta=0)
 means: [ 4.81096644e-16 -7.40148683e-17 2.2204
4605e-16 -5.92118946e-16]
 stds: [0.99999995 0.99999999 1.
                                        0.9999
99691
After layer normalization (gamma= [3. 3. 3.], bet
a = [5.5.5.]
 means: [5. 5. 5. 5.]
 stds: [2.99999985 2.99999998 2.9999999 2.9999
99071
```

```
# 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
print('dx error: ', rel error(dx num, dx))
print('dgamma error: ', rel_error(da_num, dgamma))
print('dbeta error: ', rel_error(db_num, dbeta))
```

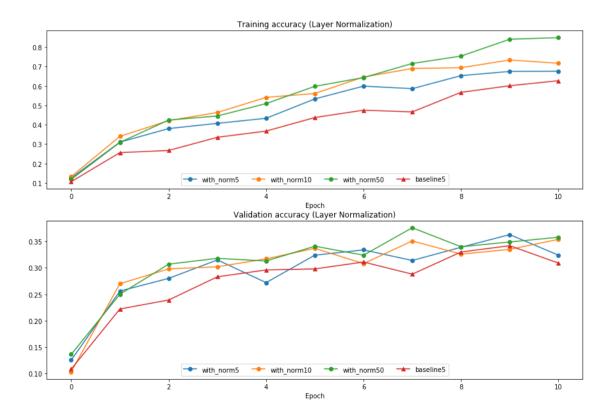
```
dx error: 1.433615146847572e-09
dgamma error: 4.519489546032799e-12
dbeta error: 2.276445013433725e-12
```

# 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!

#### In [16]:

No normalization: batch size = 5 Normalization: batch size = 5 Normalization: batch size = 10 Normalization: batch size = 50



# **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:**

- [INCORRECT] In the previous example, the network had five layers and it can be considered as a deep network. Thus, using layer normalization in deep networks works correctly.
- 2. [CORRECT] Having a small dimension of features affects the performance of layer normalization. The problem is very similar to that of batch normalization with small batch size because in layer normalization we calculate the statistics according to the number of hidden units, which represent the features that the network is learning. Thus, the smaller the hidden size the noisier the statistics used in layer normalization.
- 3. [CORRECT] Having a high regularization term affects the performance of layer normalization. In general, when the regularization term is very high, the model learns very simple functions (underfitting).

Tests showing the performance of layer normalization with different hidden sizes and regularization values can be seen in the next cells.

	In [ ]:				
ı					