
DEEP LEARNING STUDY NOTES

ALL CREDITS GO TO **L. Fei-Fei, A. Karpathy, J. Johnson** TEACHERS OF THE CS231N COURSE. THANK YOU FOR THIS AMAZING COURSE!!

BY

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Part I

DATA

Chapter 1

Data Preprocessing

There are different techniques used in learning in order to improve the accuracy of the model by preprocessing the data:

There are three common forms of data preprocessing a data matrix X , where we will assume that X is of size $[N \times D]$ (N is the number of data, D is their dimensionality).

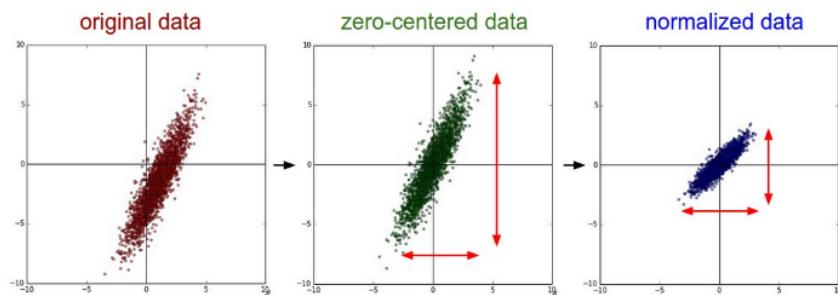


FIGURE 1.1: Preprocessing example 1

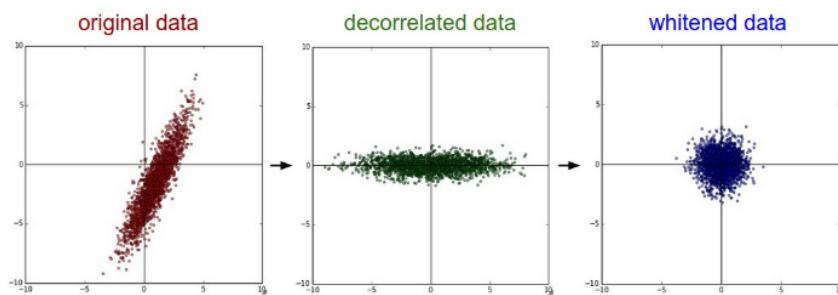


FIGURE 1.2: Preprocessing example 2

Mean subtraction Most common form of preprocessing. It involves subtracting the mean across every individual feature in the data, and has the geometric interpretation of centring the cloud of data around the origin along every dimension. In NumPy, this operation would be implemented as: `X -= np.mean(X, axis = 0)`. With images specifically, for convenience it

can be common to subtract a single value from all pixels (e.g. `X -= np.mean(X)`), or to do so separately across the three color channels.

Normalization Refers to normalizing the data dimensions so that they are of approximately the same scale. There are two common ways of achieving this normalization. One is to divide each dimension by its standard deviation, once it has been zero-centered: (`X /= np.std(X, axis = 0)`). Another form of this preprocessing normalizes each dimension so that the min and max along the dimension is -1 and 1 respectively. It only makes sense to apply this preprocessing if you have a reason to believe that different input features have different scales (or units), but they should be of approximately equal importance to the learning algorithm. In case of images, the relative scales of pixels are already approximately equal (and in range from 0 to 255), so it is not strictly necessary to perform this additional preprocessing step.

PCA The reason why one would want to use PCA is if one expects that many of the features are in fact dependent. This would be particularly handy for Naive Bayes where independence is assumed. Most datasets are far too large to use PCA. Attention: PCA complexity is $O(n^3)$, so more sophisticated methods are required. But if your dataset is small, and you don't have the time to investigate more sophisticated methods, then by all means go ahead and apply an out-of-box PCA for feature selection.

Whitening Takes the data in the eigenbasis and divides every dimension by the eigenvalue to normalize the scale. The geometric interpretation of this transformation is that if the input data is a multi-variable Gaussian, then the whitened data will be a Gaussian with zero mean and identity covariance matrix.

Deep Learning with images

For Deep Learning for images we will only use: center our data to zero. To do so, for each pixel, compute its mean across all the dataset and subtract the resulting mean image to all the training samples. If you have more than one channel (e.g. RGB) do it for each of the channels separately.

Chapter 2

Making the most of your data - Data Augmentation and Transfer Learning

2.1 Data Augmentation

Horizontal Flip

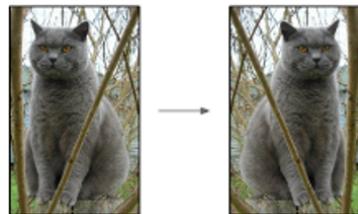


FIGURE 2.1: Horizontal flip. Flip image 180 degrees in the horizontal direction.

Random Crops/Scales

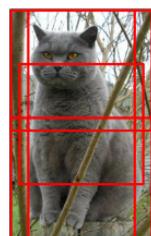


FIGURE 2.2: Random Crops/Scales

Training: sample random crops / scales

Specific example. How ResNet trains:

1. Pick random L in range [256, 480]
2. Resize training image, short side = L
3. Sample random 224 x 224 patch

Testing: average a fixed set of crops

Specific example. How ResNet tests:

1. Resize image at 5 scales: 224, 256, 384, 480, 640
2. For each size, use 10 224 x 224 crops: 4 corners + center, + flips

Color Jitter

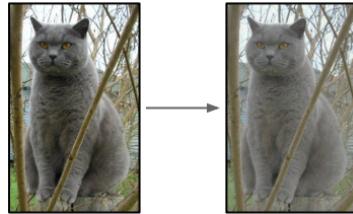


FIGURE 2.3: Color Jitter

- Simple: Randomly jitter contrast
- Complex:
 1. Apply PCA to all [R, G, B] pixels in training set
 2. Sample a color offset along principal component directions
 3. Add offset to all pixels of a training image

Get Creative

At the end you have to think for your specific case, what transformations you want to be robust to and add them to the training set: translation, rotation, stretching, shearing, lens distortions, ...

2.2 Transfer Learning

In practice, very few people train an entire Deep Network from scratch (with random initialization), because it is relatively rare to have a dataset of sufficient size. Instead, it is common to pretrain a ConvNet on a very large dataset (e.g. ImageNet, which contains 1.2 million images with 1000 categories), and then use the ConvNet either as an initialization or a fixed feature extractor for the task of interest. Pretrained DeepNets are usually trained with large computer clusters and downloaded by others. The three major Transfer Learning scenarios look as follows:

ConvNet as fixed feature extractor Take a ConvNet pretrained on ImageNet, remove the last fully-connected layer (this layers outputs are the 1000 class scores for a different task like ImageNet), then treat the rest of the ConvNet as a fixed feature extractor for the new dataset. In an AlexNet, this would compute a 4096-D vector for every image that contains the activations of the hidden layer immediately before the classifier. We call these features CNN codes.

Fine-tuning the ConvNet The second strategy is not only to replace and retrain the classifier on top of the ConvNet on the new dataset, but to also fine-tune the weights of the pretrained network by continuing the backpropagation. It is possible to fine-tune all the layers of the ConvNet, or its possible to keep some of the earlier layers fixed (due to overfitting concerns) and only fine-tune some higher-level portion of the network. This is motivated by the observation that the earlier features of a ConvNet contain more generic features (e.g. edge detectors or color blob detectors) that should be useful to many tasks, but later layers of the ConvNet becomes progressively more specific to the details of the classes contained in the original dataset. In case of ImageNet for example, which contains many dog breeds, a significant portion of the representational power of the ConvNet may be devoted to features that are specific to differentiating between dog breeds.

Check points of pretrained models Since modern ConvNets take 2-3 weeks to train across multiple GPUs on ImageNet, it is common to see people release their final ConvNet checkpoints for the benefit of others who can use the networks for fine-tuning. For example, the Caffe library has a Model Zoo where people share their network weights.

When and how to fine-tune?

IMPORTANT: When doing transfer learning if you make changes in the green and orange part do the following:

- Train green, freeze everything else
- When green is starting to converge unfreeze the orange part that you want to modify

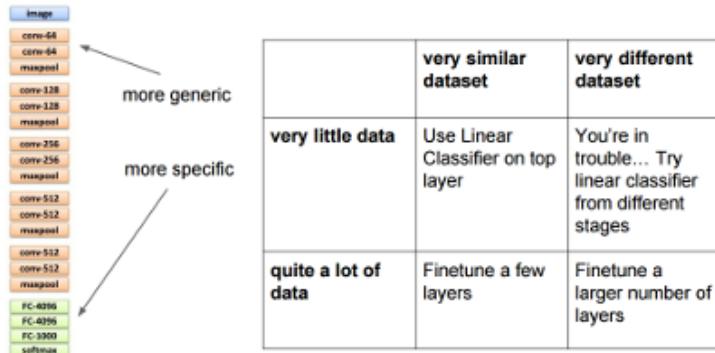


FIGURE 2.4: Random Crops/Scales

This is necessary because green starts with random values and it may produce very big gradients that may destroy your previous layers

Practical advice

There are a few additional things to keep in mind when performing Transfer Learning:

Constraints from pretrained models Note that if you wish to use a pretrained network, you may be slightly constrained in terms of the architecture you can use for your new dataset. For example, you can't arbitrarily take out Conv layers from the pretrained network. However, some changes are straight-forward: Due to parameter sharing, you can easily run a pretrained network on images of different spatial size. This is clearly evident in the case of Conv/Pool layers because their forward function is independent of the input volume spatial size (as long as the strides fit). In case of FC layers, this still holds true because FC layers can be converted to a Convolutional Layer: For example, in an AlexNet, the final pooling volume before the first FC layer is of size [6x6x512]. Therefore, the FC layer looking at this volume is equivalent to having a Convolutional Layer that has receptive field size 6x6, and is applied with padding of 0.

Learning rates It's common to use a smaller learning rate for ConvNet weights that are being fine-tuned, in comparison to the (randomly-initialized) weights for the new linear classifier that computes the class scores of your new dataset. This is because we expect that the ConvNet weights are relatively good, so we don't wish to distort them too quickly and too much (especially while the new Linear Classifier above them is being trained from random initialization).

Part II

LEARNING

Chapter 3

Neural Network

Imagine a linear classification which computes scores for different visual categories given the image using the formula $s = Wx$, where W is a matrix and x was an input column vector containing all pixel data of the image. In the case of CIFAR-10, x is a 3072×1 column vector, and W is a 10×3072 matrix, so that the output scores is a vector of 10 class scores.

An example neural network would instead compute $s = W_2 \max(0, W_1 x)$. Here, W_1 could be, for example, a 100×3072 matrix transforming the image into a 100-dimensional intermediate vector. The function $\max(0, -)$ is a non-linearity that is applied elementwise. There are several choices we could make for the non-linearity (which we will see later), but this one is a common choice and simply thresholds all activations that are below zero to zero. These are the so called *activation functions*. Finally, the matrix W_2 would then be of size 10×100 , so that we again get 10 numbers out that we interpret as the class scores. Notice that the non-linearity is critical computationally - if we left it out, the two matrices could be collapsed to a single matrix, and therefore the predicted class scores would again be a linear function of the input. The non-linearity is where we get the wiggle. The parameters W_2, W_1 are learned with stochastic gradient descent, and their gradients are derived with chain rule (and computed with backpropagation).

A three-layer neural network could analogously look like $s = W_3 \max(0, W_2 \max(0, W_1 x))$, where all of W_3, W_2, W_1 are parameters to be learned. The sizes of the intermediate hidden vectors are hyperparameters of the network and well see how we can set them later. Lets now look into how we can interpret these computations from the neuron/network perspective.

An example code for forward-propagating a single neuron might look as follows:

```
class Neuron(object):
    # ...
    def forward(inputs):
        """ assume inputs and weights are 1-D numpy arrays and bias is a number """
        cell_body_sum = np.sum(inputs * self.weights) + self.bias
        firing_rate = 1.0 / (1.0 + math.exp(-cell_body_sum)) # sigmoid activation function
        return firing_rate
```

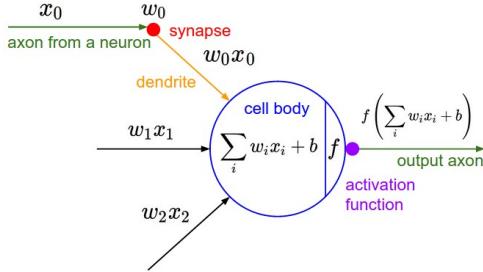


FIGURE 3.1: Cell diagram

In other words, each neuron performs a dot product with the input and its weights, adds the bias and applies the non-linearity (or activation function), in this case the sigmoid $\sigma(x) = 1/(1+e^{-x})$. We will go into more details about different activation functions at the chapter *Activation Function*.

Single neuron as a linear classifier

The mathematical form of the model Neurons forward computation might look familiar to you. As with linear classifiers, a neuron has the capacity to “like (activation near one) or “dislike (activation near zero) certain linear regions of its input space. Hence, with an appropriate loss function on the neurons output, we can turn a single neuron into a linear classifier:

Binary Softmax classifier For example, we can interpret $\sigma(\sum_i w_i x_i + b)$ to be the probability of one of the classes $P(y_i = 1 | x_i; w)$. The probability of the other class would be $P(y_i = 0 | x_i; w) = 1 - P(y_i = 1 | x_i; w)$, since they must sum to one. With this interpretation, we can formulate the cross-entropy loss as we have seen in the Linear Classification section, and optimizing it would lead to a binary Softmax classifier (also known as logistic regression). Since the sigmoid function is restricted to be between 0-1, the predictions of this classifier are based on whether the output of the neuron is greater than 0.5.

Binary SVM classifier Alternatively, we could attach a max-margin hinge loss to the output of the neuron and train it to become a binary Support Vector Machine.

Regularization interpretation The regularization loss in both SVM/Softmax cases could, in this biological view, be interpreted as gradual forgetting, since it would have the effect of driving all synaptic weights w towards zero after every parameter update.

Layer-wise organization

Neural Networks as neurons in graphs. Neural Networks are modeled as collections of neurons that are connected in an acyclic graph. In other words, the outputs of some neurons can become

inputs to other neurons. Cycles are not allowed since that would imply an infinite loop in the forward pass of a network. Instead of an amorphous blobs of connected neurons, Neural Network models are often organized into distinct layers of neurons. For regular neural networks, the most common layer type is the **fully-connected layer** in which neurons between two adjacent layers are fully pairwise connected, but neurons within a single layer share no connections.

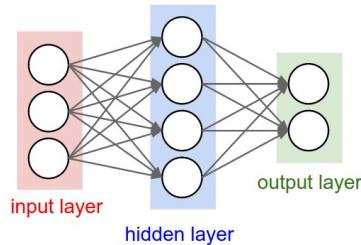


FIGURE 3.2: Layer-wise organization. A 2-layer Neural Network (one hidden layer of 4 neurons (or units) and one output layer with 2 neurons), and three inputs

Example feed-forward computation

```

# forward-pass of a 3-layer neural network:
f = lambda x: 1.0/(1.0 + np.exp(-x)) # activation function (use sigmoid)
x = np.random.randn(3, 1) # random input vector of three numbers (3x1)
h1 = f(np.dot(W1, x) + b1) # calculate first hidden layer activations (4x1)
h2 = f(np.dot(W2, h1) + b2) # calculate second hidden layer activations (4x1)
out = np.dot(W3, h2) + b3 # output neuron (1x1)

```

Setting number of layers and their sizes

How do we decide on what architecture to use when faced with a practical problem? Should we use no hidden layers? One hidden layer? Two hidden layers? How large should each layer be? First, note that as we increase the size and number of layers in a Neural Network, the capacity of the network increases. That is, the space of representable functions grows since the neurons can collaborate to express many different functions. For example, suppose we had a binary classification problem in two dimensions. We could train three separate neural networks, each with one hidden layer of some size and obtain different classifiers (Fig. 3.3).

In the diagram above, we can see that Neural Networks with more neurons can express more complicated functions. However, this is both a blessing (since we can learn to classify more complicated data) and a curse (since it is easier to overfit the training data). **Overfitting** occurs when a model with high capacity fits the noise in the data instead of the (assumed) underlying relationship. For example, the model with 20 hidden neurons fits all the training data but at the cost of segmenting the space into many disjoint red and green decision regions. The model with 3 hidden neurons only has the representational power to classify the data in broad strokes. It models the data as two blobs and interprets the few red points inside the green

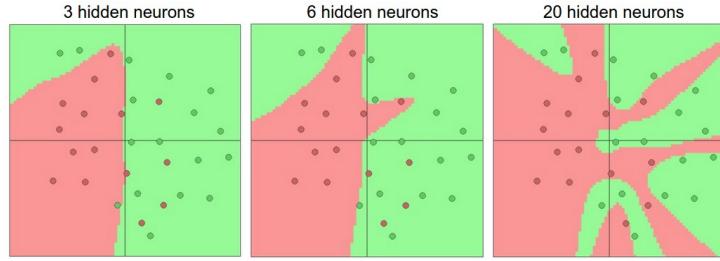


FIGURE 3.3: Larger Neural Networks can represent more complicated functions. You can play with these examples in [this ConvNetsJS demo](#).

cluster as **outliers** (noise). In practice, this could lead to better **generalization** on the test set. To reiterate, the **regularization** strength is the preferred way to control the overfitting of a neural network. We can look at the results achieved by three different settings (Fig. 3.4).

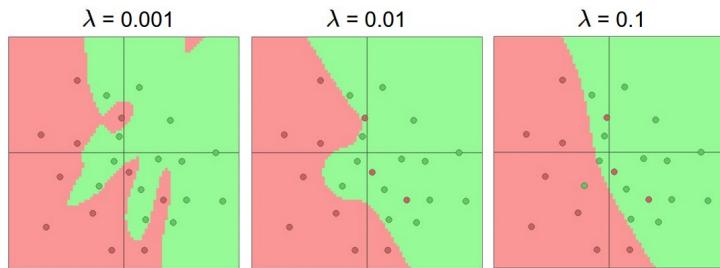


FIGURE 3.4: The effects of regularization strength: Each neural network above has 20 hidden neurons, but changing the regularization strength makes its final decision regions smoother with a higher regularization. You can play with these examples in [this ConvNetsJS demo](#).

Chapter 4

Parameters Initialization

4.1 Weights

We normally want the weights to be close to zero because it is reasonable to think that half of the weights will be positive and half of them will be negative.

Warning: It's not necessarily the case that smaller numbers will work strictly better. For example, a Neural Network layer that has very small weights will during backpropagation compute very small gradients on its data (since this gradient is proportional to the value of the weights). This could greatly diminish the gradient signal flowing backward through a network, and could become a concern for deep networks. It would bring all the activations towards 0.

If the weights are initialized with a too high value the activations will explode and saturate to 1.

W=0

There is no balance break. If every neuron in the network computes the same output, then they will also all compute the same gradients during backpropagation and undergo the exact same parameter updates. In other words, there is no source of asymmetry between neurons if their weights are initialized to be the same.

Small random numbers

Every neuron's weight vector is initialized as a random vector sampled from a multi-dimensional gaussian, so the neurons point in random direction in the input space. It is also possible to use small numbers drawn from a uniform distribution, but this seems to have relatively little impact on the final performance in practice.

One of the problems of this method is to decide how small should the gradients be. Example assuiming tanh:

- $W = 0.01 * np.random.randn(D, H)$: might be too small, all activations become zero, therefore, no gradient accumulation.
- $W = 1 * np.random.randn(D, H)$: might be too large, almost all neurons completely saturated, thus gradients will be zero.

Another problem is that the distribution of the outputs from a randomly initialized neuron has a variance that grows with the number of inputs.

Xavier initialization

eq: $w = np.random.randn(n) / \sqrt{n}$

eq for ReLU: $w = np.random.randn(n) / \sqrt{2n}$

It is possible to solve the distribution problem of the above method by normalizing the variance of each neurons output to 1 by scaling its weight vector by the square root of its fan-in (i.e. its number of inputs). This ensures that all neurons in the network initially have approximately the same output distribution and empirically improves the rate of convergence.

Mathematical reasoning

Consider the inner product $s = \sum_i^n w_i x_i$ between the weights and input , which gives the raw activation of a neuron before the non-linearity. We can examine the variance of s :

$$\begin{aligned}
\text{Var}(s) &= \text{Var}\left(\sum_i^n w_i x_i\right) \\
&= \sum_i^n \text{Var}(w_i x_i) \\
&= \sum_i^n [E(w_i)]^2 \text{Var}(x_i) + E[(x_i)]^2 \text{Var}(w_i) + \text{Var}(x_i) \text{Var}(w_i) \\
&= \sum_i^n \text{Var}(x_i) \text{Var}(w_i) \\
&= (n \text{Var}(w)) \text{Var}(x)
\end{aligned} \tag{4.1}$$

where in the first 2 steps we have used properties of variance. In third step we assumed zero mean inputs and weights, so $E[x_i] = E[w_i] = 0$. Note that this is not generally the case: For example ReLU units will have a positive mean. In the last step we assumed that all x_i, w_i are

identically distributed. From this derivation we can see that if we want s to have the same variance as all of its inputs xx , then during initialization we should make sure that the variance of every weight w is $1/n$. And since $\text{Var}(aX) = a^2\text{Var}(X)$ for a random variable X and a scalar a , this implies that we should draw from unit Gaussian and then scale it by $a = \sqrt{1/n}$, to make its variance $1/n$. This gives the initialization `w = np.random.randn(n) / sqrt(n)`.

A similar analysis is carried out for ReLUs, reaching the conclusion that the variance of neurons in the network should be $2/n$. This gives the initialization `w = np.random.randn(n) / sqrt(2n)`, and is the current recommendation for use in practice in the specific case of neural networks with ReLU neurons.

Sparse Initialization

Another way to address the uncalibrated variances problem is to set all weight matrices to zero, but to break symmetry every neuron is randomly connected (with weights sampled from a small gaussian as above) to a fixed number of neurons below it. A typical number of neurons to connect to may be as small as 10.

4.2 Biases

It is possible and common to initialize the biases to be zero, since the asymmetry breaking is provided by the small random numbers in the weights. For ReLU non-linearities, some people like to use small constant value such as 0.01 for all biases because this ensures that all ReLU units fire in the beginning and therefore obtain and propagate some gradient. However, it is not clear if this provides a consistent improvement (in fact some results seem to indicate that this performs worse) and it is more common to simply use 0 bias initialization.

Summary

The current recommendation is to use ReLU units and use the Xavier initialization (`w = np.random.randn(n) * sqrt(2.0/n)`) or `w = np.random.randn(n) * sqrt(2.0/n)`; and all biases equal to 0 or 0.01. Also batch normalization helps having less headaches with weight initialization.

Chapter 5

Activation Function

Activation functions are needed in order to increase the complexity of the NN, without them it would just be a linear sandwich of linear functions (multiplying linear functions = linear function)

5.1 Sigmoid

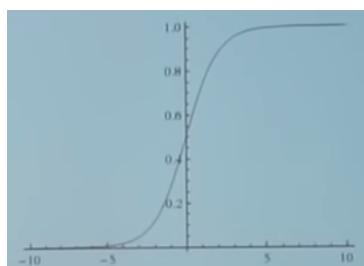


FIGURE 5.1: Sigmoid

$$\sigma(x) = \frac{1}{1+e^{-x}}$$

- Saturated neurons "kill" the gradient
- No zero-centered output
- $\exp()$ is computationally expensive

5.2 Tanh

$$\tanh(x) = \frac{2}{e^{-2x}+1} - 1$$

- Zero centered output

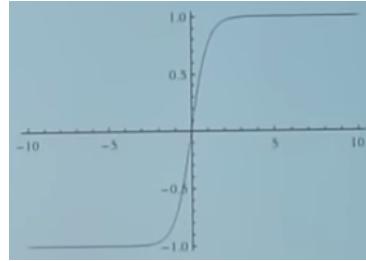


FIGURE 5.2: Tanh

- Saturated neurons "kill" the gradient

5.3 ReLU

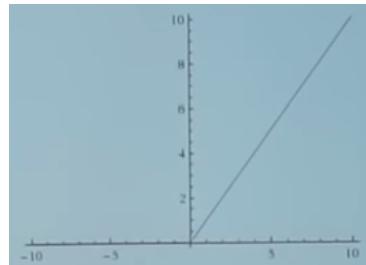


FIGURE 5.3: ReLU

$f(x) = \max(0, x)$ If a neuron has $x < 0$ the ReLU kills it in backprop. The gradient will be 0 so that neuron will not backprop downwards and its weights will not be updated. Its like a boolean gate. The gradient in the case of $x = 0$ is undefined (unlike case).

- Does not saturate
- Computationally efficient
- Converges much faster than sigmoid/tanh in practice ($6x$)
- Not zero centred output
- If you're unlucky a neuron may be never active because the initialization has put it outside the manifold.
- When the learning rate is high it's easy to kill a lot of neurons. Imagine the activation function as a threshold which is moving while training. If the learning rate is too high it may move out of the data manifold. In that case the neuron dies and will never be able to recover because it will never update again.

It is a good practice to initialize them with a slightly positive initial bias to avoid "dead neurons"

5.4 Leaky ReLU

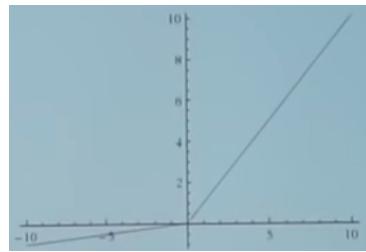


FIGURE 5.4: Leaky ReLU

$f(x) = \max(\alpha x, x) = \begin{cases} \alpha x & \text{if } x \leq 0 \\ x & \text{if } x > 0 \end{cases}$ where α is a very small number 0.01 which can be a hyperparameter or learned through .

- Does not saturate
- Computationally efficient
- Converges much faster than sigmoid/tanh in practice ($6x$)
- Does not die
- Not zero centred output
- Consistency of the benefits across tasks not clear

5.5 ELU - Exponential Linear Units

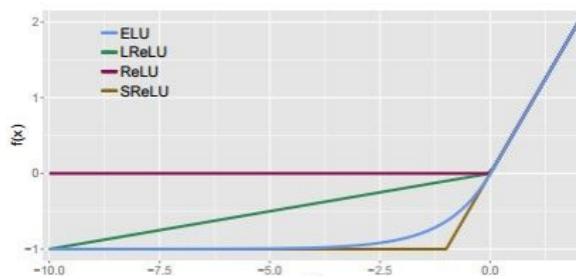


FIGURE 5.5: ELU

$$f(x) = \begin{cases} \alpha(e^x - 1) & \text{if } x \leq 0 \\ x & \text{if } x > 0 \end{cases}$$

- All benefits of ReLU
- Does not die
- Closer to zero mean outputs

- $\exp()$ is computationally expensive
- There is controversy if it actually trains better with this than with ReLU

5.6 Maxout

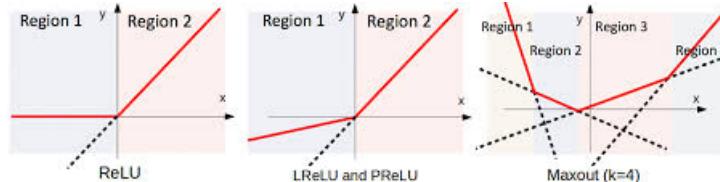


FIGURE 5.6: Maxout

$$f(x) = \max(w_1^T x + b_1, w_2^T x + b_2)$$

- Generalized ReLU and Leaky ReLU
- Does not saturate
- Does not die
- Linear Regime
- Doubles the number of parameters/neurons

Conclusion

- Use ReLU but be careful with your learning rates. So for example a 3-layer neural network will look like this: $f(x) = W_3 \max(0, W_2 \max(0, W_1 x))$
- Try out Leaky ReLU/Maxout/ELU
- Try out tanh but don't expect much
- Don't use sigmoid

Chapter 6

Loss function

The objective function is composed by the data loss and a regularization term. The data loss takes the form of an average over the data losses for every individual example. That is, $L = \frac{1}{N} \sum_i L_i$ where N is the number of training data. Lets abbreviate $f = f(x_i; W)$ to be the activations of the output layer in a Neural Network. There are several types of problems you might want to solve in practice:

Classification

We assume a dataset of examples and a single correct label (out of a fixed set) for each example. One of two most commonly seen cost functions in this setting are the *SVM* (e.g. the Weston Watkins formulation):

$$L_i = \sum_{j \neq y_i} \max(0, f_j - f_{y_i} + 1) \quad (6.1)$$

where f_{y_i} is the activation of the correct class, and f_j the activation of the class j . As we briefly alluded to, some people report better performance with the squared hinge loss (i.e. instead using $\max(0, f_j - f_{y_i} + 1)^2$). The second common choice is the *Softmax classifier* that uses the cross-entropy loss. It transforms the *logits* (scores) output of the network and transforms them into probabilities.

$$L_i = -\log \left(\frac{e^{f_{y_i}}}{\sum_j e^{f_j}} \right) \quad (6.2)$$

In some rough sense, the cross-entropy is measuring how inefficient our predictions are for describing the truth.

Probabilistic interpretation. Looking at the expression, we see that

$$P(y_i | x_i; W) = \frac{e^{f_{y_i}}}{\sum_j e^{f_j}} \quad (6.3)$$

can be interpreted as the (normalized) probability assigned to the correct label y_i given the image x_i and parameterized by W . To see this, remember that the Softmax classifier interprets the scores inside the output vector f as the unnormalized log probabilities (log probabilities means representing probabilities in logarithmic space, instead of the standard [0,1] interval. This has practical advantages because computers can perform addition more efficiently than multiplication). Exponentiating these quantities therefore gives the (unnormalized) probabilities, and the division performs the normalization so that the probabilities sum to one. In the probabilistic interpretation, we are therefore minimizing the negative log likelihood of the correct class, which can be interpreted as performing *Maximum Likelihood Estimation* (MLE). A nice feature of this view is that we can now also interpret the regularization term $R(W)$ in the full loss function as coming from a Gaussian prior over the weight matrix W , where instead of MLE we are performing the *Maximum a posteriori* (MAP) estimation. We mention these interpretations to help your intuitions, but the full details of this derivation are beyond the scope of this class.

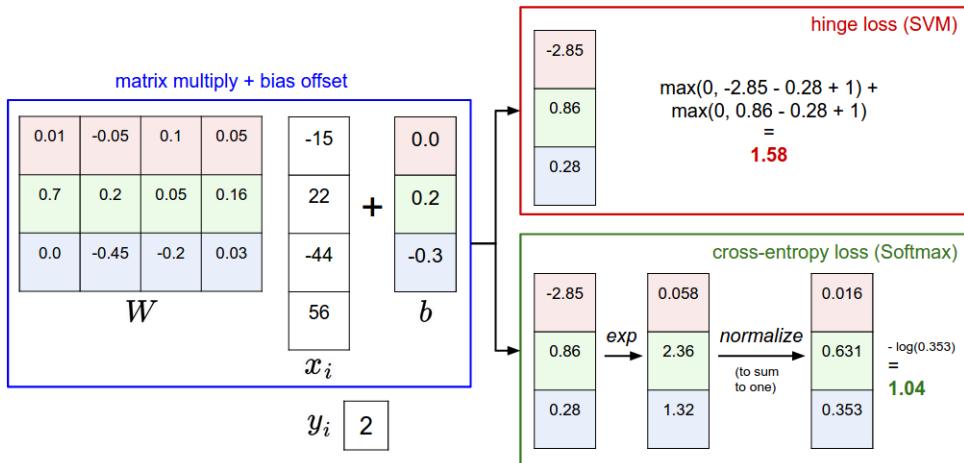


FIGURE 6.1: Example of the difference between the SVM and Softmax classifiers for one datapoint. In both cases we compute the same score vector f (e.g. by matrix multiplication in this section). The difference is in the interpretation of the scores in f : The SVM interprets these as class scores and its loss function encourages the correct class (class 2, in blue) to have a score higher by a margin than the other class scores. The Softmax classifier instead interprets the scores as (unnormalized) log probabilities for each class and then encourages the (normalized) log probability of the correct class to be high (equivalently the negative of it to be low). The final loss for this example is 1.58 for the SVM and 1.04 (note this is 1.04 using the natural logarithm, not base 2 or base 10) for the Softmax classifier, but note that these numbers are not comparable; They are only meaningful in relation to loss computed within the same classifier and with the same data.

Problem

Large number of classes. When the set of labels is very large (e.g. words in English dictionary, or ImageNet which contains 22,000 categories), it may be helpful to use *Hierarchical Softmax*. The hierarchical softmax decomposes labels into a tree. Each label is then represented as a path along the tree, and a Softmax classifier is trained at every node of the tree to disambiguate between the left and right branch. The structure of the tree strongly impacts the performance and is generally problem-dependent.

Attribute Classification

Both losses above assume that there is a single correct answer y_i . But what if y_i is a binary vector where every example may or may not have a certain attribute, and where the attributes are not exclusive? For example, images on Instagram can be thought of as labeled with a certain subset of hashtags from a large set of all hashtags, and an image may contain multiple. A sensible approach in this case is to build a binary classifier for every single attribute independently. For example, a binary classifier for each category independently would take the form:

$$L_i = \sum_j \max(0, 1 - y_{ij} f_j) \quad (6.4)$$

where the sum is over all categories j , and y_{ij} is either +1 or -1 depending on whether the i -th example is labeled with the j -th attribute, and the score vector f_j will be positive when the class is predicted to be present and negative otherwise. Notice that loss is accumulated if a positive example has score less than +1, or when a negative example has score greater than -1.

An alternative to this loss would be to train a logistic regression classifier for every attribute independently. A binary logistic regression classifier has only two classes (0,1), and calculates the probability of class 1 as:

$$P(y = 1 | x; w, b) = \frac{1}{1 + e^{-(w^T x + b)}} = \sigma(w^T x + b) \quad (6.5)$$

Since the probabilities of class 1 and 0 sum to one, the probability for class 0 is $P(y = 0 | x; w, b) = 1 - P(y = 1 | x; w, b)$. Hence, an example is classified as a positive example ($y = 1$) if $\sigma(w^T x + b) > 0.5$, or equivalently if the score $w^T x + b > 0$. The loss function then maximizes the log likelihood of this probability. You can convince yourself that this simplifies to:

$$L_i = \sum_j y_{ij} \log(\sigma(f_j)) + (1 - y_{ij}) \log(1 - \sigma(f_j)) \quad (6.6)$$

where the labels y_{ij} are assumed to be either 1 (positive) or 0 (negative), and $\sigma(\cdot)$ is the sigmoid function. The expression above can look scary but the gradient on f is in fact extremely simple and intuitive: $\partial L_i / \partial f_j = y_{ij} - \sigma(f_j)$ (as you can double check yourself by taking the derivatives).

Regression

Regression is the task of predicting real-valued quantities, such as the price of houses or the length of something in an image. For this task, it is common to compute the loss between the predicted quantity and the true answer and then measure the L2 squared norm, or L1 norm of the difference. The L2 norm squared would compute the loss for a single example of the form:

$$L_i = \|f - y_i\|_2^2 \quad (6.7)$$

The reason the L2 norm is squared in the objective is that the gradient becomes much simpler, without changing the optimal parameters since squaring is a monotonic operation. The L1 norm would be formulated by summing the absolute value along each dimension:

$$L_i = \|f - y_i\|_1 = \sum_j |f_j - (y_i)_j| \quad (6.8)$$

where the sum \sum_j is a sum over all dimensions of the desired prediction, if there is more than one quantity being predicted. Looking at only the j -th dimension of the i -th example and denoting the difference between the true and the predicted value by δ_{ij} , the gradient for this dimension (i.e. $\partial L_i / \partial f_j$) is easily derived to be either δ_{ij} with the L2 norm, or $\text{sign}(\delta_{ij})$. That is, the gradient on the score will either be directly proportional to the difference in the error, or it will be fixed and only inherit the sign of the difference.

Outliers L2 norm is not that good dealing with outliers in the training samples. You may consider L1 or variations of L2.

Word of caution It is important to note that the L2 loss is much harder to optimize than a more stable loss such as Softmax. Intuitively, it requires a very fragile and specific property from the network to output exactly one correct value for each input (and its augmentations). Notice that this is not the case with Softmax, where the precise value of each score is less important: It only matters that their magnitudes are appropriate. Additionally, the L2 loss is less robust because outliers can introduce huge gradients. When faced with a regression problem, first consider if it is absolutely inadequate to quantize the output into bins. For example, if you are predicting star rating for a product, it might work much better to use 5 independent classifiers for ratings of 1-5 stars instead of a regression loss. Classification has the additional benefit

that it can give you a distribution over the regression outputs, not just a single output with no indication of its confidence. If you're certain that classification is not appropriate, use the L2 but be careful: For example, the L2 is more fragile and applying dropout in the network (especially in the layer right before the L2 loss) is not a great idea.

When faced with a regression task, first consider if it is absolutely necessary. Instead, have a strong preference to discretizing your outputs to bins and perform classification over them whenever possible.

Structured prediction

The structured loss refers to a case where the labels can be arbitrary structures such as graphs, trees, or other complex objects. Usually it is also assumed that the space of structures is very large and not easily enumerable. The basic idea behind the structured SVM loss is to demand a margin between the correct structure y_i and the highest-scoring incorrect structure. It is not common to solve this problem as a simple unconstrained optimization problem with gradient descent. Instead, special solvers are usually devised so that the specific simplifying assumptions of the structure space can be taken advantage of. We mention the problem briefly but consider the specifics to be outside of the scope of the class.

Chapter 7

Backpropagation

Introduction

Motivation In this section we will develop expertise with an intuitive understanding of backpropagation, which is a way of computing gradients of expressions through recursive application of **chain rule**. Understanding of this process and its subtleties is critical for you to understand, and effectively develop, design and debug Neural Networks.

Problem statement The core problem studied in this section is as follows: We are given some function $f(x)$ where x is a vector of inputs and we are interested in computing the gradient of f at x (i.e. $\nabla f(x)$).

Motivation Recall that the primary reason we are interested in this problem is that in the specific case of Neural Networks, f will correspond to the loss function (L) and the inputs x will consist of the training data and the neural network weights. For example, the loss could be the SVM loss function and the inputs are both the training data $(x_i, y_i), i = 1 \dots N$ and the weights and biases W, b . Note that (as is usually the case in Machine Learning) we think of the training data as given and fixed, and of the weights as variables we have control over. Hence, even though we can easily use backpropagation to compute the gradient on the input examples x_i , in practice we usually only compute the gradient for the parameters (e.g. W, b) so that we can use it to perform a parameter update. However, as we will see later in the class the gradient on x_i can still be useful sometimes, for example for purposes of visualization and interpreting what the Neural Network might be doing.

Simple expressions and interpretation of the gradient

Lets start simple so that we can develop the notation and conventions for more complex expressions. Consider a simple multiplication function of two numbers $f(x, y) = xy$. It is a matter of simple calculus to derive the partial derivative for either input:

$$f(x, y) = xy \quad \rightarrow \quad \frac{\partial f}{\partial x} = y \quad \frac{\partial f}{\partial y} = x \quad (7.1)$$

Interpretation Keep in mind what the derivatives tell you: They indicate the rate of change of a function with respect to that variable surrounding an infinitesimally small region near a particular point:

$$\frac{df(x)}{dx} = \lim_{h \rightarrow 0} \frac{f(x+h) - f(x)}{h} \quad (7.2)$$

A technical note is that the division sign on the left-hand sign is, unlike the division sign on the right-hand sign, not a division. Instead, this notation indicates that the operator $\frac{d}{dx}$ is being applied to the function f , and returns a different function (the derivative). A nice way to think about the expression above is that when h is very small, then the function is well-approximated by a straight line, and the derivative is its slope. In other words, **the derivative on each variable tells you the sensitivity of the whole expression on its value**. For example, if $x = 4, y = -3$ then $f(x, y) = -12$ and the derivative on x $\frac{\partial f}{\partial x} = -3$. This tells us that if we were to increase the value of this variable by a tiny amount, the effect on the whole expression would be to decrease it (due to the negative sign), and by three times that amount. This can be seen by rearranging the above equation ($f(x+h) = f(x) + h \frac{df(x)}{dx}$). Analogously, since $\frac{\partial f}{\partial y} = 4$, we expect that increasing the value of y by some very small amount h would also increase the output of the function (due to the positive sign), and by $4h$.

The derivative on each variable tells you the sensitivity of the whole expression on its value.

As mentioned, the gradient ∇f is the vector of partial derivatives, so we have that $\nabla f = [\frac{\partial f}{\partial x}, \frac{\partial f}{\partial y}] = [y, x]$. Even though the gradient is technically a vector, we will often use terms such as “the gradient on x ” instead of the technically correct phrase “the partial derivative on x ” for simplicity.

We can also derive the derivatives for the addition operation:

$$f(x, y) = x + y \quad \rightarrow \quad \frac{\partial f}{\partial x} = 1 \quad \frac{\partial f}{\partial y} = 1 \quad (7.3)$$

that is, the derivative on both x, y is one regardless of what the values of x, y are. This makes sense, since increasing either x, y would increase the output of f , and the rate of that increase would be independent of what the actual values of x, y are (unlike the case of multiplication above). The last function well use quite a bit in the class is the max operation:

$$f(x, y) = \max(x, y) \quad \rightarrow \quad \frac{\partial f}{\partial x} = (x \geq y) \quad \frac{\partial f}{\partial y} = (y \geq x) \quad (7.4)$$

That is, the (sub)gradient is 1 on the input that was larger and 0 on the other input. Intuitively, if the inputs are $x = 4, y = 2$, then the max is 4, and the function is not sensitive to the setting of y . That is, if we were to increase it by a tiny amount h , the function would keep outputting 4, and therefore the gradient is zero: there is no effect. Of course, if we were to change y by a large amount (e.g. larger than 2), then the value of f would change, but the derivatives tell us nothing about the effect of such large changes on the inputs of a function; They are only informative for tiny, infinitesimally small changes on the inputs, as indicated by the $\lim_{h \rightarrow 0}$ in its definition.

Compound expressions with chain rule

Lets now start to consider more complicated expressions that involve multiple composed functions, such as $f(x, y, z) = (x+y)z$. This expression is still simple enough to differentiate directly, but well take a particular approach to it that will be helpful with understanding the intuition behind backpropagation. In particular, note that this expression can be broken down into two expressions: $q = x + y$ and $f = qz$. Moreover, we know how to compute the derivatives of both expressions separately, as seen in the previous section. f is just multiplication of q and z , so $\frac{\partial f}{\partial q} = z, \frac{\partial f}{\partial z} = q$ and q is addition of x and y so $\frac{\partial q}{\partial x} = 1, \frac{\partial q}{\partial y} = 1$. However, we dont necessarily care about the gradient on the intermediate value q - the value of $\frac{\partial f}{\partial q}$ is not useful. Instead, we are ultimately interested in the gradient of f with respect to its inputs x, y, z . The chain rule tells us that the correct way to “chain” these gradient expressions together is through multiplication. For example, $\frac{\partial f}{\partial x} = \frac{\partial f}{\partial q} \frac{\partial q}{\partial x}$. In practice this is simply a multiplication of the two numbers that hold the two gradients. Lets see this with an example:

```
# set some inputs
x = -2; y = 5; z = -4

# perform the forward pass
q = x + y # q becomes 3
f = q * z # f becomes -12

# perform the backward pass (backpropagation) in reverse order:
# first backprop through f = q * z
dfdz = q # df/dz = q, so gradient on z becomes 3
dfdq = z # df/dq = z, so gradient on q becomes -4
# now backprop through q = x + y
dfdx = 1.0 * dfdq # dq/dx = 1. And the multiplication here is the chain rule!
dfdy = 1.0 * dfdq # dq/dy = 1
```

At the end we are left with the gradient in the variables `[dfdx, dfdy, dfdz]`, which tell us the sensitivity of the variables `x, y, z` on `f`. This is the simplest example of backpropagation. Going

forward, we will want to use a more concise notation so that we don't have to keep writing the df part. That is, for example instead of dfdq we would simply write dq , and always assume that the gradient is with respect to the final output.

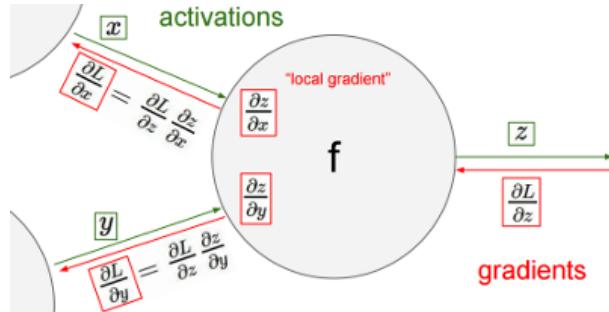


FIGURE 7.1: Neuron backprop

This computation can also be nicely visualized with a circuit diagram in figure 7.2. The real-valued “circuit” shows the visual representation of the computation. The forward pass computes values from inputs to output (shown in green). The backward pass then performs backpropagation which starts at the end and recursively applies the chain rule to compute the gradients (shown in red) all the way to the inputs of the circuit. The gradients can be thought of as flowing backwards through the circuit.

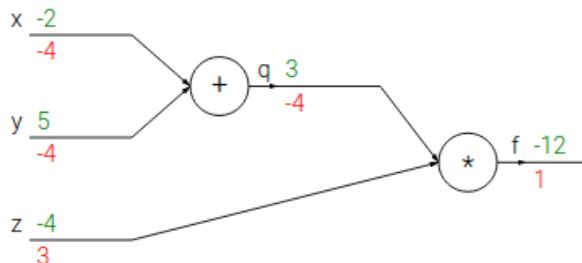


FIGURE 7.2: Neuron backprop diagram

Intuitive understanding of backpropagation

Notice that backpropagation is a beautifully local process. Every gate in a circuit diagram gets some inputs and can right away compute two things: 1. its output value and 2. the local gradient of its inputs with respect to its output value. Notice that the gates can do this completely independently without being aware of any of the details of the full circuit that they are embedded in. However, once the forward pass is over, during backpropagation the gate will eventually learn about the gradient of its output value on the final output of the entire circuit. Chain rule says that the gate should take that gradient and multiply it into every gradient it normally computes for all of its inputs.

This extra multiplication (for each input) due to the chain rule can turn a single and relatively useless gate into a cog in a complex circuit such as an entire neural network.

Lets get an intuition for how this works by referring again to the example. The add gate received inputs [-2, 5] and computed output 3. Since the gate is computing the addition operation, its local gradient for both of its inputs is +1. The rest of the circuit computed the final value, which is -12. During the backward pass in which the chain rule is applied recursively backwards through the circuit, the add gate (which is an input to the multiply gate) learns that the gradient for its output was -4. If we anthropomorphize the circuit as wanting to output a higher value (which can help with intuition), then we can think of the circuit as wanting the output of the add gate to be lower (due to negative sign), and with a force of 4. To continue the recurrence and to chain the gradient, the add gate takes that gradient and multiplies it to all of the local gradients for its inputs (making the gradient on both x and y $1 * -4 = -4$). Notice that this has the desired effect: If x, y were to decrease (responding to their negative gradient) then the add gates output would decrease, which in turn makes the multiply gates output increase.

Backpropagation can thus be thought of as gates communicating to each other (through the gradient signal) whether they want their outputs to increase or decrease (and how strongly), so as to make the final output value higher.

Modularity: Sigmoid example

The gates we introduced above are relatively arbitrary. Any kind of differentiable function can act as a gate, and we can group multiple gates into a single gate, or decompose a function into multiple gates whenever it is convenient. Lets look at another expression that illustrates this point:

$$f(w, x) = \frac{1}{1 + e^{-(w_0x_0 + w_1x_1 + w_2)}} \quad (7.5)$$

as we will see later in the class, this expression describes a 2-dimensional neuron (with inputs x and weights w) that uses the sigmoid activation function. But for now lets think of this very simply as just a function from inputs w, x to a single number. The function is made up of multiple gates. In addition to the ones described already above (add, mul, max), there are four more:

$$f(x) = \frac{1}{x} \rightarrow \frac{df}{dx} = -1/x^2, \quad f_c(x) = c+x \rightarrow \frac{df}{dx} = 1, \quad f(x) = e^x \rightarrow \frac{df}{dx} = e^x, \quad f_a(x) = ax \rightarrow \frac{df}{dx} = a \quad (7.6)$$

Where the functions f_c, f_a translate the input by a constant of c and scale the input by a constant of a , respectively. These are technically special cases of addition and multiplication, but we introduce them as (new) unary gates here since we do need the gradients for the constants c, a . The full circuit then looks as follows:

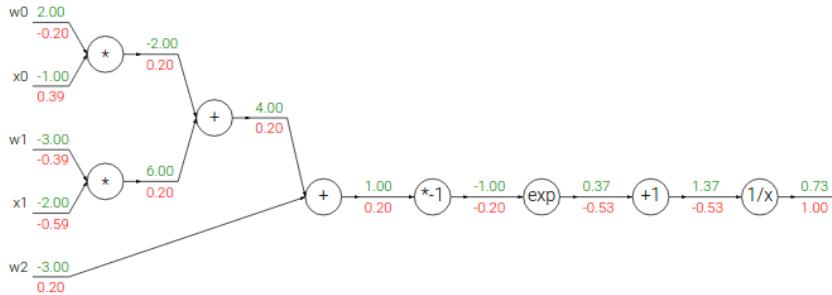


FIGURE 7.3: Example circuit for a 2D neuron with a sigmoid activation function. The inputs are $[x_0, x_1]$ and the (learnable) weights of the neuron are $[w_0, w_1, w_2]$. As we will see later, the neuron computes a dot product with the input and then its activation is softly squashed by the sigmoid function to be in range from 0 to 1.

In the example above, we see a long chain of function applications that operates on the result of the dot product between w, x . The function that these operations implement is called the sigmoid function $\sigma(x)$. It turns out that the derivative of the sigmoid function with respect to its input simplifies if you perform the derivation (after a fun tricky part where we add and subtract a 1 in the numerator):

$$\sigma(x) = \frac{1}{1 + e^{-x}} \rightarrow \frac{d\sigma(x)}{dx} = \frac{e^{-x}}{(1 + e^{-x})^2} = \left(\frac{1 + e^{-x} - 1}{1 + e^{-x}} \right) \left(\frac{1}{1 + e^{-x}} \right) = (1 - \sigma(x)) \sigma(x) \quad (7.7)$$

As we see, the gradient turns out to simplify and becomes surprisingly simple. For example, the sigmoid expression receives the input 1.0 and computes the output 0.73 during the forward pass. The derivation above shows that the local gradient would simply be $(1 - 0.73) * 0.73 = 0.2$, as the circuit computed before (see the image above), except this way it would be done with a single, simple and efficient expression (and with less numerical issues). Therefore, in any real practical application it would be very useful to group these operations into a single gate. Lets see the backprop for this neuron in code:

```
w = [2,-3,-3] # assume some random weights and data
x = [-1, -2]

# forward pass
dot = w[0]*x[0] + w[1]*x[1] + w[2]
f = 1.0 / (1 + math.exp(-dot)) # sigmoid function

# backward pass through the neuron (backpropagation)
ddot = (1 - f) * f # gradient on dot variable, using the sigmoid gradient derivation
dx = [w[0] * ddot, w[1] * ddot] # backprop into x
dw = [x[0] * ddot, x[1] * ddot, 1.0 * ddot] # backprop into w
# we're done! we have the gradients on the inputs to the circuit
```

Implementation protip staged backpropagation. As shown in the code above, in practice it is always helpful to break down the forward pass into stages that are easily backpropogated through. For example here we created an intermediate variable `dot` which holds the output of the dot product between `w` and `x`. During backward pass we then successively compute (in reverse order) the corresponding variables (e.g. `ddot`, and ultimately `dw`, `dx`) that hold the gradients of those variables.

The point of this section is that the details of how the backpropagation is performed, and which parts of the forward function we think of as gates, is a matter of convenience. It helps to be aware of which parts of the expression have easy local gradients, so that they can be chained together with the least amount of code and effort.

Backprop in practice: Staged computation

Lets see this with another example. Suppose that we have a function of the form:

$$f(x, y) = \frac{x + \sigma(y)}{\sigma(x) + (x + y)^2} \quad (7.8)$$

To be clear, this function is completely useless and its not clear why you would ever want to compute its gradient, except for the fact that it is a good example of backpropagation in practice. It is very important to stress that if you were to launch into performing the differentiation with respect to either x or y , you would end up with very large and complex expressions. However, it turns out that doing so is completely unnecessary because we dont need to have an explicit function written down that evaluates the gradient. We only have to know how to compute it. Here is how we would structure the forward pass of such expression:

```
x = 3 # example values
y = -4

# forward pass
sigy = 1.0 / (1 + math.exp(-y)) # sigmoid in numerator #(1)
num = x + sigy # numerator #(2)
sigx = 1.0 / (1 + math.exp(-x)) # sigmoid in denominator #(3)
xpy = x + y #(4)
xpysqr = xpy**2 #(5)
den = sigx + xpysqr # denominator #(6)
invden = 1.0 / den #(7)
f = num * invden # done! #(8)
```

Phew, by the end of the expression we have computed the forward pass. Notice that we have structured the code in such way that it contains multiple intermediate variables, each of which are only simple expressions for which we already know the local gradients. Therefore, computing the backprop pass is easy: Well go backwards and for every variable along the way in the forward pass (`sigy`, `num`, `sigx`, `xpy`, `xpysqr`, `den`, `invden`) we will have the same variable, but one

that begins with a `d`, which will hold the gradient of the output of the circuit with respect to that variable. Additionally, note that every single piece in our backprop will involve computing the local gradient of that expression, and chaining it with the gradient on that expression with a multiplication. For each row, we also highlight which part of the forward pass it refers to:

```
# backprop f = num * invden
dnum = invden # gradient on numerator #(8)
dinvden = num #(8)
# backprop invden = 1.0 / den
dden = (-1.0 / (den**2)) * dinvden #(7)
# backprop den = sigx + xpy**2
dsigx = (1) * dden #(6)
dxpy = (1) * dden #(6)
# backprop xpy = x + y
dxpy = (2 * xpy) * dxpy #(5)
# backprop xpy = x + y
dx = (1) * dxpy #(4)
dy = (1) * dxpy #(4)
# backprop sigx = 1.0 / (1 + math.exp(-x))
dx += ((1 - sigx) * sigx) * dsigx # Notice += !! See notes below #(3)
# backprop num = x + sigy
dx += (1) * dnum #(2)
dsigy = (1) * dnum #(2)
# backprop sigy = 1.0 / (1 + math.exp(-y))
dy += ((1 - sigy) * sigy) * dsigy #(1)
# done! phew
```

Notice a few things:

Cache forward pass variables To compute the backward pass it is very helpful to have some of the variables that were used in the forward pass. In practice you want to structure your code so that you cache these variables, and so that they are available during backpropagation. If this is too difficult, it is possible (but wasteful) to recompute them.

Gradients add up at forks The forward expression involves the variables x, y multiple times, so when we perform backpropagation we must be careful to use `+=` instead of `=` to accumulate the gradient on these variables (otherwise we would overwrite it). This follows the multivariable chain rule in Calculus, which states that if a variable branches out to different parts of the circuit, then the gradients that flow back to it will add.

Patterns in backward flow

It is interesting to note that in many cases the backward-flowing gradient can be interpreted on an intuitive level. For example, the three most commonly used gates in neural networks

(add,mul,max), all have very simple interpretations in terms of how they act during backpropagation. Consider this example circuit:

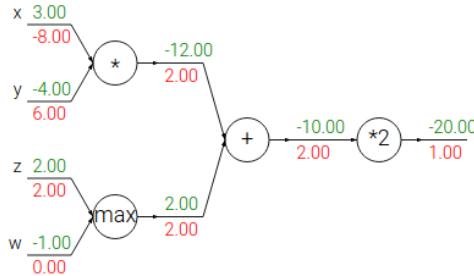


FIGURE 7.4: An example circuit demonstrating the intuition behind the operations that back-propagation performs during the backward pass in order to compute the gradients on the inputs. Sum operation distributes gradients equally to all its inputs. Max operation routes the gradient to the higher input. Multiply gate takes the input activations, swaps them and multiplies by its gradient.

Looking at the diagram above as an example, we can see that:

add gate The add gate always takes the gradient on its output and distributes it equally to all of its inputs, regardless of what their values were during the forward pass. This follows from the fact that the local gradient for the add operation is simply +1.0, so the gradients on all inputs will exactly equal the gradients on the output because it will be multiplied by x1.0 (and remain unchanged). In the example circuit above, note that the + gate routed the gradient of 2.00 to both of its inputs, equally and unchanged.

max gate The max gate routes the gradient. Unlike the add gate which distributed the gradient unchanged to all its inputs, the max gate distributes the gradient (unchanged) to exactly one of its inputs (the input that had the highest value during the forward pass). This is because the local gradient for a max gate is 1.0 for the highest value, and 0.0 for all other values. In the example circuit above, the max operation routed the gradient of 2.00 to the z variable, which had a higher value than w, and the gradient on w remains zero.

multiply gate The multiply gate is a little less easy to interpret. Its local gradients are the input values (except switched), and this is multiplied by the gradient on its output during the chain rule. In the example above, the gradient on x is -8.00, which is -4.00 x 2.00.

Unintuitive effects and their consequences. Notice that if one of the inputs to the multiply gate is very small and the other is very big, then the multiply gate will do something slightly unintuitive: it will assign a relatively huge gradient to the small input and a tiny gradient to the large input. Note that in linear classifiers where the weights are dot producted $w^T x_i$ (multiplied) with the inputs, this implies that the scale of the data has an effect on the magnitude of the gradient for the weights. For example, if you multiplied all input data examples x_i by 1000

during preprocessing, then the gradient on the weights will be 1000 times larger, and you'd have to lower the learning rate by that factor to compensate. This is why preprocessing matters a lot, sometimes in subtle ways! And having intuitive understanding for how the gradients flow can help you debug some of these cases.

Gradients for vectorized operations

The above sections were concerned with single variables, but all concepts extend in a straightforward manner to matrix and vector operations. However, one must pay closer attention to dimensions and transpose operations.

Matrix-Matrix multiply gradient Possibly the most tricky operation is the matrix-matrix multiplication (which generalizes all matrix-vector and vector-vector) multiply operations:

```
# forward pass
W = np.random.randn(5, 10)
X = np.random.randn(10, 3)
D = W.dot(X)

# now suppose we had the gradient on D from above in the circuit
dD = np.random.randn(*D.shape) # same shape as D
dW = dD.dot(X.T) #.T gives the transpose of the matrix
dX = W.T.dot(dD)
```

Tip: use dimension analysis! Note that you do not need to remember the expressions for dW and dX because they are easy to re-derive based on dimensions. For instance, we know that the gradient on the weights dW must be of the same size as W after it is computed, and that it must depend on matrix multiplication of X and dD (as is the case when both X, W are single numbers and not matrices). There is always exactly one way of achieving this so that the dimensions work out. For example, X is of size [10 x 3] and dD of size [5 x 3], so if we want dW and W has shape [5 x 10], then the only way of achieving this is with $dD \cdot dot(X.T)$, as shown above.

Work with small, explicit examples. Some people may find it difficult at first to derive the gradient updates for some vectorized expressions. Our recommendation is to explicitly write out a minimal vectorized example, derive the gradient on paper and then generalize the pattern to its efficient, vectorized form.

Erik Learned-Miller has also written up a longer related document on taking matrix/vector derivatives which you might find helpful. Find it [here](#).

Summary

- We developed intuition for what the gradients mean, how they flow backwards in the circuit, and how they communicate which part of the circuit should increase or decrease and with what force to make the final output higher.
- We discussed the importance of staged computation for practical implementations of backpropagation. You always want to break up your function into modules for which you can easily derive local gradients, and then chain them with chain rule. Crucially, you almost never want to write out these expressions on paper and differentiate them symbolically in full, because you never need an explicit mathematical equation for the gradient of the input variables. Hence, decompose your expressions into stages such that you can differentiate every stage independently (the stages will be matrix vector multiplies, or max operations, or sum operations, etc.) and then backprop through the variables one step at a time.

In the next section we will start to define Neural Networks, and backpropagation will allow us to efficiently compute the gradients on the connections of the neural network, with respect to a loss function. In other words, we're now ready to train Neural Nets, and the most conceptually difficult part of this class is behind us! ConvNets will then be a small step away.

Chapter 8

Parameters Update

IMPORTANT NOTATION!!: Through all this chapter I am saying parameters = weights, but remember that there are other ones like the bias.

First Order Optimization Methods

The pseudo-code for updating the parameters:

```
while True:  
    data_batch = dataset.sample_data_batch()  
    loss = network.forward(data_batch)  
    dw = network.backward()  
    w += -learning_rate * dw
```

This update ($w += -lr * dw$) is an stochastic gradient descent. Let's see better ways of updating the parameters. The plot below shows different update strategies to optimize the loss function. In DNN, as you scale your model, local minima become less and less of an issue. The loss function shape we should have in mind is like a landscape with a lot of local minima but all with more or less the same loss. As you scale the model the distance between the bigger and lower local minima scales down (there is no bad local minima in big networks).

Stochastic gradient descent

$$w_t = w_{t-1} - \alpha \Delta L(w_{t-1}) \quad (8.1)$$

where $\Delta L(w_{t-1})$ is the derivative of the loss function and the learning rate

Problem: Suppose a 2dimensions loss function (NN with 2 layers and 2 weights) which is steep vertically and shallow horizontally (weight represented in the vertical axis has a higher impact

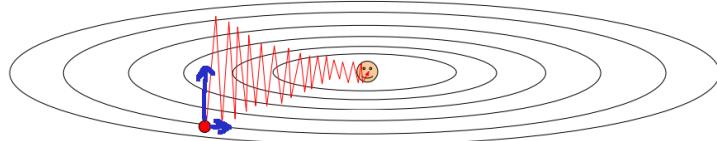


FIGURE 8.1: SGD example

over the loss function). The horizontal gradient is small because it is a shallow function in the horizontal direction, and a big gradient in the vertical direction because it is steep in this direction. Then, the update is way too slow in the horizontal direction, and too fast in the vertical direction. This causes an ugly jerk shape.

Notice that it is not an option to just decrease the learning rate because then it would take forever to cross all the shallow surface.

IMPORTANT NOTATION!!: Through all this chapter I am saying parameters = weights, but remember that there are other ones like the bias.

Momentum update

$$\begin{aligned} v_0 &= 0 \\ v_t &= \mu v_{t-1} - \alpha \Delta L(w_{t-1}) \\ w_t &= w_{t-1} + v_t \end{aligned} \tag{8.2}$$

where $\mu \in [0, 1]$ and usually takes values around 0.5, 0.9 or 0.99 (sometimes annealed over time 0.5 -> 0.99)

It can be interpreted as a ball rolling across the loss function starting at velocity 0. Notice that would represent position, velocity, friction (decreasing velocity over time). Now you will speed up in shallow surfaces directions and in steep directions you will be jiggly but you will have a force pushing you to the center (like a ball rolling in a convex function).

In the first plot we can see that first it overshoots because of all the accumulated velocity but it ends converging to the minimum quicker than SVG.

Nesterov Momentum update (or Nesterov accelerated gradient descent)

$$\begin{aligned} v_0 &= 0 \\ v_t &= \mu v_{t-1} - \alpha \Delta L(w_{t-1} + \mu v_{t-1}) \\ w_t &= w_{t-1} + v_t \end{aligned} \tag{8.3}$$

A slightly different version of the momentum update has recently been gaining popularity. It enjoys stronger theoretical converge guarantees for convex functions and in practice it also consistently works slightly better than standard momentum.

The core idea behind Nesterov momentum is that when the current parameter vector is at some position x (weights), then looking at the momentum update above, we know that the momentum term alone (i.e. ignoring the second term with the gradient) is about to nudge the parameter vector by $\mu * v$. Therefore, if we are about to compute the gradient, we can treat the future approximate position $x + \mu * v$ as a lookahead - this is a point in the vicinity of where we are soon going to end up. Hence, it makes sense to compute the gradient at $x + \mu * v$ instead of at the old/stale position x .

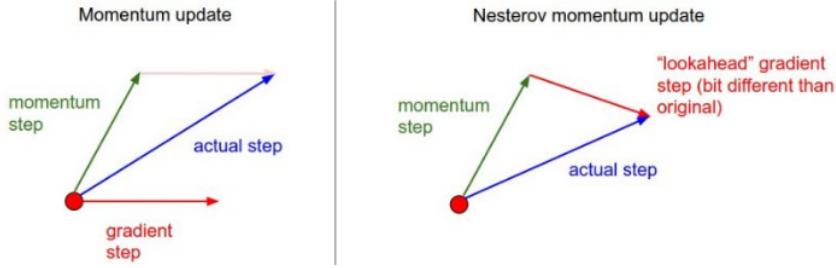


FIGURE 8.2: Nesterov Momentum update

IMPORTANT NOTATION!!: Through all this chapter I am saying parameters = weights, but remember that there are other ones like the bias.

AdaGrad

$$\begin{aligned} c_t &= c_{t-1} + \Delta L(w_{t-1})^2 \\ w_t &= w_{t-1} - \frac{\alpha \Delta L(w_{t-1})}{\sqrt{c_t} + 1^{-7}} \end{aligned} \tag{8.4}$$

where is the cache is a giant vector of the same size of your parameter vector . So this cache keeps track of the sum of squares (or also called the uncentered second moment) of your parameters in each direction (array). So now, with the cache, each dimension of the parameter vector has a different dynamic learning rate which is dynamically scaled based on what kind of gradients are you seeing in terms of their scale.

In the first case explained steep vertically and shallow horizontally loss function (2 weights). We have a large gradient vertically which will be added to cache. So in the vertical direction, the parameter update will be getting smaller and smaller (= smaller steps) because the denominator keeps increasing (= learning rate keeps decreasing). In the horizontal direction the surface is shallow, so will be small and as a consequence the learning rate will not decrease. The horizontal direction will have faster progress relative to the vertical direction.

So the cache is acting as an equalizer accounting for the stiffness () of the loss function in the parameters direction. Keep in mind that this equalizer effect over a specific parameter (=weight) is independent of the rest of parameters. It is not like a normalization.

Problem: When training over long time the step size (= learning rate) goes to 0 and the network stops learning. To solve this, the next method is proposed.

RMSProp

$$\begin{aligned} c_t &= \gamma c_{t-1} + (1 - \gamma) \Delta L(w_{t-1})^2 \\ w_t &= w_{t-1} - \frac{\alpha \Delta L(w_{t-1})}{\sqrt{c} + 1^{-7}} \end{aligned} \tag{8.5}$$

The idea is that instead of keeping the sum of squares in every direction of the parameters, we make the counter leaky by introducing the decay rate . So we maintain the nice effect of equalizing the directions but it will not converge to 0. The term is there just to prevent the division by 0.

You can also imagine the decay rate as a sort of a forgetting factor that makes the cache only a function of the last few gradient but in an exponential weighted sum rate. Also, notice that we could not do something like having a sliding window just to track the previous n gradients because it would take too much memory.

Comparing RMSProp vs AdaGrad in DNNs, in practice AdaGrad usually ends up stopping too early.

IMPORTANT NOTATION!!: Through all this chapter I am saying parameters = weights, but remember that there are other ones like the bias.

Adam update - USE THIS ONE

$$\begin{aligned} v_0 = c_0 = 0 \\ v_t &= \frac{\mu v_{t-1} + (1 - \mu) \Delta L(w_{t-1})}{1 - \mu^t} \\ c_t &= \frac{\gamma c_{t-1} + (1 - \gamma) \Delta L(w_{t-1})^2}{1 - \gamma^t} \\ w_t &= w_{t-1} - \frac{\alpha v_t}{\sqrt{c} + 1^{-7}} \end{aligned} \tag{8.6}$$

Adam update is a combination of adaptive scaling (AdaGrad) and momentum (Momentum update). v_t is the velocity and the cache. Sometimes v_t and c_t are denoted first and second momentum, respectively. To compensate the fact that the momentums are initialized to zero () and need some time to "warm up", we introduce a bias correction in order to scale up and in the first iterations so you do not get a very biased estimate of the first and second momentum. bias correction: $\frac{1}{1-\mu^t}, \frac{1}{1-\gamma^t}$,

So in DNNs you are in a stochastic setting, you are sampling from a mini-batch, there is going to be a lot of randomness in the forward pass. In other words, you are getting a lot of noisy gradients.

- Momentum contribution: So instead of using every gradient at every single time step we only use the decaying sum of previous gradients which helps use stabilize your gradient direction a bit.
- Cache contribution: To make sure that the steps size workout in steep and shallow directions.

Usually $\mu = 0.9, \gamma = 0.995$ but if you cross-validate them it can help a bit.

Why using momentum update instead of Nesterov Momentum update? It could be also done but it does not make such a difference.

IMPORTANT NOTATION!!: Through all this chapter I am saying parameters = weights, but remember that there are other ones like the bias.

Learning Rate - Decay over time

So which is the best learning rate for training DNNs? None. The best one is to start with high learning first because it optimizes faster than the "good learning rate", in this way you do really fast progress. But at some point you are going to be too stochastic and you can not converge to a minimum. So, at this point you must start decreasing the learning rate.

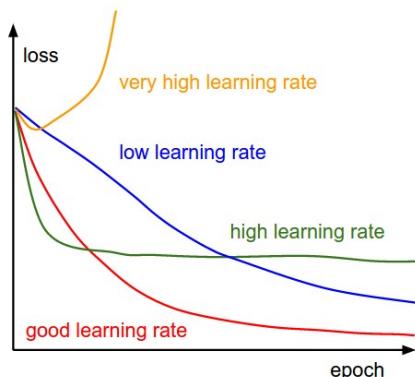


FIGURE 8.3: Learning rate selection

Another way of seeing it is to have in mind is that with a high learning rate, the system contains too much kinetic energy and the parameter vector bounces around chaotically, unable to settle down into deeper, but narrower parts of the loss function.

Knowing when to decay the learning rate can be tricky: Decay it slowly and you'll be wasting computation bouncing around chaotically with little improvement for a long time. But decay it too aggressively and the system will cool too quickly, unable to reach the best position it can. There are three common types of implementing the learning rate decay:

- **Step decay:** Reduce the learning rate by some factor every few epochs. Typical values might be reducing the learning rate by a half every 5 epochs, or by 0.1 every 20 epochs.

These numbers depend heavily on the type of problem and the model. One heuristic you may see in practice is to watch the validation error while training with a fixed learning rate, and reduce the learning rate by a constant (e.g. 0.5) whenever the validation error stops improving.

- **Exponential decay** has the mathematical form $\alpha = \alpha_0 e^{-kt}$, where α_0, k are hyperparameters and t is the iteration number (but you can also use units of epochs).
- **$1/t$ decay** has the mathematical form $\alpha = \frac{\alpha_0}{1+Kt}$ where α_0, k are hyperparameters and t is the iteration number.

In practice, we find that the step decay dropout is slightly preferable because the hyperparameters it involves (the fraction of decay and the step timings in units of epochs) are more interpretable than the hyperparameter k . Lastly, if you can afford the computational budget, err on the side of slower decay and train for a longer time.

Notice that Adagrad, RMSProp and Adam already have a learning decay. When using these methods it may help a little bit also adding a learning decay method. For SGD and SGD+Momentum you should always add the learning decay.

Second Order Optimization Methods

The advantages of second order optimization methods are: faster convergence and no hyperparameters. However they are not normally used because they are too heavy, too much stuff happening, and they use too much memory. It is preferable just to use first order methods.

The idea of second order methods is that they approximate the loss function in a more precise way. As first order methods, they approximate with a hyperplane of like which way are we sloping. Also they also approximate with Hessian (H) which is telling you how is this surface curving. So you need the gradient for the sloping estimation and the Hessian for the surface curvature.

For example in the Newton method, once you have formed this bowl like Hessian approximation to your objective function, you can use this information to jump directly to the minimum of that approximation, and you keep iterating this strategy.

$$J(\theta) \approx J(\theta_0) + (\theta - \theta_0)^T \nabla_{\theta} J(\theta_0) + \frac{1}{2} (\theta - \theta_0)^T H(\theta - \theta_0) \theta^* = \theta_0 - H^{-1} \nabla_{\theta} J(\theta_0) \quad (8.7)$$

This method is impractical to train DNNs because it would take too much memory. So imagine you have 10 million parameters, then your Hessian matrix would be of size 10 million by 10 million and then you want to invert it, so good luck.

There is a variant of the Newton method called BGFS which approximates the inverse Hessian with rank 1 over time so you do not need to compute the inverse of the Hessian but still requires to store the gigantic matrix in memory.

Another extension called L-BGFS does not require you to store all the full inverse matrix. This method is used some time to train DNNs. It usually works very good in full batch and deterministic functions with no stochastic settings. However, it does not transfer very well to mini-batch settings because the approximation it is doing of the Hessian matrix are incorrect when you switch the batch. Also, it does not work well with stochastic settings, so you have to remove all sources of randomness (e.g. dropout). Still, it is too heavy, too much stuff is happening, and it is better just to use first order methods.

Summary

- Adam is a good default choice in most cases. Some times it helps adding learning rate decay like step decay or exponential decay even-though Adam already has inside a learning decay.
- If you can afford to do full batch updates then try out L-BFGS (do not forget to disable all sources of noise/randomness)
- Decay your learning rate over the period of the training. For example, halve the learning rate after a fixed number of epochs, or whenever the validation accuracy tops off.

IMPORTANT NOTATION!!: Through all this chapter I am saying parameters = weights, but remember that there are other ones like the bias.

Chapter 9

Dropout

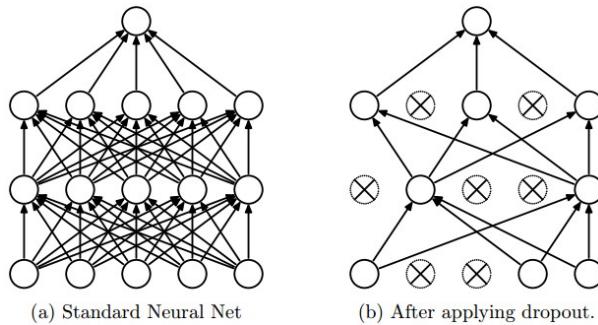


FIGURE 9.1: Dropout

“Randomly set %50 some neurons to zero in the forward pass.” There two ways of understanding why this works:

- This makes the network unable to rely on a single feature. Thus, forcing it to generate redundant representation. In this way, you will be representing an object with redundant descriptors, detectors, etc. So in test time, if some features can not be detected, you still can rely on the other ones.



FIGURE 9.2: Relay on multiple features

- It can also be seen as training a large ensemble of models that share parameters. When applying a mask to shutdown some randomly neurons at the forward pass this mask stays there for the backward pass. Thus, in the backprop step, no gradient will flow through

the neurons that were shut off to zero; so its weights to the previous layer will not be updated. In other words, the neurons that were dropout do not update their connections to the previous layer, just as if they were not there. With dropout you are subsampeling a part of your neural network and training it with the current example. So each binary mask is one model which gets trained on only one datapoint (or more in the case that two random mask are equal during training).

WARNING: If dropout is not working for you, you should probably be using a bigger network.

Train time

During the forward pass set 50% of the neurons to 0. To get an idea the code below shows a naive implementation of dropout

```
""" Vanilla Dropout: Not recommended implementation (see notes below) """
p = 0.5 # probability of keeping a unit active. higher = less dropout

def train_step(X):
    """ X contains the data """

    # forward pass for example 3-layer neural network
    H1 = np.maximum(0, np.dot(W1, X) + b1)
    U1 = np.random.rand(*H1.shape) < p # first dropout mask
    H1 *= U1 # drop!
    H2 = np.maximum(0, np.dot(W2, H1) + b2)
    U2 = np.random.rand(*H2.shape) < p # second dropout mask
    H2 *= U2 # drop!
    out = np.dot(W3, H2) + b3

    # backward pass: compute gradients... (not shown)
    # perform parameter update... (not shown)
```

Test time

At test time ideally we would like to integrate out all the noise (not possible). A Monte Carlo approximation would be: do many forward passes with different dropout masks, average all predictions. Unfortunately its not time efficient. Instead, you can approximate this method without not droping out any unit in the forward pass during testing but we have to be careful.

Example

Imagine this linear NN and a dropout $prob = 0.5$. During training time the expected value is the average between all the possible masks activation value. However, if we use all the neurons

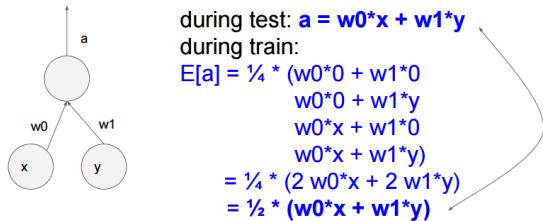


FIGURE 9.3: Example

in test time, the activation value is going to be twice the expected value in train time. So it is easy to see that this half comes from the fact that at train time we dropout neurons with probability 50%.

With $p = 0.5$ dropout prob, using all inputs in the forward pass would inflate the activations by $2x$ from what the network was "used to" during training. Thus, you have to compensate by scaling the activations back down by $1/2$. In this way, the output at test time is equal to the expected output at training time. Let's see a naive implementation:

```
""" Vanilla Dropout: Not recommended implementation (see notes below) """
p = 0.5 # probability of keeping a unit active. higher = less dropout

def predict(X):
    # ensembled forward pass
    H1 = np.maximum(0, np.dot(W1, X) + b1) * p # NOTE: scale the activations
    H2 = np.maximum(0, np.dot(W2, H1) + b2) * p # NOTE: scale the activations
    out = np.dot(W3, H2) + b3
```

A better way of implementing dropout is using "inverted dropout". The idea is very simple, instead of multiplying the activation values by the dropout probability during test time, divide the training activation values by the dropout probability during train time. In this way, test time is unchanged.

```
"""
Inverted Dropout: Recommended implementation example.
We drop and scale at train time and don't do anything at test time.
"""

p = 0.5 # probability of keeping a unit active. higher = less dropout

def train_step(X):
    # forward pass for example 3-layer neural network
    H1 = np.maximum(0, np.dot(W1, X) + b1)
    U1 = (np.random.rand(*H1.shape) < p) / p # first dropout mask. Notice /p!
    H1 *= U1 # drop!
    H2 = np.maximum(0, np.dot(W2, H1) + b2)
    U2 = (np.random.rand(*H2.shape) < p) / p # second dropout mask. Notice /p!
    H2 *= U2 # drop!
    out = np.dot(W3, H2) + b3
```

```
# backward pass: compute gradients... (not shown)
# perform parameter update... (not shown)

def predict(X):
    # ensembled forward pass
    H1 = np.maximum(0, np.dot(W1, X) + b1) # no scaling necessary
    H2 = np.maximum(0, np.dot(W2, H1) + b2)
    out = np.dot(W3, H2) + b3
```

There is a method to dropout single weights instead of full neurons call dropconnect

Chapter 10

Hyper-Parameters selection and babysitting

Sanity test

Make sure that you can overfit very small portion of the training data. Take 20 samples, turn off regularization and make sure that you can get a loss of 0. If you can't overfit there is a problem: or there is something broken or you have to scale up your network.

Learning Rate

Start with small regularization (0.00001) and find learning rate that makes the loss not go down. Then, find learning rate that makes the loss go down but at some point it explodes. The learning rate is in between this range.

The screenshot shows two code cells in a Jupyter notebook. The left cell contains code for initializing a two-layer model and training it with a learning rate of 3e-3. The output shows the loss increasing rapidly across all 10 epochs, indicating that the learning rate is too high. The right cell contains similar code but with a learning rate of 1e-6, resulting in a loss that remains nearly constant across all 10 epochs, indicating that the learning rate is too low.

```
model = init_two_layer_model(32*32*3, 50, 10) # input size, hidden size, number of classes
trainer = ClassifierTrainer()
best_model, stats = trainer.train(X_train, y_train, X_val, y_val,
                                  model, two_layer_net,
                                  num_epochs=10, reg=0.00001,
                                  update='sgd', learning_rate_decay=1,
                                  sample_batches=True,
                                  learning_rate=3e-3, verbose=True)

Finished epoch 1 / 10: cost 2.186654, train: 0.308000, val 3.000000e-03
Finished epoch 2 / 10: cost 2.176230, train: 0.330000, val 0.350000, lr 3.000000e-03
Finished epoch 3 / 10: cost 1.942257, train: 0.376000, val 0.352000, lr 3.000000e-03
Finished epoch 4 / 10: cost 1.827868, train: 0.329000, val 0.310000, lr 3.000000e-03
Finished epoch 5 / 10: cost inf, train: 0.128000, val 0.128000, lr 3.000000e-03
Finished epoch 6 / 10: cost inf, train: 0.144000, val 0.147000, lr 3.000000e-03

3e-3 is still too high. Cost explodes....
```

```
model = init_two_layer_model(32*32*3, 50, 10) # input size, hidden size, number of classes
trainer = ClassifierTrainer()
best_model, stats = trainer.train(X_train, y_train, X_val, y_val,
                                  model, two_layer_net,
                                  num_epochs=10, reg=0.00001,
                                  update='sgd', learning_rate_decay=1,
                                  sample_batches=True,
                                  learning_rate=1e-6, verbose=True)

Finished epoch 1 / 10: cost 2.302576, train: 0.080000, val 0.103000, lr 1.000000e-06
Finished epoch 2 / 10: cost 2.302582, train: 0.121000, val 0.124000, lr 1.000000e-06
Finished epoch 3 / 10: cost 2.302558, train: 0.119000, val 0.138000, lr 1.000000e-06
Finished epoch 4 / 10: cost 2.302519, train: 0.127000, val 0.151000, lr 1.000000e-06
Finished epoch 5 / 10: cost 2.302517, train: 0.158000, val 0.171000, lr 1.000000e-06
Finished epoch 6 / 10: cost 2.302518, train: 0.179000, val 0.172000, lr 1.000000e-06
Finished epoch 7 / 10: cost 2.302466, train: 0.180000, val 0.176000, lr 1.000000e-06
Finished epoch 8 / 10: cost 2.302452, train: 0.175000, val 0.185000, lr 1.000000e-06
Finished epoch 9 / 10: cost 2.302459, train: 0.206000, val 0.192000, lr 1.000000e-06
Finished epoch 10 / 10: cost 2.302420, train: 0.190000, val 0.192000, lr 1.000000e-06
finished optimization. best validation accuracy: 0.192000
```

Loss barely changing

FIGURE 10.1: **Left:** Loss exploding, **Right:** Loss not going down

- Loss not going down: learning rate too low. Something funny can happen in this case. Loss not going down but accuracy improving until 20%. How is that possible? So because

of the way softmax is computed, small changes in the lost can cause small changes in the scores which make the correct class has a tiny bigger score than the others. This, makes the softmax classifier get more samples correct.

- Loss exploding (NaN happens): learning rate too high

Hyperparameter Optimization - Coarse to fine serach

Tuning regularization and learning rate. Do coarse \rightarrow fine for cross-validation in stages. In other words, first do a rough search, see what it works, and keep iterating to longer narrow in ranges that are working. First, only a few epochs to get rough idea of what parameters work (few minutes is enough). Second, longer running time, finer search.

Also, it is better to optimize parameters in log space because reg and learning rate work manipulatively in the dynamics of your back propagation.

For example: run coarse search for 5 epochs

```

max_count = 100
for count in xrange(max_count):
    reg = 10**uniform(-5, 5)
    lr = 10**uniform(-3, -6)

    trainer = ClassifierTrainer()
    model = init two layer model(32*32*3, 50, 10) # input size, hidden size, number of classes
    trainer = ClassifierTrainer()
    best_model_local, stats = trainer.trainX(train, y_train, X_val, y_val,
                                              model, two_layer_net,
                                              num_epochs=5, reg=reg,
                                              update='momentum', learning_rate_decay=0.9,
                                              sample_batches=True, batch_size=100,
                                              learning_rate=lr, verbose=False)

    [val acc: 0.412000, lr: 1.405206e-04, reg: 4.793564e-01, (1 / 100)]
    [val acc: 0.214000, lr: 7.231888e-06, reg: 2.321281e-04, (2 / 100)]
    [val acc: 0.208000, lr: 2.119571e-06, reg: 8.011857e+01, (3 / 100)]
    [val acc: 0.196000, lr: 1.551131e-05, reg: 4.374936e-05, (4 / 100)]
    [val acc: 0.079000, lr: 1.753300e-05, reg: 1.200424e+03, (5 / 100)]
    [val acc: 0.223000, lr: 4.215128e-05, reg: 4.196174e+01, (6 / 100)]
    [val acc: 0.441000, lr: 1.750259e-04, reg: 2.110807e-04, (7 / 100)]
    [val acc: 0.241000, lr: 6.749231e-05, reg: 4.226413e+01, (8 / 100)]
    [val acc: 0.482000, lr: 4.296863e-04, reg: 6.642555e-01, (9 / 100)]
    [val acc: 0.079000, lr: 5.401602e-06, reg: 1.599828e+04, (10 / 100)]
    [val acc: 0.154000, lr: 1.618500e-06, reg: 4.925252e-01, (11 / 100)]

```

nice → note it's best to optimize in log space!

FIGURE 10.2: First pass - coarse search

Now run finer search...

```

max count = 100
for count in xrange(max_count):
    reg = 10**uniform(-5, 5)
    lr = 10**uniform(-3, -6)

    [val acc: 0.527000, lr: 5.340517e-04, reg: 4.897824e-01, (1 / 100)]
    [val acc: 0.492000, lr: 1.726484e-04, reg: 7.991248e-01, (2 / 100)]
    [val acc: 0.468000, lr: 6.608423e-04, reg: 1.349727e-02, (3 / 100)]
    [val acc: 0.461000, lr: 1.928377e-04, reg: 1.228193e-02, (4 / 100)]
    [val acc: 0.469000, lr: 1.113730e-04, reg: 5.244309e-02, (5 / 100)]
    [val acc: 0.498000, lr: 9.477776e-04, reg: 2.001293e-03, (6 / 100)]
    [val acc: 0.469000, lr: 1.000000e-04, reg: 1.000000e-04, (7 / 100)]
    [val acc: 0.536000, lr: 5.586201e-04, reg: 2.312689e-04, (8 / 100)]
    [val acc: 0.536000, lr: 5.808183e-04, reg: 8.259964e-02, (9 / 100)]
    [val acc: 0.489000, lr: 1.979168e-04, reg: 1.810889e-04, (10 / 100)]
    [val acc: 0.496000, lr: 2.036031e-04, reg: 2.406271e-03, (11 / 100)]
    [val acc: 0.475000, lr: 2.021622e-04, reg: 2.287807e-01, (12 / 100)]
    [val acc: 0.518000, lr: 6.947668e-04, reg: 1.562780e-02, (13 / 100)]
    [val acc: 0.518000, lr: 9.471549e-04, reg: 1.433895e-03, (14 / 100)]
    [val acc: 0.509000, lr: 3.140888e-04, reg: 2.857518e-01, (15 / 100)]
    [val acc: 0.514000, lr: 6.438349e-04, reg: 3.035781e-01, (16 / 100)]
    [val acc: 0.509000, lr: 9.752279e-04, reg: 2.705984e-01, (17 / 100)]
    [val acc: 0.509000, lr: 2.952279e-04, reg: 2.858685e-03, (18 / 100)]
    [val acc: 0.509000, lr: 2.412048e-04, reg: 4.997821e-04, (19 / 100)]
    [val acc: 0.466000, lr: 1.119314e-04, reg: 1.189935e-02, (20 / 100)]
    [val acc: 0.516000, lr: 8.039527e-04, reg: 1.528291e-02, (21 / 100)]

```

adjust range → 53% - relatively good for a 2-layer neural net with 50 hidden neurons.

FIGURE 10.3: Second pass - finer search. Notice that there is a problem with this result. The best result (last red box) has a lr close to the boundary of search that we have set (-3). So it may we better results waiting for lower values of lr. **Careful with best values on border**

Hyper-parameter Optimization - NEVER do grid (iterative) search of parameters, do random search

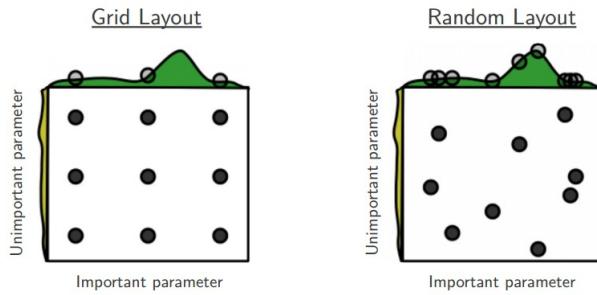


FIGURE 10.4: Second pass - finer search. Notice that there is a problem with this result. The best result (last red box) has a lr close to the boundary of search that we have set (-3). So it may be better results waiting for lower values of lr. **Careful with best values on border**

For cross-validation, use random search in stead of grid search. The issue is that one of the parameters may be much important than another one.

In this figure in particular its more important the x than the y dimension. Then, with random sampling, you are going to evaluate more different samples of the x parameter space (9 with random vs 3 with grid).

Hyper-parameter Optimization - Monitor and visualize the loss curve

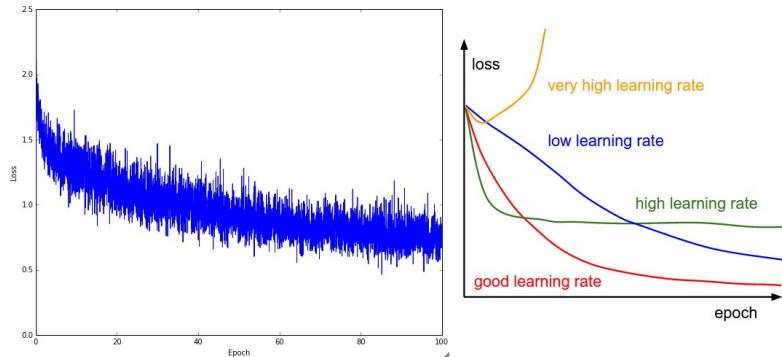


FIGURE 10.5: In this left case the loss is too slow... The learning rate is too low

Hyper-parameter Optimization - Monitor and visualize the accuracy

The gap between the training and validation accuracy indicates the amount of overfitting. Two possible cases are shown in the diagram on the left. The blue validation error curve shows very

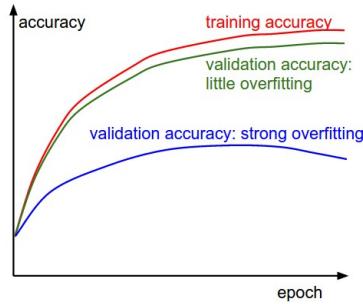


FIGURE 10.6: Monitor and visualize the accuracy

small validation accuracy compared to the training accuracy, indicating strong overfitting (note, it's possible for the validation accuracy to even start to go down after some point). When you see this in practice you probably want to increase regularization (stronger L2 weight penalty, more dropout, etc.) or collect more data. The other possible case is when the validation accuracy tracks the training accuracy fairly well. This case indicates that your model capacity is not high enough: make the model larger by increasing the number of parameters.

- gap between trainval is too big: overfitting, increase regularization
- gap between trainval too small: increase model capacity

In this figure case there is a big gap so it is probably over-fitting. We should increase the regularization factor.

Hyper-parameter Optimization - Track the ratio of weight updates / weight magnitudes

You want `weight_updates / weight_magnitudes = 0.001`.

- If this is too high to decrease learning rate
- If it is too low to increase learning rate

Dropout not working

If dropout is not working for you, you should probably be using a bigger network.

Chapter 11

Other definitions

11.1 Embedding

An embedding maps an input representation, such as a word or sentence, into a vector. A popular type of embedding are word embeddings such as word2vec or GloVe. We can also embed sentences, paragraphs or images. For example, by mapping images and their textual descriptions into a common embedding space and minimizing the distance between them, we can match labels with images. Embeddings can be learned explicitly, such as in word2vec, or as part of a supervised task, such as Sentiment Analysis. Often, the input layer of a network is initialized with pre-trained embeddings, which are then fine-tuned to the task at hand.

11.2 Gradient Clipping

Gradient Clipping is a technique to prevent exploding gradients in very deep networks, typically Recurrent Neural Networks. There exist various ways to perform gradient clipping, but the a common one is to normalize the gradients of a parameter vector when its L2 norm exceeds a certain threshold according to `new_gradients = gradients * threshold / l2_norm(gradients)`.

11.3 Vanishing Gradient Problem

The vanishing gradient problem arises in very deep Neural Networks, typically Recurrent Neural Networks, that use activation functions whose gradients tend to be small (in the range of 0 from 1). Because these small gradients are multiplied during backpropagation, they tend to vanish throughout the layers, preventing the network from learning long-range dependencies. Common ways to counter this problem is to use activation functions like ReLUs that do not suffer from small gradients, or use architectures like LSTMs that explicitly combat vanishing gradients. The opposite of this problem is called the exploding gradient problem.

Chapter 12

Tricks

Real Ensambles

- Train multiple independent models
- At test time average their results

Enjoy $\sim 2\%$ extra performance.

Simulated Ensambles

Fake ensambles. As you are training, normally, you save a checkpoint after each epoch to figure out what was your validation performance at that point. At test time average multiple model checkpoints of a single model. Enjoy $\sim 1\%$ extra performance

Average parameter vector

Keep track of (and use at test time) a running average parameter vector. `params_test` is a running sum exponentially decaying. One way to visualize it is the following. Think in a case of optimizing a bowl like function and you are bouncing around the minimum, then taking the average of all this steps gets you closer to the minimum.

```
while True:  
    data_batch = dataset.sample_data_batch()  
    loss = network.forward(data_batch)  
    dparams = network.backward()  
    params += update_params() //instead of using this parameters at test time  
    params_test = 0.995*params_test + 0.005*params //use this ones
```

This can give you a small boost.

Use float32 instead of float64

Just that. In fact if you can work with float16 do it.

ReLU bias initialization

It is a good practice to initialize them with a slightly positive initial bias to avoid "dead neurons".

Chapter 13

Visualization

13.1 t-SNE visualization

Embed high-dimensional points so that locally, pairwise distances are conserved i.e. similar things end up in similar places. Dissimilar things end up wherever. Pronounced "tisni".

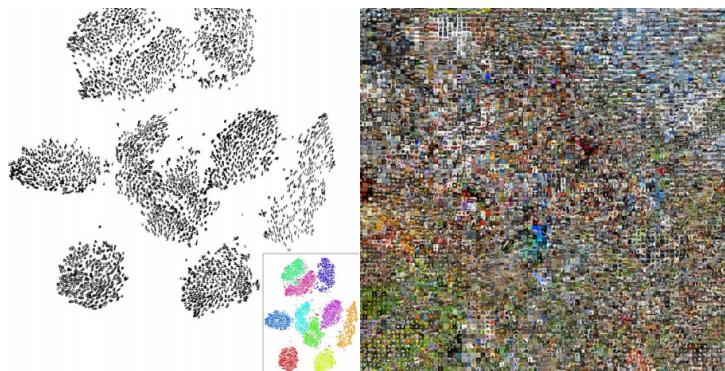


FIGURE 13.1: **Left:** Example embedding of MNIST digits (0-9) in 2D, **Right:** CIFAR-10 representation warped in a square shape

More CIFAR-10 examples [here](#)

13.2 Occlusions

The idea is to create a hit map of the output probability of the CNN of the image in the left corresponding to the correct label as you slide an occlusion across the image.

This helps you understand what are the regions which contains the most valuable features, for the CNN, in the image.

For example, in the first row, we can observe that the probability of the left image being a Pomeranian decreases when we cover its face.

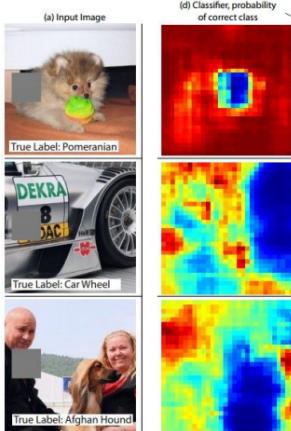
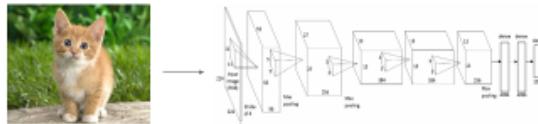


FIGURE 13.2: Visualize with occlusions

13.3 Visualizing Activations

How can we compute the gradient of any arbitrary neuron in the network w.r.t. the image?

1. Feed image into net



2. Pick a layer, set the gradient there to be all zero except for one 1 for some neuron of interest
3. Backprop to image:

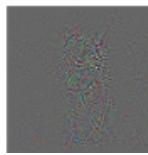


FIGURE 13.3: Visualize with occlusions

The output image gradients is not that good to understand what is going on. We will now see two strategies to obtain "better looking" gradients: Deconv approaches and

13.4 Deconv approaches

We are trying to see what parts of the input image are exciting a specific neuron. The idea to get a better looking gradient image is only to take into account the positive gradients. In this way we remove the noise caused by mixing positive and negative gradients.

The ReLU layer only backprop the gradient of the activations that where positive at the forward pass.

With Deconv, we must hack this behavior and change it so the ReLU layers only backprop positive gradients and do not care if the forward activations where positive or negative.

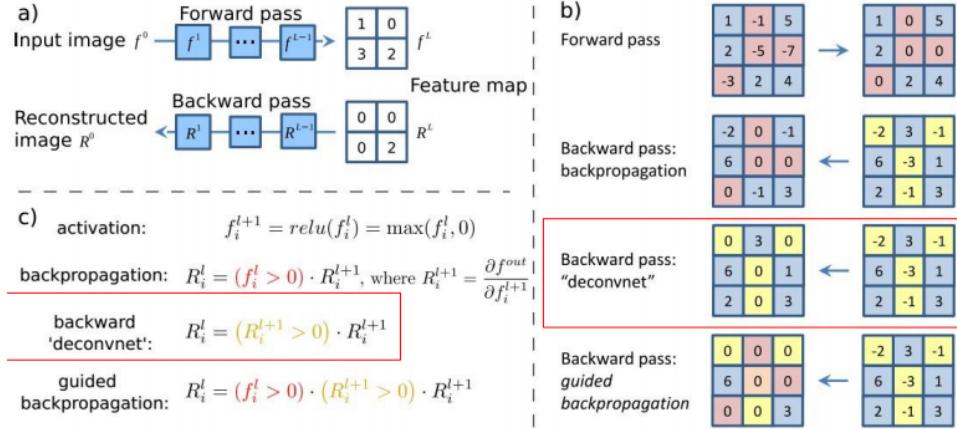


FIGURE 13.4: Deconv approaches

Now we can show what excites a specific neuron. The figures below show a grid of neurons at each layer. For each neuron we can see the image that arrived to that neuron and what parts of this image excited the neuron. So the deeper the neuron the bigger the area of the original image it observes. Also, it can be seen that the deeper the neuron the higher feature representation.

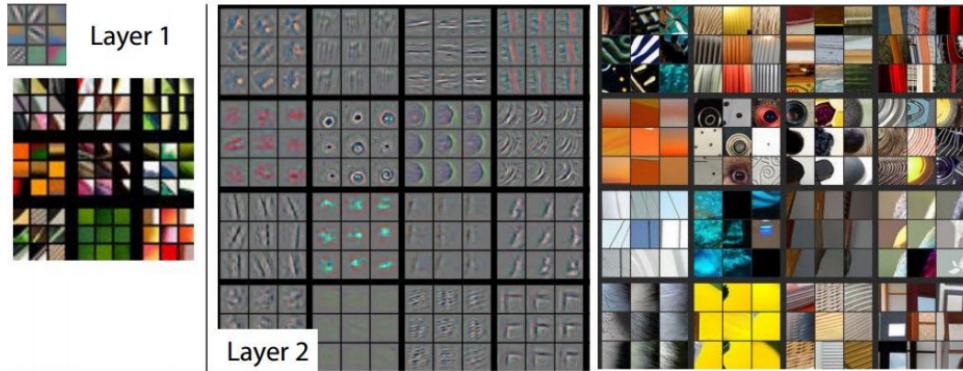


FIGURE 13.5: Deconv approaches layers 1 and 2

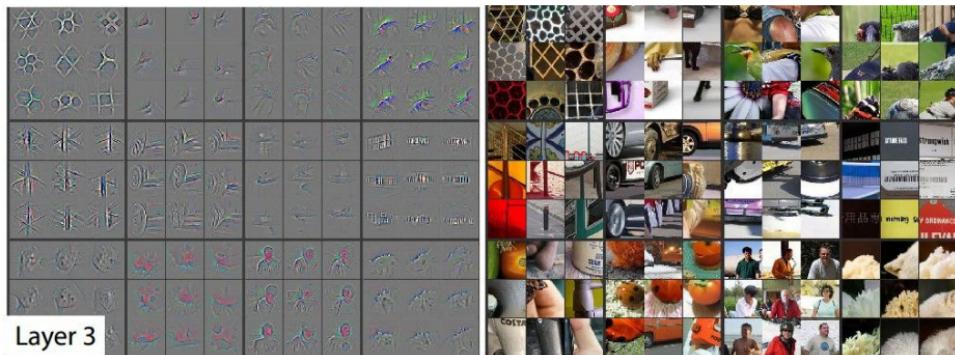


FIGURE 13.6: Deconv approaches layer 3

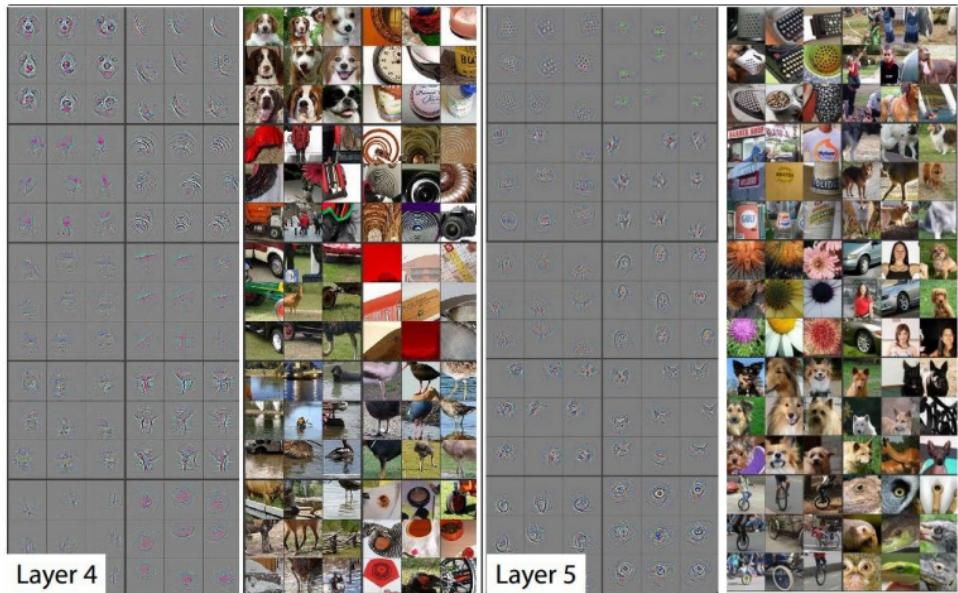


FIGURE 13.7: Deconv approaches layers 4 and 5

13.5 Optimization to image

How can we find an image that maximizes some class score?

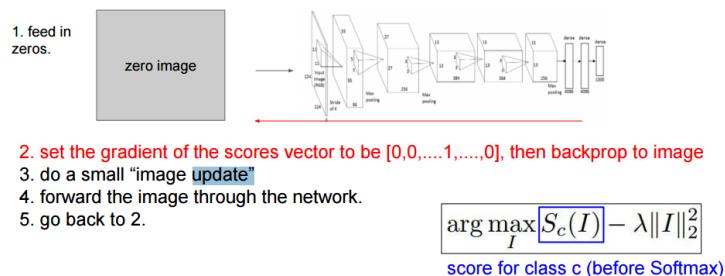


FIGURE 13.8: Optimize an image to maximize the score of a specific class

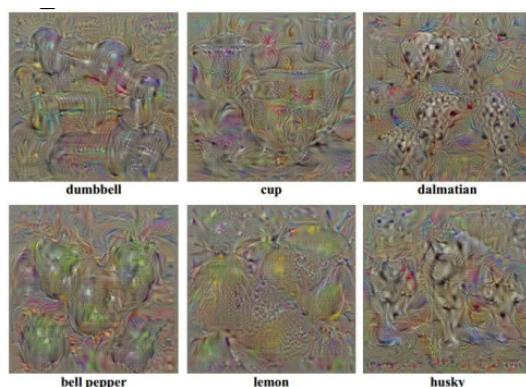


FIGURE 13.9: Optimize an image to maximize the score of a specific class output

13.6 Visualizing Data Gradients

1. Feed an input image
2. Set the gradient of the scores vector be $[0,0,\dots,0,1,0,\dots,0]$ where 1 is the ground-truth label class of the image
3. Backprop to the image
4. For each pixel of the image store the max gradient value across all the channels
5. Plot the gradients

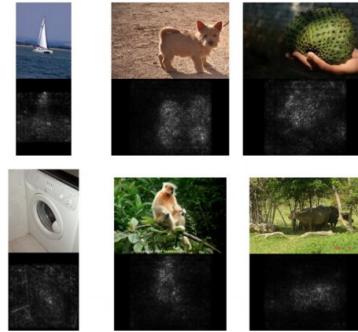


FIGURE 13.10: Data gradients output

The interpretation of the results are that the stuff that is black is not influencing the score of the image. So you can wiggle them and the score will not change. So at the end it is telling you the area of the image that is influencing the network.

13.7 DeepDream

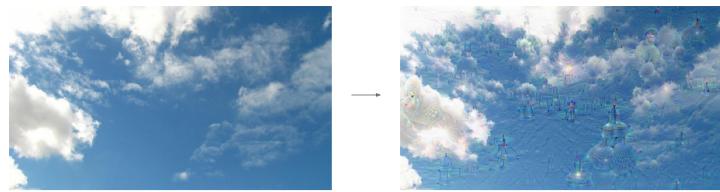


FIGURE 13.11: DeepDream output

DeepDream [link](#)

It normally hallucinates a lot of dogs because there are a lot of dog samples in ImageNet (because there are a lot of different dog classes).

At the end, the only thing DeepDream is doing is: given a layer immediately after a ReLU layer, it sets the gradients equal to the activation values. So if you iteratively update the image with the gradients, you are modifying the image to increase whatever it is exiting the selected layer.

```

def objective_L2(dst):
    dst.diff[:] = dst.data # DeepDream: set dx = x :)

def make_step(net, step_size=1.5, end='inception_4c/output',
             jitter=32, clip=True, objective=objective_L2):
    '''Basic gradient ascent step.'''
    src = net.blobs['data'] # input image is stored in Net's 'data' blob
    dst = net.blobs[end]

    ox, oy = np.random.randint(-jitter, jitter+1, 2)
    src.data[0] = np.roll(np.roll(src.data[0], ox, -1), oy, -2) # apply jitter shift

    net.forward(end=end)
    objective(dst) # specify the optimization objective
    net.backward(start=end)
    g = src.diff[0]
    # apply normalized ascent step to the input image
    src.data[:] += step_size/np.abs(g).mean() * g # "image update"

    src.data[0] = np.roll(np.roll(src.data[0], -ox, -1), -oy, -2) # unshift image

    if clip:
        bias = net.transformer.mean['data']
        src.data[:] = np.clip(src.data, -bias, 255-bias)

```

FIGURE 13.12: DeepDream code

In other words, DeepDream modifies the image in a way that “boosts all activations. This creates a feedback loop: e.g. any slightly detected dog face will be made more and more dog like over time.

13.8 DeepArt

Explained at cs231n lecture 9

13.9 Adversarial data - Fooling networks

It is very simple to fool a CNN. Images are super high dimensional objects (one dimension for each pixel). The training images are constrained to a small manifold and we are training ConvNets over it. The trained ConvNet works extremely well inside that manifold, but outside of it is complete random.

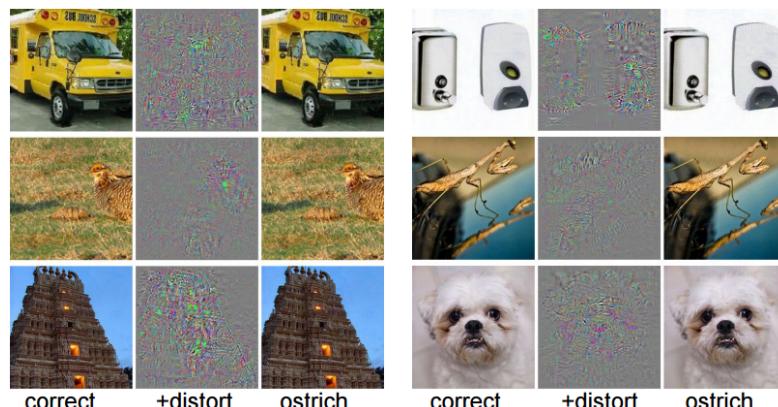


FIGURE 13.13: Adversarial data examples

The think is that during the forward pass we are applying dot product over all the dimensions of the data (one per pixel) so if we change the input a tiny bit but all in the correct way, the dot product output is going to change a lot.

X	2	-1	3	-2	2	2	1	-4	5	1	← input example
W	-1	-1	1	-1	1	-1	1	1	-1	1	← weights
adversarial x	1.5	-1.5	3.5	-2.5	2.5	1.5	1.5	-3.5	4.5	1.5	

class 1 score before:
 $-2 + 1 + 3 + 2 + 2 - 2 + 1 - 4 - 5 + 1 = -3$
 $\Rightarrow \text{probability of class 1 is } 1/(1+e^{(-3)}) = 0.0474$

$-1.5+1.5+3.5+2.5-1.5+1.5-3.5-4.5+1.5 = 2$
 $\Rightarrow \text{probability of class 1 is now } 1/(1+e^{(-2)}) = 0.88$
i.e. we improved the class 1 probability from 5% to 88%

This was only with 10 input dimensions. A 224x224 input image has 150,528.
 (It's significantly easier with more numbers, need smaller nudge for each)

FIGURE 13.14: Adversarial data case example

This is not a problem of deep learning, its is a problem in learning in general. There are some ways of decreasing this effect. One is to train also with adversarial examples; it will be more robust to them but the classifier accuracy will decrease. Another one is to instead of classifying the hole image, classify chunks.

Part III

LAYERS

Chapter 14

Input Layer

The input layer (that contains the image) should be divisible by 2 many times. Common numbers include 32 (e.g. CIFAR-10), 64, 96 (e.g. STL-10), or 224 (e.g. common ImageNet ConvNets), 384, and 512.

Chapter 15

Convolutional layer

Structure

- Accepts a volume of size $W_1 \times H_1 \times D_1$
- Requires four hyperparameters:
 - Number of filters K
 - their spatial extent F
 - the stride S
 - the amount of zero padding P
- Produces a volume of size $W_2 \times H_2 \times D_2$ where:
 - $W_2 = \frac{W_1 - F + 2P}{S + 1}$
 - $H_2 = \frac{H_1 - F + 2P}{S + 1}$
 - $D_2 = K$
- With parameter sharing, it introduces FFD_1 weights per filter, for a total of FFD_1K weights and K biases.
- The filters depth is always equal to the input volume depth.
- The output volume depth is equal to the number of filters
- In the output volume, the d -th depth slice (of size $W_2 \times H_2$) is the result of performing a valid convolution of the d -th filter over the input volume with a stride of S , and then offset by d -th bias.
- Normally, in the same conv layer, all filters have the same dimensions so that special optimized routines can be invoked.

Explanation

In a convolution layer we have filters. Each filter is a weight matrix which is convoluted over the input volume. Each time the filter is applied it outputs a single value. The result of convolving the filter over all the input volume is another volume called activation map. A filter's depth is always equal to the input volume depth. You can imagine a filter as a neuron.

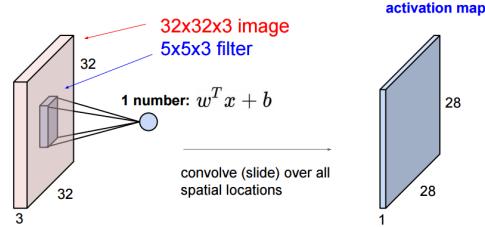


FIGURE 15.1: Convolution filter

Instead of having only one filter, we have a stack of filter which produce a stack of activation maps (one for each filter). In this example this conv layer has a stack of 6 filters ($5 \times 5 \times 3$). All positions in the same activation map share the same weights (filter)

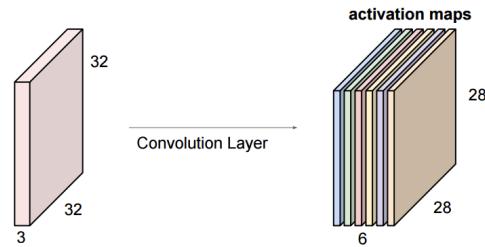


FIGURE 15.2: 6 convolution filters output

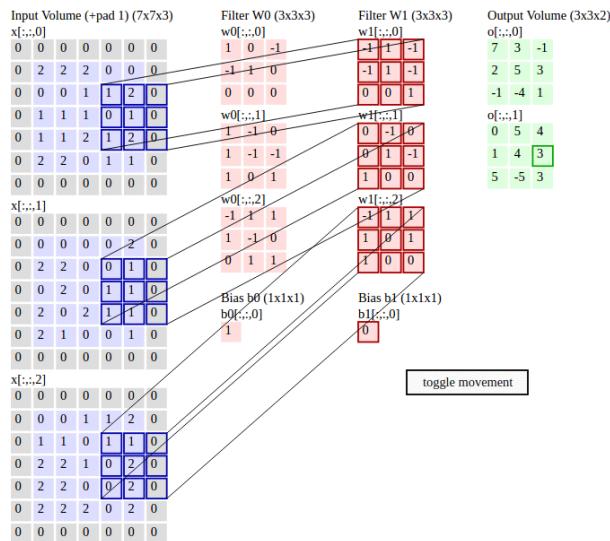


FIGURE 15.3: Example of a conv layer with 2 filters ($3 \times 3 \times 3$) over an input volume of $(7 \times 7 \times 3)$ with 1 padding, which produces and output map of $(3 \times 3 \times 2)$

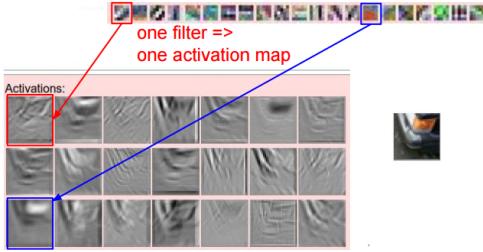


FIGURE 15.4: Example of the activation maps produced in a convolution layer. The small images on top are the stack of filters, the image on the right the input volume, and the grey-scale images the activation maps. White corresponds to high activations and black to low activations. Notice for example, that the filter marked in blue has an orange part, so when you slide this filter through the input image there are a lot of high activations in the area corresponding to the orange part of the input image. The output of this convolutional layer is going to be a stack of all these activation matrices.

Padding Add zeros around the input volume. The output volume decreases after each conv layer, so if we would not pad, the size of the activation maps will decrease very fast and after a few layers the volume would be $1 \times 1 \times ?$. This behavior is not desired because in deep learning we want to have a lot of layers. Why padding with zeros and not an extension of the image for example? Because in this way the padded cells do not contribute to the filter.

Choosing hyperparameters Should be using small filters (e.g. 3×3 or at most 5×5), using a stride of $S = 1$, and crucially, padding the input volume with zeros in such way that the conv layer does not alter the spatial dimensions of the input. A common setting of the hyperparameters are: $F = 3, S = 1, P = 1$; $F = 5, S = 1, P = 2$; $F = 5, S = 2, P = ?$ and $F = 1, S = 1, P = 0$. K is usually a power of two value (e.g. 32,64,128,512) because libraries normally have optimized routines. If you must use bigger filter sizes (such as 7×7 or so), it is only common to see this on the very first conv layer that is looking at the input image. For a general F , it can be seen that $P = (F - 1)/2$ preserves the input size.

Parameter sharing In an activation map, all the positions (neurons) share the same weights (filter). It makes sense to share weights because for example if the filter is searching for edges, it makes sense that it search edges in all the image. Moreover, sharing weights spatially avoids over-fitting the filter. It also helps reducing the number of parameters.

Backpropagation The backward pass for a convolution operation (for both the data and the weights) is also a convolution (but with spatially-flipped filters). This is easy to derive in the 1-dimensional case with a toy example (not expanded on for now).

1×1 convolution As an aside, several papers use 1×1 convolutions, as first investigated by Network in Network. Some people are at first confused to see 1×1 convolutions especially when they come from signal processing background. Normally signals are 2-dimensional so 1×1

convolutions do not make sense (its just pointwise scaling). However, in ConvNets this is not the case because one must remember that we operate over 3-dimensional volumes, and that the filters always extend through the full depth of the input volume. For example, if the input is $[32 \times 32 \times 3]$ then doing 1×1 convolutions would effectively be doing 3-dimensional dot products (since the input depth is 3 channels).

Dilated convolutions A recent development (e.g. see paper by Fisher Yu and Vladlen Koltun) is to introduce one more hyperparameter to the CONV layer called the dilation. So far weve only dicussed CONV filters that are contiguous. However, its possible to have filters that have spaces between each cell, called dilation. As an example, in one dimension a filter w of size 3 would compute over input x the following: $w[0]*x[0] + w[1]*x[1] + w[2]*x[2]$. This is dilation of 0. For dilation 1 the filter would instead compute $w[0]*x[0] + w[1]*x[2] + w[2]*x[4]$; In other words there is a gap of 1 between the applications. This can be very useful in some settings to use in conjunction with 0-dilated filters because it allows you to merge spatial information across the inputs much more agressively with fewer layers. For example, if you stack two 3×3 CONV layers on top of each other than you can convince yourself that the neurons on the 2nd layer are a function of a 5×5 patch of the input (we would say that the effective receptive field of these neurons is 5×5). If we use dilated convolutions then this effective receptive field would grow much quicker.

The power of small filters

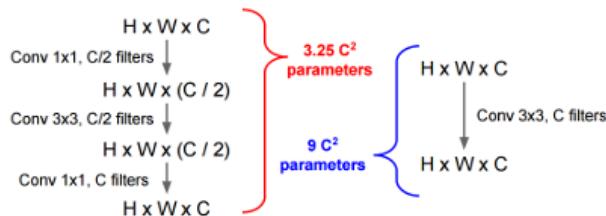


FIGURE 15.5: Network in network

3 Conv layers of 3×3 filters have the same receptive field as a 1 Conv layer of 7×7 filters. But lets compare both options:

Suppose input is $H \times W \times C$ and we use convolutions with C filters to preserve depth (stride 1, padding to preserve H, W)

- three CONV with 3×3 filters
 - Number of weights: $3 \times C \times (3 \times 3 \times C) = 27C^2$
 - Number of multiply-adds: $3 \times (H \times W \times C) \times (3 \times 3 \times C) = 27HWC^2$
- one CONV with 7×7 filters

- Number of weights: $C \times (7 \times 7 \times C) = 49C^2$
- Number of multiply-adds: $(Hx \times W \times C) \times (7 \times 7 \times C) = 49HWC^2$

So 3 Conv layers of 3×3 filters have the same receptive field of a larger Conv layer of 7×7 filters but have a lower computation and memory complexity. Moreover, if we have ReLU layers after each Conv layer the 3 Conv layer option is more nonlinear (good).

Lets get crazy. ResNet and GoogLeNet go one step further and use 1x1 filters all over the place to even reduce further the number of parameters. This is commonly known as "Network in Network"

Chapter 16

Pooling layer

Structure

- Accepts a volume of size
- Requires two hyperparameters: $W_1 \times H_1 \times D_1$
 - their spatial extent F
 - the stride S
- Produces a volume of size where: $W_2 \times H_2 \times D_2$
 - $W_2 = \frac{W_1 - F}{S} + 1$
 - $H_2 = \frac{H_1 - F}{S} + 1$
 - $D_2 = D_1$
- Introduces zero parameters since it computes a fixed function of the input
- Note that it is not common to use zero-padding for Pooling layers
- Has **NO** parameters

Explanation

It is common to periodically insert a Pooling layer in-between successive Conv layers in a ConvNet architecture. Its function is to progressively reduce the spatial size of the representation to reduce the amount of parameters and computation in the network, and hence to also control overfitting. This down sampling operations happens on each activation map independently and the volume depth is preserved. The most common way of pooling is max pooling, another one is average pooling but it does not work that well.

The most common form is a pooling layer with filters of size 2×2 applied with a stride of 2 downsamples every depth slice in the input by 2 along both width and height, discarding 75% of the activations. Every MAX operation would in this case be taking a max over 4 numbers (little 2×2 region in some depth slice). The depth dimension remains unchanged.

Notice that the pooling layer has NO parameters!!

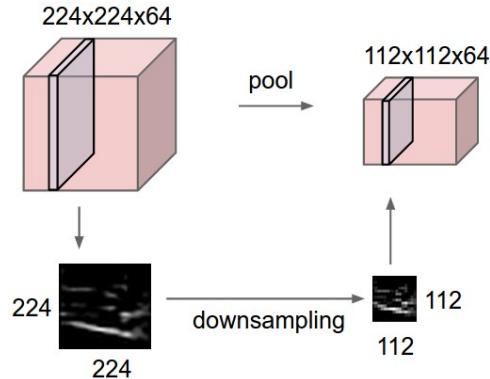


FIGURE 16.1: In this example, the input volume of size $[224 \times 224 \times 64]$ is pooled with filter size 2, stride 2 into output volume of size $[112 \times 112 \times 64]$. Notice that the volume depth is preserved.

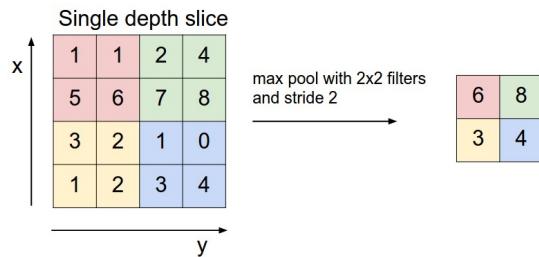


FIGURE 16.2: The most common downsampling operation is max, giving rise to max pooling, here shown with a stride of 2. That is, each max is taken over 4 numbers (little 2×2 square).

Choosing hyperparameters There are 2 common setting of the hyperparameters. $F = 2, S = 2$ that this discards exactly 75% of the activations in an input volume (due to downsampling by 2 in both width and height). Another slightly less common setting is t . It is very uncommon to see receptive field sizes for max pooling that are larger than 3 because the pooling is then too lossy and aggressive. This usually leads to worse performance.

Backpropagation Recall from the backpropagation chapter that the backward pass for a $\text{max}(x, y)$ operation has a simple interpretation as only routing the gradient to the input that had the highest value in the forward pass. Hence, during the forward pass of a pooling layer it is common to keep track of the index of the max activation (sometimes also called the switches) so that gradient routing is efficient during backpropagation.

Getting rid of pooling Many people dislike the pooling operation and think that we can get away without it. For example, Striving for Simplicity: The All Convolutional Net proposes to discard the pooling layer in favor of architecture that only consists of repeated CONV layers. To reduce the size of the representation they suggest using larger stride in CONV layer once in a while. Discarding pooling layers has also been found to be important in training good generative models, such as variational autoencoders (VAEs) or generative adversarial networks (GANs). It seems likely that future architectures will feature very few to no pooling layers.

Chapter 17

Batch Normalization layer

You have a minibatch of data and you are taking it through your network. You have inserted batch normalization layers through the layer which take your input x and make sure that every feature dimension across the batch you have unit gaussian activations.

Basic Idea

Batch normalization potentially helps in two ways: faster learning and higher overall accuracy. The improved method also allows you to use a higher learning rate, potentially providing another boost in speed.

Why does this work? Well, we know that normalization (shifting inputs to zero-mean and unit variance) is often used as a pre-processing step to make the data comparable across features. As the data flows through a deep network, the weights and parameters adjust those values, sometimes making the data too big or too small again - a problem the authors refer to as "internal covariate shift". By normalizing the data in each mini-batch, this problem is largely avoided.

Basically, rather than just performing normalization once in the beginning, you're doing it all over place. Of course, this is a drastically simplified view of the matter (since for one thing, I'm completely ignoring the post-processing updates applied to the entire network), but hopefully this gives a good high-level overview.

Adding batch normalization normally slows 30%.

Extended explanation

So image you have N samples in your minibatch and D **features / neuron activations** at some points. So this matrix of X is the input to the batch normalization layer. Evaluates

empirical mean and variance along every feature. So it makes sure that each column of X is a unit gaussian. You can do this because its perfectly fine to transform it to unit Gaussian because it is differentiable so you can backpropagate.

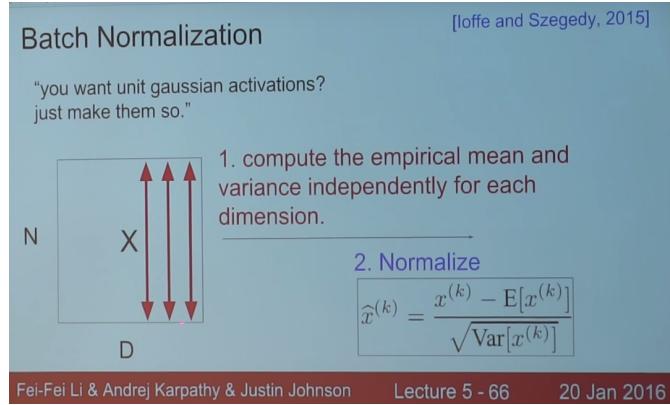


FIGURE 17.1: Compute empirical mean and variance for each dimension

So what you usually have are fully connected or convolutional layers followed by batch normalization layer before the non-linearity. So they ensure that everything is roughly unit Gaussian at each step of the neural net. One problem is that its not clear that tanh wants exactly unit Gaussian. Because you want tanh to be able to make its outputs more or less defused (more or less saturated) so right now it would not be able to do that.

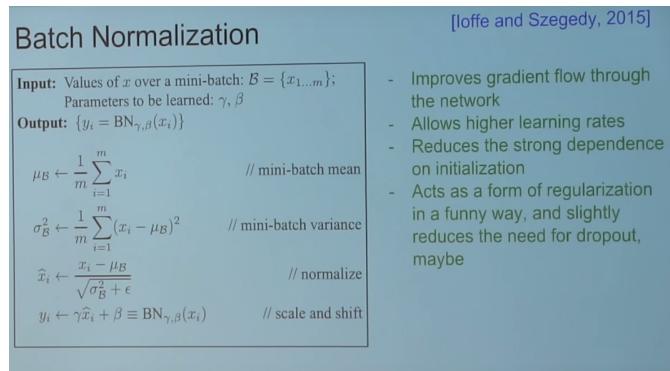


FIGURE 17.2: In network

To solve this you do not only normalize X but you also allow the network to shift by gamma and add B . So after you have centred your data you are allowing the network through the backprop to shift and scale the distribution. Also note that the network can learn to undo this layer (it can learn to have the batch normalization layer to be an identity)

[2 good (from fig 17.4)] As you are swiping through different choices of initialization values with and without batch norm you'll see a huge difference. With batch norm it will work with much bigger settings of the initial scale so you don't have to worry as much, it really helps.

[3 good (from fig 17.4)] It acts somehow as a way of regularization because with batch norm when you have some kind of input x and it goes through the network its representation in some

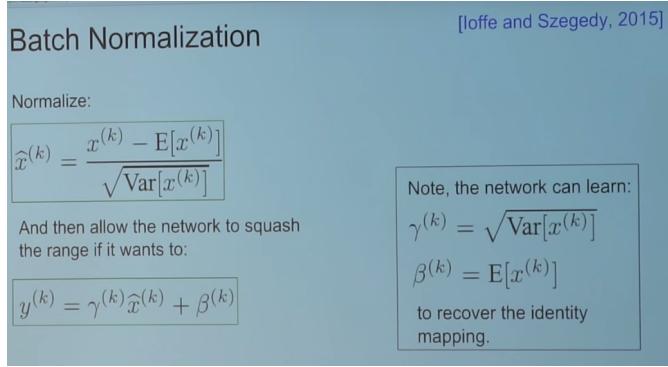


FIGURE 17.3: Equations

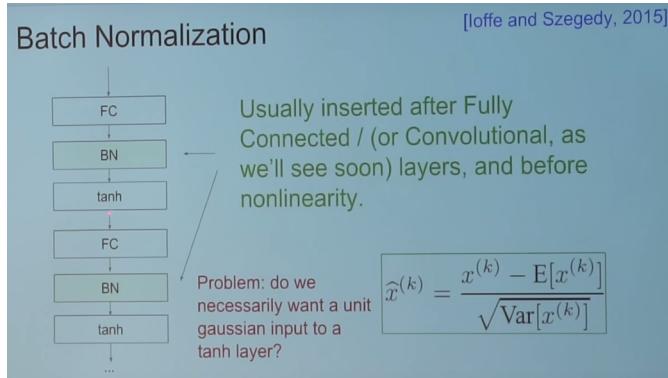


FIGURE 17.4: Algorithm

layer of the network is basically not only function of it but also whatever other examples are with x in the batch. Normally all examples are processed independently in parallel, but batch norm tides them together and so your representation at some layer is a function on whatever batch you happen to be sampled in at what it does is to jigger your place in the representation space in that layer. Which is a nice regularizer effect.

During testing it works a little different. During test you what this to be a deterministic function. μ and σ are the once that you used during training.

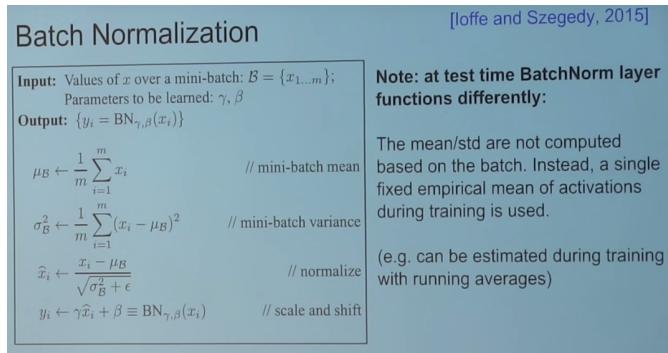


FIGURE 17.5: During testing it works a little different

Chapter 18

Upsampling Layer (“Transposed Convolution”)

Learnable Upsampling

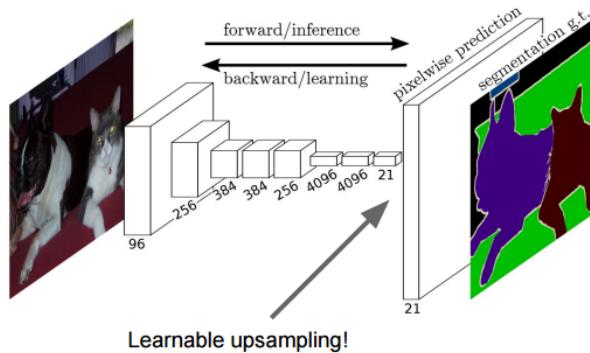


FIGURE 18.1: From the paper: “Fully Convolutional Networks for Semantic Segmentation”

Transposed Convolution

Transposed convolutions – also called fractionally strided convolutions – work by swapping the forward and backward passes of a convolution. One way to put it is to note that the kernel defines a convolution, but whether its a direct convolution or a transposed convolution is determined by how the forward and backward passes are computed.

The transposed convolution is implemented as the backward pass of a corresponding non-transposed convolution. It can be thought of as dilating the input (by adding “stride - 1” zeros between adjacent input elements), padding it with the needed number of zeros so it is not out. And then, apply the convolution with the filter flipped 180 degrees.

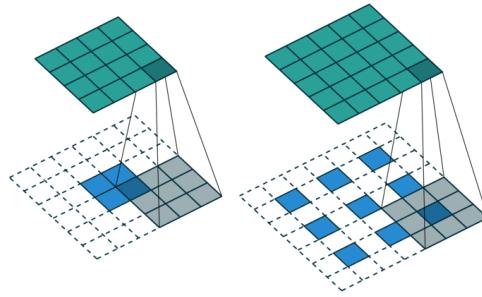


FIGURE 18.2: **Left:** Visually, for a transposed convolution with stride one and no padding, we just pad the original input (blue entries) with zeroes (white entries). **Right:** Stride two and padding, the transposed convolution would look like this

Skip Connections

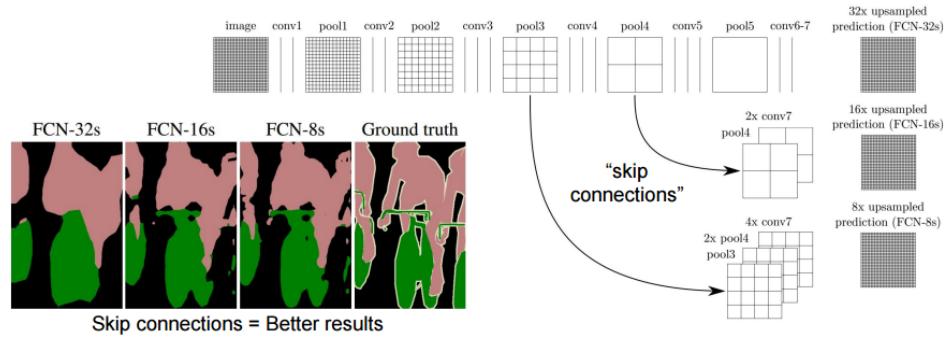


FIGURE 18.3: Skip Connections

Chapter 19

Fully Connected Layer

FC layers, also called affine layers, produce the high-level reasoning in the DNN. Neurons in a fully connected layer have full connections to all activations in the previous layer, as in regular Neural Networks. Their activations can hence be computed with a matrix multiplication followed by a bias offset.

Chapter 20

Highway Layer

A Highway Layer is a type of Neural Network layer that uses a gating mechanism to control the information flow through a layer. Stacking multiple Highway Layers allows for training of very deep networks. Highway Layers work by learning a gating function that chooses which parts of the inputs to pass through and which parts to pass through a transformation function, such as a standard affine layer for example. The basic formulation of a Highway Layer is $T * h(x) + (1 - T) * x$, where T is the learned gating function with values between 0 and 1, $h(x)$ is an arbitrary input transformation and x is the input. Note that all of these must have the same size.

Part IV

NETWORKS

Chapter 21

Recurrent Neural Networks (RNN)

A recurrent neural network (RNN) is a class of artificial neural network where connections between units form a directed cycle. This creates an internal state of the network which allows it to exhibit dynamic temporal behavior. Unlike feedforward neural networks, RNNs can use their internal memory to process arbitrary sequences of inputs. This makes them applicable to tasks such as unsegmented connected handwriting recognition or speech recognition.

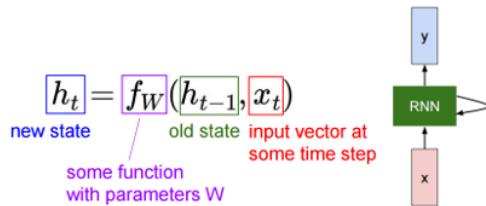


FIGURE 21.1: RNN

Notice that the same function and the same set of parameters are used at every time step! We can not have RNNs of enormous length because we have to store in memory the hidden state of each time step to be able to back-propagate. Normally we will have RNNs of max 25 length. To process bigger sequences we will divide the sequence in chunks of 25 and the last hidden state of a chunk is the initial hidden state of the next chunk.

Let's see some wire examples (fig 21.2):

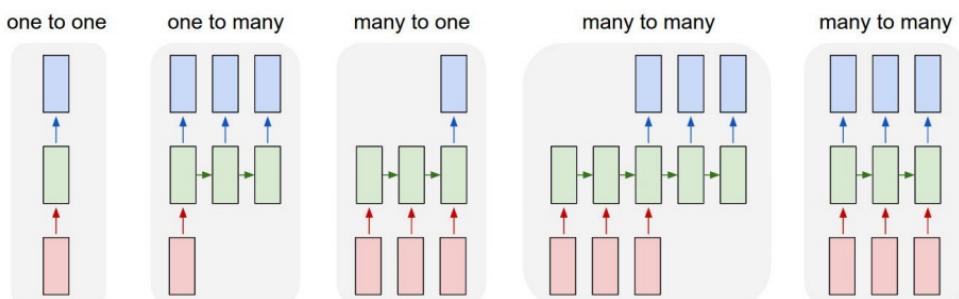


FIGURE 21.2: RNN examples

1. Vanilla Neural Networks
2. Image Captioning (image to sequence of words)
3. Sentiment Classification (sequence of words to sentiment)
4. Machine Translation (seq of words to seq of words)
5. Video classification on frame level

We can stack RNNs together to produce deeper RNN. In figure 21.4 case we have stack 3 RNNs. It still works the same as before but now we have 3 weights.

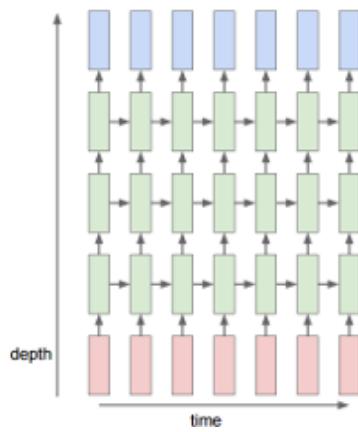


FIGURE 21.3: Stacked RNNs

Practical Example

Lets create a RNN that given a sequence of characters predicts the next one. The output is always the prob of each letter in the vocabulary to be the next character:

- Vocabulary: [h,l,e,o]
- Example training sequence: "hello"

For this example we will use a Vanilla RNN:

$$h_t = \tanh(W_{hh}h_{t-1} + W_{xh}x_t) \quad (21.1)$$

$$y_t = W_{hy}h_t \quad (21.2)$$

$$h_0 = 0 \quad (21.3)$$

A 100 lines of code implementation of a character recognition in python is implemented here:
<https://gist.github.com/karpathy/d4dee566867f8291f086>

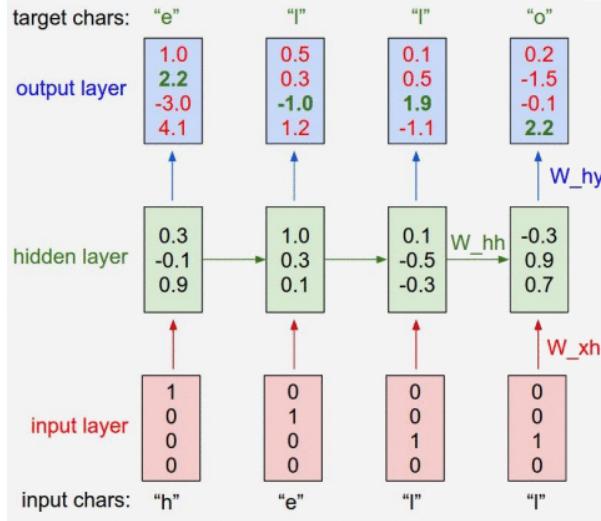


FIGURE 21.4: Practical example. You can see this blue boxes being softmax classifier, in other words, at each time step there is a softmax classifier.

Image Captioning

Example using RNN for image captioning

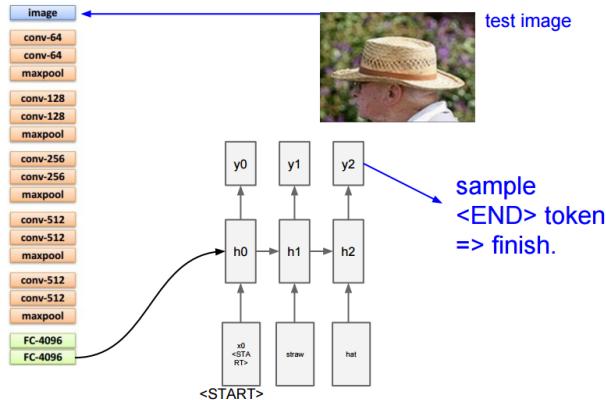


FIGURE 21.5: "Deep Visual-Semantic Alignments for Generating Image Descriptions", Karpathy and Fei-Fei

21.1 LSTM

Normally we will not use Vanilla RNNs, instead, all papers use LSTM. It is very similar, it stills take into account the input and the last state but now the combination of both is more complex and works better. With RNN we have one vector h at each time step. But with LSTM we have two vectors at each time step h_t and c_t . Moreover in RNNs the repeating module only has one layer, \tanh . Instead, LSTM has four.

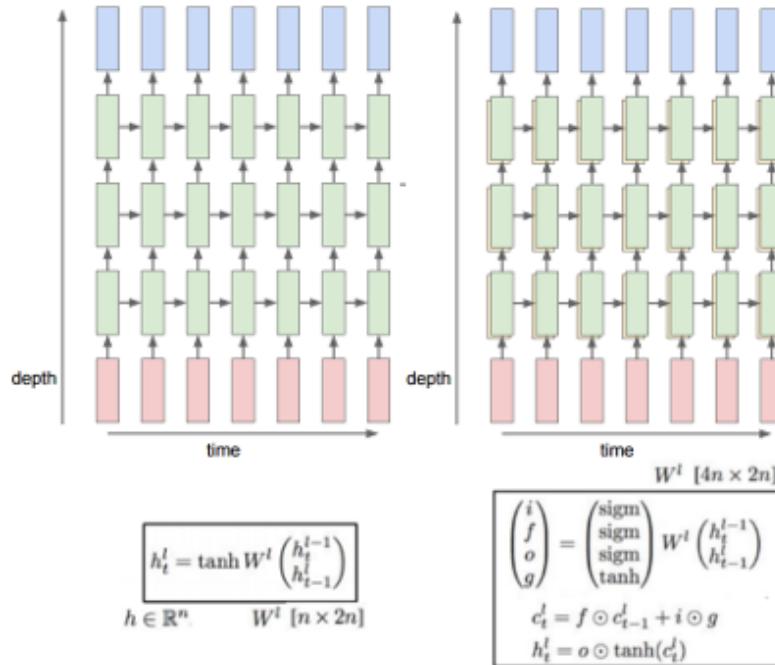


FIGURE 21.6: **Left:** RNN with h (green), **Right:** LSTM with h (hidden vector, green) and c (cell state vector, yellow)

How do they work?

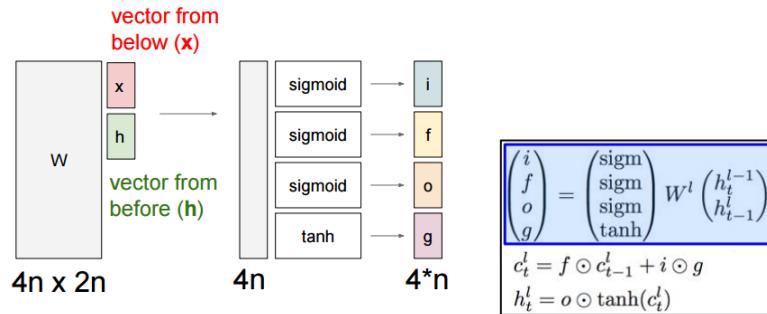


FIGURE 21.7: LSTM first equation diagram

where:

- c cells. Are best though as counters
- h hidden states
- h_t^{l-1} vector from below of size $n \times 1$ (x in the figure)
- h_{t-1}^l vector from before of size $n \times 1$ (h in the figure)
- $i \in [0, 1]$ to chose if we want to add, or not, to a cell
- $f \in [0, 1]$ forget gate to reset cells to 0

- $o \in [0, 1]$ to choose which cells are used to produce
- $g \in [-1, 1]$ to add -1 or 1 to a cell

The key to LSTMs is the cell state, the horizontal line running through the top of the diagram. The cell state is kind of like a conveyor belt. It runs straight down the entire chain, with only some minor linear interactions. Its very easy for information to just flow along it unchanged. The LSTM does have the ability to remove or add information to the cell state, carefully regulated by structures called gates.

Think of i, f, o as boolean variables, we want them to have an interpretation like a gate. They are a result of a sigmoid to make them differentiable. They allow us to reset and add to counters, as well as to choose what cells should be used to update h_t^l . We can do two operations to cells (counters):

- reset them with $f \odot c_{t-1}^l$
- add -1 or 1 with $i \odot g$
- choose what cells should be used with o to update h_t^l : $o \odot \tanh(c_t^l)$

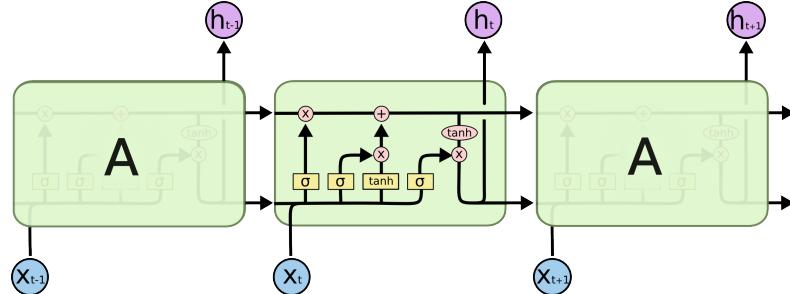


FIGURE 21.8: LSTM diagram

Step by step

First Decide what information were going to throw away from the cell state. This decision is made by a sigmoid layer called the “forget gate layer. It looks at h_{t-1} and x_t , and outputs a number between 0 and 1 for each number in the cell state C_{t-1} . A 1 represents “completely keep this while a 0 represents “completely get rid of this. For example, in a language model trying to predict the next word based on all the previous ones the cell state might include the gender of the present subject, so that the correct pronouns can be used. When we see a new subject, we want to forget the gender of the old subject.

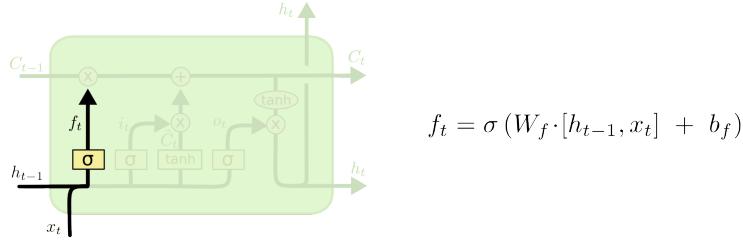


FIGURE 21.9: First, decide what information were going to throw away from the cell state

Second Decide what new information were going to store in the cell state. This has two parts. First, a sigmoid layer called the “input gate layer decides which values well update. Next, a *tanh* layer creates a vector of new candidate values, \tilde{C}_t , that could be added to the state. In the next step, well combine these two to create an update to the state. For example, in the example of our language model, wed want to add the gender of the new subject to the cell state, to replace the old one were forgetting.

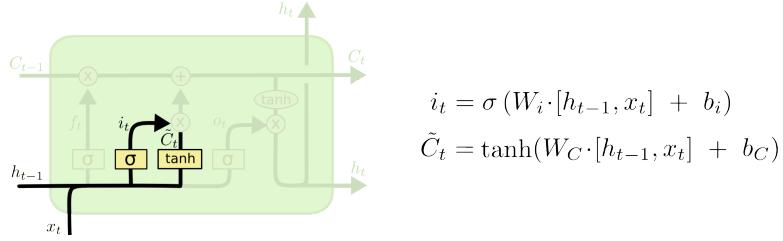


FIGURE 21.10: Second, decide what information were going to store in the cell state

Third Update the old cell state, C_{t-1} , into the new cell state C_t . The previous steps already decided what to do, we just need to actually do it. We multiply the old state by f_t , forgetting the things we decided to forget earlier. Then we add $i_t \tilde{C}_t$. This is the new candidate values, scaled by how much we decided to update each state value. In the case of the language model, this is where wed actually drop the information about the old subjects gender and add the new information, as we decided in the previous steps.

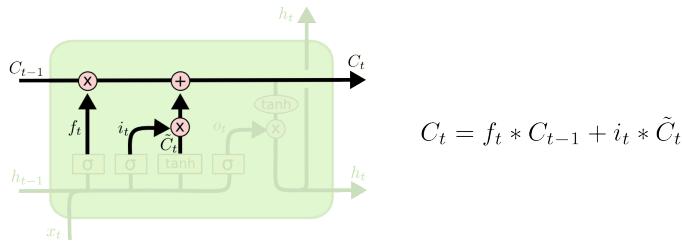


FIGURE 21.11: Third, update the old cell state

Fourth Finally, we decide what were going to output. This output will be based on our cell state, but will be a filtered version. First, we run a sigmoid layer which decides what parts of the cell state were going to output. Then, we put the cell state through *tanh* (to push the values

to be between -1 and 1) and multiply it by the output of the sigmoid gate, so that we only output the parts we decided to. For the language model example, since it just saw a subject, it might want to output information relevant to a verb, in case that's what is coming next. For example, it might output whether the subject is singular or plural, so that we know what form a verb should be conjugated into if that's what follows next.

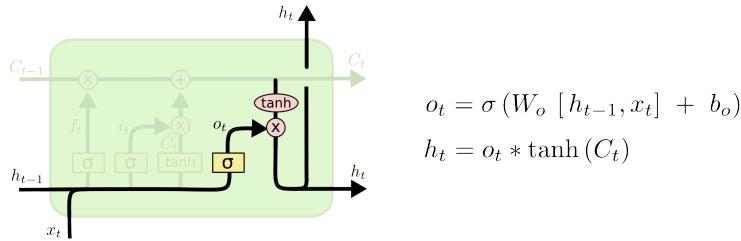


FIGURE 21.12: Fourth, decide what were going to output

Variants on LSTMs

But not all LSTMs are the same as the above. In fact, it seems like almost every paper involving LSTMs uses a slightly different version. The differences are minor, but it's worth mentioning some of them. Greff, et al. (2015) do a nice comparison of popular variants, finding that they're all about the same.

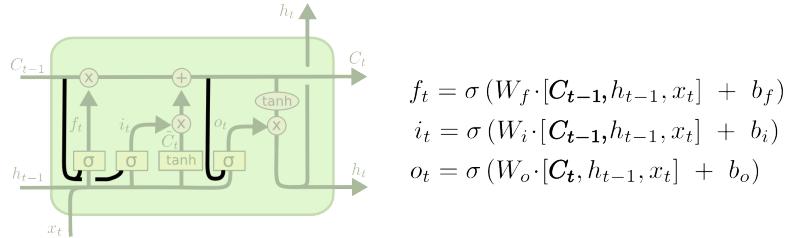


FIGURE 21.13: One popular LSTM variant, introduced by Gers & Schmidhuber (2000), is adding “peephole connections. This means that we let the gate layers look at the cell state. The diagram adds peepholes to all the gates, but many papers will give some peepholes and not others.

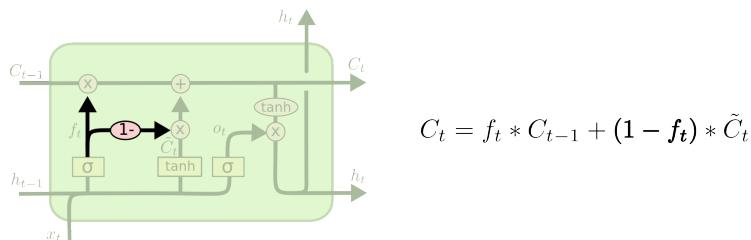


FIGURE 21.14: Another variation is to use coupled forget and input gates. Instead of separately deciding what to forget and what we should add new information to, we make those decisions together. We only forget when we're going to input something in its place. We only input new values to the state when we forget something older.

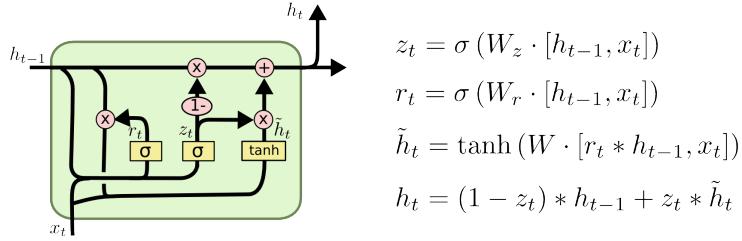


FIGURE 21.15: A slightly more dramatic variation on the LSTM is the Gated Recurrent Unit, or GRU, introduced by Cho, et al. (2014). It combines the forget and input gates into a single “update gate”. It also merges the cell state and hidden state, and makes some other changes. The resulting model is simpler than standard LSTM models, and has been growing increasingly popular.

These are only a few of the most notable LSTM variants. There are lots of others, like Depth Gated RNNs by Yao, et al. (2015). There's also some completely different approach to tackling long-term dependencies, like Clockwork RNNs by Koutnik, et al. (2014).

Difference between RNN and LSTM

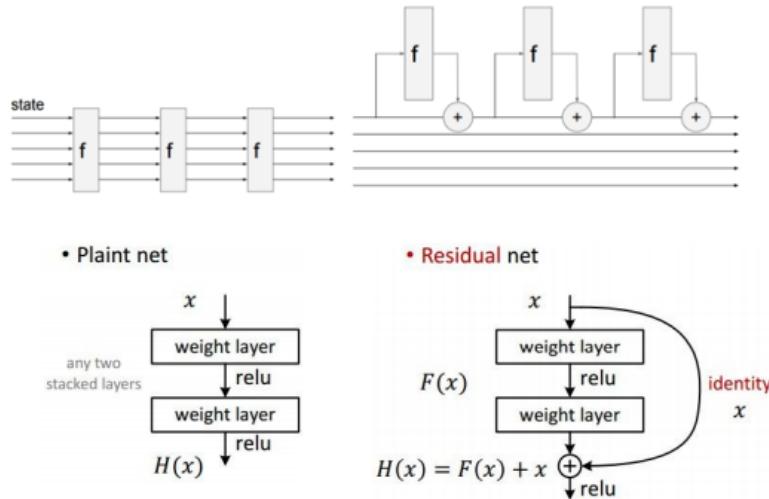


FIGURE 21.16: Difference between RNN (left) and LSTM (right)

- RNN transformative interaction of the state, LSTM additive interaction of the state.
- In RNN you are operating and transforming the state vector. So you are changing your hidden state from time step to time step. Instead, LSTM has these cells states flowing through. A subset of cells (or all of them) to compute the hidden state. Then, based on hidden state, we decide how to operate over the cell. We can reset it and/or adding interaction.
- RNNs look identical to plain nets, LSTMs to ResNets.
- RNNs has vanishing gradients, so you can not learn dependencies between distant time steps. LSTMs do not have the problem of vanishing gradients. In a video in evernote

they introduce gradient noise to a layer and show how it evolves. We can see that RNN automatically kills it while LSTM maintains it more time. This means that RNN can not learn long term relationships.

21.1.1 Bidirectional RNN

A Bidirectional Recurrent Neural Network is a type of Neural Network that contains two RNNs going into different directions. The forward RNN reads the input sequence from start to end, while the backward RNN reads it from end to start. The two RNNs are stacked on top of each others and their states are typically combined by appending the two vectors. Bidirectional RNNs are often used in Natural Language problems, where we want to take the context from both before and after a word into account before making a prediction.

Summary

- RNNs allow a lot of flexibility in architecture design
- Vanilla RNNs are simple but don't work very well
- Common to use LSTM or GRU: their additive interactions improve gradient flow
- Backward flow of gradients in RNN can explode or vanish. Exploding is controlled with gradient clipping. Vanishing is controlled with additive interactions (LSTM)
- Better/simpler architectures are a hot topic of current research
- Better understanding (both theoretical and empirical) is needed.

Chapter 22

Autoencoders

The idea of autoencoders is that we want to force a network to try to reconstruct our data and hopefully it will learn a useful representation of the data. For traditional autoencoders this is use for feature learning and for variational autoencoders to generate fake samples.

22.1 Traditional Autoencoder

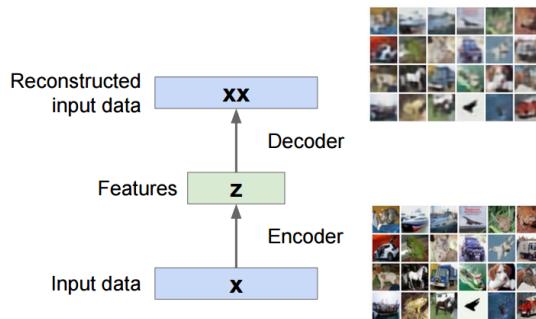


FIGURE 22.1: Traditional Autoencoder

Normally encoder will be a 4-layer conv, and the decoder a 4-layer upconv. Sometimes they share weights (in fact they are the same, we only have to transpose them): $\dim(x) = D, \dim(z) = H, w_e : H \times D, w_d : D \times H = w_e^T$.

The idea is to pass the input data through an encoder network that will produce features z . This step can be thought as PCA. We are transforming the input data and transforming it into another feature representation. The encoder network is usually a ReLU CNN. Normally z is smaller than x (dimensionality reduction).

The problem is that we do not have any explicit labels to use to know which information is relevant and which one its not to decide how to make the dimensionality reduction. To decide which is the most relevant data we reconstruct the input data only with the information in the z features and compare the reconstruction with the original image.

Specially if the z features are small, hopefully, it will force the network to summarize the useful statistics and to discover useful features of the input image.

How do we train them?

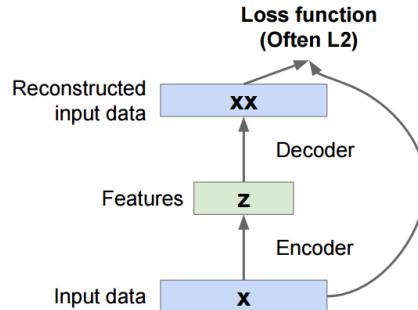


FIGURE 22.2: Traditional Autoencoder

Normally it is trained as a regular NN with euclidean distance as the loss function comparing the input data vs the reconstructed data.

Normally, after training, the decoder will be thrown away.

Unsupervised + Supervised learning

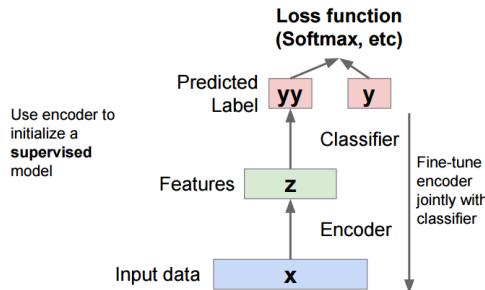


FIGURE 22.3: Traditional Autoencoder

The idea is first to train the network with unsupervised data, and then fine tune with a small labeled dataset. This idea is beautiful in the sense that we could train our networks with millions of unlabeled data from the internet, and then fine tune the network with a labeled dataset such as KITTI. The problem is that it is yet not working that well.

22.2 Variational Autoencoder

Transforming traditional autoencoders to Bayesian, allows us to generate data.

We assume that it exists a distribution with parameters ϕ^* that is generating the latent states z from x (given x we can sample z): $q_{\phi^*}(z|x)$

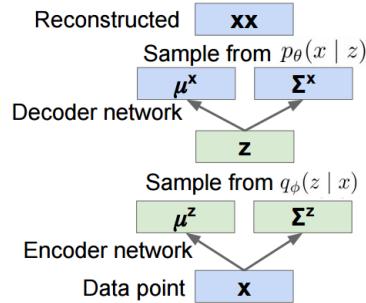


FIGURE 22.4: Variational Autoencoder

We also assume a conditional distribution with parameters θ^* that given the latent states we can sample x : $p_{\theta^*}(x|z^{(i)})$

We assume that these distributions are diagonal Gaussian and we will try to approximate them with two neural nets with parameters ϕ and θ . Also, we assume $p_{\theta^*}(z)$ unit Gaussian. We could assume more fancy distributions, but we use these ones because they are easy to work with.

1. Input data x
2. Pass x through the encoder network which will spit out a distribution of the latent states z (the mean and (diagonal) covariance of $q_\phi(z|x)$).
3. Sample latent states of high probability from the obtained distribution $q_\phi(z|x)$ given our input data x .
4. Once we have some samples of z , pass them through the decoder which will spit out the distribution of the input data x (the mean and (diagonal) covariance of $p_{\theta^*}(x|z^{(i)})$)
5. Finally, we can sample data xx of high probability from the obtained distribution $p_\theta(x|z)$ given the latent variables z . xx should be similar to x .

Generate fake data

We train Variational Autoencoders as a normal autoencoder: reconstruction loss at the end, regularization toward prior in the middle. After the network is trained we can use it to generate fake data.

1. We first sample from the prior distribution $p_\theta(z)$ which we have assumed that it is unit Gaussian so it is really easy to draw random samples from this distribution.
2. Pass a sample through the decoder network which we have trained during training. The decoder will spit out a distribution over our data points x in terms of both a mean and covariance

3. Once we have a mean and a covariance, we have a the $p_\theta(x|z)$ which is also easy to sample from because we have assumed that it is diagonal Gaussian. These samples are going to be new fake data xx

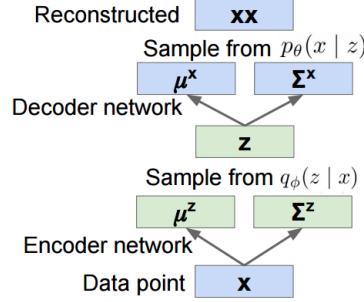


FIGURE 22.5: Generate data with Variational Autoencoder

To try figuring out what the decoder network has learned we can densely sample all the latent space.

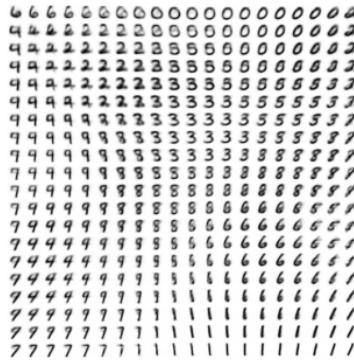


FIGURE 22.6: In this example, the encoder was trained with a latent space of two dimensions. This image was generated by passing each point of this 2D latent space through the decoder. It can be seen that the net has learned to interpolate the digits. For example, we can see 6s transforming to 0s.

One of the reasons that is making this nice separation of classes between axes, is that we are assuming diagonal distribution. Thus, we are assuming as a prior to our system that latent variables (axis in the latent space) are independent (there is no relation between them).

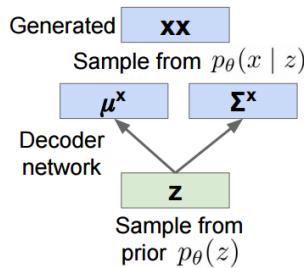


FIGURE 22.7: Example of dense sampling of the latent space

In the CS231n course - lecture 14 there is more information explaining that it is not possible to use maximum likelihood and explaining how to solve this situation.

All together

A recent paper (Dosovitskiy and Brox, Generating Images with Perceptual Similarity Metrics based on Deep Networks, arXiv 2016) proposes a method to mix all together to make the fake images more realistic.

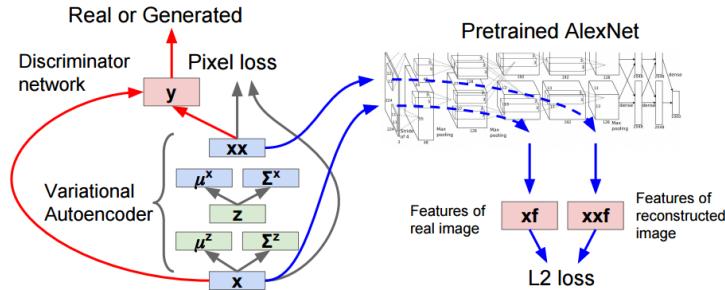


FIGURE 22.8: All together to generate better images

Overview

The idea of autoencoders is that we want to force a network to try to reconstruct our data and hopefully it will learn a useful representation of the data. For traditional autoencoders this is use for feature learning and for variational autoencoders to generate fake samples.

- Traditional Autoencoders
 - Try to reconstruct input
 - Used to learn features, initialize supervised model
 - Not used much anymore
- Variational Autoencoders
 - Bayesian meets deep learning
 - Sample from model to generate images

Chapter 23

Generative Adversarial Nets

A strategy to generate images with less math than Variational Autoencoders.

The idea is to start with a random noise that is normally drawn from a unit Gaussian (or something similar) and we will pass it through a Generator network. The Generator network looks very much like the decoder on the autoencoder and spits fake images. Then, a discriminator network is responsible for deciding/estimating if the input image is fake or a real.

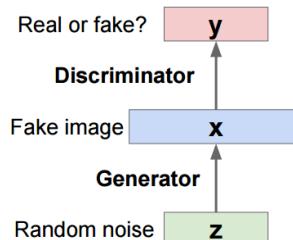


FIGURE 23.1: GAN

How to we train them?

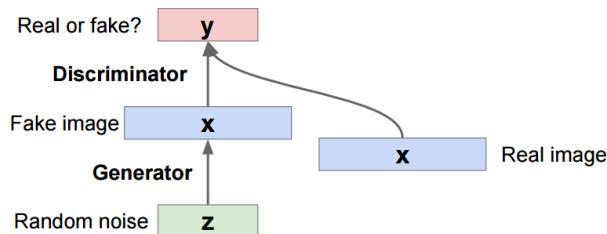


FIGURE 23.2: GAN

We train this network all together. The Generator will receive minibatches of random noise, at it will spit fake images. The Discriminator will receive batches partially fake and real; and it will try to decide which ones are real and which ones are fake.

The generator will improve its fake images by trying to trick the discriminator.

Examples

Interpolating random noise vectors. In figure 23.3 the images in the left and right extremes are obtained by sampling two random points (vectors) of the z space and passing them through the generative net. Then the interpolation between pairs of extreme images is performed by interpolating the z vectors and passing them through the generative net.



FIGURE 23.3: Interpolating random noise vectors

Another experiment shows that the generator is learning a useful representation of the data. In this experiment they randomly generated multiple images of faces and then manually organized in the classes "smiling woman", "neutral woman", "neutral man". Then, they produce the mean z average representation of each image and perform the arithmetic operation of $z_{\text{smiling woman}} - z_{\text{neutral woman}} + z_{\text{neutral man}}$ produces a smiling man.

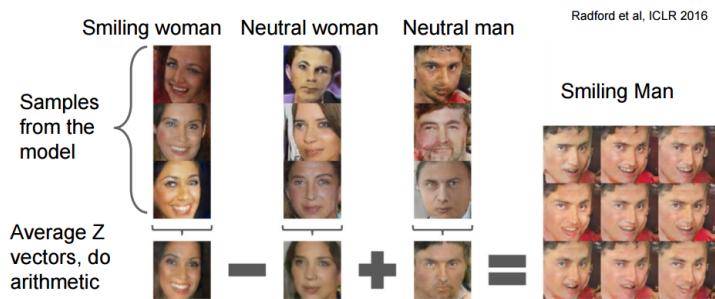


FIGURE 23.4: Generator learning a useful representation of the data

Chapter 24

Region Based CNN (R-CNN)

R-CNN (Region based CNN) are used for object detection. Another network also used for objection detection is YOLO (You only look once)

24.1 R-CNN (do not use this method, deprecated)

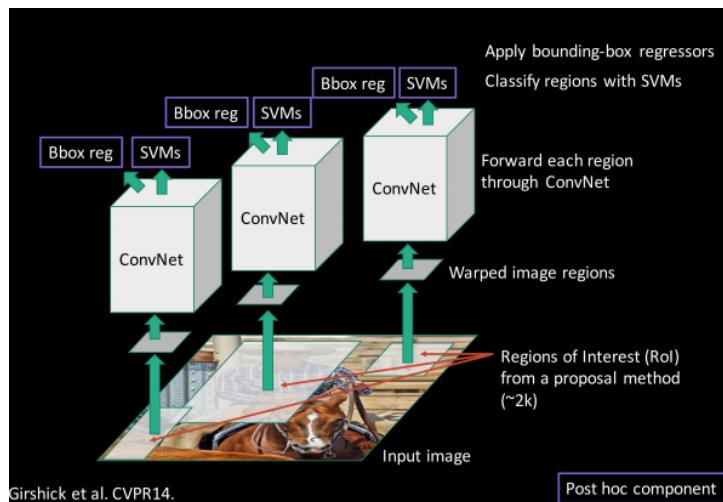


FIGURE 24.1: R-CNN

So the idea for object detection is to use R-CNN. First run a region proposal algorithm to obten regions of interets. Then, warp this regions into a fixed size and run the ConvNet with regression head and classification head. The regression head objective is to output an offset to correct "slightly wrong" region proposals.

R-CNN probelms:

- Slow at test-time: need to run full forward pass of CNN for each region proposal

- SVMs and regressors are post-hoc: CNN features not updated in response to SVMs and regressors
- Complex multistage training pipeline

Train a R-CNN

1.- **Train** (or download) a classification model (e.g. AlexNet). 2.- **Fine-tune model for**

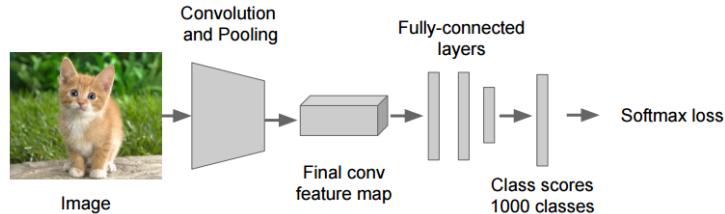


FIGURE 24.2: Train

detection. Instead of X classes you want Y classes + background. To do so, throw away the final fully-connected layer, add the new one and retrain using positive/negative regions from detection images. 3.- **Extract features.** Extract region proposals for all images. For each

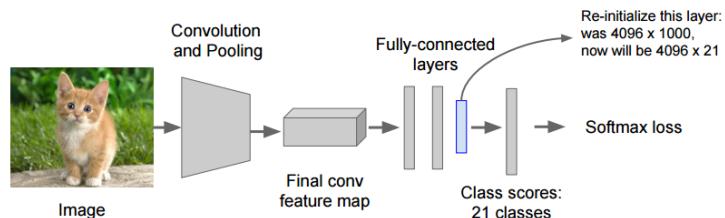


FIGURE 24.3: Fine-tune

region: warp to CNN input size, run forward through CNN, save last pool features to disk. 4.-



FIGURE 24.4: Extract features

Train classification head. Train one binary SVM per class using as input the saved features from last step.

5- **Train regression head.** For each class, train a linear regression model to map from cached features to offset to GT boxes to make up for "slightly wrong" proposals of the region proposals.



FIGURE 24.5: Train regression head

	PASCAL VOC (2010)	ImageNet Detection (ILSVRC 2014)	MS-COCO (2014)
Number of classes	20	200	80
Number of images (train + val)	~20k	~470k	~120k
Mean objects per image	2.4	1.1	7.2

FIGURE 24.6: Usual datasets for training

24.2 Fast R-CNN (do not use this method, deprecated)

A new proposal called Fast R-CNN appeared trying to solve R-CNN problems. The idea is just to swap the order of extracting regions of interest and features. Still, no sliding windows.

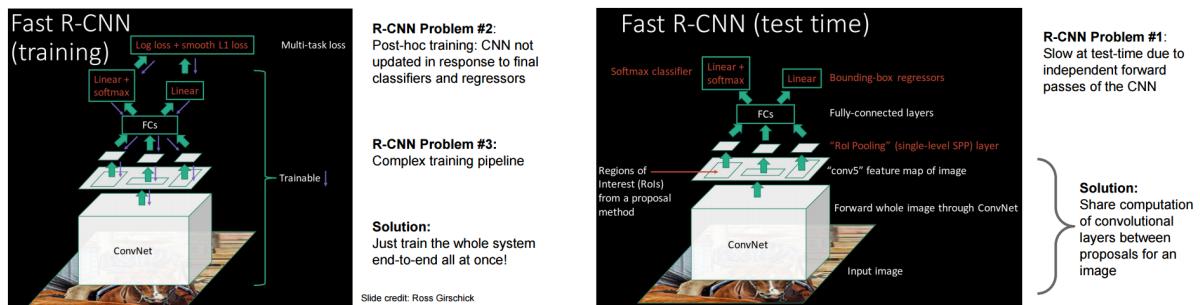


FIGURE 24.7: Fast R-CNN training/testing time

The only mystery here are region of interest pooling.

The HxW image is processed by the convolution and pooling part and regions of interest are proposed (with an external algorithm). The problem now is that the FC layer is expecting a fixed size input. To solve this, given the region proposal, we project it onto the conv feature map, divide it into hwx grid and do max-pool within each grid cell.

With this simple straggly the FC layer will always receive the same input size. Moreover, we can backprop through this strategy with no problem because it only uses max-pool.

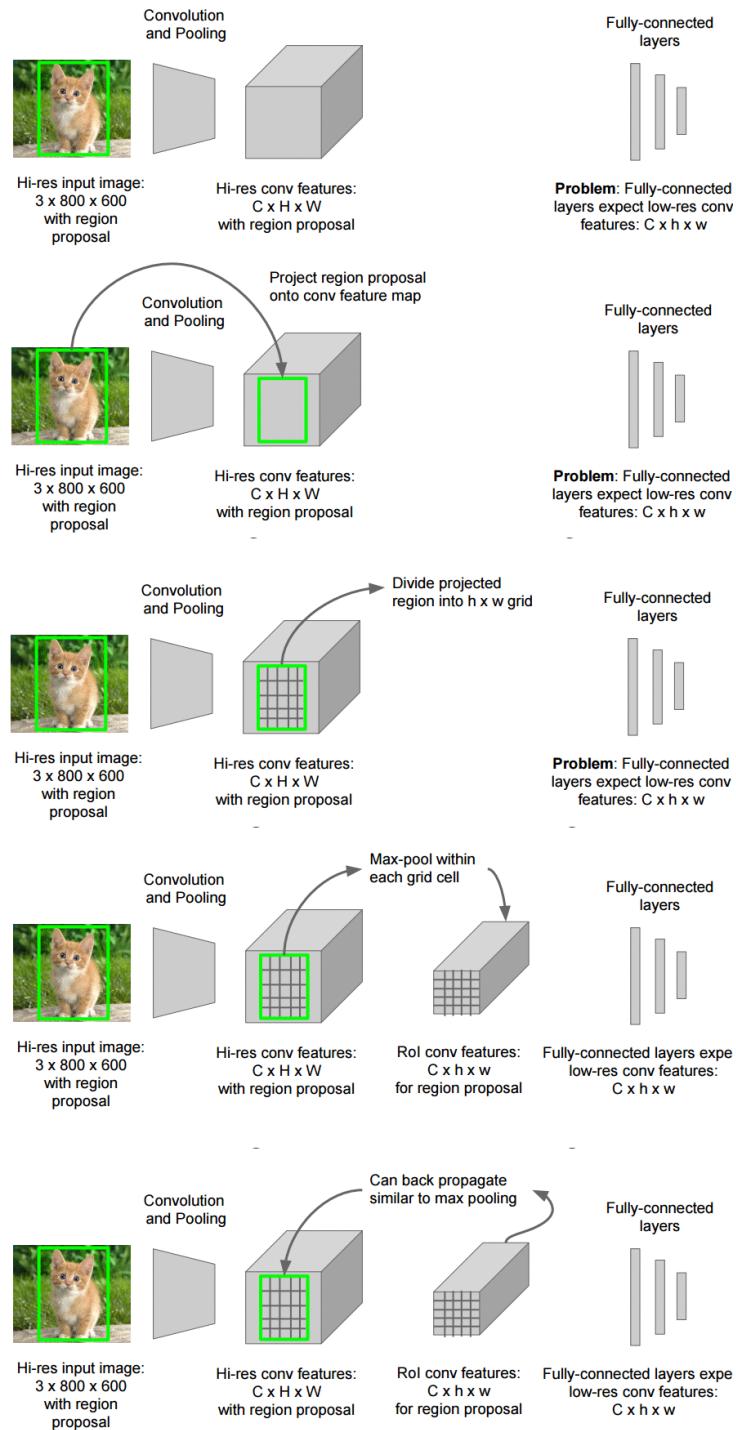


FIGURE 24.8: process

In terms of mAP (accuracy) is more or less the same. The computational time improvement of Fast R-CNN w.r.t. R-CNN is impressive:

- x8.8 training time speed up
- x25 testing time speed up (50s to 2s)

However, the bottleneck is extracting the regions with a region proposals algorithm. Without it, the speed would be:

- x146 testing time speed up (47s to 0.32s)

so why not extracting the ROIs with the CNN? this is what Faster R-CNN proposes

24.3 Faster R-CNN

To solve the problem of region proposals algorithm bottleneck in Fast R-CNN, Faster R-CNN proposes to extract the regions of interest with another network using the information of the last layer of the CNN net.

Insert a Region Proposal Network (RPN) after the last convolutional layer.

RPN trained to produce region proposals directly; no need for external region proposals!

After RPN, use ROI Pooling and an upstream classifier and bbox regressor just like Fast R-CNN

Still, the ROI pooling does not allow to introduce rotations into your regions of interests. To introduce rotations, a DeepMind paper proposes to do bilinear interpolation similar computer graphics.

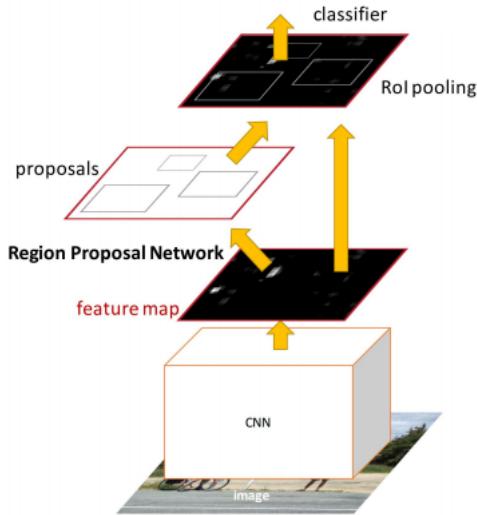


FIGURE 24.9: Faster R-CNN

The cool thing here is the region proposal network, how does it work?

Build a small network for:

- Classifying object or no-object, and

- Regressing bbox locations

Use N anchor boxes at each location. Anchors are translation invariant: use the same ones at every location. For all the feature map points, at the original input image apply all anchors to the point corresponding to the current feature map point. So we take as features the convolutional feature map region corresponding to the anchor region in the original image.

For each of this anchor boxes it produces a score whether there is an object or not; and a finer localization with reference to the anchor (they are an offset to the anchor boxes)

So you are learning different weights for each anchor.

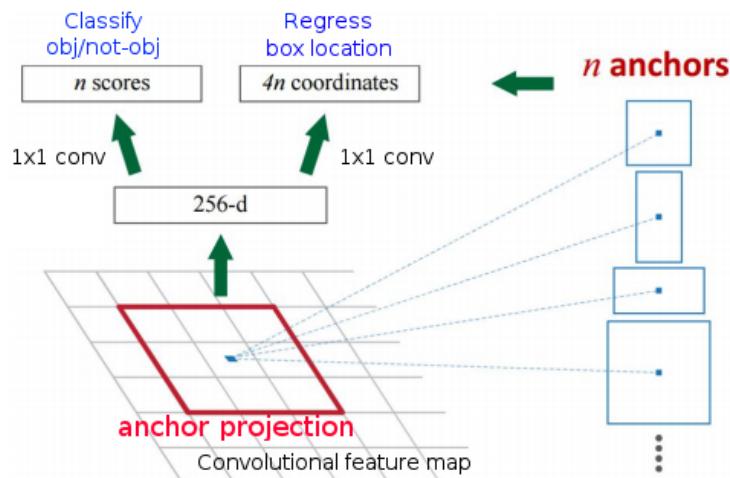


FIGURE 24.10: The cool thing here is the region proposal network, how does it work?

How to train it?

To train it, in some unpublished work, it trains the net as a net with 4 losses

One network, four losses

- RPN classification (anchor good / bad)
- RPN regression (anchor to proposal)
- Fast R-CNN classification (over classes)
- Fast R-CNN regression (proposal to box)

In terms of mAP (accuracy) is more or less the same than with Fast R-CNN. The computational time improvement of Faster R-CNN w.r.t. R-CNN is impressive: x250 testing time speed up (50s to 0.2s)

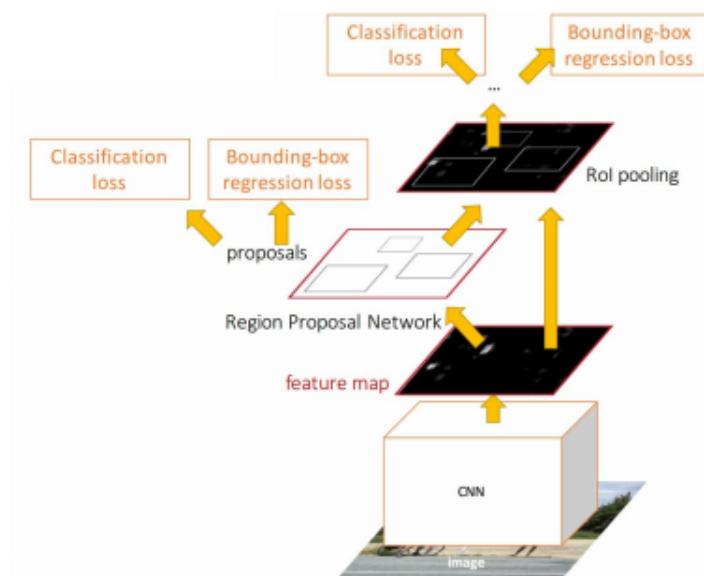


FIGURE 24.11: How to train it?

Chapter 25

YOLO

YOLOs are used for object detection. Other networks also used for objection detection are R-CNN, Fast R-CNN & Faster R-CNN.

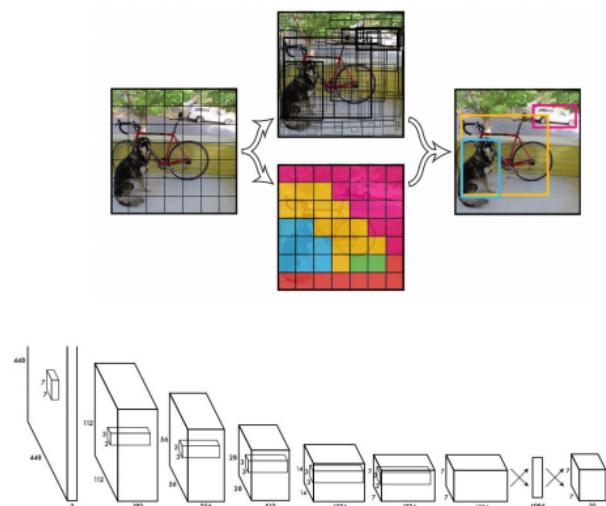


FIGURE 25.1: YOLO

Another detection strategy that is also commonly used is YOLO. The idea is to solve detection as only a regression problem. Direct prediction using a CNN.

Divide image into $S \times S$ grid, they use $S = 7$. Then, within each grid cell predict:

- B boxes: 4 coordinates + confidence, they use $B = 2$
- Class scores: C numbers

Regression from input image to output $S \times S \times (5 * B + C)$ tensor.

It can go at real-time at the expense of lower mAP than Faster R-CNN.

Chapter 26

RNN ConvNet

Local motion

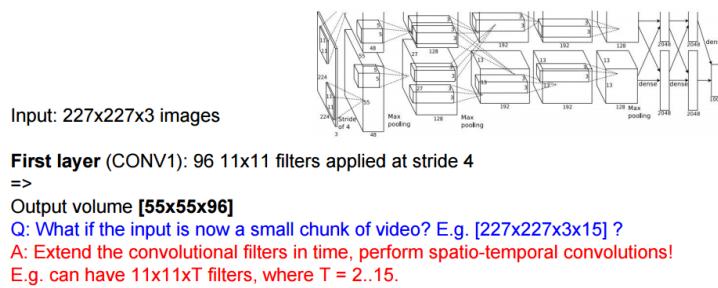


FIGURE 26.1: Getting there

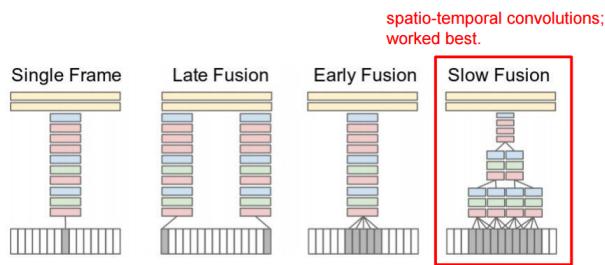


FIGURE 26.2: Large-scale Video Classification with Convolutional Neural Networks, Karpathy et al., 2014]

Global motion

Delving Deeper into Convolutional Networks for Learning Video Representations, Ballas et al., 2016

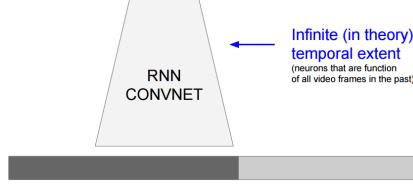


FIGURE 26.3: Large-scale Video Classification with Convolutional Neural Networks, Karpathy et al., 2014]

GRU $\mathbf{z}_t = \sigma(\mathbf{W}_z \mathbf{x}_t + \mathbf{U}_z \mathbf{h}_{t-1}),$ $\mathbf{r}_t = \sigma(\mathbf{W}_r \mathbf{x}_t + \mathbf{U}_r \mathbf{h}_{t-1}),$ $\tilde{\mathbf{h}}_t = \tanh(\mathbf{W} \mathbf{x}_t + \mathbf{U} (\mathbf{r}_t \odot \mathbf{h}_{t-1}))$ $\mathbf{h}_t = (1 - \mathbf{z}_t) \mathbf{h}_{t-1} + \mathbf{z}_t \tilde{\mathbf{h}}_t,$	Matrix multiply \Rightarrow CONV	$\mathbf{z}_t^l = \sigma(\mathbf{W}_z^l * \mathbf{x}_t^l + \mathbf{U}_z^l * \mathbf{h}_{t-1}^l),$ $\mathbf{r}_t^l = \sigma(\mathbf{W}_r^l * \mathbf{x}_t^l + \mathbf{U}_r^l * \mathbf{h}_{t-1}^l),$ $\tilde{\mathbf{h}}_t^l = \tanh(\mathbf{W}^l * \mathbf{x}_t^l + \mathbf{U} * (\mathbf{r}_t^l \odot \mathbf{h}_{t-1}^l)),$ $\mathbf{h}_t^l = (1 - \mathbf{z}_t^l) \mathbf{h}_{t-1}^l + \mathbf{z}_t^l \tilde{\mathbf{h}}_t^l,$
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FIGURE 26.4: The idea is to modify GRU adding convolutions. The Gated Recurrent Unit (GRU) is a simplified version of an LSTM unit with fewer parameters. Just like an LSTM cell, it uses a gating mechanism to allow RNNs to efficiently learn long-range dependency by preventing the vanishing gradient problem. The GRU consists of a reset and update gate that determine which part of the old memory to keep vs. update with new values at the current time step.

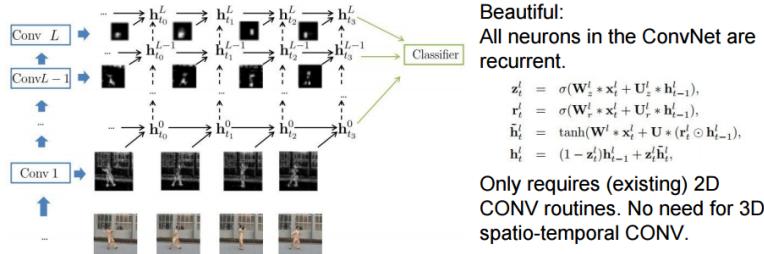


FIGURE 26.5: Original paper diagram

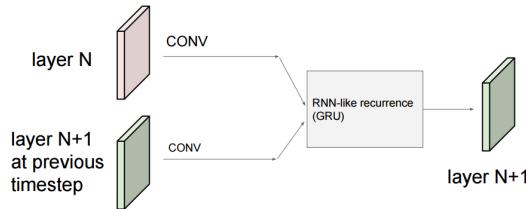


FIGURE 26.6: A more clear diagram

Summary

1. You think you need a Spatio-Temporal Fancy VideoConvNet
2. STOP. Do you really?
3. Okay fine: do you want to model:
 - local motion? (use 3D CONV), or
 - global motion? (use LSTM).

4. Try out using Optical Flow in a second stream (can work better sometimes)
5. Try out GRU-RCN! (imo best model)

Chapter 27

Attention Models

First recall RNN for Captioning. It would be better if it could look at the image more than once and focus its attention to specific parts of the image.

Soft Attention for Captioning

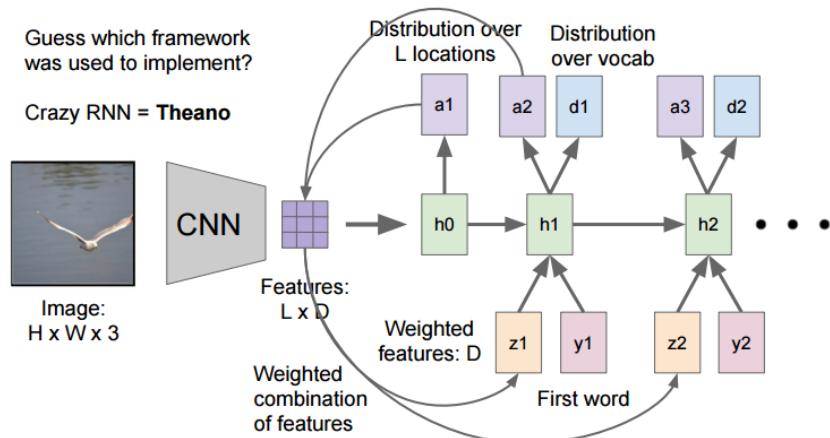


FIGURE 27.1: Soft Attention for Captioning - Xu et al, Show, Attend and Tell: Neural Image Caption Generation with Visual Attention, ICML 2015

1. Compute the features with a CNN. Extract not the last ones but from an early layer. This is why it is a grid of features and not a single vector. In this way we have information of the localization of the input image features
2. Use these features to initialize the h_0 hidden state
3. Now things get different with respect to RNN for captioning. Instead of using h_0 to compute distribution over words, we use it to compute a prob distribution over L locations. This would be implemented with a couple of affine layers (= FC) and softmax to give a distribution.

4. We produce a weighted sum of features using the prob distribution over locations. This can be seen as taking the feature vector and summarizing it to a vector. This gives us a weighted features that is used to decide where to focus.
5. The next hidden state has as inputs the past hidden state and a word (like CNN) but now we also add the weighted features.
6. The hidden state is used to produce a prob distribution over words and also a new distribution over locations. These are implemented with a couple of FC layers on top of the hidden state.
7. Go to 4

Soft vs Hard Attention

Lets see how these summarization vectors z are produced

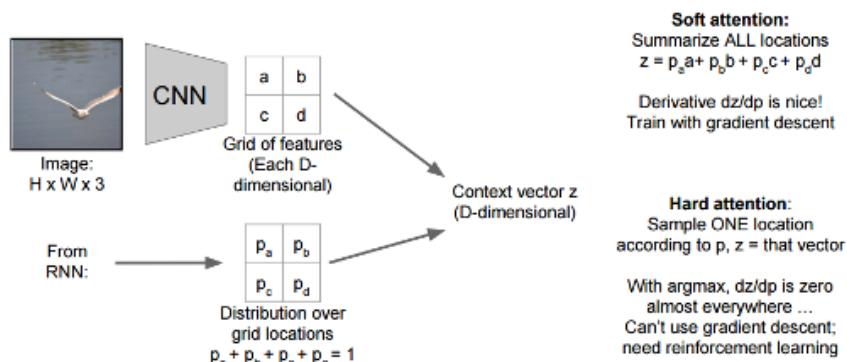


FIGURE 27.2: Soft Attention vs Hard Attention

Both soft and hard attention model produce the same output. Notice that the soft attention is more diffuse because it is averaging prob from the image. And Hard attention it is only focusing in one element.

Hard attention is normally faster at test time because it is only focusing on a specific thing at every step instead of looking at big regions of the image.

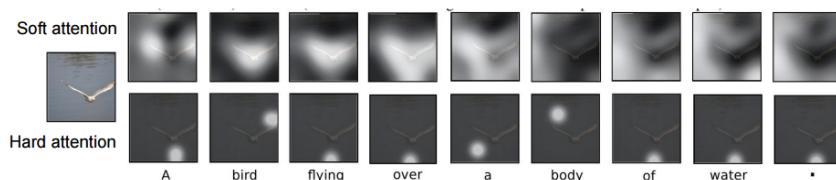


FIGURE 27.3: Example Soft Attention vs Hard Attention

Both have the same problem, they are constrained by a grid over the input image which makes them more blurry. z is not continuous. To work in a continuous way it exists the so called "Spatial Transformed Networks"

Recap

Soft attention:

- Easy to implement: produce distribution over input locations, reweight features and feed as input
- Attend to arbitrary input locations using spatial transformer networks

Hard attention:

- Attend to a single input location
- Can't use gradient descent!
- Need reinforcement learning (because they are not differentiable)

Chapter 28

Spatial Transformer Networks

The use of spatial transformers results in models which learn invariance to translation, scale, rotation and more generic warping, resulting in state-of-the-art performance on several benchmarks, and for a number of classes of transformations.

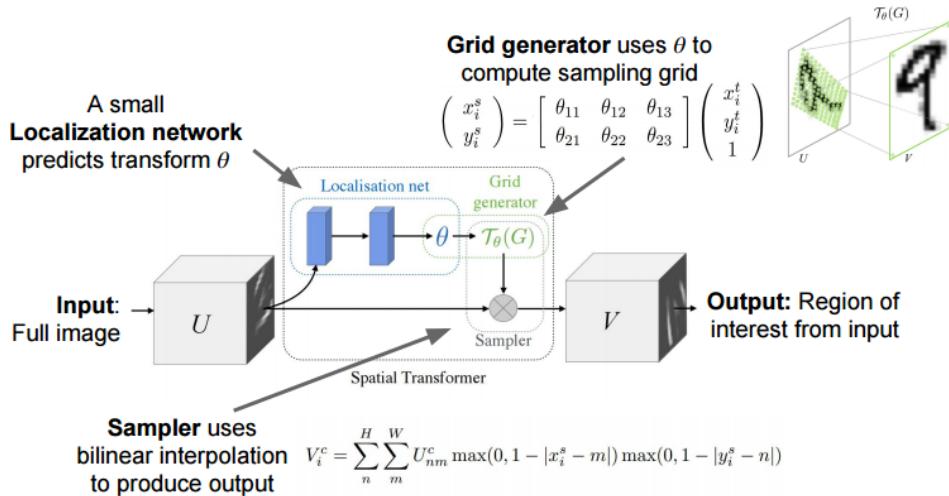


FIGURE 28.1: Spatial Transformer Networks

We would like a function that given a continuous box coordinates cropped the input image. The problem is that it is not possible to crop in a continuous way because we are constraint, at least, by pixels.

The idea is to have a function that maps output pixels coord to input pixels coord. This function is differentiable w.r.t the affine parameters θ .

With this function, the network will attend to input regions by predicting θ .

They put the info explained on top row in a self contained module which they call "Spatial Transformer" which is divided in three parts:

- Localisation net: Outputs the affine parameters θ .

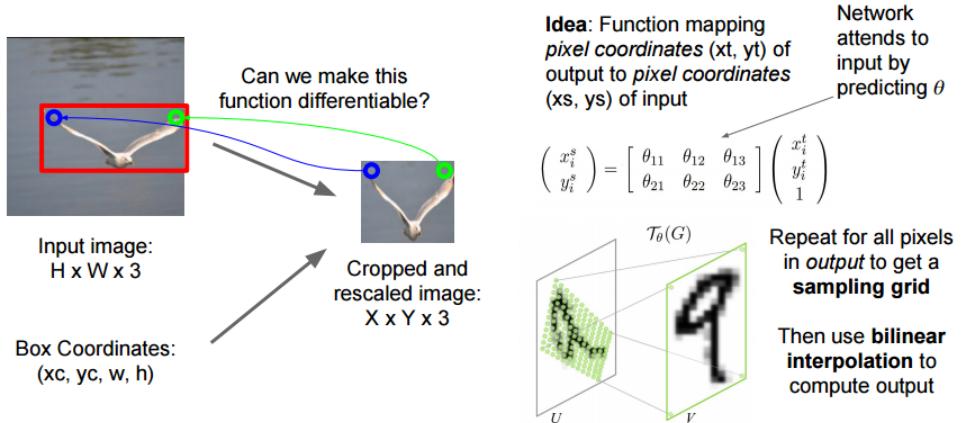


FIGURE 28.2: continuous coordinates

- Grid Generator: Use θ to compute the sampling grid
- Sampler: With bilinear interpolation produces outputs.

Notice that all this modules are continuous and differentiable.

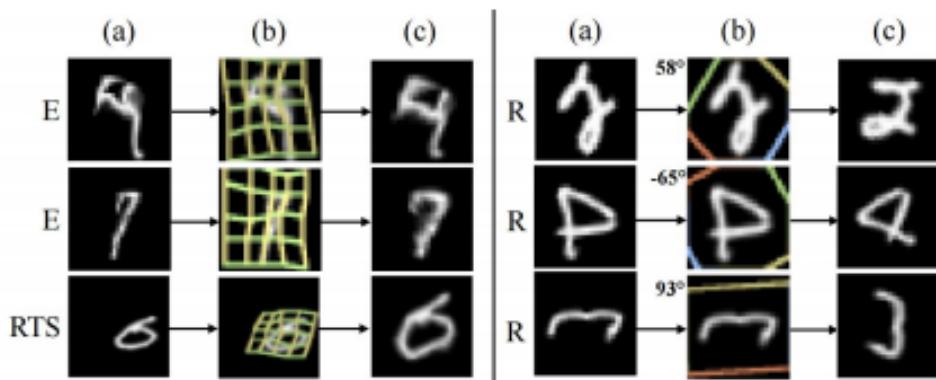


FIGURE 28.3: Deformation examples

Chapter 29

Famous Networks

LeNet-5

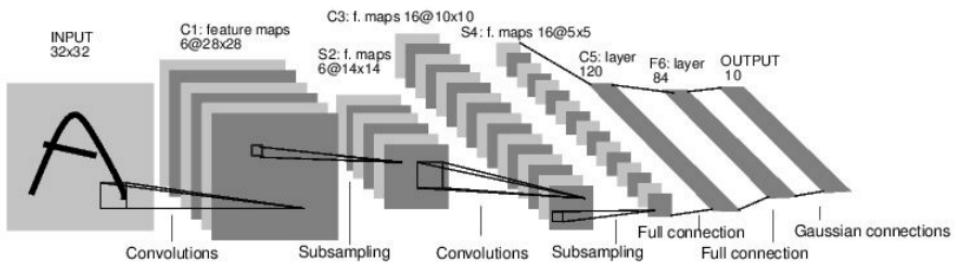


FIGURE 29.1: LeNet

The first successful applications of Convolutional Networks were developed by Yann LeCun in 1990s. Of these, the best known is the LeNet architecture that was used to read zip codes, digits, etc.

AlexNet

The first work that popularized Convolutional Networks in Computer Vision was the AlexNet, developed by Alex Krizhevsky, Ilya Sutskever and Geoff Hinton. The AlexNet was submitted to the ImageNet ILSVRC challenge in 2012 and significantly outperformed the second runner-up (top 5 error of 16% compared to runner-up with 26% error). The Network had a very similar architecture to LeNet, but was deeper, bigger, and featured Convolutional Layers stacked on top of each other (previously it was common to only have a single CONV layer always immediately followed by a POOL layer). 60M parameters.

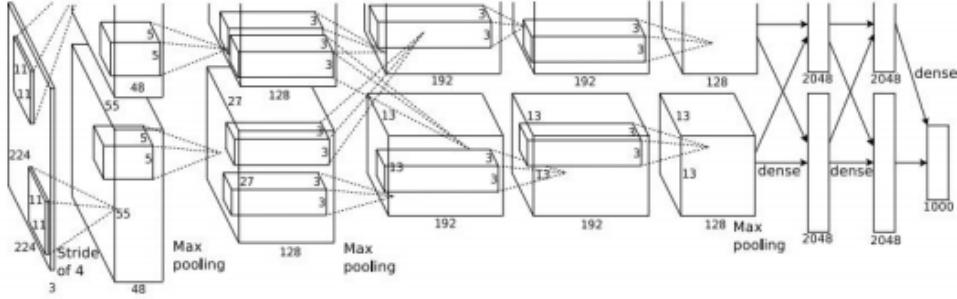


FIGURE 29.2: AlexNet. There is an error in the paper figure, the input image must be of 227 so that the rest of volumes are coherent

VGGNet

The runner-up in ILSVRC 2014 was the network from Karen Simonyan and Andrew Zisserman that became known as the VGGNet. Its main contribution was in showing that the depth of the network is a critical component for good performance. Their final best network contains 16 CONV/FC layers and, appealingly, features an extremely homogeneous architecture that only performs 3×3 convolutions and 2×2 pooling from the beginning to the end. Their pretrained model is available for plug and play use in Caffe. A downside of the VGGNet is that it is more expensive to evaluate and uses a lot more memory and parameters (140M). Most of these parameters are in the first fully connected layer, and it was since found that these FC layers can be removed with no performance downgrade, significantly reducing the number of necessary parameters.

The most of the network parameters are in the FC layer and most of the memory required by the network is used in the first 2 ConvLayers.

ConvNet Configuration			
B	C	D	19
13 weight layers	16 weight layers	16 weight layers	
put [224 x 224 RGB image]			
conv3-64	conv3-64	conv3-64	cc
conv3-64	conv3-64	conv3-64	cc
maxpool			
conv3-128	conv3-128	conv3-128	co
conv3-128	conv3-128	conv3-128	co
maxpool			
conv3-256	conv3-256	conv3-256	co
conv3-256	conv3-256	conv3-256	co
conv1-256	conv3-256	conv3-256	co
maxpool			
conv3-512	conv3-512	conv3-512	co
conv3-512	conv3-512	conv3-512	co
conv1-512	conv3-512	conv3-512	co
maxpool			
conv3-512	conv3-512	conv3-512	co
conv3-512	conv3-512	conv3-512	co
conv1-512	conv3-512	conv3-512	co
maxpool			
FC-4096			
FC-4096			
FC-1000			
soft-max			

INPUT: [224x224x3] memory: 224*224*3=150K params: 0 (not counting biases)
 CONV3-64: [224x224x64] memory: 224*224*64=3.2M params: (3*3*3)*64 = 1,728
 CONV3-64: [224x224x64] memory: 224*224*64=3.2M params: (3*3*64)*64 = 36,864
 POOL2: [112x112x64] memory: 112*112*64=800K params: 0
 CONV3-128: [112x112x128] memory: 112*112*128=1.6M params: (3*3*64)*128 = 73,728
 CONV3-128: [112x112x128] memory: 112*112*128=1.6M params: (3*3*128)*128 = 147,456
 POOL2: [56x56x128] memory: 56*56*128=400K params: 0
 CONV3-256: [56x56x256] memory: 56*56*256=800K params: (3*3*128)*256 = 294,912
 CONV3-256: [56x56x256] memory: 56*56*256=800K params: (3*3*256)*256 = 589,824
 CONV3-256: [56x56x256] memory: 56*56*256=800K params: (3*3*256)*256 = 589,824
 POOL2: [28x28x256] memory: 28*28*256=200K params: 0
 CONV3-512: [28x28x512] memory: 28*28*512=400K params: (3*3*256)*512 = 1,179,648
 CONV3-512: [28x28x512] memory: 28*28*512=400K params: (3*3*512)*512 = 2,359,296
 CONV3-512: [28x28x512] memory: 28*28*512=400K params: (3*3*512)*512 = 2,359,296
 POOL2: [14x14x512] memory: 14*14*512=100K params: 0
 CONV3-512: [14x14x512] memory: 14*14*512=100K params: (3*3*512)*512 = 2,359,296
 CONV3-512: [14x14x512] memory: 14*14*512=100K params: (3*3*512)*512 = 2,359,296
 CONV3-512: [14x14x512] memory: 14*14*512=100K params: (3*3*512)*512 = 2,359,296
 POOL2: [7x7x512] memory: 7*7*512=25K params: 0
 FC: [1x1x4096] memory: 4096 params: 77*512*4096 = 102,760,448
 FC: [1x1x4096] memory: 4096 params: 4096*4096 = 16,777,216
 FC: [1x1x1000] memory: 1000 params: 4096*1000 = 4,096,000
 TOTAL memory: 24M * 4 bytes ~ 93MB / image (only forward! ~*2 for bwd)
 TOTAL params: 138M parameters

FIGURE 29.3: VGGNet - 7.3% top 5 error in ImageNet

GoogLeNet

The ILSVRC 2014 winner was a Convolutional Network from Szegedy et al. from Google. Its main contribution was the development of an Inception Module that dramatically reduced the number of parameters in the network (4M, compared to AlexNet with 60M). Additionally, this paper uses Average Pooling instead of Fully Connected layers at the top of the ConvNet, eliminating a large amount of parameters that do not seem to matter much. There are also several followup versions to the GoogLeNet, most recently Inception-v4.

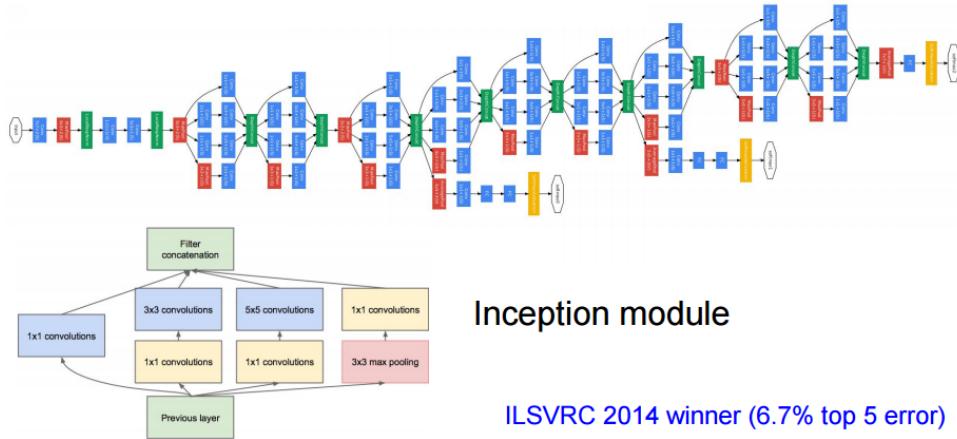


FIGURE 29.4: VGGNet - 6.7% top 5 error in ImageNet

type	patch size/stride	output size	depth	#1x1	#3x3 reduce	#3x3	#5x5 reduce	#5x5	pool proj	params	ops
convolution	7x7/2	112x112x64	1							2.7K	34M
max pool	3x3/2	56x56x64	0								
convolution	3x3/1	56x56x192	2		64	192				112K	360M
max pool	3x3/2	28x28x192	0								
inception (3a)		28x28x256	2	64	96	128	16	32	32	159K	128M
inception (3b)		28x28x480	2	128	128	192	32	96	64	380K	304M
max pool	3x3/2	14x14x480	0								
inception (4a)		14x14x512	2	192	96	208	16	48	64	364K	73M
inception (4b)		14x14x512	2	160	112	224	24	64	64	437K	88M
inception (4c)		14x14x512	2	128	128	256	24	64	64	463K	100M
inception (4d)		14x14x528	2	112	144	288	32	64	64	580K	119M
inception (4e)		14x14x832	2	256	160	320	32	128	128	840K	170M
max pool	3x3/2	7x7x832	0								
inception (5a)		7x7x832	2	256	160	320	32	128	128	1072K	54M
inception (5b)		7x7x1024	2	384	192	384	48	128	128	1388K	71M
avg pool	7x7/1	1x1x1024	0								
dropout (40%)		1x1x1024	0								
linear		1x1x1000	1							1000K	1M
softmax		1x1x1000	0								

Fun features:

- Only 5 million params! (Removes FC layers completely)

Compared to AlexNet:

- 12X less params
- 2x more compute
- 6.67% (vs. 16.4%)

FIGURE 29.5: VGGNet structure

ResNet

Let $H(x)$ be a function that you desire to obtain. In a typical net you would compute a sequence of steps $\text{ReLU}(\text{ReLU}(xw_1+b_1)*w_2+b_2)$ to transform x to $H(x)$. Instead, in a ResNet you compute a delta to be added to the original input to obtain $H(x)$.

What is nice about it is that in *plain nets*, gradients must flow through all the transformations. Instead, in *residual nets* because it is addition (distributes the gradient equally to all its

children), the gradient will flow through the (weights, ReLU) but will also skip this transformations and will go directly to the previous part and flow directly to the previous block. So the gradients can skip all the transformations and go directly to the first layer. In this way, you can train very fast the first layer which is doing simple statistics, and the rest of layers will learn to add to the single in between to make it work at the end.

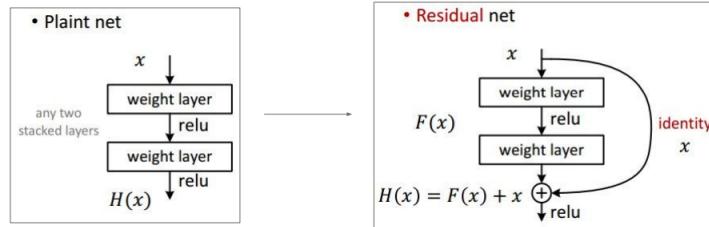


FIGURE 29.6: Plain vs Residual Net

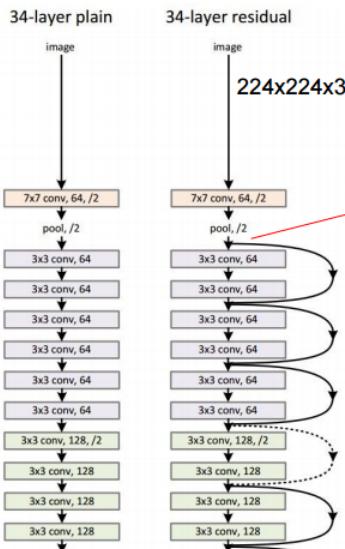


FIGURE 29.7: ResNet (much more layers than the ones on the diagram)

Another way of seeing it, is that ResNets are only computing a delta on top of the identity. So it makes it nice to optimize.

Residual Network developed by Kaiming He et al. was the winner of ILSVRC 2015. It features special skip connections and a heavy use of batch normalization. The architecture is also missing fully connected layers at the end of the network. ResNets are currently by far state of the art Convolutional Neural Network models and are the default choice for using ConvNets in practice (as of May 10, 2016). In particular, also see more recent developments that tweak the original architecture from Kaiming He et al. Identity Mappings in Deep Residual Networks (published March 2016). It is interesting that after the first layer they do polling (the only polling in all the net) and they scale the input image of 244×244 to 56×56 , and the net works that well. Its crazy that all the layer (except the first one) work with $56 \times 56 \times ?$ and even compressing the data this much it has a high accuracy.

layer name	output size	18-layer	34-layer	50-layer	101-layer	152-layer
conv1	112×112			7×7, 64, stride 2		
conv2_x	56×56	$\left[\begin{array}{l} 3 \times 3, 64 \\ 3 \times 3, 64 \end{array} \right] \times 2$	$\left[\begin{array}{l} 3 \times 3, 64 \\ 3 \times 3, 64 \end{array} \right] \times 3$	$\left[\begin{array}{l} 1 \times 1, 64 \\ 3 \times 3, 64 \\ 1 \times 1, 256 \end{array} \right] \times 3$	$\left[\begin{array}{l} 1 \times 1, 64 \\ 3 \times 3, 64 \\ 1 \times 1, 256 \end{array} \right] \times 3$	$\left[\begin{array}{l} 1 \times 1, 64 \\ 3 \times 3, 64 \\ 1 \times 1, 256 \end{array} \right] \times 3$
conv3_x	28×28	$\left[\begin{array}{l} 3 \times 3, 128 \\ 3 \times 3, 128 \end{array} \right] \times 2$	$\left[\begin{array}{l} 3 \times 3, 128 \\ 3 \times 3, 128 \end{array} \right] \times 4$	$\left[\begin{array}{l} 1 \times 1, 128 \\ 3 \times 3, 128 \\ 1 \times 1, 512 \end{array} \right] \times 4$	$\left[\begin{array}{l} 1 \times 1, 128 \\ 3 \times 3, 128 \\ 1 \times 1, 512 \end{array} \right] \times 4$	$\left[\begin{array}{l} 1 \times 1, 128 \\ 3 \times 3, 128 \\ 1 \times 1, 512 \end{array} \right] \times 8$
conv4_x	14×14	$\left[\begin{array}{l} 3 \times 3, 256 \\ 3 \times 3, 256 \end{array} \right] \times 2$	$\left[\begin{array}{l} 3 \times 3, 256 \\ 3 \times 3, 256 \end{array} \right] \times 6$	$\left[\begin{array}{l} 1 \times 1, 256 \\ 3 \times 3, 256 \\ 1 \times 1, 1024 \end{array} \right] \times 6$	$\left[\begin{array}{l} 1 \times 1, 256 \\ 3 \times 3, 256 \\ 1 \times 1, 1024 \end{array} \right] \times 23$	$\left[\begin{array}{l} 1 \times 1, 256 \\ 3 \times 3, 256 \\ 1 \times 1, 1024 \end{array} \right] \times 36$
conv5_x	7×7	$\left[\begin{array}{l} 3 \times 3, 512 \\ 3 \times 3, 512 \end{array} \right] \times 2$	$\left[\begin{array}{l} 3 \times 3, 512 \\ 3 \times 3, 512 \end{array} \right] \times 3$	$\left[\begin{array}{l} 1 \times 1, 512 \\ 3 \times 3, 512 \\ 1 \times 1, 2048 \end{array} \right] \times 3$	$\left[\begin{array}{l} 1 \times 1, 512 \\ 3 \times 3, 512 \\ 1 \times 1, 2048 \end{array} \right] \times 3$	$\left[\begin{array}{l} 1 \times 1, 512 \\ 3 \times 3, 512 \\ 1 \times 1, 2048 \end{array} \right] \times 3$
	1×1			average pool, 1000-d fc, softmax		
FLOPs		1.8×10^9	3.6×10^9	3.8×10^9	7.6×10^9	11.3×10^9

FIGURE 29.8: ResNet structure. 3.6% top 5 error in ImageNet, 152 layers, 2-3 weeks training on 8 GPU machine, faster at test time than VGGNet

Should we add infinite layers?

In plot 29.9 it is clear that networks are getting deeper and deeper But we have to be careful.

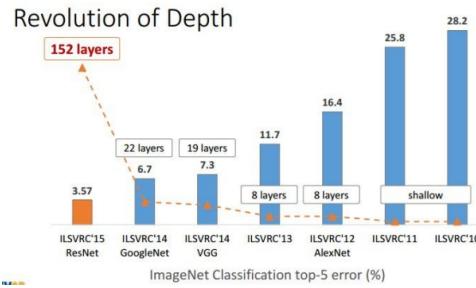


FIGURE 29.9: Depth revolution

Plot 29.10 shows CIFAR-10 training error. In the left plain nets (weighted layer + ReLU) in the right ResNet. How it is possible to get a higher training error (dashed lines) with higher number of layers? It should not happen, the model is more complex. The explanation is that we are still not capable of optimizing them good enough.

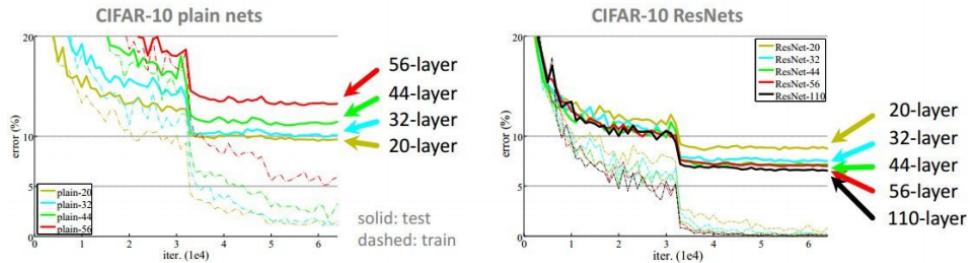


FIGURE 29.10: CIFAR-10 training error

However, ResNets always improve the test and training error.

So the answer to the question is that we should keep adding more layers but not in a naive way, we should do it in a ResNet way

Part V

Applications

Chapter 30

Applications

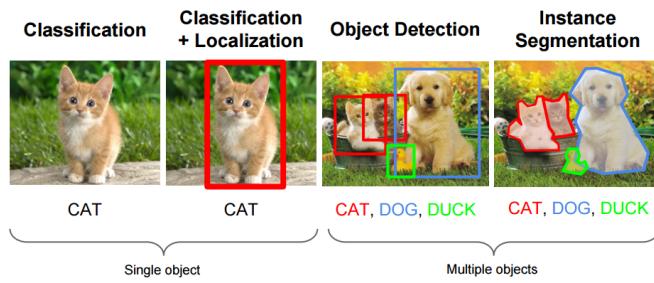


FIGURE 30.1: Some applications examples

30.1 Classification

Train a classification model with softmax loss. The input is the entire image and the output are C probabilities (one per class) of being in the image.

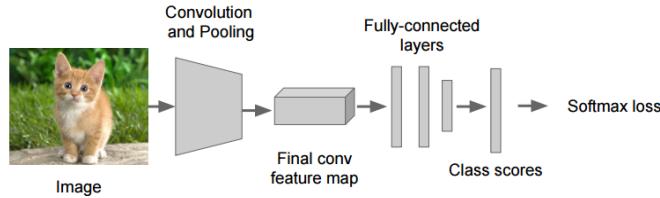


FIGURE 30.2: Classification

30.2 Classification + Localization

Goal is to find a fixed number of objects (one or many) in an image

3 easy steps recipe:

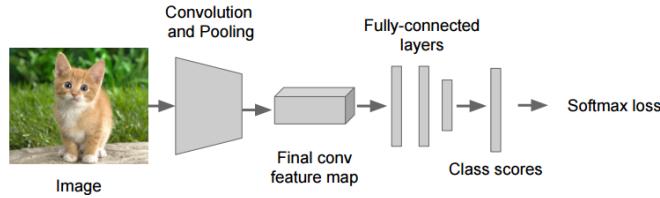


FIGURE 30.3: 1.- Train (or download) a classification model (AlexNet, VGG, GoogLeNet)

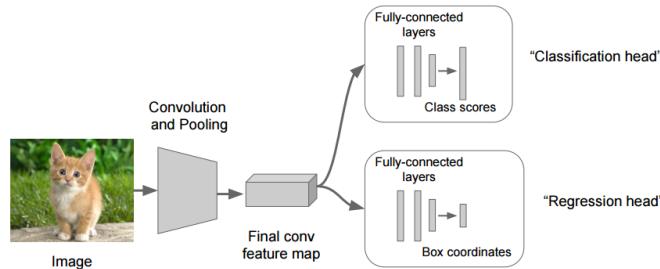


FIGURE 30.4: 2.- Attach a new fully-connected "regression head" to the network to compute bounding boxes (x, y, w, h)

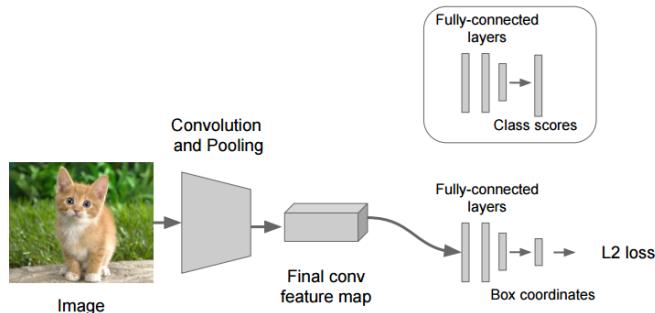


FIGURE 30.5: 3.- Train the regression head only with SGD and L2 loss with the bounding boxes as ground-truth.

When training the regression head there are two options: You can backpropagate only till the regression head or the entire network. 2nd option will improve a little bit the accuracy at a expense of higher training computation cost. If you choose option 2 you will be changing the original Conv layers on which the classification head is trained. So there are two options. Or you have two independent networks: the original one (Conv+classification head) and the other one (Modified Conv+regression head). Or you train both at the same time so you will have only one model(classification head + regression head).

The final DNN for classification + uses both heads at test time

- Classification head:
- The output are C numbers (one per class)
- Regression head:
- There are two regression heads strategies (choose one):

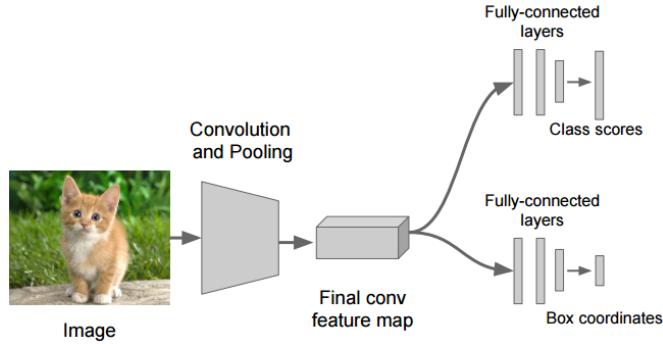


FIGURE 30.6: 4.- At test time use both heads

1. Class agnostic: Output 4 numbers (one bounding box)
2. Class specific: Output $C \times 4$ numbers (one bounding box per class)

Notice that instead of training a regression head which outputs bounding boxes, we could have train it to output a specific number of joints or whatever

Sliding windows (DO NOT DO THIS)

At test time this (classification + regression) network should be applied inside a sliding windows at multiple scales. (This would have an **enormous computation cost**, we will see later that there is a better way of doing this).

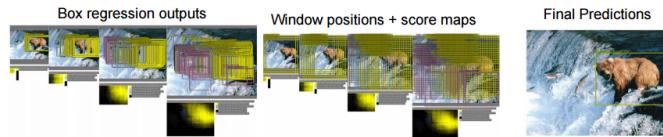


FIGURE 30.7: Classification + Localization - Box regression and Sliding windows diagram

The idea is to run the classification + regression network inside a sliding windows and at multiple scales. So for each windows position we obtain a cat bb (x, y, w, h) and the probability of a cat being inside the current sliding windows position. (This would have an enormous computation cost, we will see later that there is a better way of doing this)

Finally, we have obtained multiple bound boxes and we must merge them and estimate the prob of this final bb of containing a cat. To do so, an algorithm such us non-max suppression could be used.

Bounding Box regression (ONLY INPUT IMAGE FIXED SIZE)

Fully connected layers can only deal with input of a fixed size, because it requires a certain amount of parameters to "fully connect" the input and output.

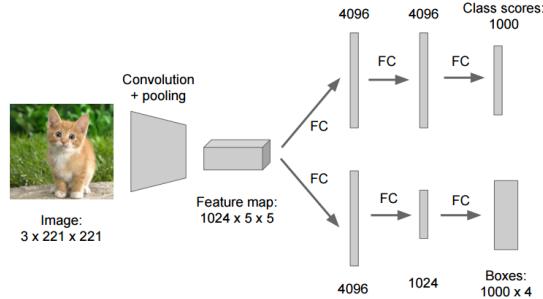


FIGURE 30.8: Classification + Localization - Box regression architecture

In the example network with fully-connected layers at the end, a 224×224 image will output a 1000 size vector of class scores. If we apply the network on a larger image, the network will fail because of the inconsistency between the input and parameters of the first fully-connected layer.

In other words, we would have to execute the DNN in each sliding windows position. This has an enormous computation cost. To solve this we can transform the FC layers to Conv layers. (see right column)

Without sliding windows (DO THIS)

Any FC layer can be converted to a CONV layer. For example, in this model, the FC layer with $K=4096$ that is looking at the input volume of size $5 \times 5 \times 1024$ can be equivalently expressed as a CONV layer with $F = 5, P = 0, S = 1, K = 4096$.

In other words, we are setting the filter size to be exactly the size of the input volume, and hence the output will simply be $1 \times 1 \times 4096$ since only a single depth column “fits” across the input volume, giving identical result as the initial FC layer.

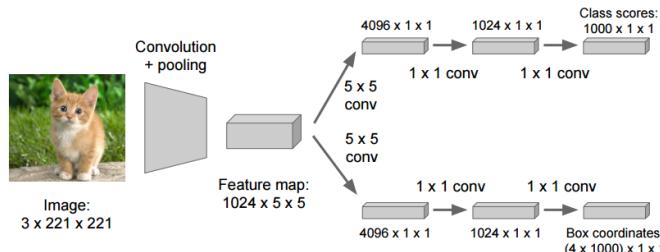


FIGURE 30.9: Classification + Localization - Without sliding windows

FC can only deal with input of a fixed size, while convolutional layers just “slide” the same filters across the input, so it can basically deal with input of an arbitrary spatial size.

At train time we train with samples of the sliding windows. At test time we can evaluate input images of higher dimension sharing computations.

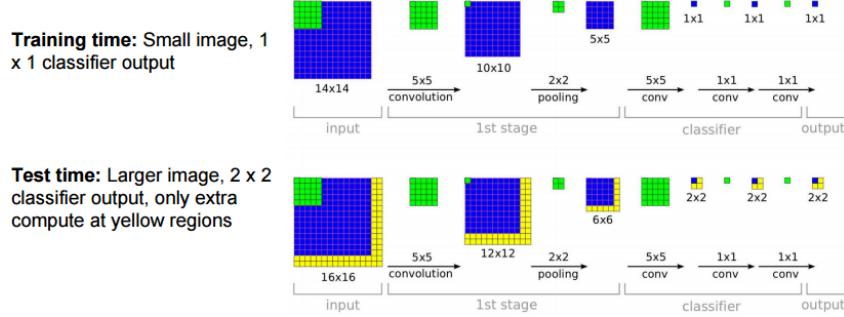


FIGURE 30.10: ConvLayers allow input of any size in an efficient way

For example, in this image, we have trained the model with input image sizes of 14×14 producing and output vector of $1 \times 1 \times C$. At test time, we are using input images of 16×16 which produces and output of $2 \times 2 \times C$. This is equivalent to execute the model 4 times (in the four corners of the image) like in the previous example of the cat. But instead of doing the whole computation 4 times, we are sharing the computation, the extra computation is only done in the yellow parts.

No more sliding windows!!

30.3 Object Detection

Object detection is a harder problem than classification + localization. Now, there can be a non fix number of instances of the same class as well as other classes. We can not use regression with detection (as within classification) because we need variable sized outputs. So we have to go back to sliding windows? It would be very costly to run a classification model in each windows position. The answer is no. We will first search for tiny subsets of possible positions. These models are called R-CNN (Region based CNN).

Evaluation. To evaluate a detection model we use "mean average precision" (mAP). Compute average precision (AP) separately for each class, then average over classes. A detection is a true positive if it has a IoU with a ground-truth box greater than some threshold (usually 0.5) (mAP@0.5). Combine all detection from all test images to draw a precision / recall curve for each class; AP is the area under the curve. mAP is a number from 0 to 100. The higher the better.

There are two main networks used for object detection explained in previous chapters:

- R-CNN (Region Based CNN) - R-CNN, Fast R-CNN & Faster R-CNN
- YOLO (You only look once)

30.4 Segmentation

30.4.1 Semantic Segmentation

Semantic Segmentation tries to label every pixel. It does not differentiate instances (e.g. cows are a massive compact blue mass)

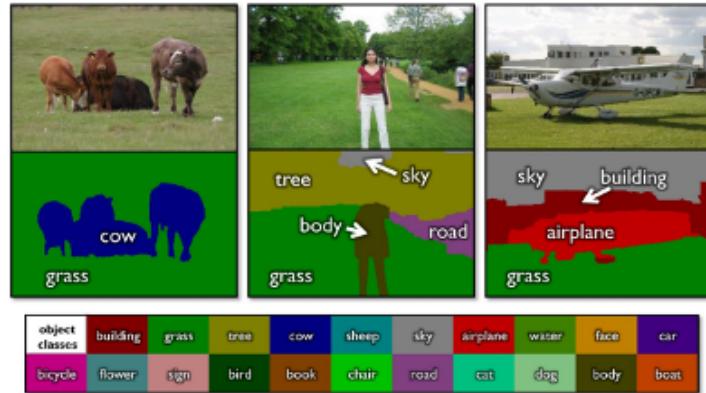


FIGURE 30.11: Semantic Segmentation

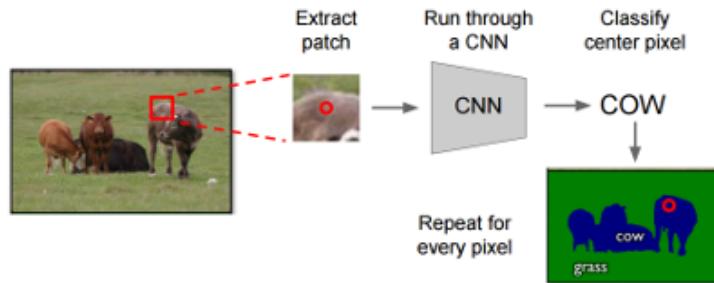


FIGURE 30.12: General idea. Problem: Smaller output due to pooling. Other more complex solutions are required

There are three common strategies used for semantic segmentation:

- Multiscale, or
- Refinement, or
- Upsampling

Multiscale

Solve problem by working in different scales and up-scaling output. In this example they are also doing a parallel process to compute super-pixels to improve the segmentation.

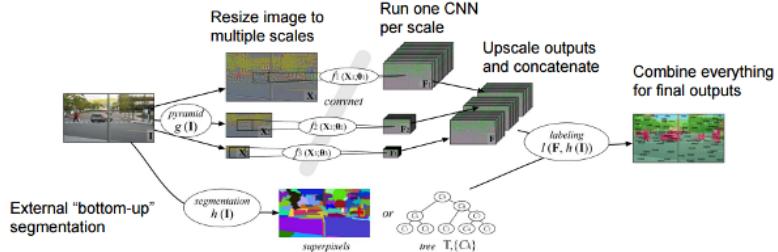


FIGURE 30.13: Farabet et al, “Learning Hierarchical Features for Scene Labeling”, TPAMI 2013

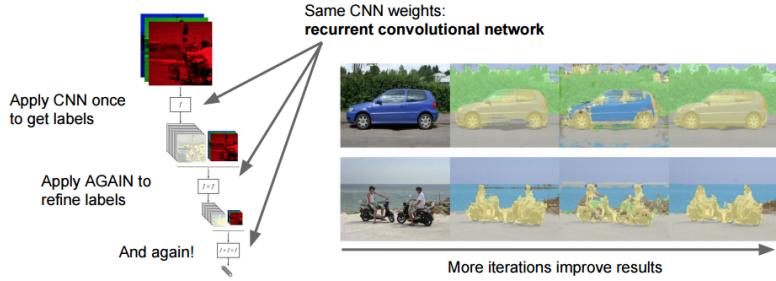


FIGURE 30.14: Pinheiro and Collobert, “Recurrent Convolutional Neural Networks for Scene Labeling”, ICML 2014

Refinement

1. Pass the input image through a CNN which produces the low res image segmentation (labels).
2. Rescale the inputs to the lower resolution and apply again the ConvNet to the lower resolution input image and results from the previous segmentation
3. Goto 2 until threshold

Upsampling

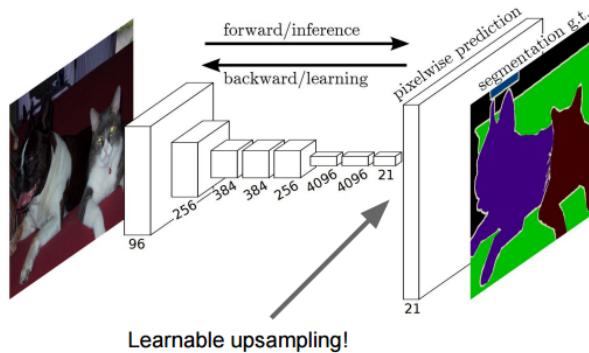


FIGURE 30.15: Long, Shelhamer, and Darrell, “Fully Convolutional Networks for Semantic Segmentation”, CVPR 2015

They pass the input image through a CNN which produces the low res image segmentation (labels). But with this method, they add an extra learnable layer at the end of the ConvNet

that learns to upsample the low resolution segmentation (Upsampling layer). They also do skip connections

30.4.2 Instance Segmentation

Detect instances, give category, label pixels. simultaneous detection and segmentation (SDS). Normally look very similar to the detection models.

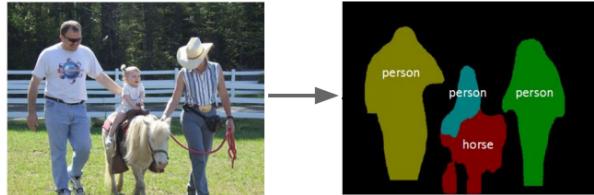


FIGURE 30.16: Instance Segmentation

Architecture similar to R-CNN

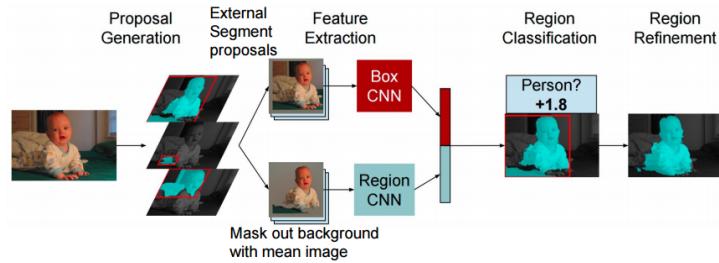


FIGURE 30.17: Architecture - Hariharan et al, “Simultaneous Detection and Segmentation”, ECCV 2014

1. External segment proposals that outputs pixels not boxes
2. Produce a BBox of the segmented region
3. Take the BBox image and run it through a CNN
4. Take the BBox image and set the non segment proposal pixels to the mean image value of the dataset and run it through a CNN
5. Concatenate both features and run it through a classifier
6. Refine the proposed region

Notice that this is very similar to R-CNN

For the refinement step there is a following paper that improves it (Hariharan et al, Hypercolumns for Object Segmentation and Fine-grained Localization, CVPR 2015). The idea is that

once we have the region cropped pass it through AlexNet and extract features of various layers. Then, upsample this feature maps and combine them together. Finally, do a logistic classifier for each of the pixels that predicts how much likely is to be background.

Architecture similar to Faster R-CNN

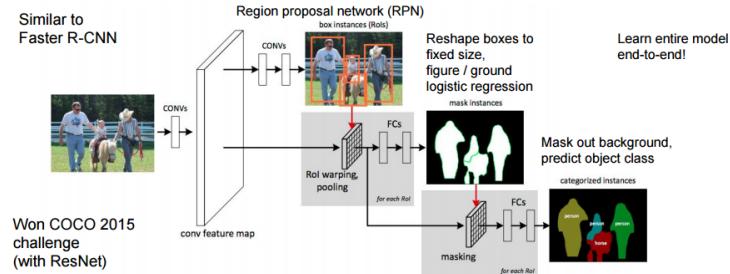


FIGURE 30.18: Architecture - Dai et al, “Instance-aware Semantic Segmentation via Multi-task Network Cascades”, arXiv 2015

Stuck the model in the left to ResNet. Similar to Faster R-CNN

From the high resolution feature map propose regions, then reshape boxes and finally mask background and predict object class.

Notice that the three steps (intermediate levels of the net) can be evaluated with ground-truth.

30.5 Image captioning

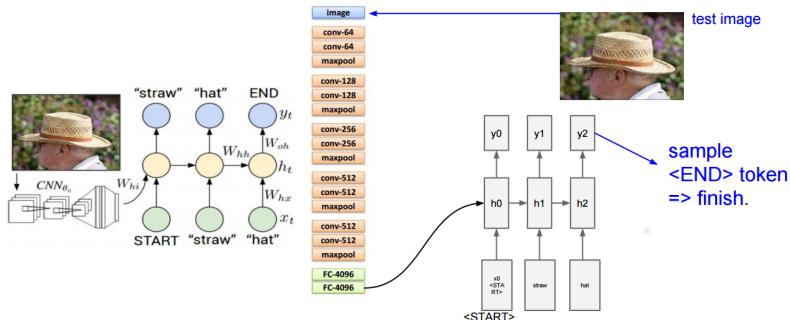


FIGURE 30.19: “Deep Visual-Semantic Alignments for Generating Image Descriptions”, Karpathy and Fei-Fei

Summary

- Localization
 - Find a fixed number of objects (one or many)

- L2 regression from CNN features to box coordinates
 - Overfeat: Regression + efficient sliding window with FC -> conv conversion
 - Deeper networks do better
- Object Detection
 - Find a variable number of objects by classifying image regions
 - Before CNNs: dense multiscale sliding window (HoG, DPM)
 - Avoid dense sliding window with region proposals
 - R-CNN: Selective Search + CNN classification / regression
 - Fast R-CNN: Swap order of convolutions and region extraction
 - Faster R-CNN: Compute region proposals within the network
 - Deeper networks do better
 - Semantic segmentation
 - Classify all pixels
 - Fully convolutional models, downsample then upsample
 - Learnable upsampling: fractionally strided convolution
 - Skip connections can help
 - Instance Segmentation
 - Detect instance, generate mask
 - Similar pipelines to object detection

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