NPFL114, Lecture 03

Training Neural Networks II





Milan Straka



Let us have a dataset with a training, validation and test sets, each containing examples (\boldsymbol{x},y) . Depending on y, consider one of the following output activation functions:

$$egin{cases} ext{none} & ext{if } y \in \mathbb{R} \ \sigma & ext{if } y ext{ is a probability of an outcome} \ ext{softmax} & ext{if } y ext{ is a gold class} \end{cases}$$

If $\mathbf{x} \in \mathbb{R}^d$, we can use a neural network with an input layer of size d, hidden layer of size h with a non-linear activation function, and an output layer of size o (either 1 or number of classification classes) with the mentioned output function.



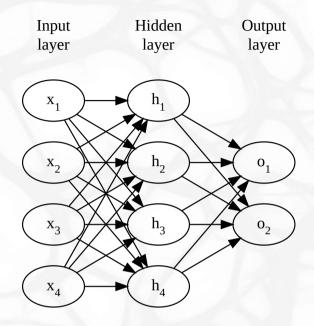
Input layer	Hidden layer	Output layer
x_1	h_1	
X_2	h ₂	01
x_3	h ₃	02
x_4	h ₄	

We have

$$h_i = f^{(1)} \left(\sum_j m{W}_{i,j}^{(1)} x_j + b_i^{(1)}
ight)$$

where $\pmb{W}^{(1)} \in \mathbb{R}^{h \times d}$ is a matrix of weights, $\pmb{b}^{(1)} \in \mathbb{R}^h$ is a vector of biases, and $f^{(1)}$ is an activation function.





Similarly

$$o_i = f^{(2)} \left(\sum_j m{W}_{i,j}^{(2)} h_j + b_i^{(2)}
ight)$$

with $\pmb{W}^{(2)} \in \mathbb{R}^{o \times h}$ another matrix of weights, $\pmb{b}^{(2)} \in \mathbb{R}^o$ another vector of biases, and $f^{(2)}$ being an output activation function.

ÚFAL

The parameters of the model are therefore $\pmb{W}^{(1)}, \pmb{W}^{(2)}, \pmb{b}^{(1)}, \pmb{b}^{(2)}$ of total size $d \times h + h \times o + h + o$.

To train the network, we repeatedly sample m training examples and perform SGD (or any its adaptive variant), updating the parameters to minimize the loss.

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We set the hyperparameters (size of the hidden layer, hidden layer activation function, learning rate, ...) using performance on the validation set and evaluate generalization error on the test set.



• Processing all input in batches.



- Processing all input in batches.
- Vector representation of the network.



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$$oldsymbol{h} = f^{(1)} \left(oldsymbol{W}^{(1)} oldsymbol{x} + oldsymbol{b}^{(1)}
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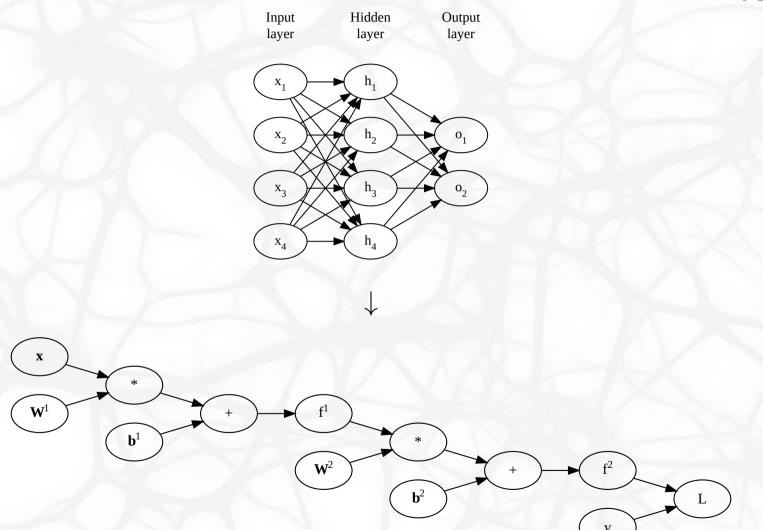
The derivatives

$$rac{\partial f^{(1)}\left(oldsymbol{W}^{(1)}oldsymbol{x}+oldsymbol{b}^{(1)}
ight)}{\partial oldsymbol{W}^{(1)}}$$

are then matrices (called Jacobians).

Computation Graph





Neural Networks Web Browser Demos



- TensorFlow Playground
- Sketch RNN Demo
- <u>DeepLearn.js Demos</u>
 - o <u>DeepLearn.js Image Style Transfer Demo</u>

High Level Overview

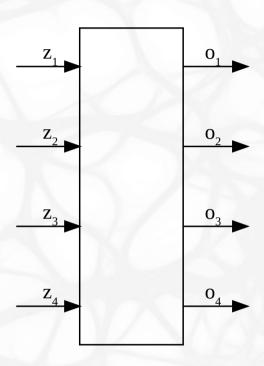


	Classical ('90s)	Deep Learning
Architecture		::::::::::::::::::::::::::::::::::::::
Activation func.	$ anh, \sigma$	tanh, ReLU, PReLU, ELU, SELU, Swish,
Output function	none, σ	none, σ , softmax
Loss function	MSE	NLL (or cross-entropy or KL-divergence)
Optimalization	SGD, momentum	SGD, RMSProp, Adam,
Regularization	L2, L1	L2, Dropout, BatchNorm, LayerNorm,

MLE Loss of Softmax



Softmax



Let us have a softmax output layer with

$$o_i = rac{e^{z_i}}{\sum_j e^{z_j}}.$$

MLE Loss of Softmax



Consider now the MLE estimation. The loss $L(\operatorname{softmax}(\pmb{z}), gold)$ for gold class index gold is then

$$L(\operatorname{softmax}(\boldsymbol{z}), gold) = -\log o_{gold}.$$

The derivation of the loss with respect to z is then

$$egin{aligned} rac{\partial L}{\partial z_i} = &rac{\partial}{\partial z_i} \left[-\log rac{e^{z_{gold}}}{\sum_j e^{z_j}}
ight] \ = &-rac{\partial z_{gold}}{\partial z_i} + rac{\partial \log(\sum_j e^{z_j})}{\partial z_i} \ = &[gold = i] + rac{1}{\sum_j e^{z_j}} e^{z_i} \ = &[gold = i] + o_i. \end{aligned}$$

Therefore, $rac{\partial L}{\partial m{z}} = - m{1}_{gold} + m{o}$, where $m{1}_{gold}$ is 1 at index gold and 0 otherwise.

MLE Loss of Softmax and Sigmoid



In the previous case, the gold distribution was *sparse*, with only one probability being 1.

In the case of general gold distribution g, we have

$$L(\operatorname{softmax}(oldsymbol{z}), oldsymbol{g}) = -\sum_i g_i \log o_i.$$

Adapting the previous procedure, we obtain

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Sigmoid

Analogously, for $o=\sigma(z)$ we get $\frac{\partial L}{\partial z}=-g+o$, where g is the target gold probability.

Regularization



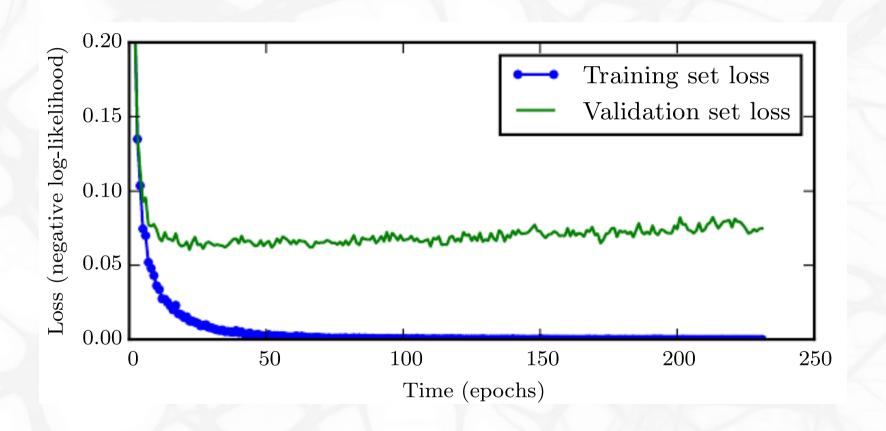
As already mentioned, regularization is any change in the machine learning algorithm that is designed to reduce generalization error but not necessarily its training error.

Regularization is usually needed only if training error and generalization error are different. That is often not the case if we process each training example only once. Generally the more training data, the better generalization performance.

- Early stopping
- L2, L1 regularization
- Dataset augmentation
- Ensembling
- Dropout

Regularization - Early Stopping





L2 Regularization



We prefer models with parameters small under L2 metric.

The L2 regularization, also called *weight decay*, *Tikhonov regularization* or *ridge regression* therefore minimizes

$$ilde{J}(oldsymbol{ heta}; \mathbb{X}) = J(oldsymbol{ heta}; \mathbb{X}) + \lambda ||oldsymbol{ heta}||_2^2$$

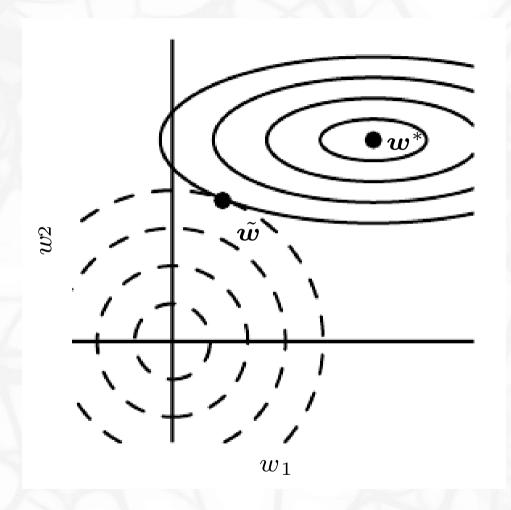
for a suitable (usually very small) λ .

During the parameter update of SGD, we get

$$heta_i \leftarrow heta_i - lpha rac{\partial J}{\partial heta_i} - 2lpha \lambda heta_i$$

L2 Regularization





L2 Regularization as MAP



Another way to arrive at L2 regularization is to utilize Bayesian inference.

With MLE we have

$$oldsymbol{ heta}_{ ext{MLE}} = rg\max_{oldsymbol{ heta}} p(\mathbb{X}; oldsymbol{ heta}).$$

Instead, we may want to maximize *maximum a posteriori (MAP)* point estimate:

$$m{ heta}_{ ext{MAP}} = rg\max_{m{ heta}} p(m{ heta}; \mathbb{X})$$

Using Bayes' theorem $[p(m{ heta};\mathbb{X})=p(\mathbb{X};m{ heta})p(m{ heta})/p(\mathbb{X})]$, we get

$$m{ heta}_{ ext{MAP}} = rg \max_{m{ heta}} p(\mathbb{X}; m{ heta}) p(m{ heta}).$$

L2 Regularization as MAP



The $p(\theta)$ are prior probabilities of the parameter values (our *preference*).

One possibility for such a prior is $\mathcal{N}(\boldsymbol{\theta}; 0, \sigma^2)$.

Then

$$egin{aligned} oldsymbol{ heta}_{ ext{MAP}} &= rg \max_{oldsymbol{ heta}} p(\mathbb{X}; oldsymbol{ heta}) p(oldsymbol{ heta}) \ &= rg \max_{oldsymbol{ heta}} \sum_{i=1}^m p(oldsymbol{x}^{(i)}; oldsymbol{ heta}) p(oldsymbol{ heta}) \ &= rg \min_{oldsymbol{ heta}} \sum_{i=1}^m -\log p(oldsymbol{x}^{(i)}; oldsymbol{ heta}) -\log p(oldsymbol{ heta}) \end{aligned}$$

By substituting the probability of the Gaussian prior, we get

$$m{ heta}_{ ext{MAP}} = rg\min_{m{ heta}} \sum_{i=1}^m -\log p(m{x}^{(i)};m{ heta}) + rac{1}{2}\log(2\pi\sigma^2) + rac{m{ heta}^2}{2\sigma^2}$$

L1 Regularization



Similar to L2 regularization, but we prefer low L1 metric of parameters. We therefore minimize

$$ilde{J}(oldsymbol{ heta}; \mathbb{X}) = J(oldsymbol{ heta}; \mathbb{X}) + \lambda ||oldsymbol{ heta}||_1$$

The corresponding SGD update is then

$$heta_i \leftarrow heta_i - lpha rac{\partial J}{\partial heta_i} - lpha \lambda.$$

Regularization – Dataset Augmentation



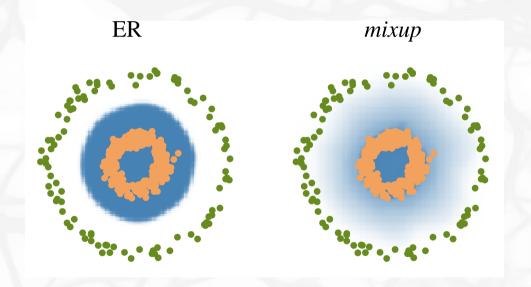
For some data, it is cheap to generate slightly modified examples.

- Image processing: translations, horizontal flips, scaling, rotations, color adjustments, ...
- Speech recognition
- More difficult for discrete domains like text.

Regularization - Dataset Augmentation



- Noise injection to input examples
- Label smoothing
- Mixup (25 Oct 2017)



(b) Effect of mixup on a toy problem.

Regularization - Ensembling

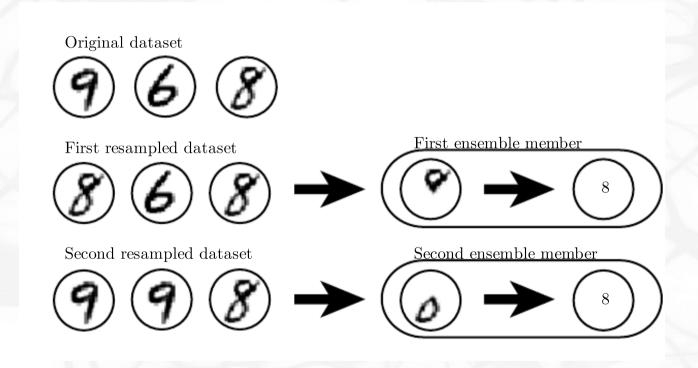


Ensembling (also called *model averaging* or *bagging*) is a general technique for reducing generalization error by combining several models. The models are usually combined by averaging their outputs (either distributions or output values in case of regression).

Regularization - Ensembling



Generate different datasets by sampling with replacement.



- Use random different initialization.
- Average models from last hours/days of training.

Regularization - Dropout



How to design good universal features?

 In reproduction, evolution is achieved using gene swapping. The genes must not be just good with combination with other genes, they need to be universally good.

Regularization - Dropout



How to design good universal features?

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Idea of dropout by (Srivastava et al., 2014), in preprint since 2012.

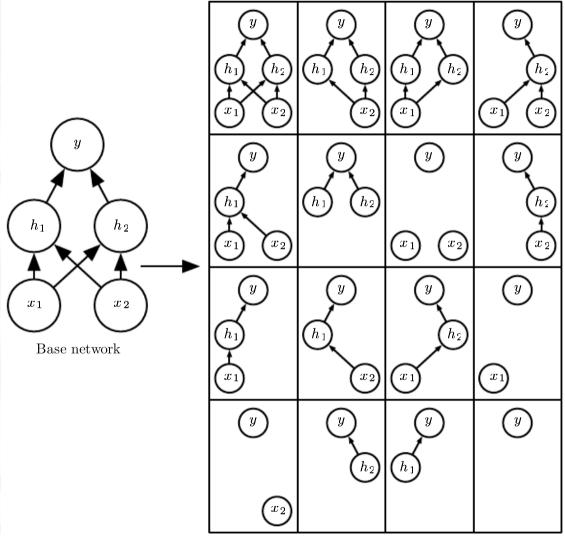
When applying dropout to a layer, we drop each neuron independently with a probability of p (usually called *dropout rate*). To the rest of the network, the dropped neurons have value of zero.

Dropout is performed only when training, during inference no nodes are dropped. However, in that case we need to scale the activations down by a factor of 1-p to account for more neurons than usual.

Alternatively, we might scale the activations up during training by a factor of 1/(1-p).

Regularization - Dropout as Ensembling





Ensemble of subnetworks

Convergence

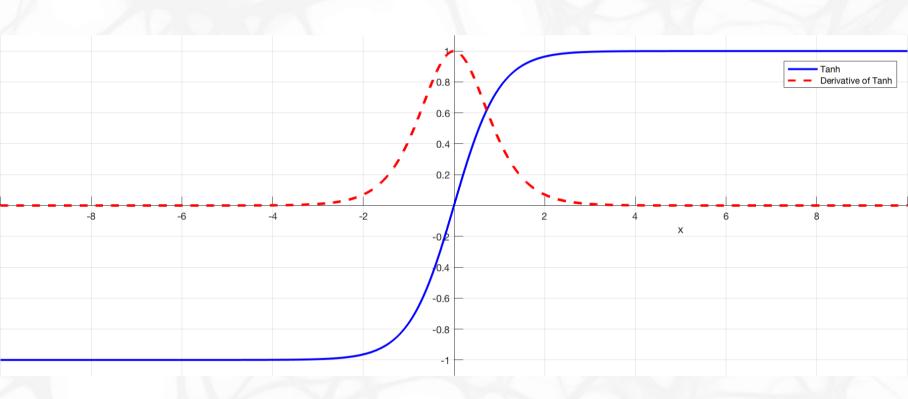


The training process might or might not converge. Even if it does, it might converge slowly or quickly.

There are *many* factors influencing convergence and its speed, we now discuss three of them.

Convergence – Saturating Non-linearities





Convergence – Parameter Initialization



Neural networks usually need random initialization to break symmetry.

• Biases are usually initialized to a constant value, usually 0.

Convergence – Parameter Initialization



Neural networks usually need random initialization to break symmetry.

- Biases are usually initialized to a constant value, usually 0.
- Weights are usually initialized to small random values, either with uniform or normal distribution.
 - The scale matters for deep networks!
 - $\circ~$ Originally, people used $U\left[-\frac{1}{\sqrt{n}},\frac{1}{\sqrt{n}}\right]$ distribution.

Convergence – Parameter Initialization



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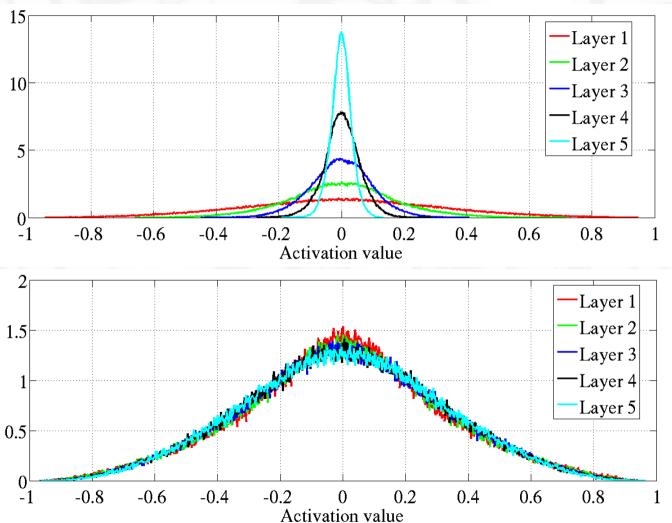
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 - Xavier Glorot and Yoshua Bengio, 2010: Understanding the difficulty of training deep feedforward neural networks.

Theoretically and experimentally showed that a suitable way to initialize a $\mathbb{R}^{n \times m}$ matrix is

$$U\left[-\sqrt{rac{6}{m+n}},\sqrt{rac{6}{m+n}}
ight].$$

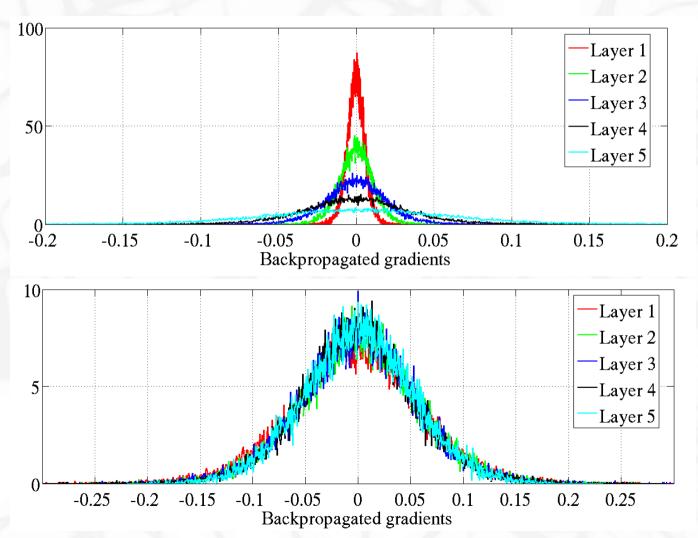
Convergence - Parameter Initialization





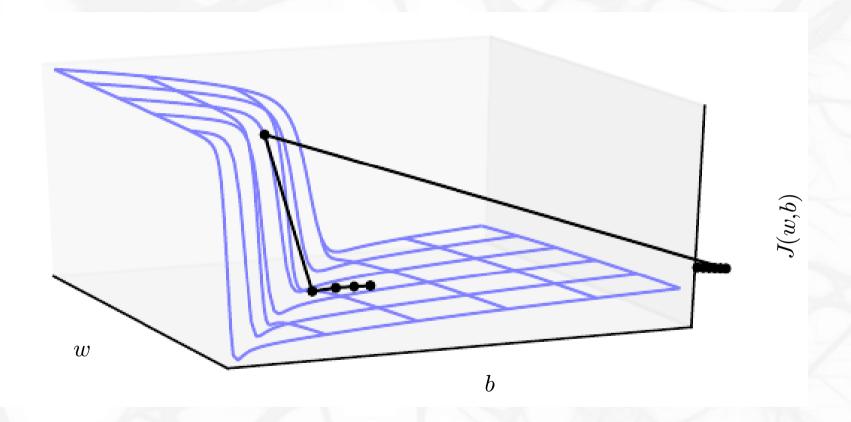
Convergence - Parameter Initialization





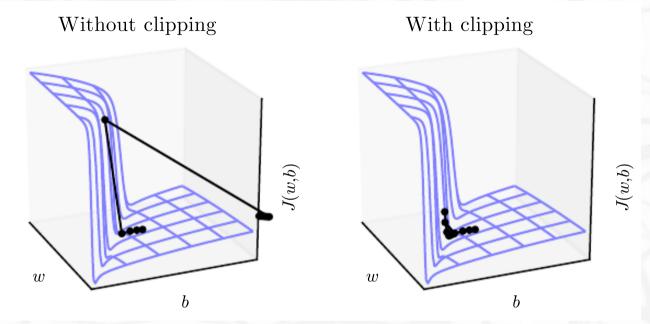
Convergence – Gradient Clipping





Convergence - Gradient Clipping





Using a given maximum norm, we may clip the gradient.

$$g \leftarrow egin{cases} g & ext{if } ||g|| \leq c \ crac{g}{||g||} & ext{if } ||g|| > c \end{cases}$$

The clipping can be per weight, per matrix or for the gradient as a whole.

Going Deeper



Going Deeper

Convolutional Networks



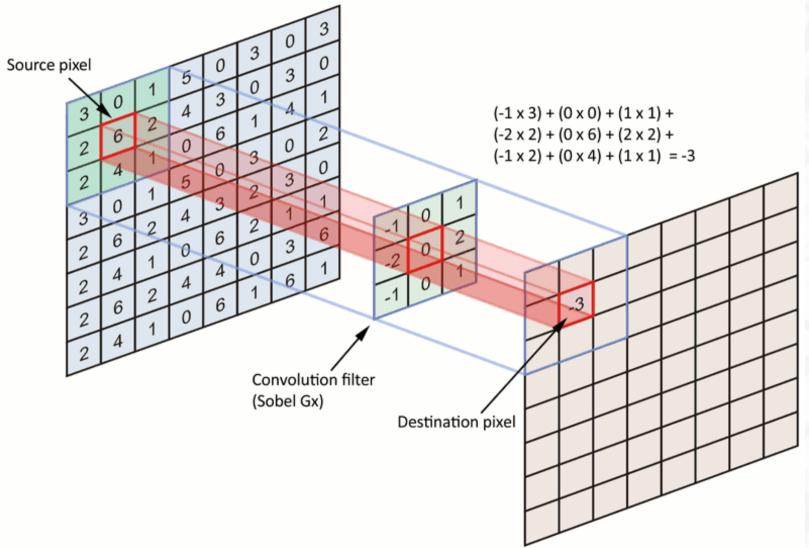
Consider data with some structure (temporal data, speech, images, ...).

Unlike densely connected layers, we might want:

- Sparse (local) interactions
- Parameter sharing (equal response everywhere)
- Shift invariance

Convolutional Networks





Convolution Operation



For a functions x and w, convolution x * w is defined as

$$(x*w)(t) = \int x(a)w(t-a)\,\mathrm{d}a.$$

For vectors, we have

$$(oldsymbol{x}*oldsymbol{w})_t = \sum_i x_i w_{t-i}.$$

Convolution operation can be generalized to two dimensions by

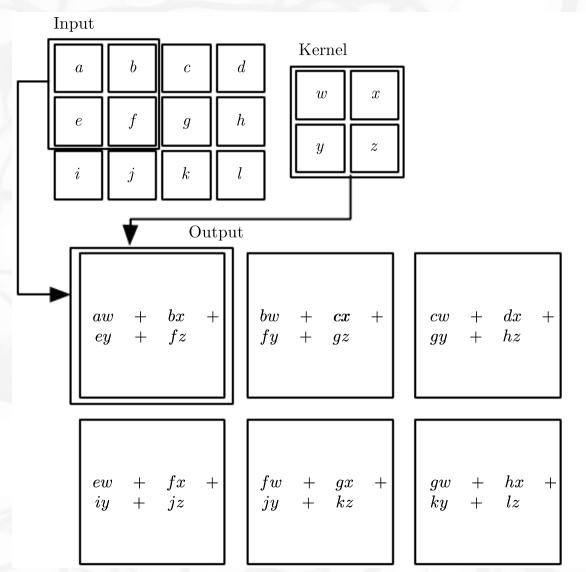
$$(oldsymbol{I} * oldsymbol{K})_{i,j} = \sum_{m,n} oldsymbol{I}_{m,n} oldsymbol{K}_{i-m,j-n}.$$

Closely related is *cross-corellation*, where K is flipped:

$$S_{i,j} = \sum_{m,n} oldsymbol{I}_{i+m,j+n} oldsymbol{K}_{m,n}.$$

Convolution





Convolution Operation



The K is usually called a kernel or a filter, and we generally apply several of them at the same time.

Consider an input image with C channels. The convolution operation with F filters of width W, height H and stride S produces an output with F channels kernels of total size $W \times H \times C \times F$ and is computed as

$$(oldsymbol{I}*oldsymbol{K})_{i,j,k} = \sum_{m,n,o} oldsymbol{I}_{i\cdot s+m,j\cdot s+n,o} oldsymbol{K}_{m,n,o,k}.$$

Convolution Operation



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There are multiple padding schemes, most common are:

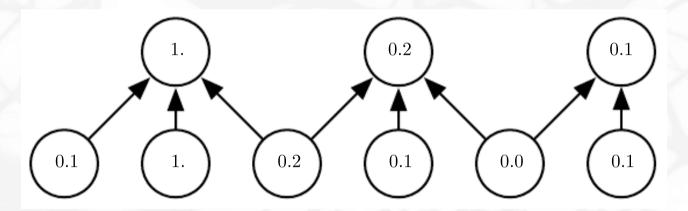
- valid: we only use valid pixels, which causes the result to me smaller
- same: we pad original image with zero pixels so that the result is exactly the size of the input

Pooling



Pooling is an operation similar to convolution, but we perform a fixed operation instead of multiplying by a kernel.

- Max pooling: minor translation invariance
- Average pooling

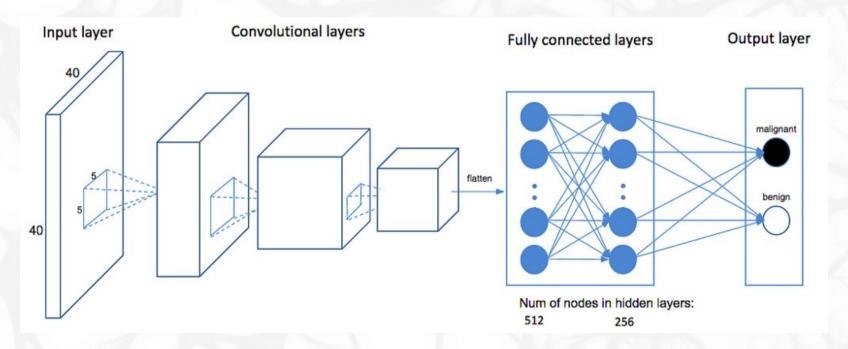


High-level CNN Architecture



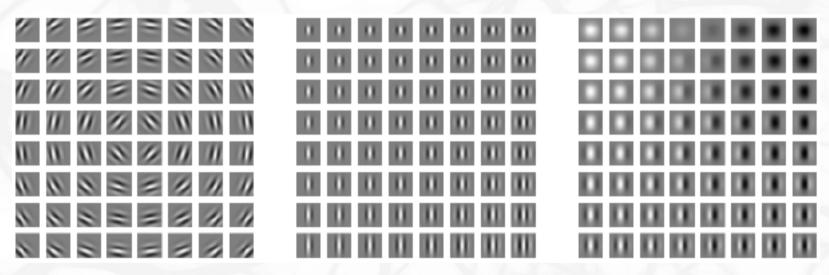
We repeatedly use the following block:

- 1. Convolution operation
- 2. Non-linear activation (usually ReLU)
- 3. Pooling



Similarities in V1 and CNNs

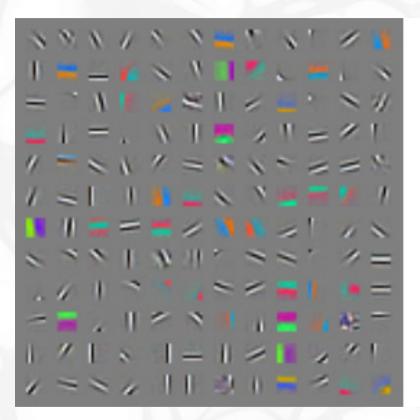


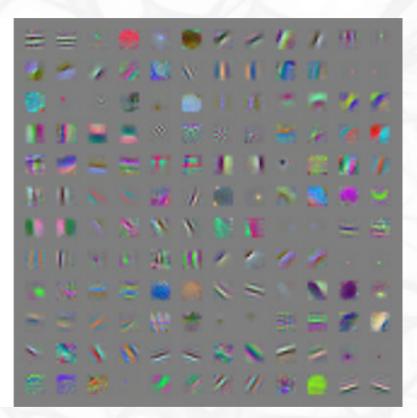


The primary visual cortex recognizes Gabor functions.

Similarities in V1 and CNNs







Similar functions are recognized in the first layer of a CNN.

AlexNet



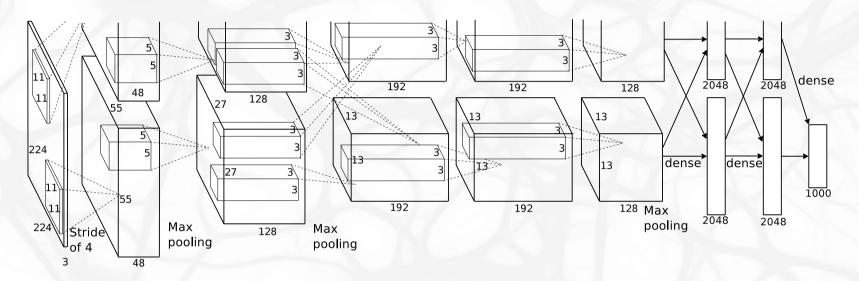
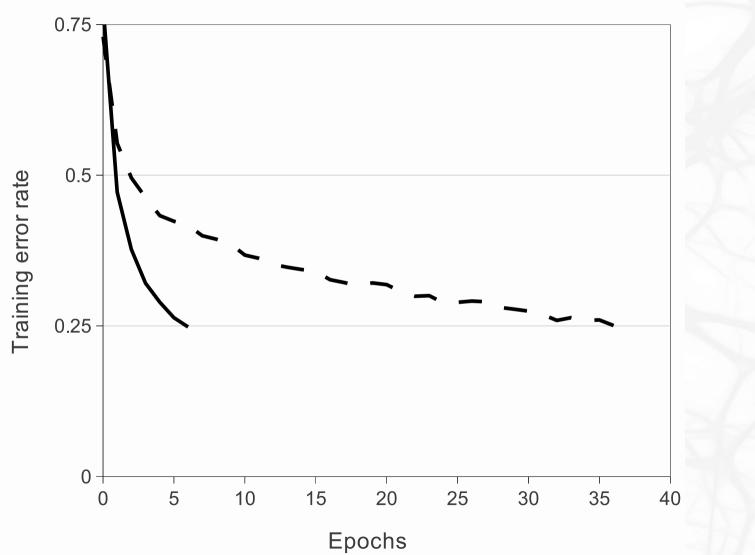


Figure 2: An illustration of the architecture of our CNN, explicitly showing the delineation of responsibilities between the two GPUs. One GPU runs the layer-parts at the top of the figure while the other runs the layer-parts at the bottom. The GPUs communicate only at certain layers. The network's input is 150,528-dimensional, and the number of neurons in the network's remaining layers is given by 253, 40–186,624–64,896–64,896–43,264–4096–4096–1000.

AlexNet





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