### NPFL114, Lecture 04

# **Convolutional Networks**



Milan Straka

### **Dropout Implementation**



```
def dropout(input, rate=0.5, training=False):
  def do_inference():
     return tf.identity(input)
  def do train():
     random_noise = tf.random_uniform(tf.shape(input))
     mask = tf.cast(tf.less(random_noise, rate),
               tf.float32)
     return x * mask / (1 - rate)
  if training == True:
     return do_train()
  if training == False:
     return do_inference()
  return tf.cond(training, do_train, do_inference)
```

# **Dropout Effect**



# **Dropout Effect**



## 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 ÚFAL



Reference: Image from

https://isaacchanghau.github.io/images/deeplearning/activationfunction/tanh.png.



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

The authors theoretically and experimentally show that a suitable way to initialize a  $R^{n \times m}$  matrix is

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





# Convergence – Gradient Clipping



### Convergence - Gradient Clipping



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

$$g \leftarrow \begin{cases} g & \text{if } ||g|| \le c \\ c \frac{g}{||g||} & \text{if } ||g|| > c \end{cases}$$

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**



Reference: Image from https://i.stack.imgur.com/YDusp.png.



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

$$(x \star w)(t) = \int x(a)w(t-a) da.$$

For vectors, we have

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

Convolution operation can be generalized to two dimensions by

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

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

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

# Convolution



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

$$(\boldsymbol{I} \star \boldsymbol{K})_{i,j,k} = \sum_{m,n,o} \boldsymbol{I}_{i \cdot S+m,j \cdot S+n,o} \boldsymbol{K}_{m,n,o,k}.$$



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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



There are two prevalent image formats:

• NHWC or channels\_last: The dimensions of the 4-dimensional image tensor are batch, height, width, and channels.

The original TensorFlow format, faster on CPU.



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- NHWC or channels\_last: The dimensions of the 4-dimensional image tensor are batch, height, width, and channels.
  - The original TensorFlow format, faster on CPU.
- NCHW or channels\_first: The dimensions of the 4-dimensional image tensor are batch, channel, height, and width.
  - Usual GPU format (used by CUDA and nearly all frameworks); on TensorFlow, not all CPU kernels are available with this layout.

### **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

Reference: Image from https://cdn-images-1.medium.com/max/1200/0\*QyXSpqpm1wc\_Dt6V... AlexNet - 2012 (16.4% error)



### AlexNet - 2012 (16.4% error)



### Training details:

- 2 GPUs for 5-6 days
- SGD with batch size 128, momentum 0.9, weight decay 0.0005
- initial learning rate 0.01, manually divided by 10 when validation error rate stopped improving
- dropout with rate 0.5 on fully-connected layers
- data augmentation using translations and horizontal reflections (choosing random  $224 \times 224$  patches from  $256 \times 256$  images)
  - during inference, 10 patches are used (four corner patches and a center patch, as well as their reflections)

# AlexNet - ReLU vs tanh



# LeNet - 1998



Achieved 0.8% test error on MNIST.

### 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.



Reference: Figure 1 of paper "Deep Prior", https://arxiv.org/abs/1712.05016



Reference: Figure 7 of paper "Deep Prior", https://arxiv.org/abs/1712.05016



Reference: Figure 5 of supplementary materials of paper "Deep Prior", https://arxiv.org/abs/1712.05016



Reference: Figure 8 of paper "Deep Prior", https://arxiv.org/abs/1712.05016

<u>Deep Prior paper website with supplementary material</u>

VGG - 2014 (6.8% error)



Inception (GoogLeNet) – 2014 (6.7% error) ÚFAL



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# Inception (GoogLeNet) – 2014 (6.7% error) ÚFAL



aux. classifiers, 0.3 weight



*Internal covariate shift* refers to the change in the distributions of hidden node activations due to the updates of network parameters during training.

Let  $\boldsymbol{x} = (x_1, \dots, x_d)$  be d-dimensional input. We would like to normalize each dimension as

$$\hat{x}_i = \frac{x_i - \mathrm{E}[x_i]}{\sqrt{\mathrm{Var}[x_i]}}.$$

Furthermore, it may be advantageous to learn suitable scale  $\gamma_i$  and shift  $\beta_i$  to produce normalized value

$$y_i = \gamma_i \hat{x}_i + \beta_i$$
.



Consider a mini-batch of m examples  $(\boldsymbol{x}^{(1)}, \dots, \boldsymbol{x}^{(m)})$ .

Batch normalizing transform of the mini-batch is the following transformation.

```
Inputs: Mini-batch (\boldsymbol{x}^{(1)}, \dots, \boldsymbol{x}^{(m)}), \varepsilon \in \mathbb{R}

Outputs: Normalized batch (\boldsymbol{y}^{(1)}, \dots, \boldsymbol{y}^{(m)})

• \boldsymbol{\mu} \leftarrow \frac{1}{m} \sum_{i=1}^{m} \boldsymbol{x}^{(i)}

• \boldsymbol{\sigma}^2 \leftarrow \frac{1}{m} \sum_{i=1}^{m} (\boldsymbol{x}^{(i)} - \boldsymbol{\mu})^2

• \boldsymbol{x}^{(i)} \leftarrow (\boldsymbol{x}^{(i)} - \boldsymbol{\mu})/\sqrt{\sigma^2 + \varepsilon}

• \boldsymbol{y}^{(i)} \leftarrow \boldsymbol{\gamma} \boldsymbol{x}^{(i)} + \boldsymbol{\beta}
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Batch normalization is commonly added just before a nonlinearity. Therefore, we replace  $f(\mathbf{W}\mathbf{x} + \mathbf{b})$  by  $f(BN(\mathbf{W}\mathbf{x}))$ .



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During inference,  $\mu$  and  $\sigma^2$  are fixed. They are either precomputed after training on the whole training data, or an exponential moving average is updated during training.

Inception with BatchNorm (4.8% error)





- Reference: Figure 1 of paper "Rethinking the Inception Architecture for Computer Vision", https://arxiv.org/abs/1512.00567.
- Reference: Figure 3 of paper "Rethinking the Inception Architecture for Computer Vision", https://arxiv.org/abs/1512.00567.



















Reference: Figure 1 of paper "Visualizing the Loss Landscape of Neural Nets", https://arxiv.org/abs/1712.09913.