COMP 448/548: Medical Image Analysis

Convolutional neural networks

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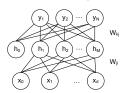
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Multilayer perceptrons

(revisited)

Contain hidden layers



Hidden units h_j's can be viewed as new "features" obtained by combining x_i's

A deeper architecture with nonlinear activations is more expressive than a shallow one

In this network

1. Each hidden unit computes its net activation $net_j^t = \sum_i x_i^t \ W_{ji}$

2. Each hidden unit emits an output that is a nonlinear function (e.g., sigmoid, ReLU) of its activation $h_i^t = non-linear-function(net_i^t)$

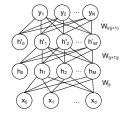
3. Each output unit computes its net activation $net_k^t = \sum_{j} h_i^t \ W_{kj}$

4. Each output units emits an output (using a linear, a sigmoid or a softmax function)

 $y_k^t = output - function(net_k^t)$

More hidden layers

(revisited)



 $\frac{\partial loss^{t}(W)}{\partial W_{ji}} = \frac{\partial loss^{t}(W)}{\partial net_{j}^{t}} \cdot \frac{\partial net_{j}^{t}}{\partial W_{ji}}$ $\frac{\partial loss^{t}(W)}{\partial t} = \delta_{i}^{t} \cdot x_{i}^{t}$

 $\delta_{j}^{t} = \sum_{(j+1)} \frac{\partial loss^{t}(W)}{\partial net^{t}_{(j+1)}} \cdot \frac{\partial net^{t}_{(j+1)}}{\partial net^{t}_{j}}$

 $\delta_j^t = \left[\sum_{i \in \mathcal{I}} \delta_{(j+1)}^t \cdot W_{(j+1)j}\right] \cdot \sigma'(net_j)$

δ_j may vanish after repeated multiplication. This makes deep architectures hard to train (when initial weights are not "good" enough)

Approaches for alleviating underfitting and overfitting problems

- Better network designs: Sparse connections, weight sharing, convolutional nets, long/short skip connections, activation functions, ...
- Better network training: Regularization, loss function definitions, larger datasets, data augmentation, ...
- Previously, layerwise pretraining (restricted Boltzmann machines, autoencoders)

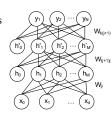
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Layerwise pretraining

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Vanishing gradient problem

- This makes hard to train deep architectures (with many hidden layers) by backpropagation
- When the initial weights are good enough, backpropagation works well
- Layerwise pretraining
 - Restricted Boltzmann machines
 - Autoencoders

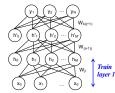


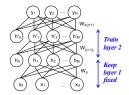
$$\begin{split} \frac{\partial loss^t(W)}{\partial W_{ji}} &= \frac{\partial loss^t(W)}{\partial net_j^t} \cdot \frac{\partial net_j^t}{\partial W_{ji}} \\ \frac{\partial loss^t(W)}{\partial W_{ji}} &= \delta_j^t \cdot x_i^t \\ \delta_j^t &= \int_{(j+1)} \frac{\partial loss^t(W)}{\partial net_{(j+1)}^t} \cdot \frac{\partial net_{(j+1)}^t}{\partial net_j^t} \\ \delta_j^t &= \left[\int_{(j+1)} \delta_{(j+1)}^t \cdot W_{(j+1)j} \right] \cdot \sigma'(net_j^t) \end{split}$$

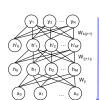
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Layerwise pretraining

- First, train one layer at a time, optimizing P(x)
- Then, fine-tune weights, optimizing P(y|x) by backpropagation







Restricted Boltzmann machines (RBMs)

■ RBM is a simple energy-based $p(x,h) = \frac{1}{Z_{\theta}} \exp(-E_{\theta}(x,h))$ model

$$\begin{array}{lll} p(x,h) & = & \frac{1}{Z_{\theta}} \exp(-E_{\theta}(x,h)) \\ \\ E_{\theta}(x,h) & = & -x^T \ W \ h - b^T \ x - d^T \ h & \text{it only allows} \\ \\ Z_{\theta} & = & \sum_{(x,h)} \exp(-E_{\theta}(x,h)) & \leftarrow & \text{normalizer} \end{array}$$

Train an RBM optimizing P(x)

$$\begin{split} \text{Maximize the log-likelihood of data} \\ \frac{\partial_{W_{ji}} \log P_W(x = x^t)}{\log P_W(x = x^t)} &= \partial_{W_{ji}} \log \sum_{h} P_W(x = x^t, h) \\ &= -\partial_{W_{ji}} \log Z_w + \partial_{W_{ji}} \log \sum_{h} \exp(-E_W(x^t, h)) \\ &= -E_{p(x, h)} \left[x_i \ h_j \right] \\ &= -E_{p(x, h)} \left[x$$

The negative phase term is expensive to calculate since it requires sampling (x, h) from the model. **Contrastive divergence** is a faster solution.

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Autoencoders

- They learn to "compress" and "reconstruct" the input data
- Encoder: $h = \sigma(Wx + b)$ Decoder: $x' = \sigma(W'h + d)$
- Learn the weights to minimize the reconstruction loss

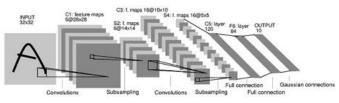
$$loss = \sum_{t} (x^t - x'^t)^2$$

- This is the same backpropagation for a network with one hidden layer, where x^t is both input and output
- They can be stacked to form a deep neural network
 - Cheaper alternatives to RBMs

Basics of convolutional neural networks

Convolutional neural networks (CNNs)

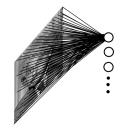
 A CNN consists of a number of convolutional and pooling (subsampling) layers optionally followed by fully connected layers



LeNet-5 by LeCun et al., 1998

Fully connected layers

■ When the input data is an image, a fully connected layer will produce a huge number of weights (parameters) to be learned



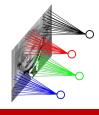
Example: 200x200 image

25K hidden units → ~1B parameters

Slide credit: M.A. Ranzalo

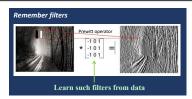
Convolutional layer

- However, spatial correlation is local and statistics is similar at different locations
- Thus, small kernels are defined and their parameters are shared by all pixels
- It is convolution with learned kernels



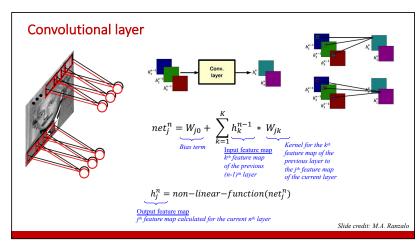
Example: 200x200 image 25K hidden units 10x10 kernels → ~2.5M parameters

(instead of 1B parameters)



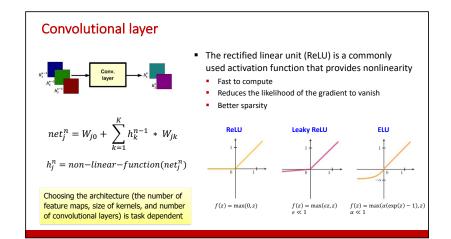
Sparse connections and weight sharing Slide credit: M.A. Ranzalo

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Pooling layer

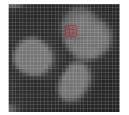
• By pooling the filter responses at different locations
• We gain robustness to the exact location of features
• Receptive field becomes larger for the next layer (the next layer will look at a larger spatial region)

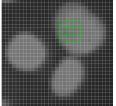
Max-pooling: $h_j^n(x,y) = \max_{\substack{\bar{x} \in N(x) \\ \bar{y} \in N(y)}} h_j^{n-1}(\bar{x},\bar{y})$ Average-pooling: $h_j^n(x,y) = 1/K \sum_{\substack{\bar{x} \in N(x) \\ \bar{y} \in N(y)}} h_j^{n-1}(\bar{x},\bar{y})$ L2-pooling: $h_j^n(x,y) = \sqrt{\sum_{\substack{\bar{x} \in N(x) \\ \bar{y} \in N(y)}}} h_j^{n-1}(\bar{x},\bar{y})^2$ Slide credit: M.A. Ranzalo

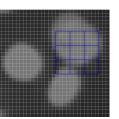
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Receptive field

■ It is the region in the input space that a convolution can "see"





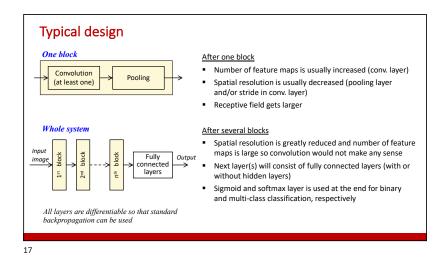


What a 3x3 convolutional filter sees What the 3x3 convolutional filter sees after 2x2 pooling

What the 3x3 convolutional filter sees after applying 2x2 pooling twice

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Example CNN architectures

C1. Income 160 Principle
Substantion Column C

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Example CNN architectures

224 x 224 x 3 224 x 224 x 64

112 x 112 x 128

56 x 56 x 256

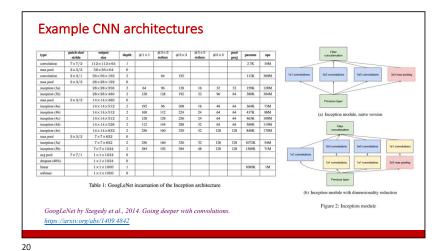
7 x 7 x 512

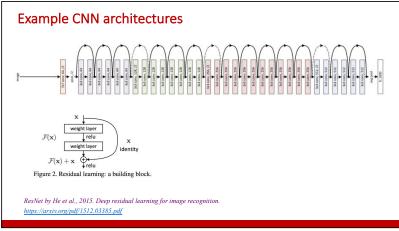
1 x 1 x 4096 1 x 1 x 1000

convolution + ReLU
max pooling
fully nected + ReLU
softmax

VGG16 by Simonyan and Zisserman, 2013. Very deep convolutional networks for large-scale image recognition.
https://arxiv.org/pdi/1409.1556.pdf

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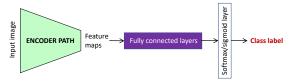
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CNNs for medical image classification

Please check the following survey paper for more about deep learning in medical images and more references. Litjens et al., a survey on deep learning in medical image analysis, Medical Image Analysis, 2017. https://www.sciencedirect.com/science/article/pii/S1361841517301135

CNNs for image classification

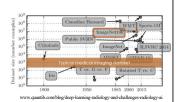
- A CNN compresses an image into a set of feature maps to capture semantic/contextual information from the image
- This compression corresponds to downsampling the image using convolution and pooling layers
- Then it puts fully connected layers on the top of the feature maps to predict a class for the entire image



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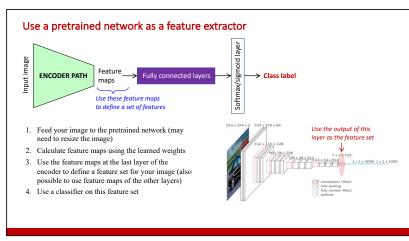
Transfer learning

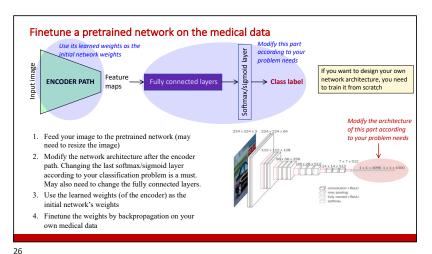
- Dataset sizes are typically small
- Thus, it is popular to use transfer learning, which employs networks (and thus, their learned weights) previously trained on large datasets

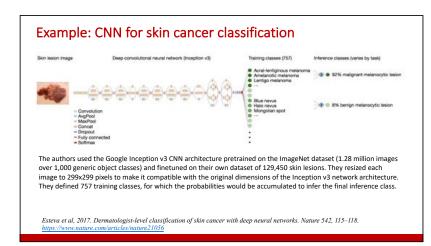


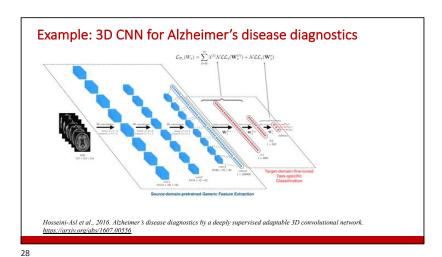
- Two main approaches
 - 1. Use a pretrained network as a feature extractor
 - 2. Finetune a pretrained network on the medical data

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CNNs for object detection and segmentation

Training:

- Small patches are cropped around individual pixels
- Each patch is labeled with the class of the pixel, around which it is cropped
- CNN is trained on these small patches

Detection/segmentation:

- For an entire (large) image, patches are obtained using a sliding window approach
- These patches are classified by the trained CNN
- Outputs (i.e., posteriors) generated by this CNN are commonly postprocessed

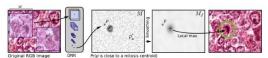
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Example: CNN for mitosis detection

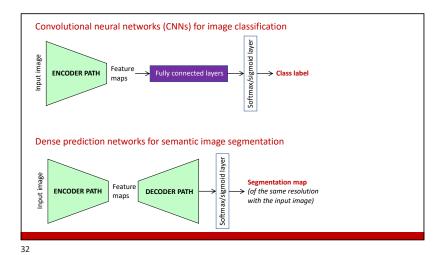
- Slide a window over an image to obtain patches
- Using the trained CNN, obtain the probability of a pixel belonging to a mitotic cell
- Find the local maxima on the smoothed probability map
- Classifying each pixel in a sliding window fashion, used by earlier studies, is expensive as it requires lots of redundant calculations
- 2. Trade-off between localization accuracy and the use of context
 - Larger patches require more max-pooling layers that reduce the localization accuracy
 - · Small patches results in seeing only little context

Dense prediction networks, used by recent studies, have greatly improved efficiency and accuracy.



Ciresan et al., 2013. Mitosis detection in breast cancer histology images with deep neural networks. MICCAI. https://link.springer.com/chapter/10.1007/978-3-642-40763-5_51

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Thank you!

Next time:

Dense prediction networks