

CS 189/289

Today's lecture:

1. From logistic to softmax.
2. Convolutional neural networks
3. Residual neural networks (resnets)

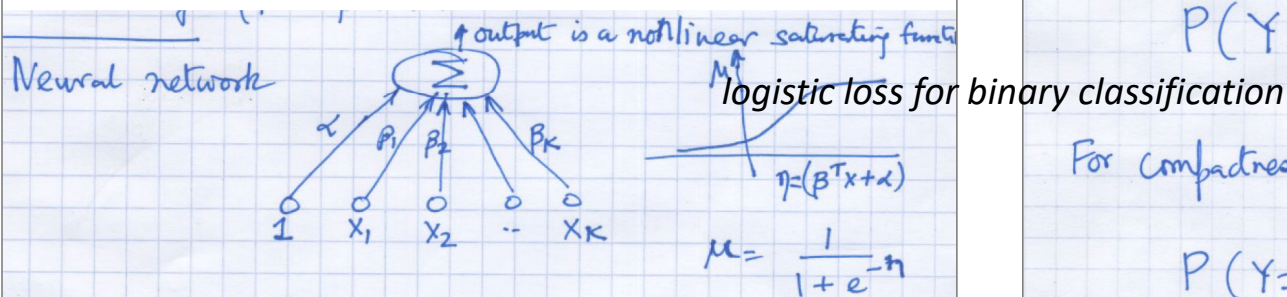
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Recall: logistic loss for binary classification

Neural networks can be modeled by logistics



Standard Trick: Add a 0^{th} component to the \underline{x} vector, which is fixed to be 1. This is connected with weight α

Modeling the probability distribution

We say that the class label Y is a Bernoulli random variable, with its probability parameter p being as above

$$P(Y=1 | X) = \frac{1}{1 + \exp(-\beta^T x)}$$

For compactness, introduce notation $\mu(x) = \frac{1}{1 + \exp(-\beta^T x)}$

$$P(Y=1 | X) = \mu(x) \quad \text{or} \quad \mu(x) = \frac{1}{1 + \exp(-\eta(x))}$$

As usual we use y to denote values taken by random variables

$$P(y|x) = \mu(x)^y (1 - \mu(x))^{1-y}$$

What if we have more than 2 classes?

From logistic regression to softmax regression

$$\begin{bmatrix} p(Y = 1|X) \\ p(Y = 0|X) \end{bmatrix} = \begin{bmatrix} \mu \\ 1 - \mu \end{bmatrix} = \begin{bmatrix} \frac{1}{1 + \exp(-\beta x)} \\ \frac{\exp(-\beta x)}{1 + \exp(-\beta x)} \end{bmatrix}$$

From logistic regression to softmax regression

$$\begin{bmatrix} p(Y=1|X) \\ p(Y=0|X) \end{bmatrix} = \begin{bmatrix} \mu \\ 1-\mu \end{bmatrix} = \begin{bmatrix} \frac{1}{1 + \exp(-\beta X)} \\ \frac{\exp(-\beta X)}{1 + \exp(-\beta X)} \end{bmatrix}$$

Instead we could write this as

$$\frac{1}{e^{\beta_1 X} + e^{\beta_2 X}} \begin{bmatrix} e^{\beta_1 X} \\ e^{\beta_2 X} \end{bmatrix} = \begin{bmatrix} \frac{1}{1 + e^{\beta_2 X - \beta_1 X}} \\ \frac{1}{1 + e^{-\beta_2 X + \beta_1 X}} \end{bmatrix}$$

Equivalent with $\beta = \beta_1 - \beta_2$

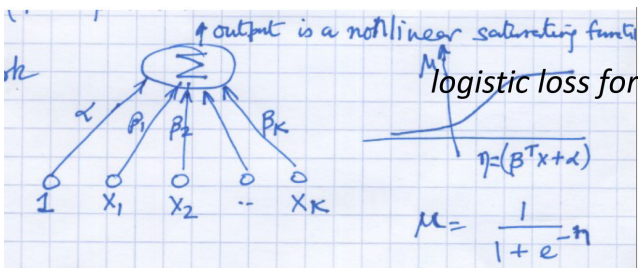
The softmax function for K-class classification

$$\begin{bmatrix} p(Y = 1|X) \\ p(Y = 2|X) \\ p(Y = 3|X) \\ \dots \\ p(Y = K|X) \end{bmatrix} =$$

$$\frac{1}{\sum_{i=1}^K e^{\beta_i X}}$$

$$\begin{bmatrix} e^{\beta_1 X} \\ e^{\beta_2 X} \\ \vdots \\ e^{\beta_K X} \end{bmatrix}$$

- Generalization of logistic regression to more than 2 classes.
- "Softmax regression" or "multinomial logistic regression", parameters β .
- Use principle of MLE to set β .
- Needs iterative optimization like gradient descent.
- Can also stick at the top of neural network to get a "softmax" loss.



The softmax function for K-class classification

$$\begin{bmatrix} p(Y = 1|X) \\ p(Y = 2|X) \\ p(Y = 3|X) \\ \dots \\ p(Y = K|X) \end{bmatrix} =$$

$$\frac{1}{\sum_{i=1}^K e^{\beta_i x}}$$

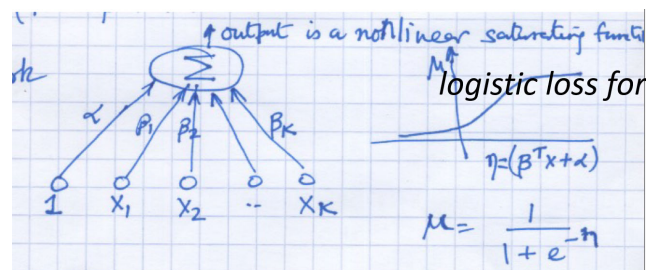
$$\begin{bmatrix} e^{\beta_1 x} \\ e^{\beta_2 x} \\ \vdots \\ e^{\beta_K x} \end{bmatrix}$$

For class `i`, the logit (log-odds) is defined as:

$$\text{logit}_i = \log \left(\frac{P(y=i|x)}{P(y \neq i|x)} \right)$$

For class `i`, the softmax function is defined as:

$$P(y = i|x) = \frac{e^{\text{logit}_i}}{\sum_{j=1}^K e^{\text{logit}_j}}$$

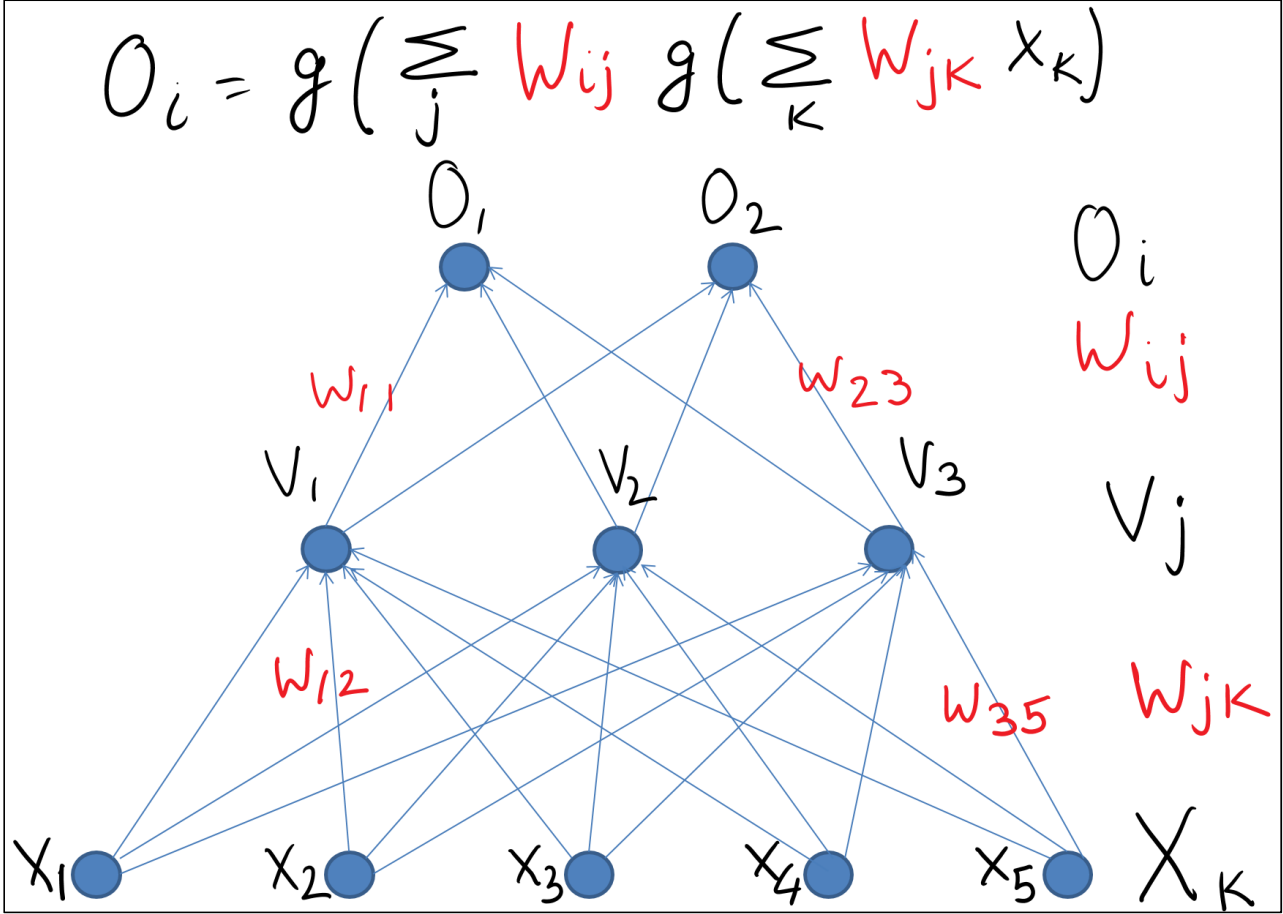


CS 189/289

Today's lecture outline:

1. From logistic to softmax.
2. Convolutional neural networks
3. Residual neural networks (resnets)

Recall: *fully connected* neural networks



Recall: feed forward, *fully connected* neural networks

Computing δ for neuron in intermediate layer

INDUCTION STEP

$$x_j^{(l)} = g \left(\sum_{i=0}^{d^{(l-1)}} w_{ij}^{(l)} x_i^{(l-1)} \right) = g(s_j^{(l)})$$

$s_j^{(l)}$ is the weighted input to node j in layer l

$$\delta_i^{(l-1)} = \frac{\partial e(w)}{\partial s_i^{(l-1)}}$$

$$= \sum_j \frac{\partial e(w)}{\partial s_j^{(l)}} \times \frac{\partial s_j^{(l)}}{\partial x_i^{(l-1)}} \times \frac{\partial x_i^{(l-1)}}{\partial s_i^{(l-1)}}$$

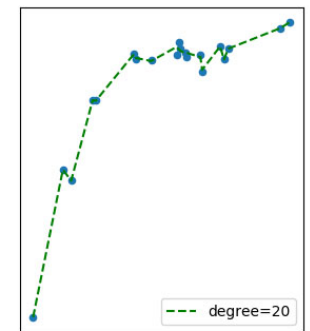
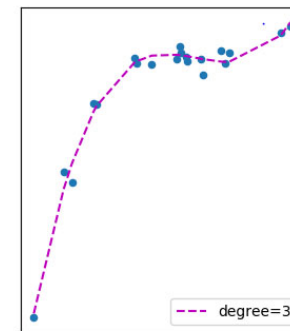
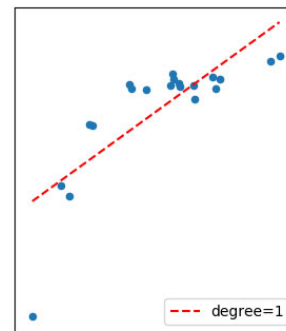
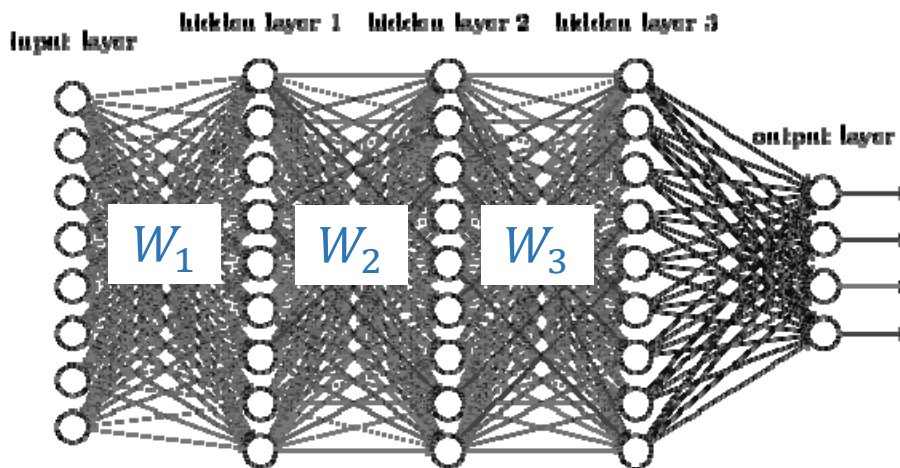
Back-propagation algorithm to compute derivatives of the parameters efficiently.

Beyond fully connected, feed-forward architectures:

1. *Convolutional*
 2. *Residual*
 3. *Recurrent* (not "feed-forward").
 4. *Attention and Transformers.*
 5. *Graph*
- As long as we have a feed-forward network, and use only differentiable components, we can apply backprop.
 - New architectures have led to break-through successes.

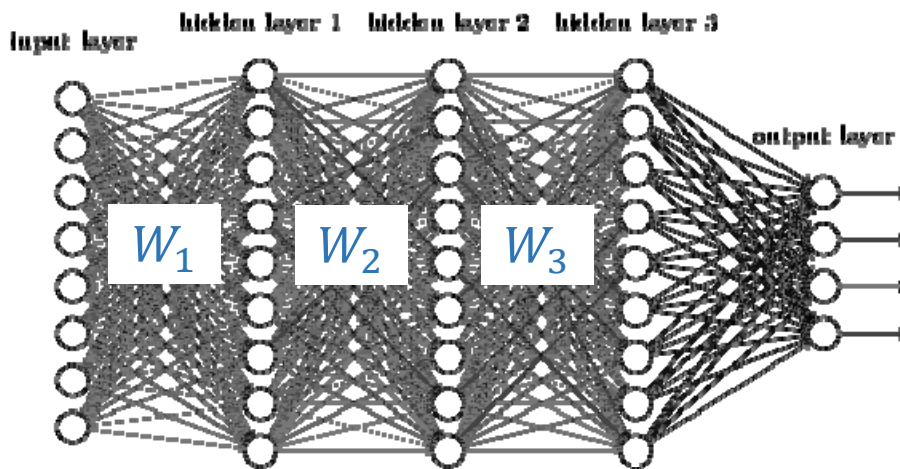
Pondering fully connected neural networks

- For “fully connected” (FC) layer, l , with $n_i(l)$ inputs and $n_o(l)$ outputs, W_l contains $n_i(l) \times n_o(l)$ parameters.
- Adds up quickly to huge #s of parameters.
- Too many parameters can contribute to problems of “overfitting”.



Pondering fully connected neural networks

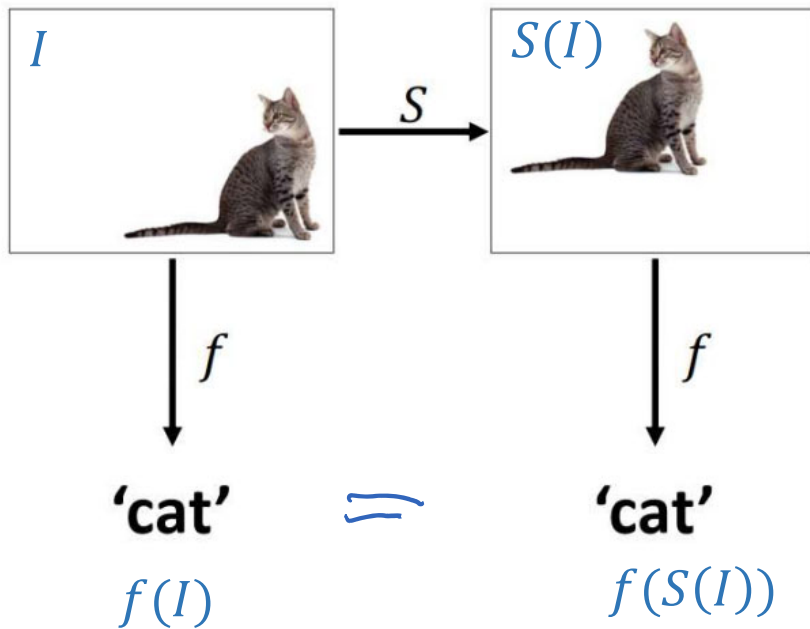
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➤ Strategy to reduce # of free parameters: “bake” in properties that encode problem symmetries.

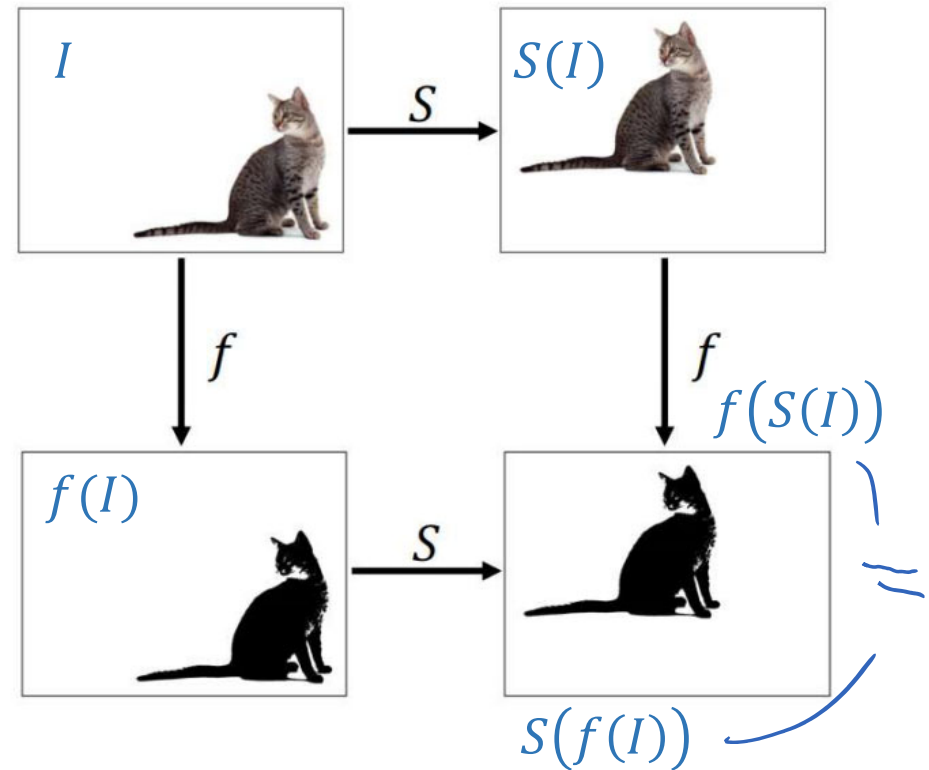
Examples of common problem symmetries

translation *in*variance



Predict: is a cat vs. not a cat

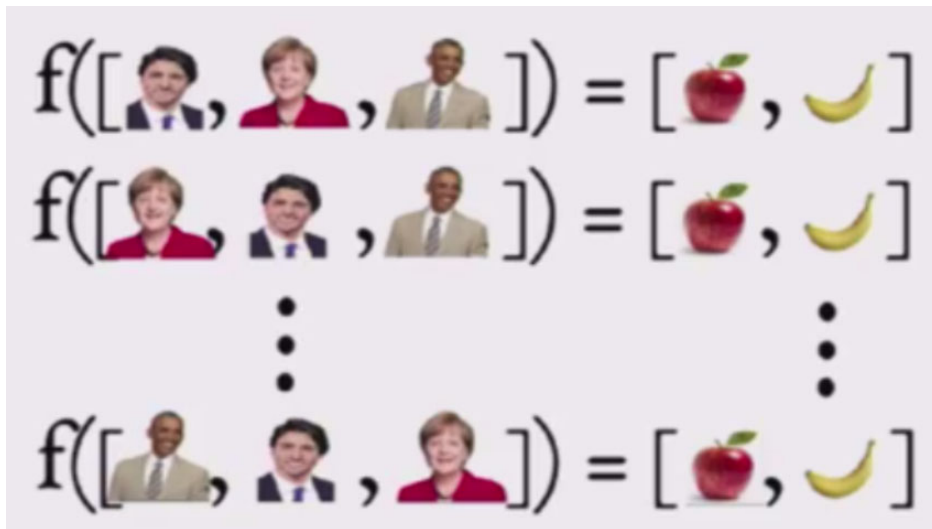
translation *equiv*ariance



Predict: which pixels are cat pixels?

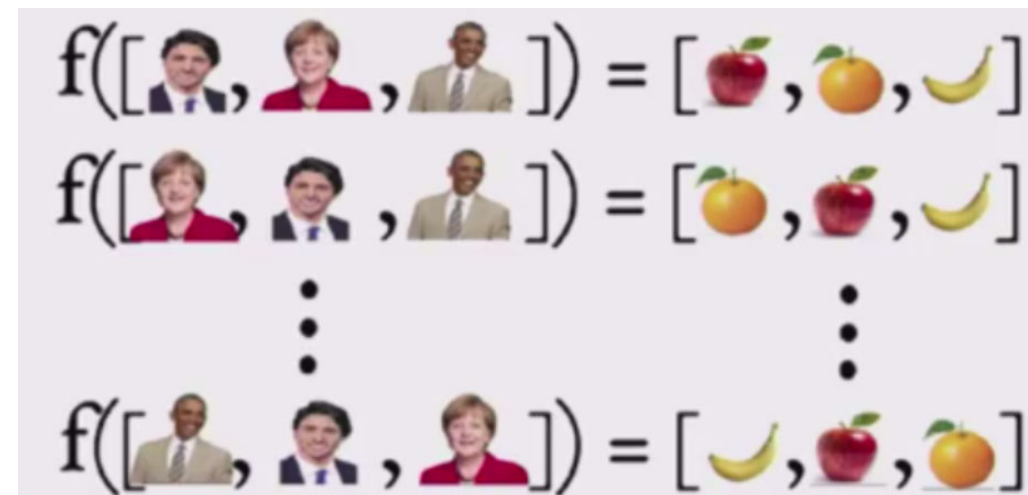
Examples of common problem symmetries

permutation *in*variance



$$f(x) = f(\text{Perm}(x))$$

permutation *equi*variance

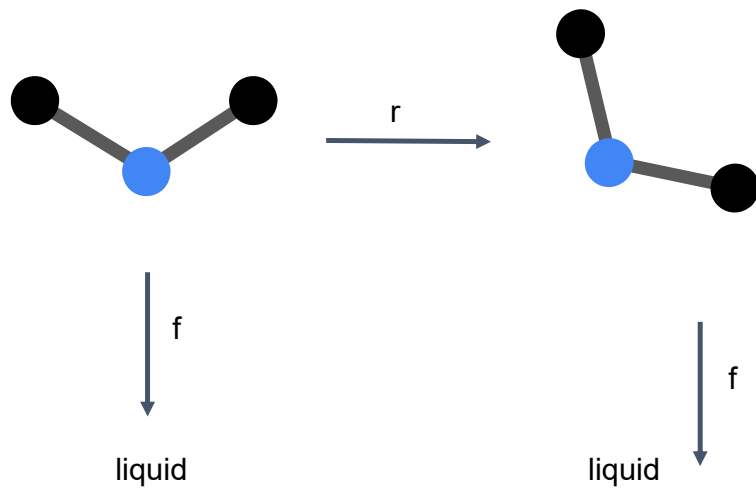


$$\text{Perm}(f(x)) = f(\text{Perm}(x))$$

Predict vector output.

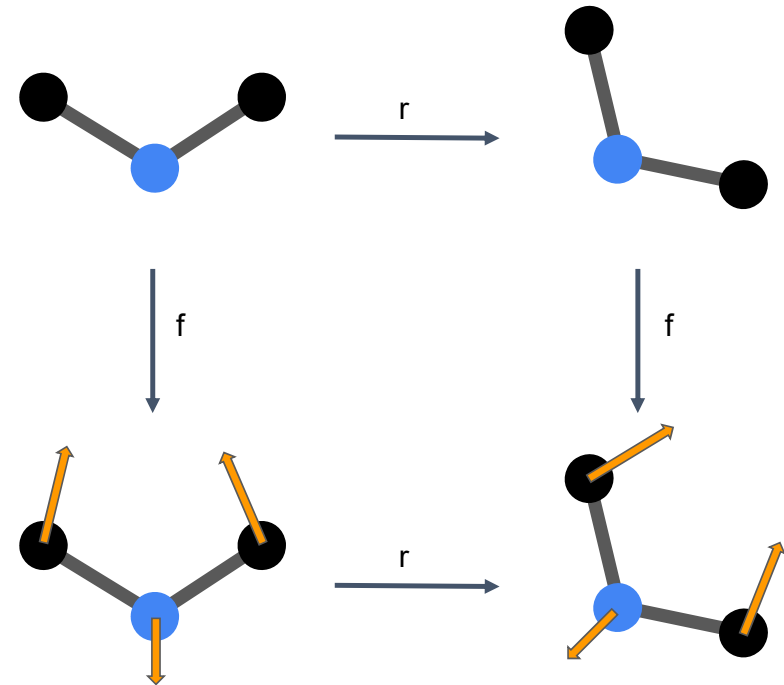
Examples of common problem symmetries

rotation *in*variance



predict phase (is liquid?)
at room temperature

rotation *equi*variance



predict **forces** (vector)

Examples of common problem symmetries

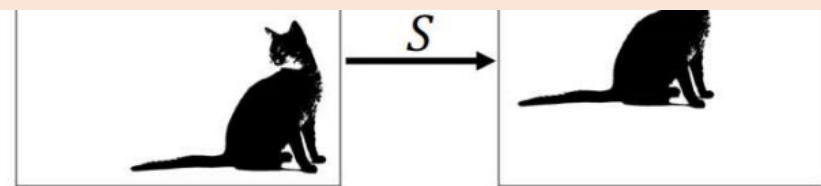
translation *in*variance

translation *equi*variance

- The convolution operation is translation equivariant.
- This operation will form the basis of *convolutional neural networks (CNNs)*.
- CNNs also be motivated by the idea of learning re-usable features (next).

'cat'

'cat'

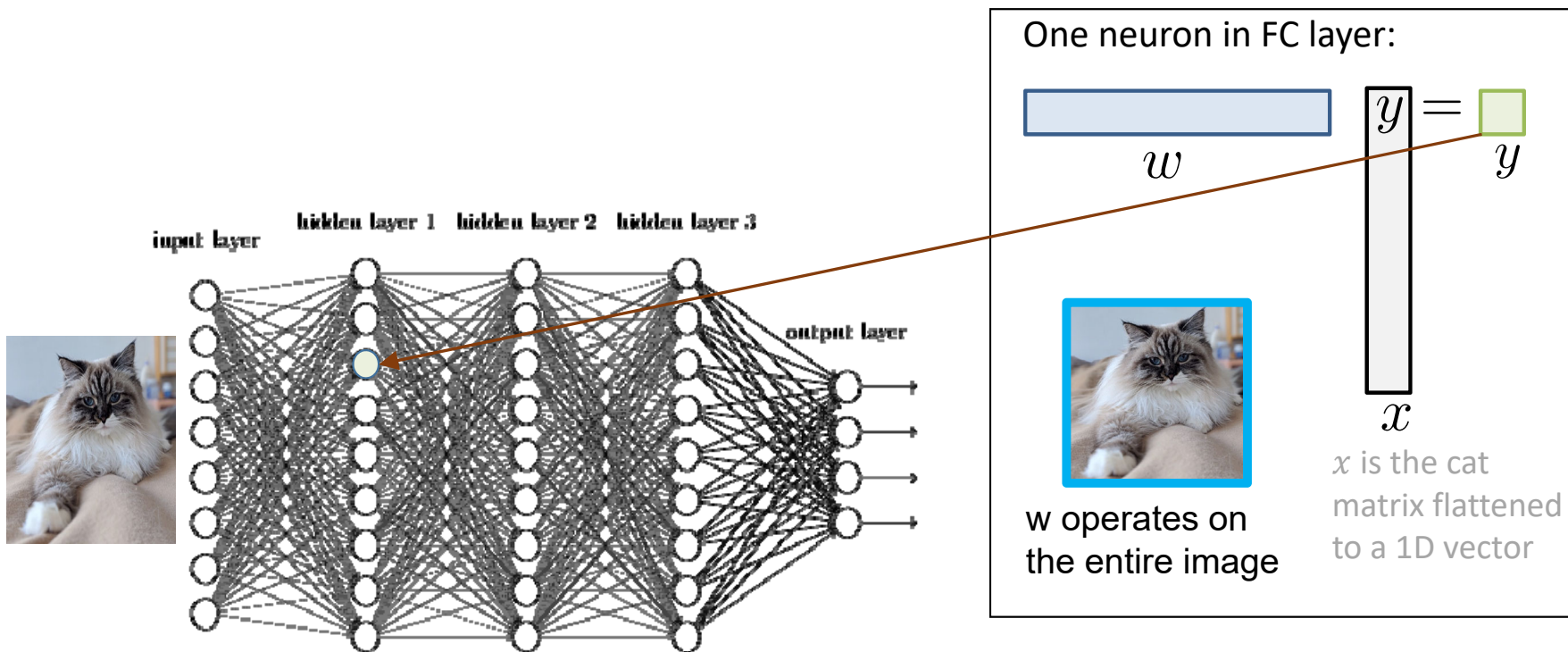


Predict: is a cat vs. not a cat

Predict: which pixels are cat pixels?

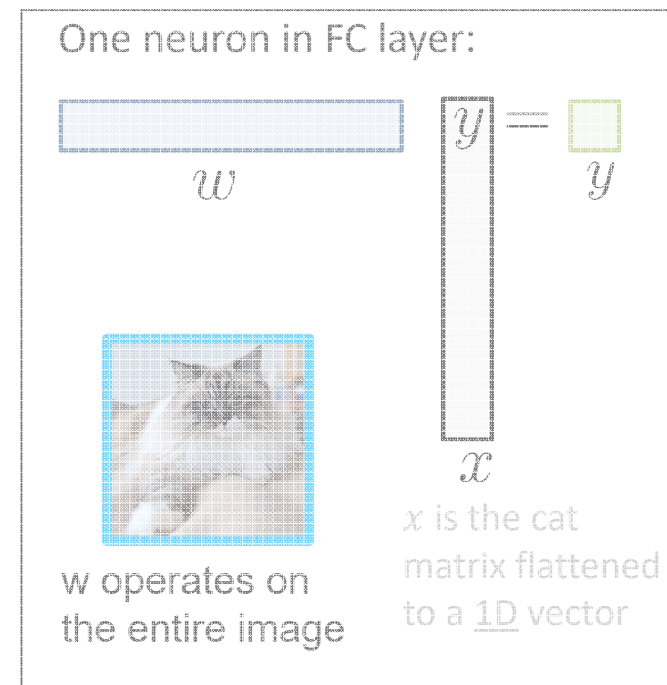
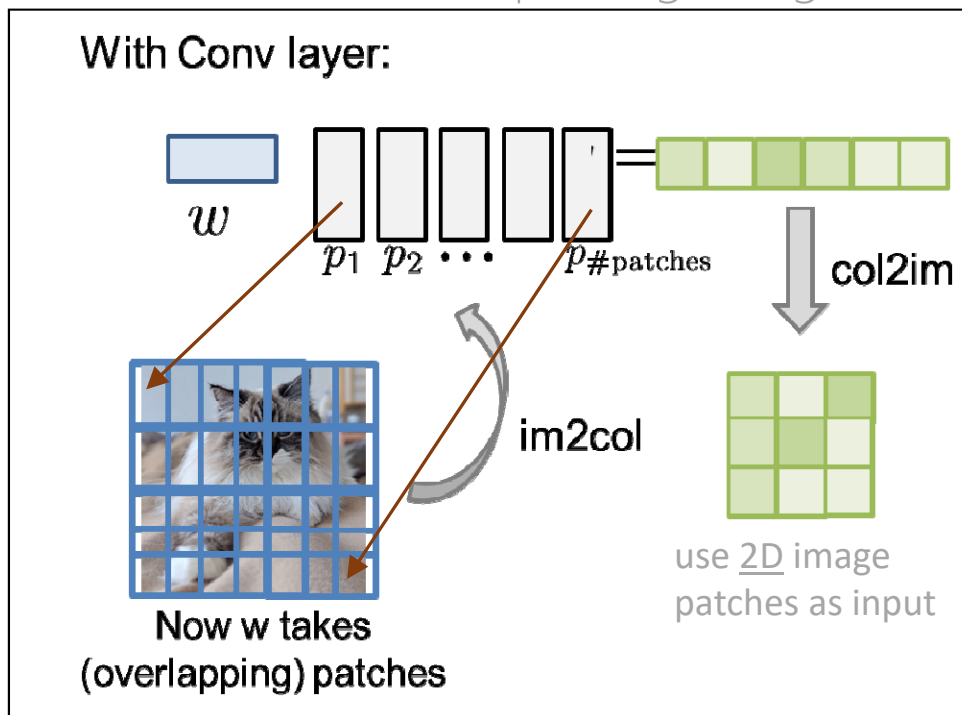
Features sharing across one input example

“Features” (e.g. is there an eye here?) constructed in fully connected layer cannot be shared across the input (e.g. image), because w is not reused across the image.



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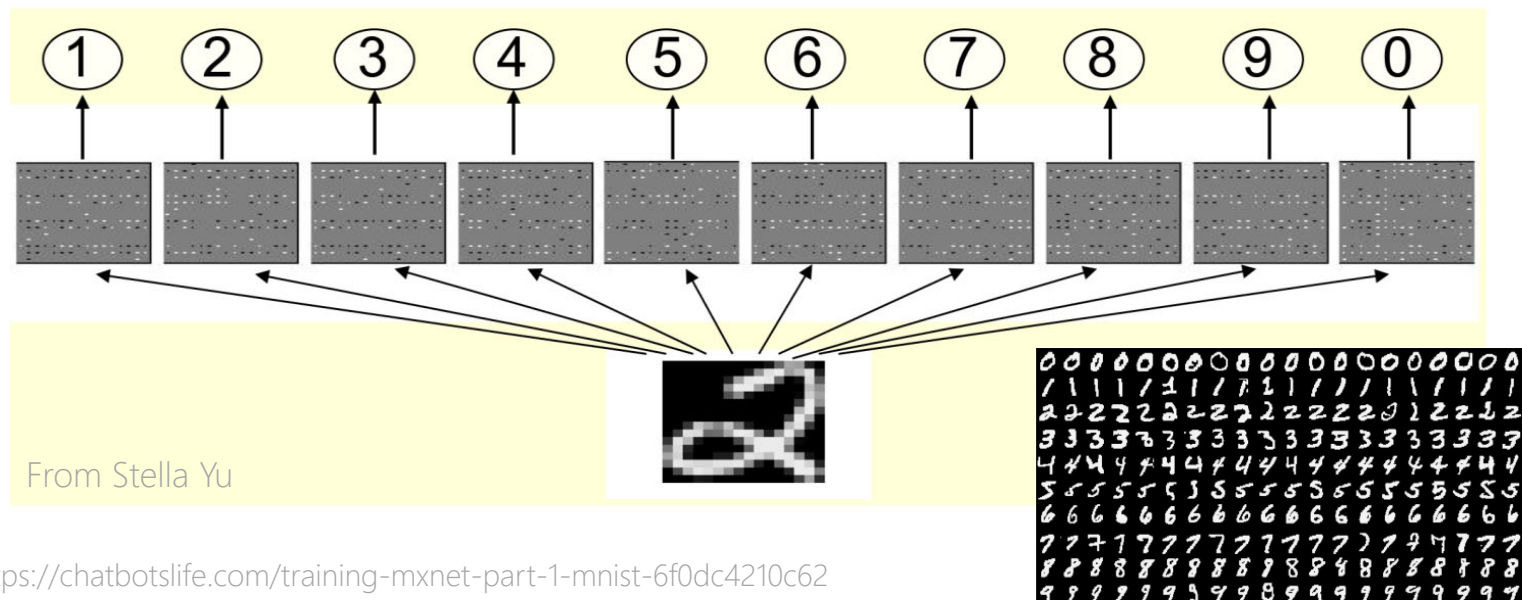


- ConvNet: learn shared features that are applied to every image patch.
- Also gives us *translational equivariance* for each filter (w) response.

Fully Connected (FC): no feature sharing

- Uses “global template matching”.
- e.g. one W matrix per class (single layer):

Iteration 1 of training:

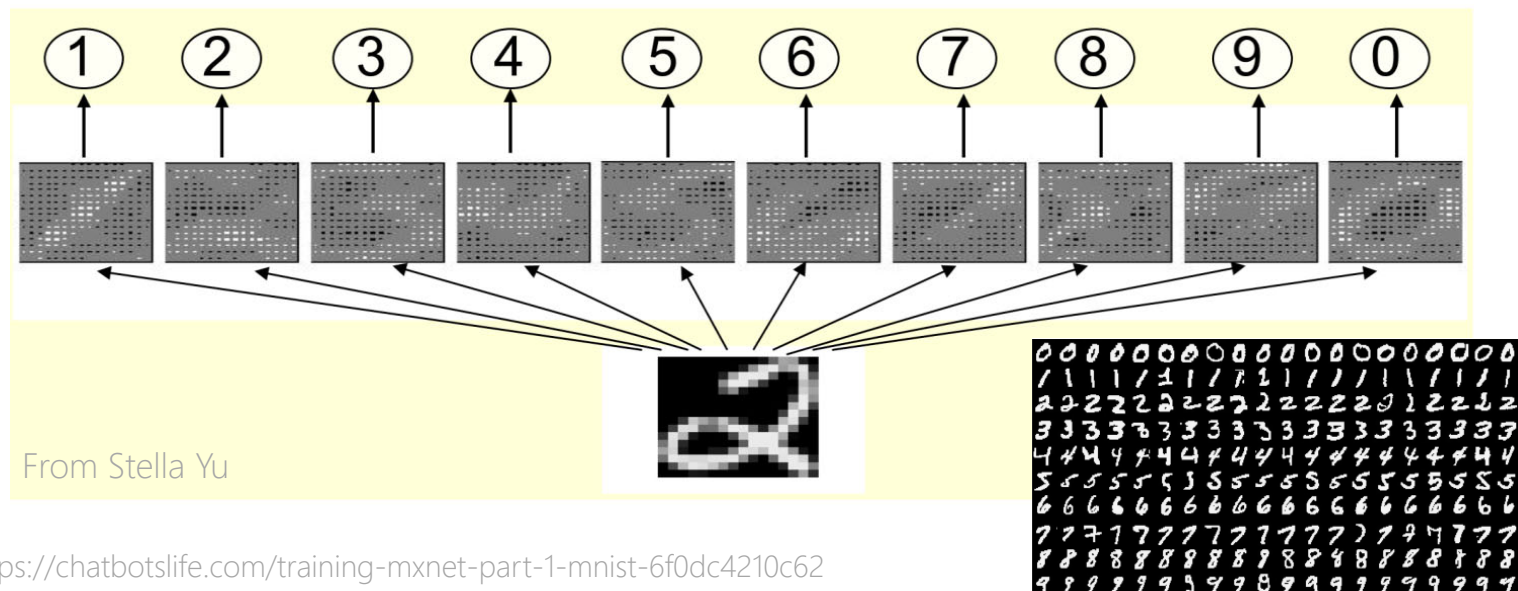


<https://chatbotslife.com/training-mxnet-part-1-mnist-6f0dc4210c62>

Fully Connected (FC): no feature sharing

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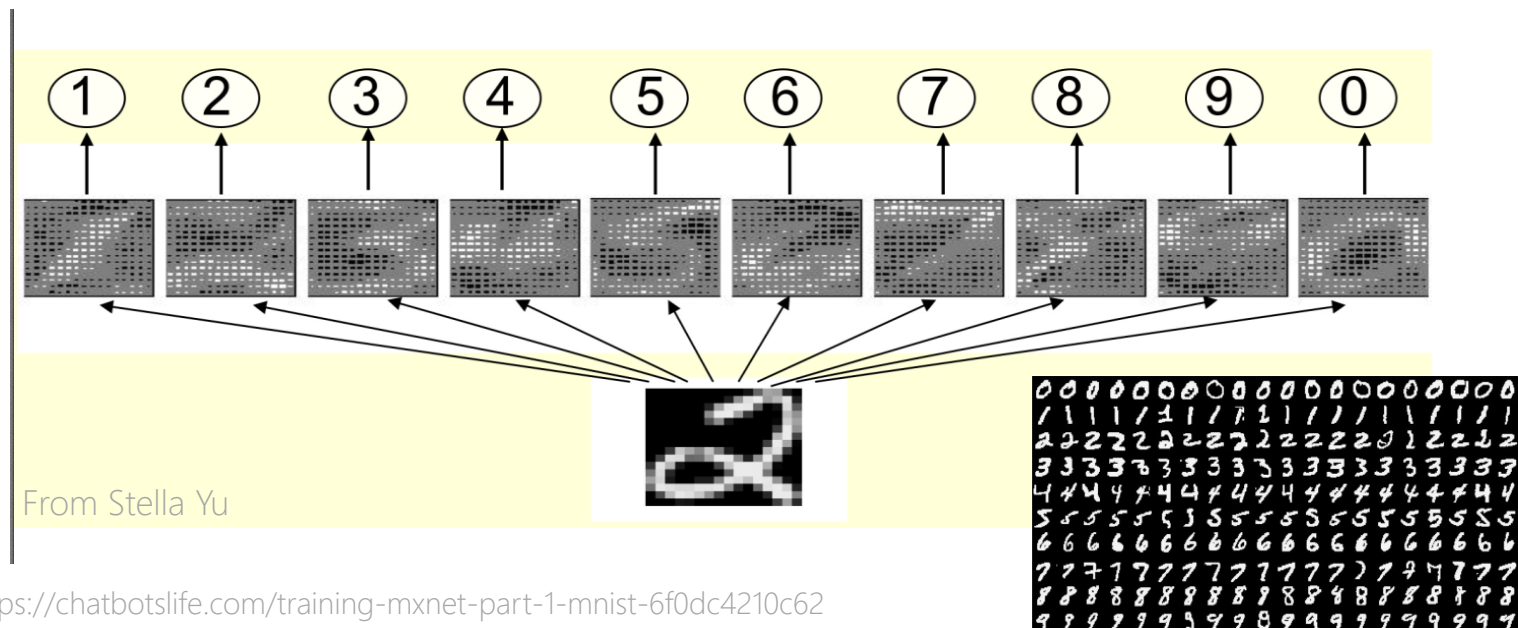
Iteration 3 of training:



Fully Connected (FC): no feature sharing

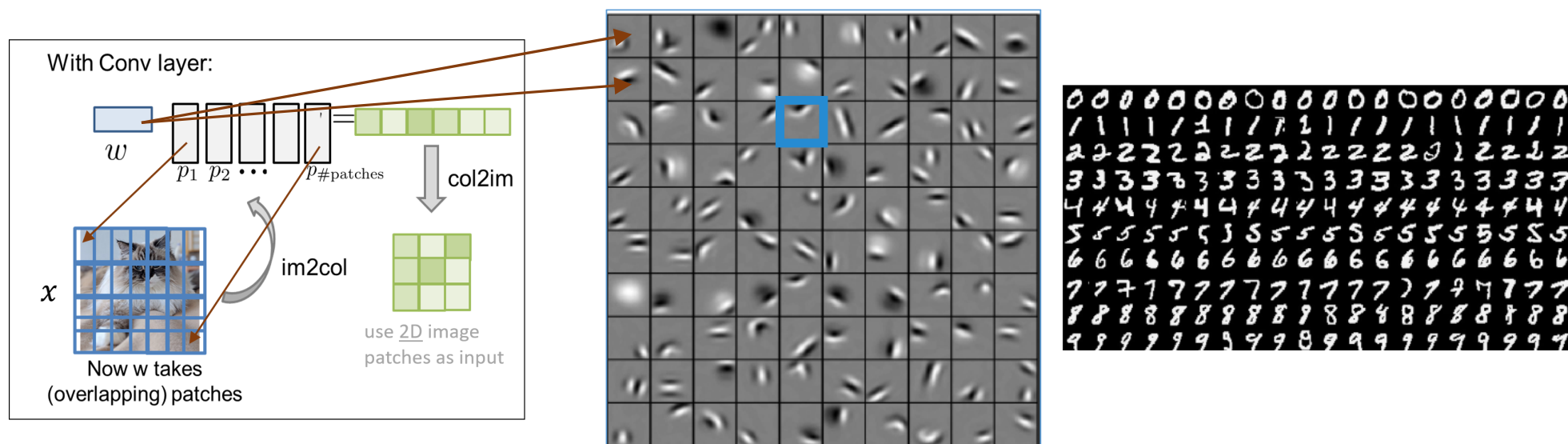
- Uses “global template matching”.
- e.g. one W matrix per class (single layer):

Iteration 7 of training:



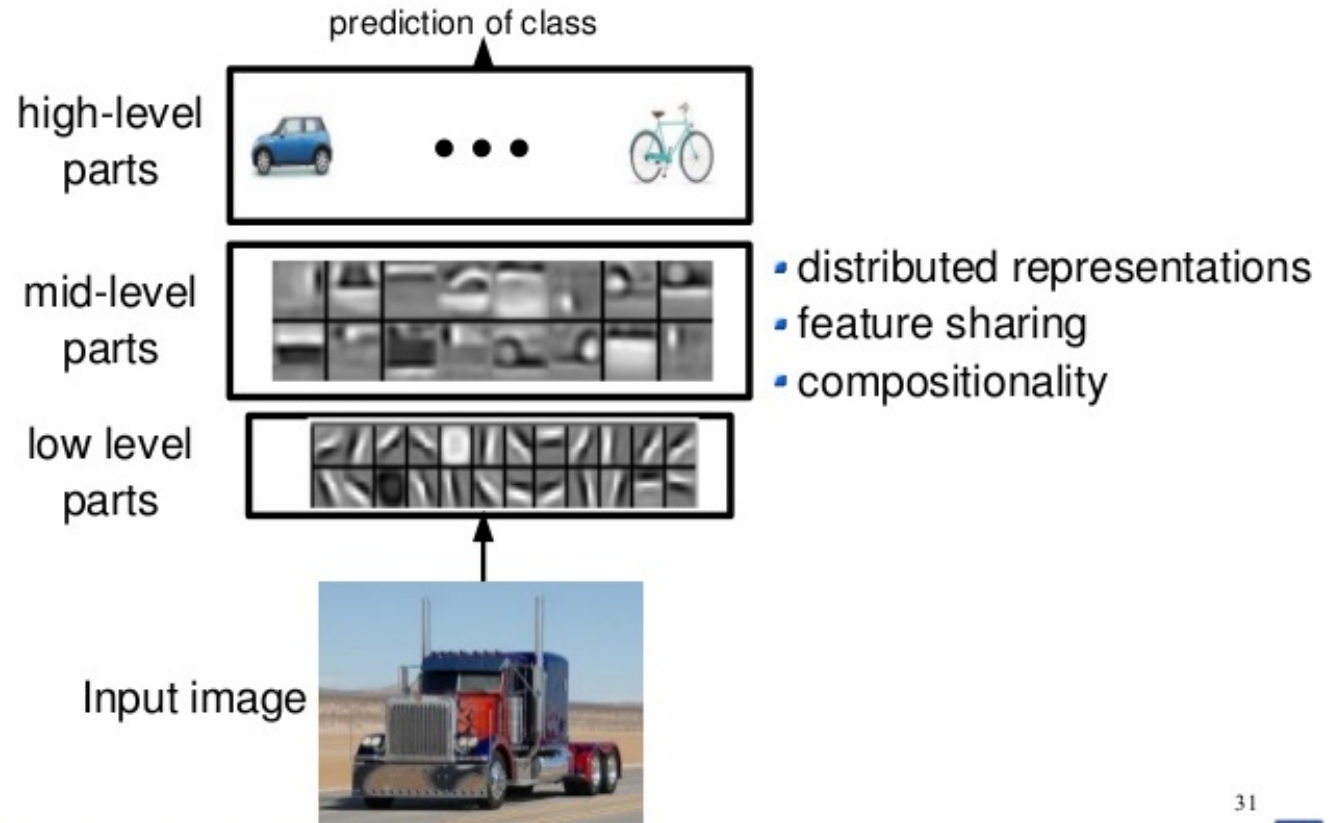
What would re-usable features look like?

- What if we could learn “local feature filters”
- Then on the next layer, learn how to combine them?



Convolutional NNs (CNNs/"Convnets")

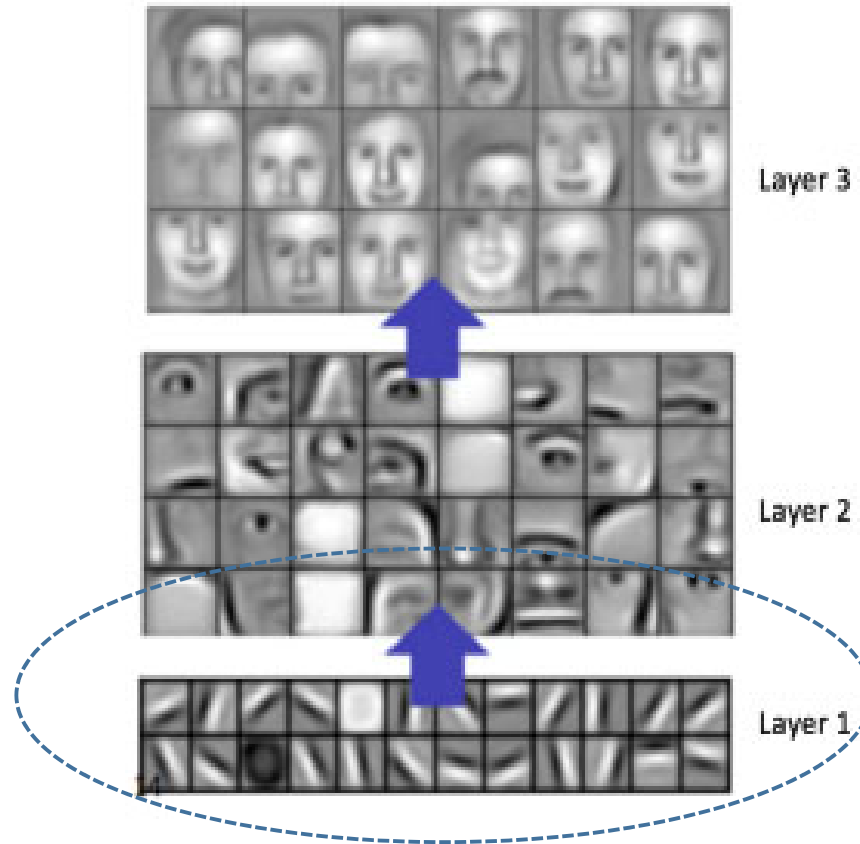
Can view CNNs as a way to construct **hierarchical features**, each of which get combined at the next level.



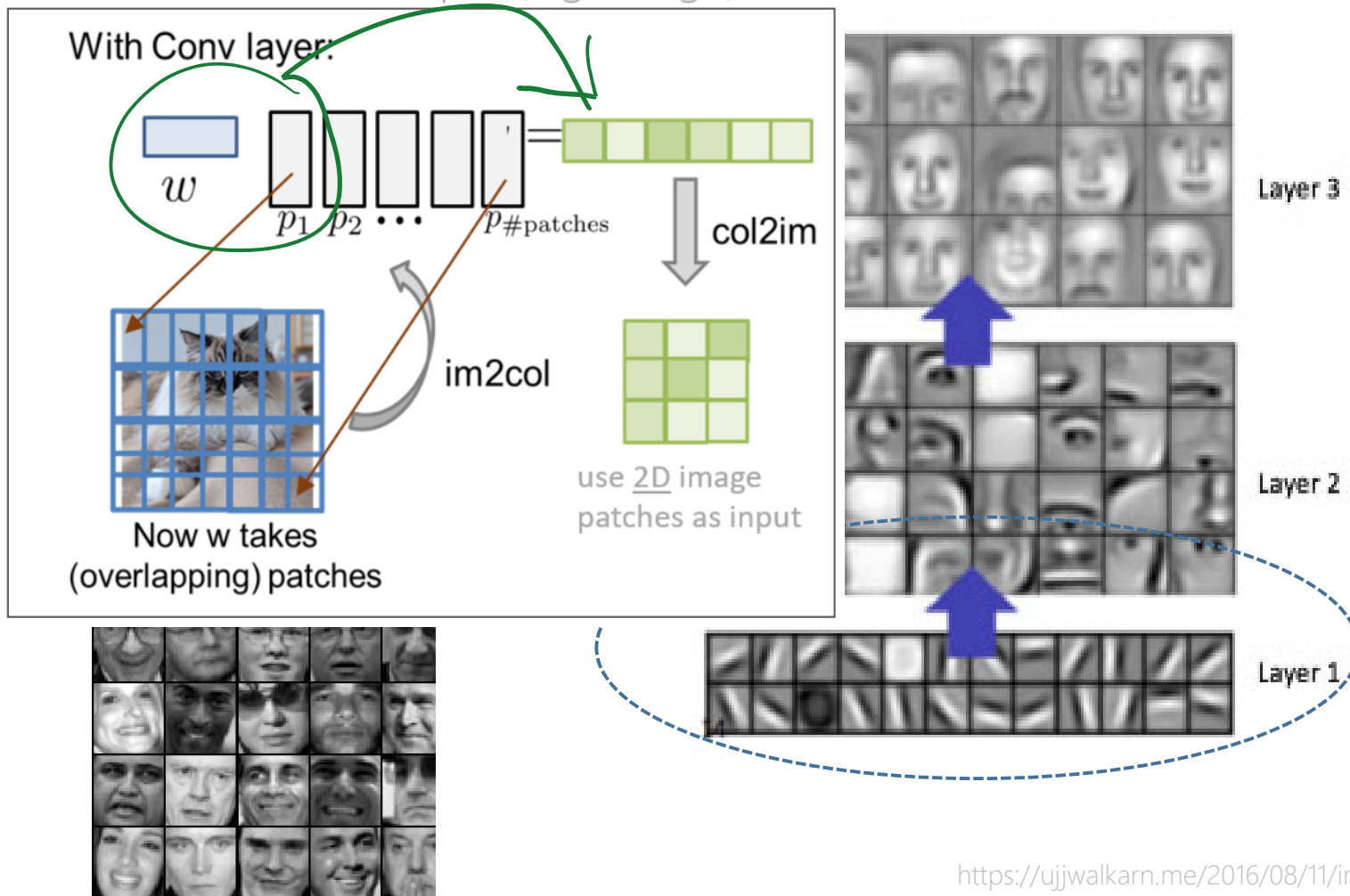
Lee et al. "Convolutional DBN's ..." ICML 2009

Convolutional NNs (CNNs/"Convnets")

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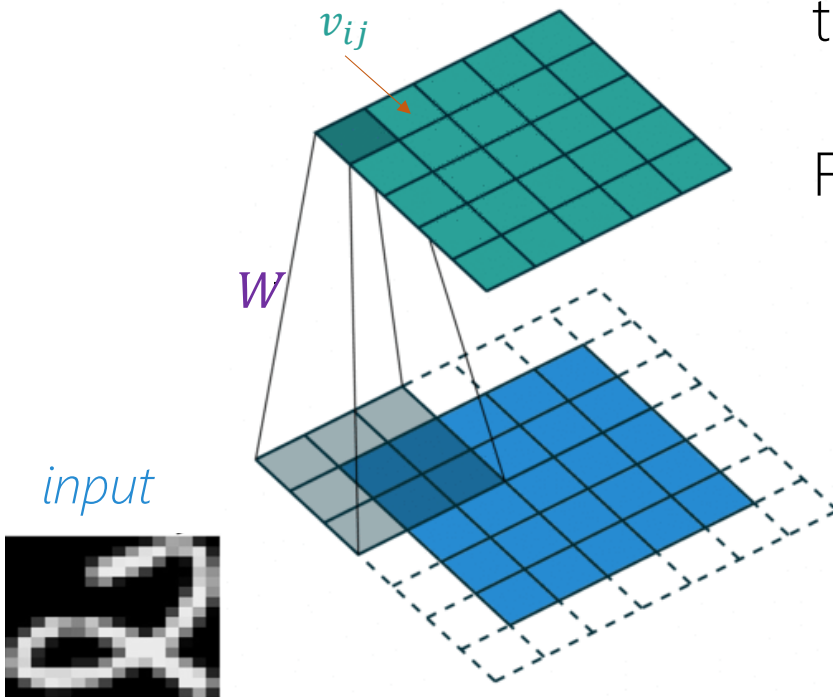


Convolutional NNs (CNNs/"Convnets")



"1D" Conv.

(2D) Convolution

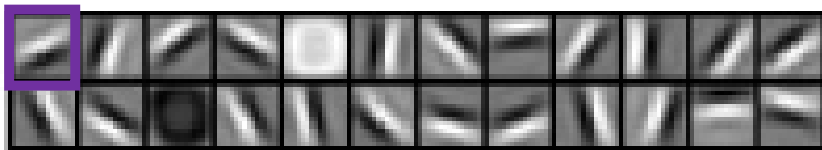


Convolve one learned “filter”, W with the input to get convolution output $\{v_{ij}\}$:

For each position, i, j :

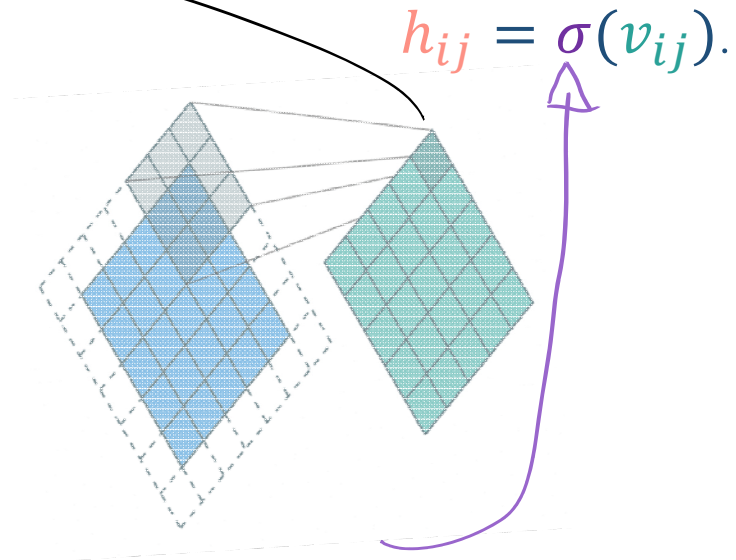
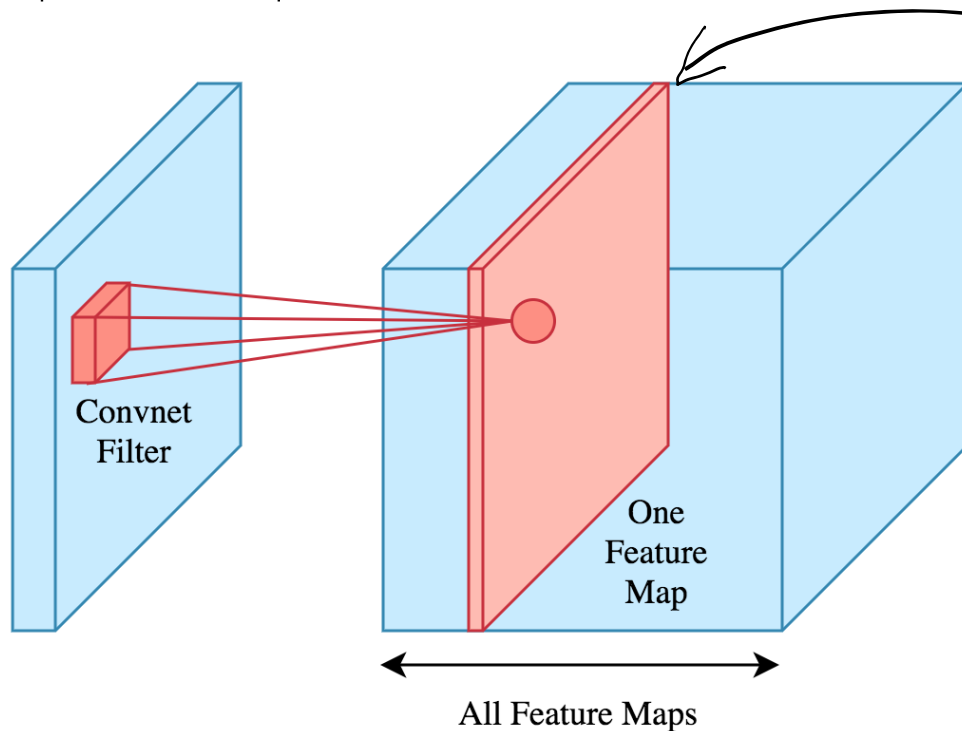
1. Element-wise product of W with image patch centered on i, j (e.g. 3×3).
2. Sum up the results to get one v_{ij} .

W called filter/template/kernel



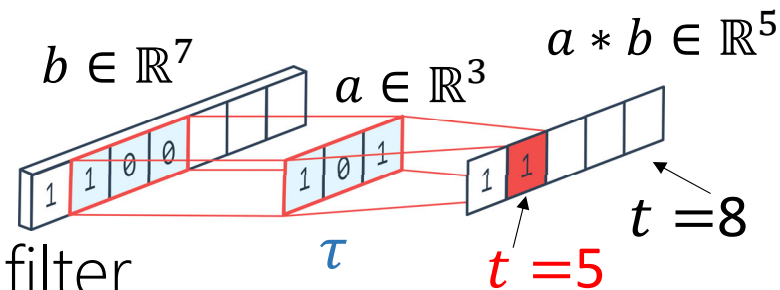
Convolutional NNs (CNNs/"Convnets")

- We will actually use multiple feature maps, $\{W_k\}_{k=1}^K$
- "Depth" of output "volume" is K :



Non-linearity to get hidden node in a hidden layer in CNN

Formally: 1D convolution



- For n-dim convolution, we use an n-dim filter.
- So 1D convolution has a 1D filter.

If a and b are two arrays,

$$(a * b) = (b * a)$$

$$a * (b * c) = (a * b) * c$$

t 'th element of the convolution $\rightarrow (a * b)_t = \sum_{\tau \in \{0,1,2,\dots\}} a_\tau b_{t-\tau} \leftarrow \text{arbitrary}$

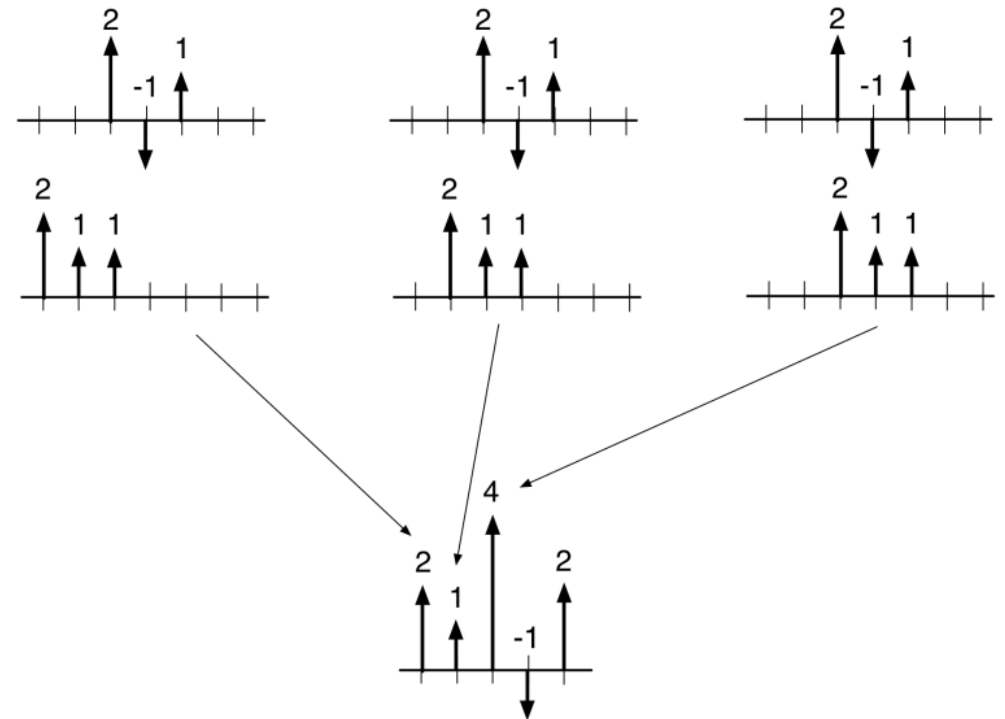
- τ is the index of the filter element ('-' means flip filter first)
- Invalid indices, e.g., $t = 1,2,3$ and $\tau = 3$, are boundaries; don't compute those t^{th} entries, or else *pad out* e.g. with zeros/mirroring input.
- No padding, size of output is $D - K + 1$ for D length input, K length filter.

Cross-correlation: $(a \otimes b)_t = \sum_{\tau} a_\tau b_{t+\tau}$

1D convolution

Method 1: flip-and-filter

$$(a * b)_t = \sum_{\tau} a_{\tau} b_{t-\tau} = \begin{array}{c} \uparrow 2 \\ | \\ \downarrow -1 \\ | \\ \uparrow 1 \end{array} * \begin{array}{c} \uparrow 1 \\ | \\ \uparrow 1 \\ | \\ \uparrow 2 \end{array} =$$



1D convolution

Method 2: translate-and-scale

$$(a * b)_t = \sum_{\tau} a_{\tau} b_{t-\tau} =$$

The diagram illustrates the convolution process using the 'translate-and-scale' method. It shows the following steps:

- Input 1:** A discrete signal with values $2, -1, 1$ at three time steps.
- Input 2:** A discrete signal with values $1, 1, 2$ at three time steps.
- Convolution:** The two signals are multiplied element-wise, resulting in $2, -1, 2$.
- Translation and Scaling:** The result is decomposed into three terms:
 - $2 \times$ (Input 2 shifted left by 1 step): $2, 2, 4$
 - $-1 \times$ (Input 2 shifted left by 0 steps): $-1, -1, -2$
 - $1 \times$ (Input 2 shifted left by 1 step): $1, 1, 2$
- Summation:** The three terms are summed to produce the final output signal: $2, 1, 4, -1, 2$.

1D convolution

Method 3

Convolution can also be viewed as matrix multiplication:

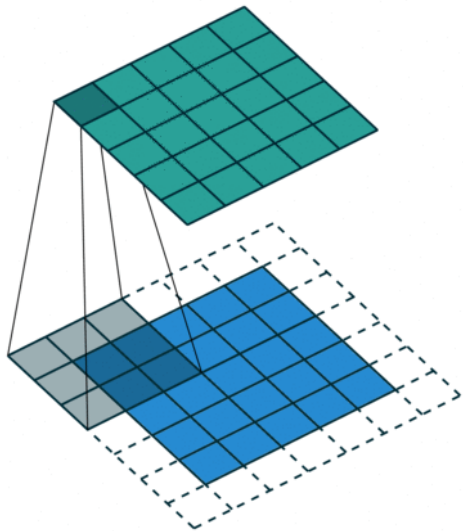
$$(a * b)_t = \sum_{\tau} a_{\tau} b_{t-\tau} = (2, -1, 1) * (1, 1, 2) = \overset{W_k}{\begin{pmatrix} 1 & & \\ 1 & 1 & \\ 2 & 1 & 1 \\ & 2 & 1 \\ & & 2 \end{pmatrix}} \begin{pmatrix} 2 \\ -1 \\ 1 \end{pmatrix} = \begin{array}{c} 2 \\ 1 \\ 4 \\ -1 \\ 2 \end{array}$$

W_k , has size 5×3 , which means it has 15 entries, yet there are only 3 parameters. Why Convnets to have relatively few parameters!

From 1D to 2D convolution

$$(A * B)_{ij} = \sum_s \sum_t A_{st} B_{i-s, j-t}$$

Method 1: Flip-and-Filter



1	3	1
0	-1	1
2	2	-1

 *

1	2
0	-1

1	3	1
0	-1	1
2	2	-1

 ×

-1	0
2	1

 →

1	5	7	2
0	-2	-4	1
2	6	4	-3
0	-2	-2	1

Note: In the original image, a blue box highlights the top 2x2 of the input grid, and a red box highlights the bottom row. Blue and red arrows point from these boxes to the corresponding rows in the output grid.

From 1D to 2D convolution

$$(A * B)_{ij} = \sum_s \sum_t A_{st} B_{i-s, j-t}$$

Method 2: Translate-and-Scale

$$\begin{array}{|c|c|c|} \hline 1 & 3 & 1 \\ \hline 0 & -1 & 1 \\ \hline 2 & 2 & -1 \\ \hline \end{array} * \begin{array}{|c|c|} \hline 1 & 2 \\ \hline 0 & -1 \\ \hline \end{array} = 1 \times \begin{array}{|c|c|c|c|} \hline 1 & 3 & 1 & \\ \hline 0 & -1 & 1 & \\ \hline 2 & 2 & -1 & \\ \hline & & & \\ \hline \end{array} + 2 \times \begin{array}{|c|c|c|c|} \hline & 1 & 3 & 1 \\ \hline & 0 & -1 & 1 \\ \hline & 2 & 2 & -1 \\ \hline & & & \\ \hline \end{array} = \begin{array}{|c|c|c|c|} \hline 1 & 5 & 7 & 2 \\ \hline 0 & -2 & -4 & 1 \\ \hline 2 & 6 & 4 & -3 \\ \hline 0 & -2 & -2 & 1 \\ \hline \end{array} + -1 \times \begin{array}{|c|c|c|c|} \hline & & & \\ \hline & 1 & 3 & 1 \\ \hline & 0 & -1 & 1 \\ \hline & 2 & 2 & -1 \\ \hline \end{array}$$

2D convolution

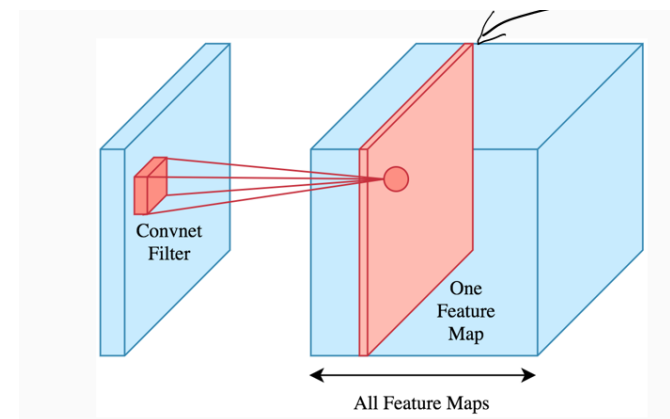
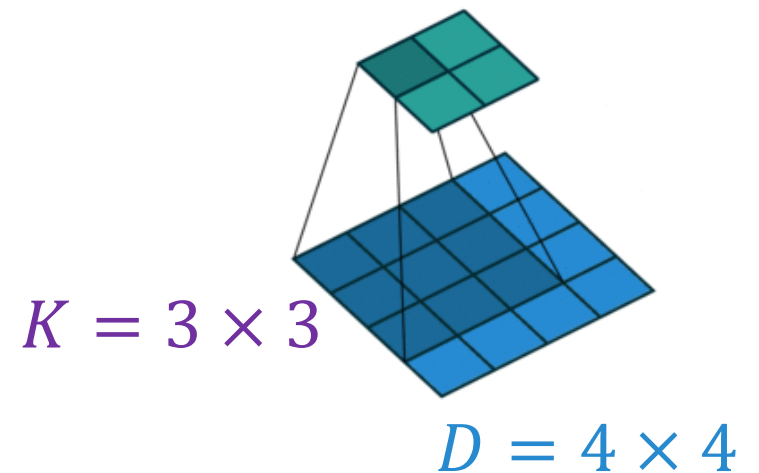
- Image is $D \times D$.
- N filters each of size $K \times K$.
- No zero-padding.

Then output from one filter has size:
 $(D - K + 1) \times (D - K + 1)$

For all N filters,
 $N \times (D - K + 1) \times (D - K + 1)$

https://github.com/vdumoulin/conv_arithmetic

convolution is 2×2

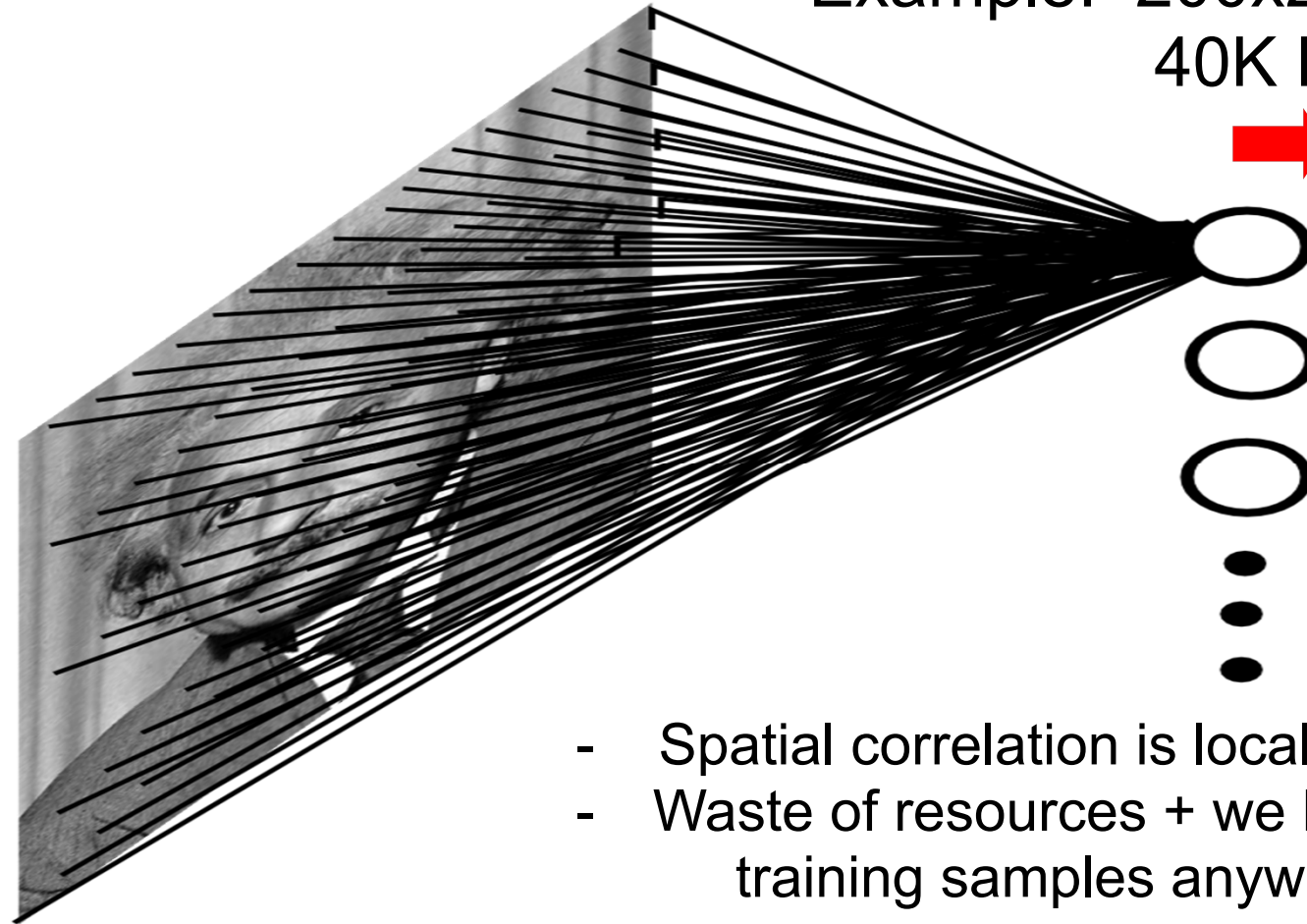


Fully-connected layer (no shared features)

Example: 200x200 image

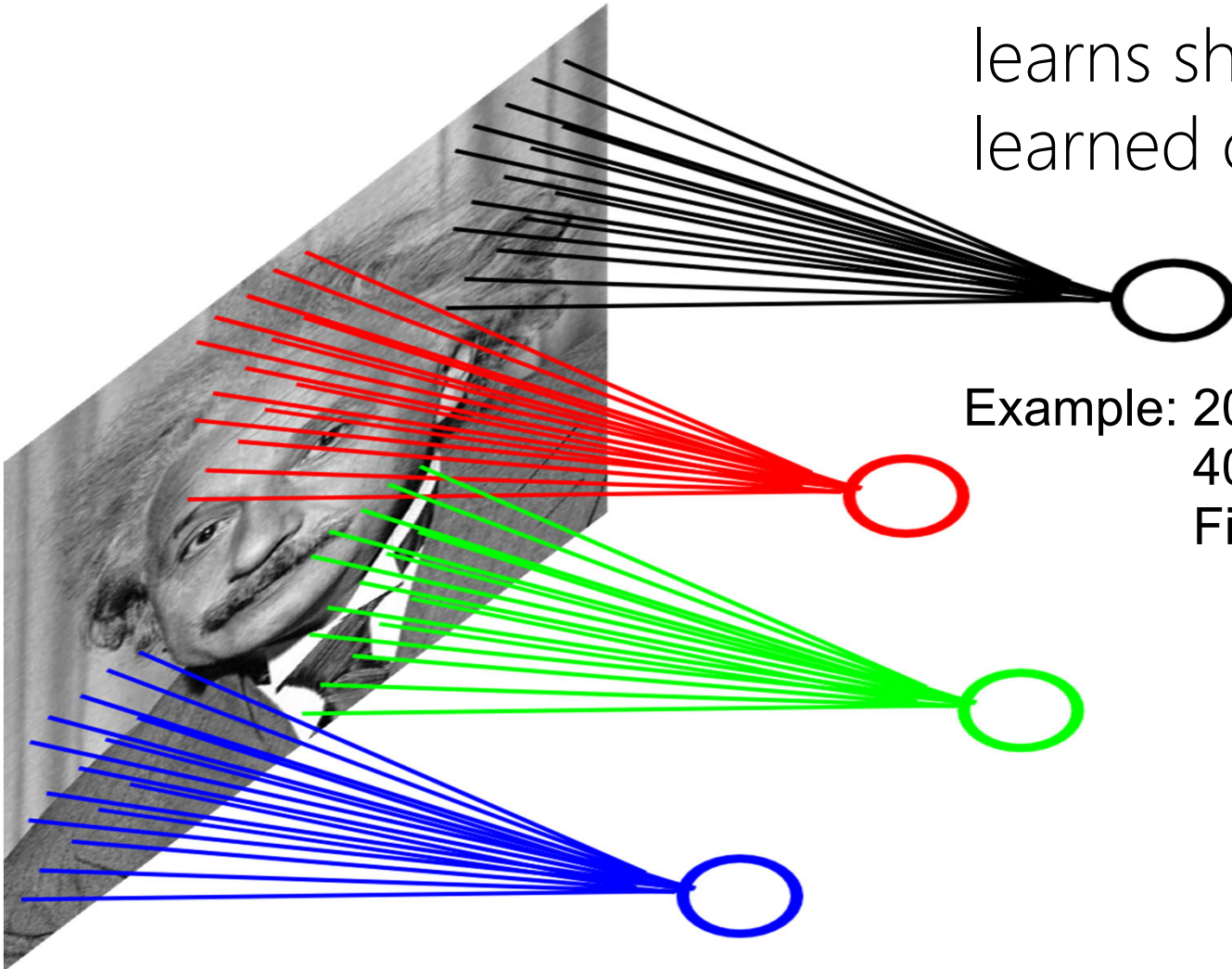
40K hidden units

→ ~2B parameters!!!



- Spatial correlation is local
- Waste of resources + we have not enough training samples anyway..

Convolutional layer



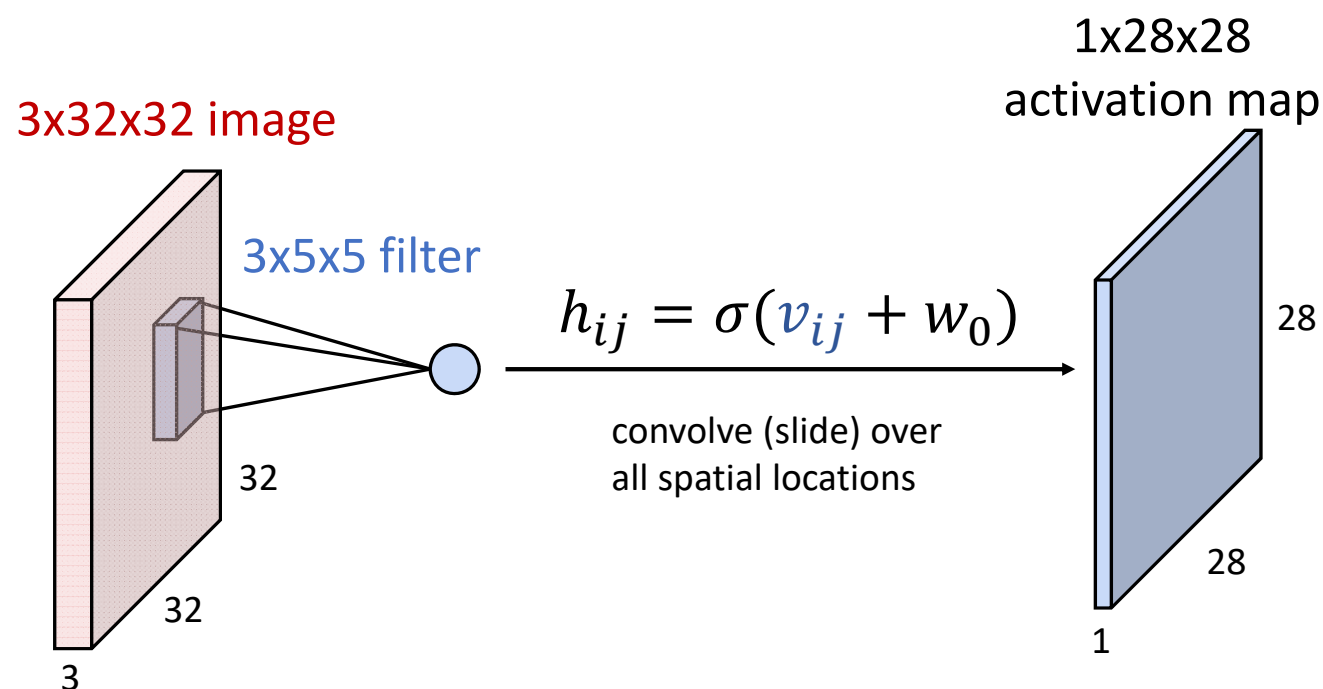
learns shared features via learned convolution kernels

Example: 200x200 image
40K hidden units
Filter size: 10x10

4M parameters

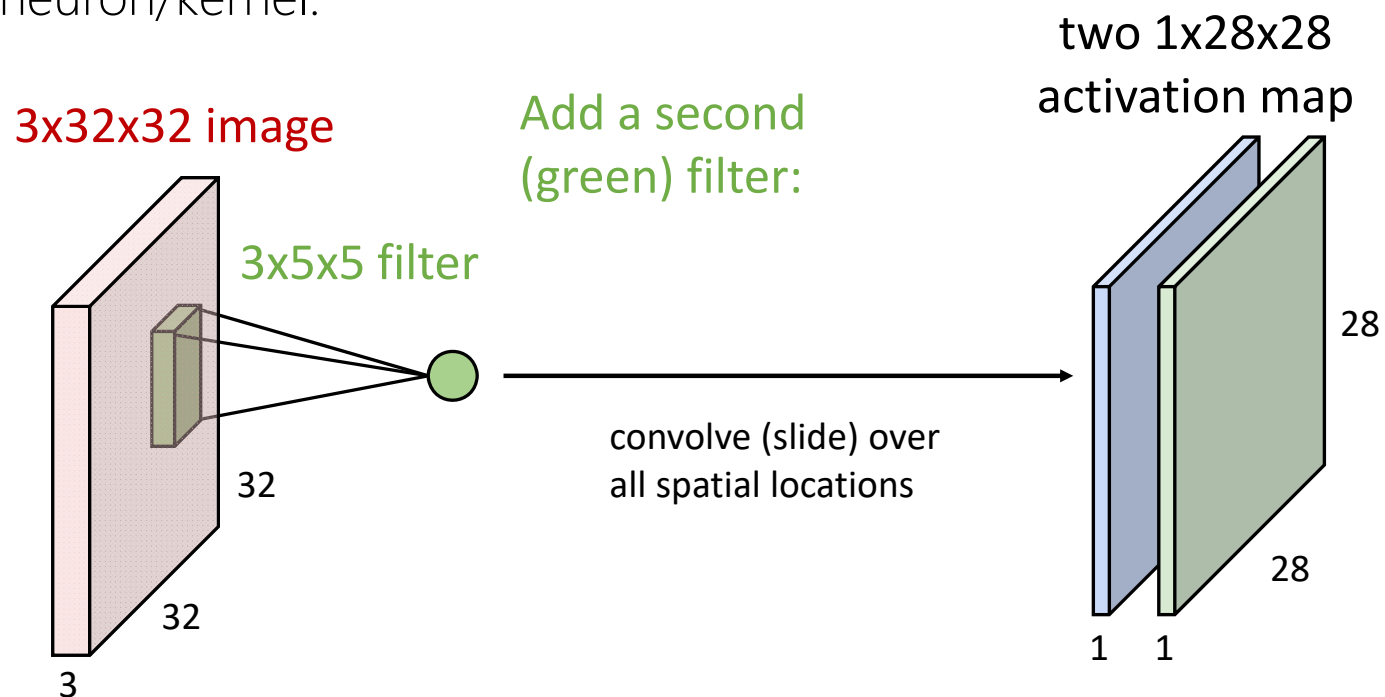
Convolution Layer

One "neuron"/kernel that "looks at" 5x5 region and outputs a sheet of *activation map*

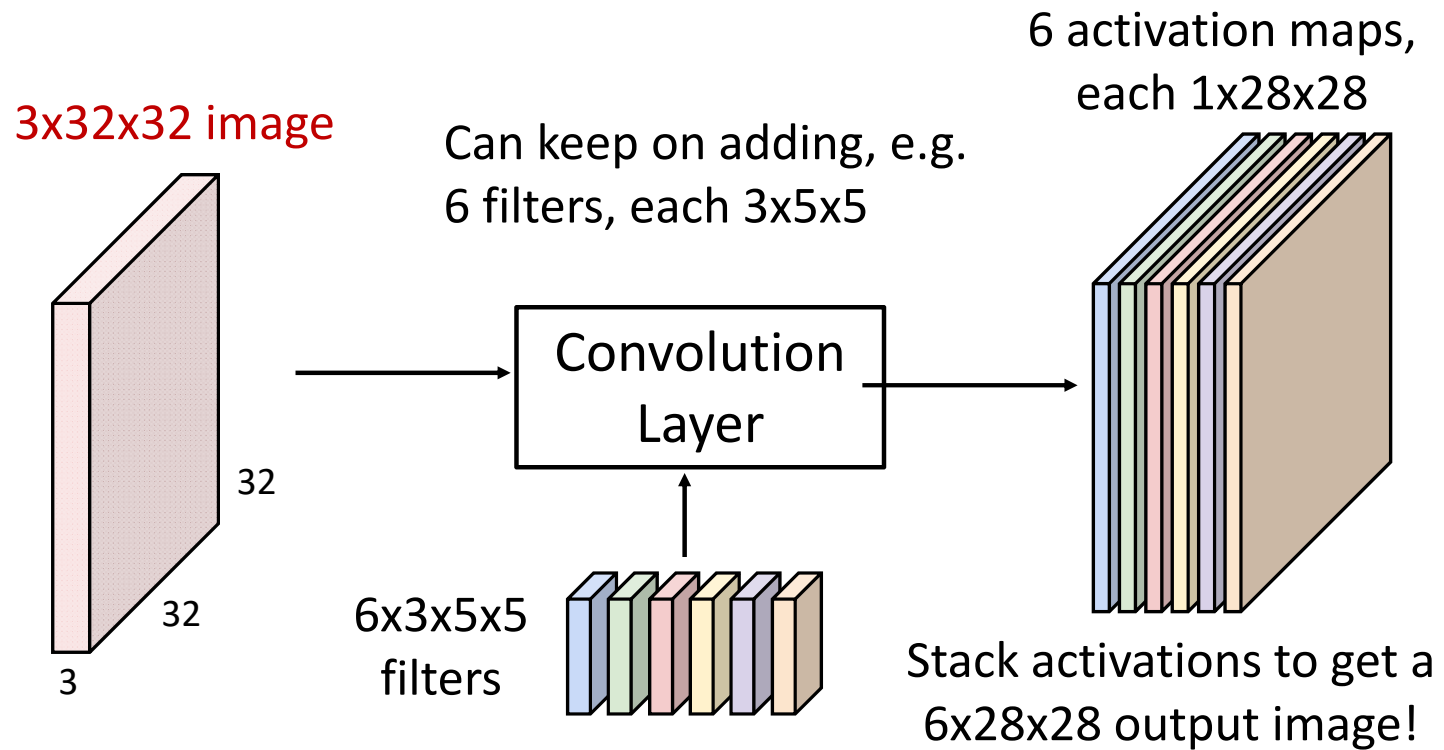


Convolution Layer

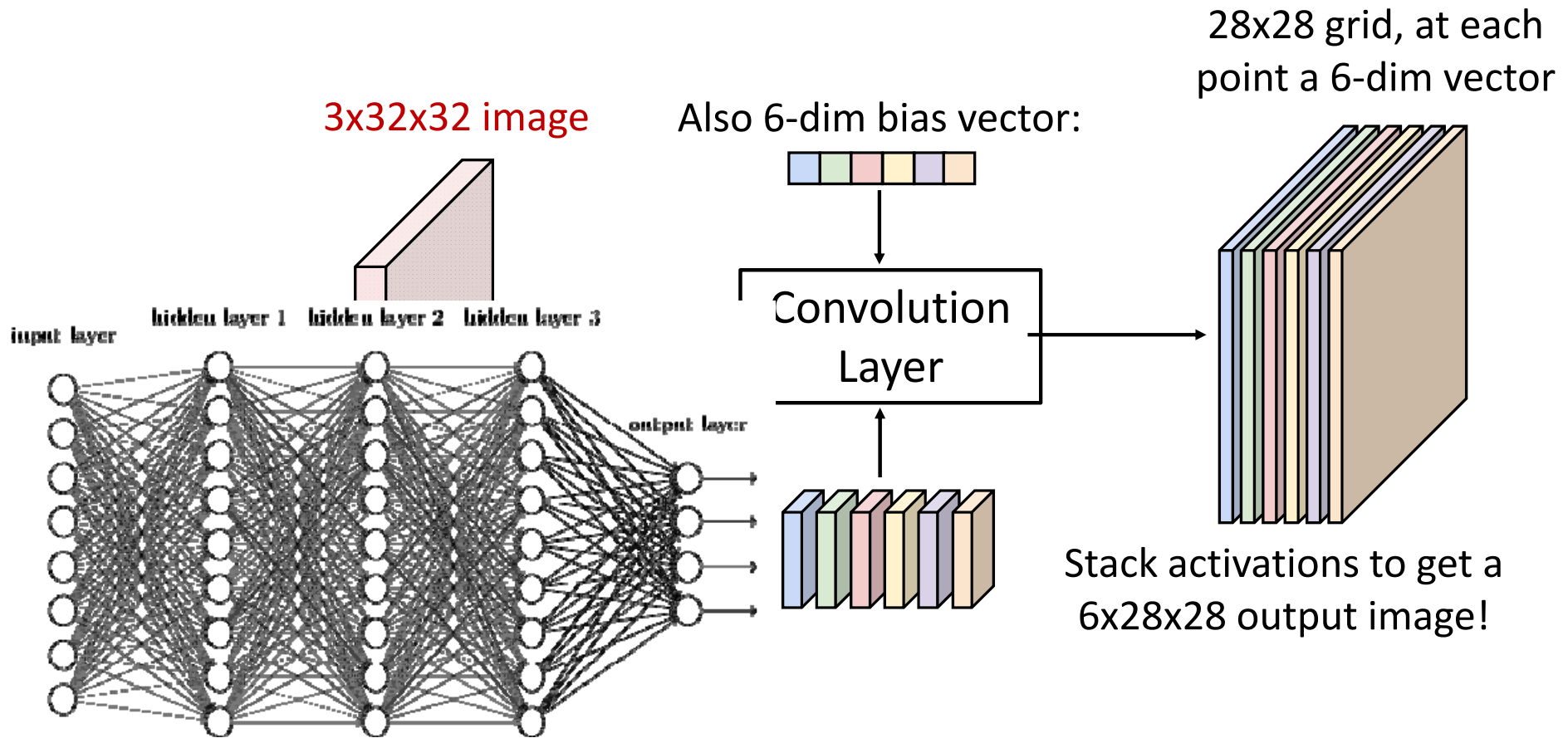
Add a second neuron/kernel.



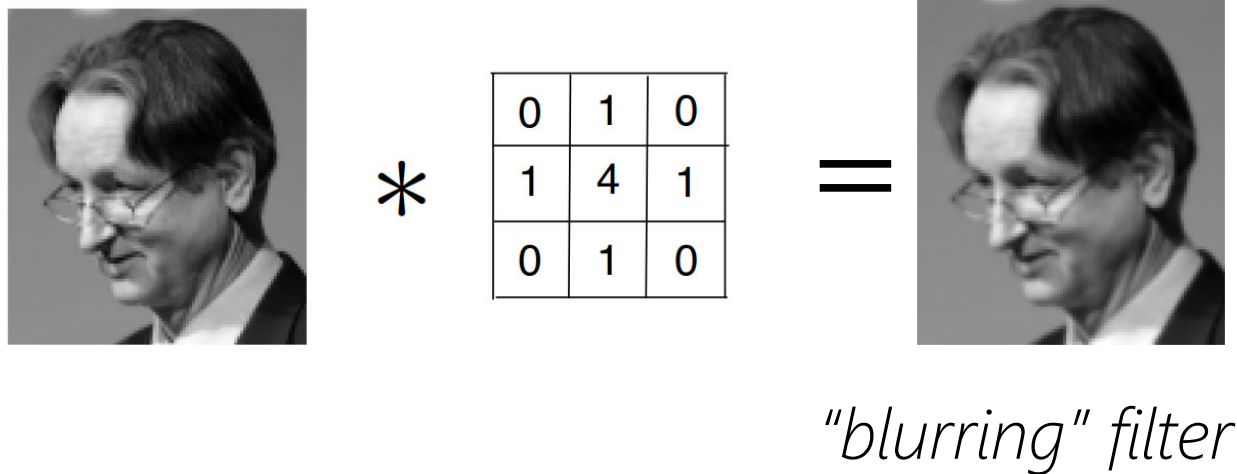
Convolution Layer



Convolution Layer



Intuition of 2D convolution kernels



Intuition of 2D convolution kernels



*

1	0	-1
2	0	-2
1	0	-1

=



"oriented edges"

Intuition of 2D convolution kernels



*

0	-1	0
-1	4	-1
0	-1	0

=



"sharpen"

Intuition of 2D convolution kernels



*

w_{11}	w_{12}	w_{13}
w_{21}	w_{22}	w_{23}
w_{31}	w_{32}	w_{33}

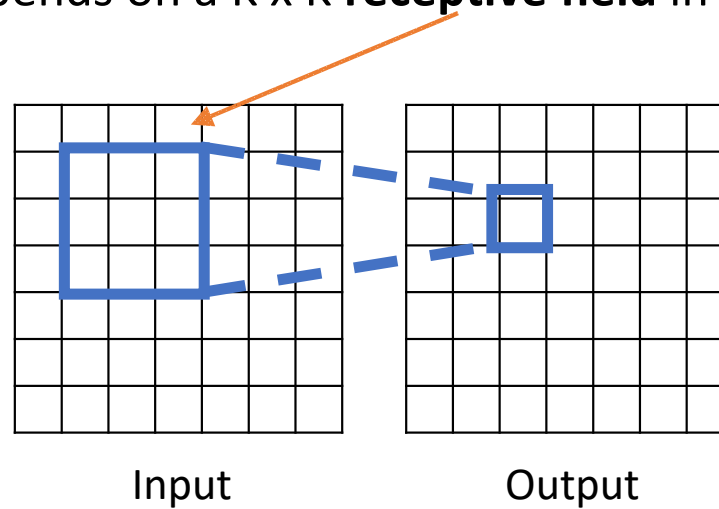
=



Gradient descent on loss will decide.

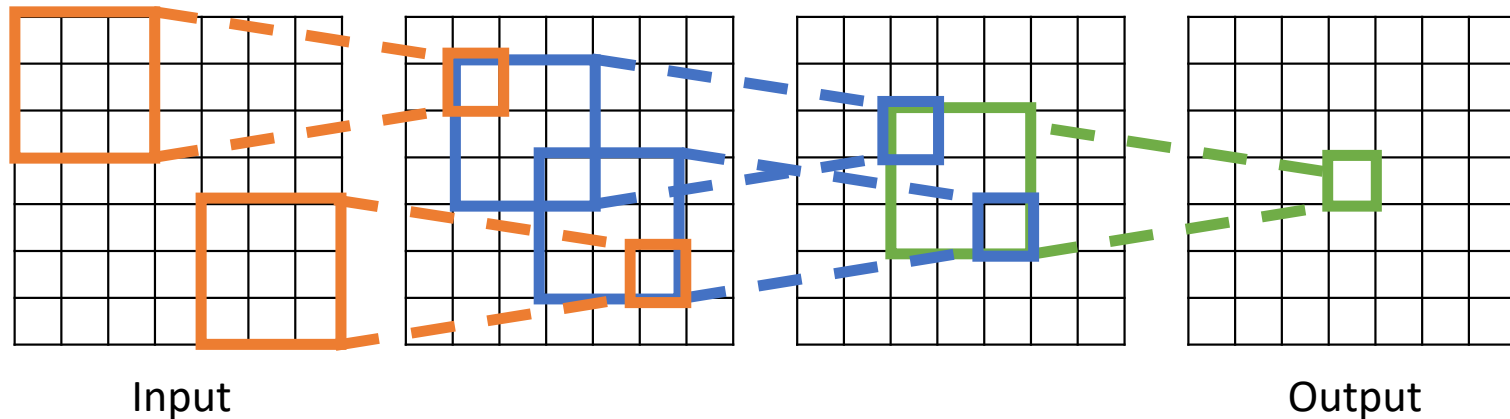
Receptive Fields

For convolution with kernel size K , each element in the output depends on a $K \times K$ **receptive field** in the input



Receptive Fields

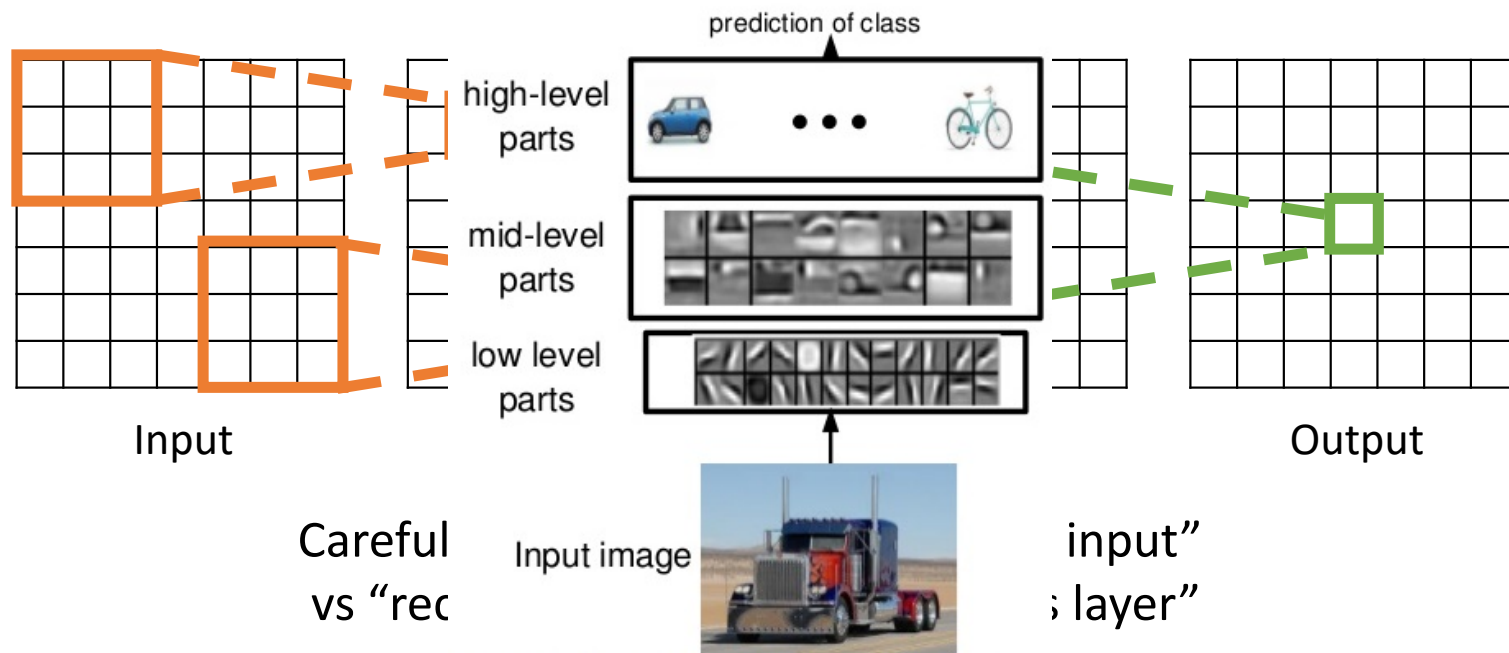
Each successive convolution adds $K - 1$ to the receptive field size
With L layers the receptive field size is $1 + L \times (K - 1)$



Careful – “receptive field wrt to the input”
vs “receptive field wrt the previous layer”

Receptive Fields

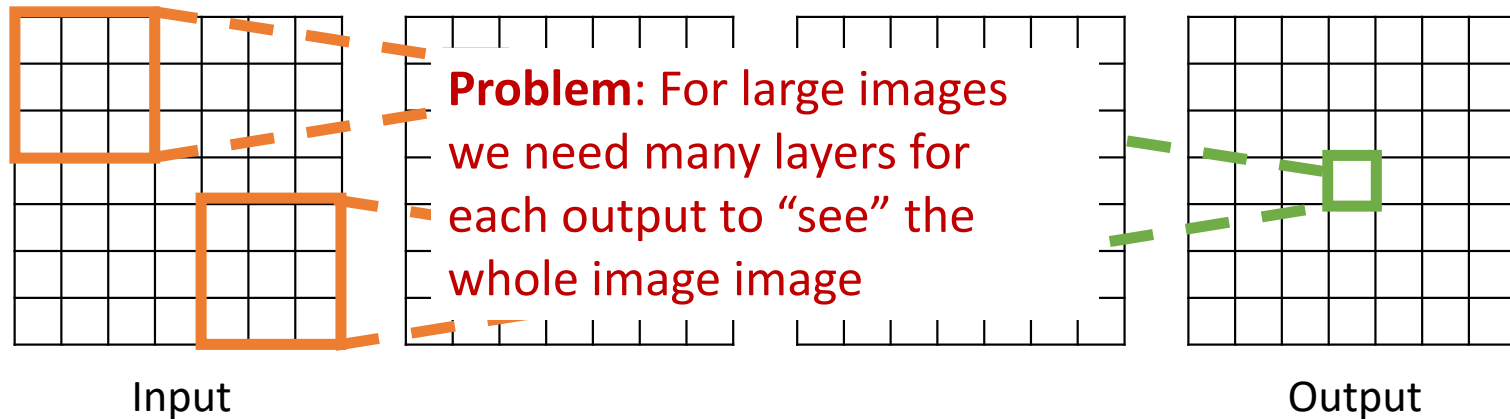
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Lee et al. "Convolutional DBN's ..." ICML 2009

Receptive Fields

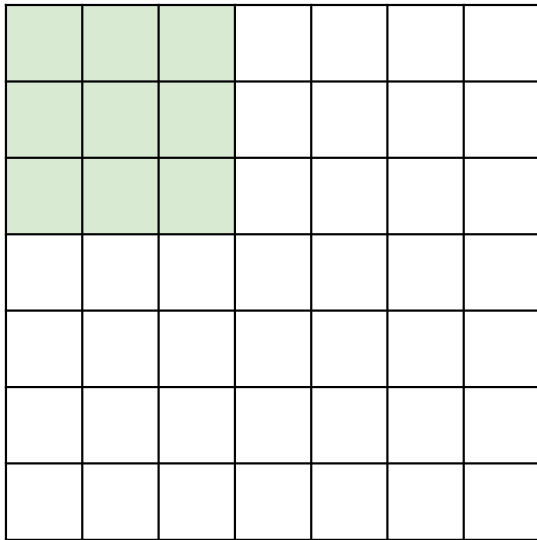
Each successive convolution adds $K - 1$ to the receptive field size
With L layers the receptive field size is $1 + L \times (K - 1)$



Solution: downsample inside the network

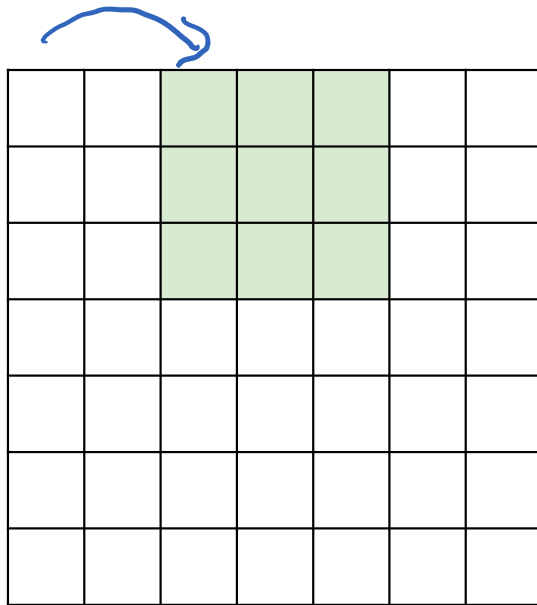
1. “Strided” convolution
2. Pooling

1. Strided Convolution



Input: 7x7
Filter: 3x3
Stride: 2

1. Strided Convolution

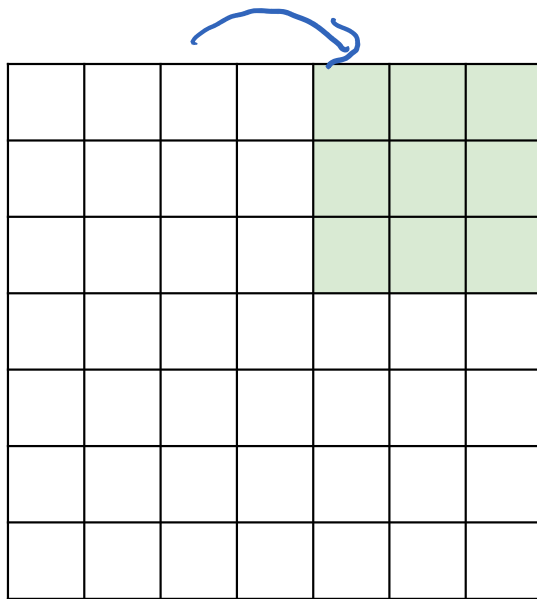


Input: 7x7

Filter: 3x3

Stride: 2

1. Strided Convolution



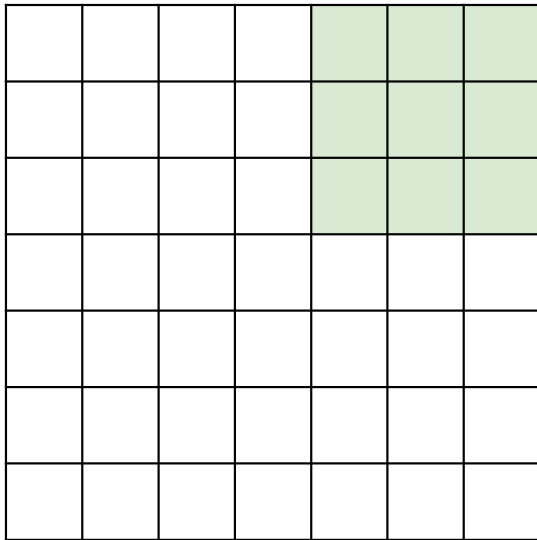
Input: 7x7

Filter: 3x3

Stride: 2

Output: 3x3

1. Strided Convolution



Input: 7x7

Filter: 3x3

Stride: 2

Output: 3x3

In general:

Input: W

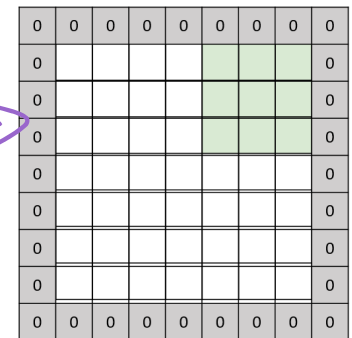
Filter: K

Padding: P

Stride: S

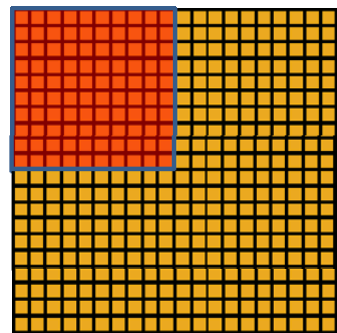
Output dimension: $(W - K + 2P) / S + 1$

(one dimension of the output square)

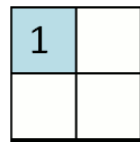


2. Pooling layers downsample its inputs

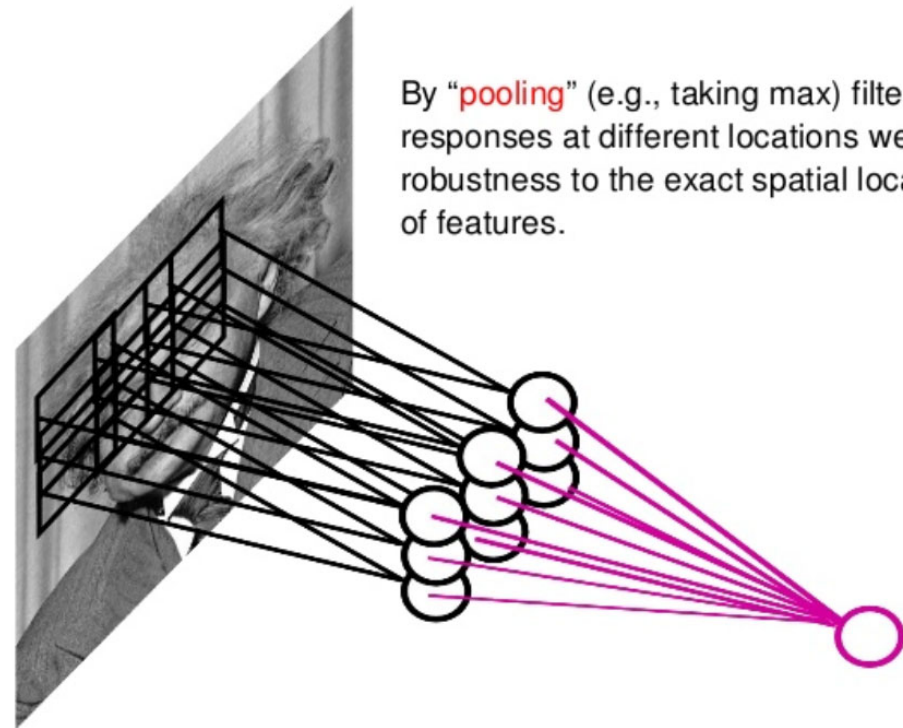
Also adds some local translational *invariance* (by summing/averaging):



Convolved
feature

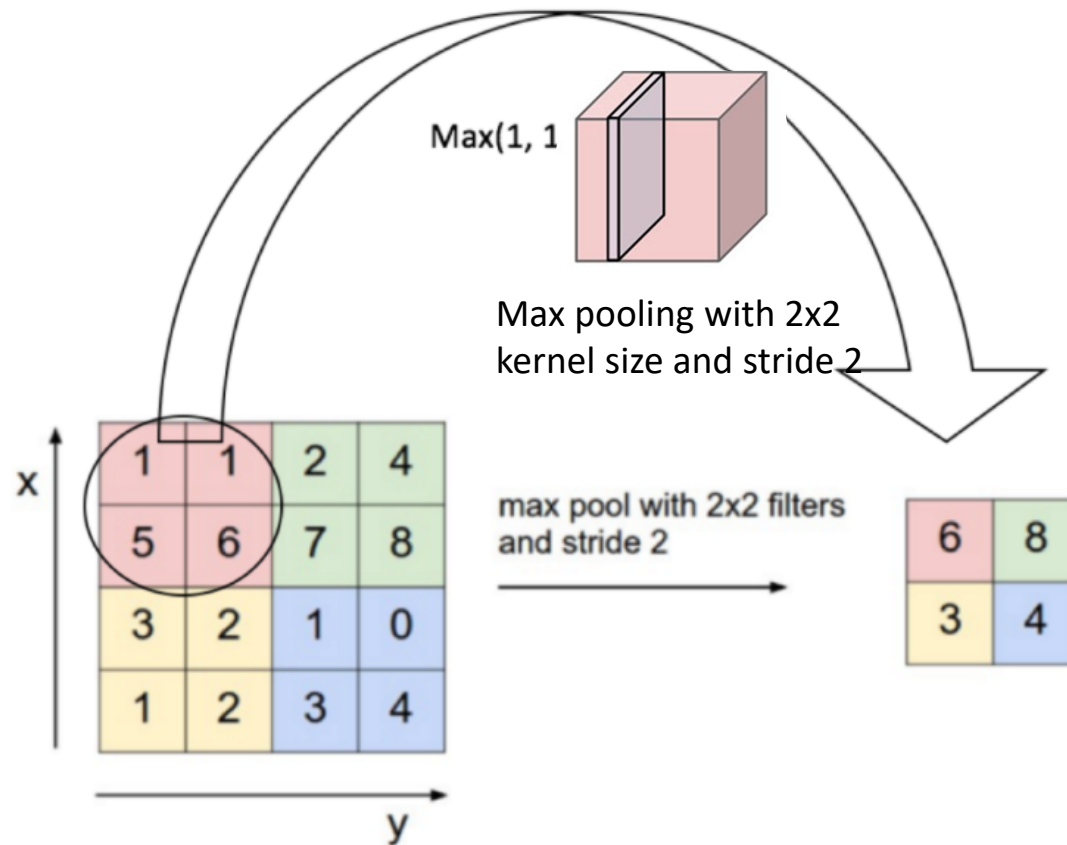


Pooled
feature



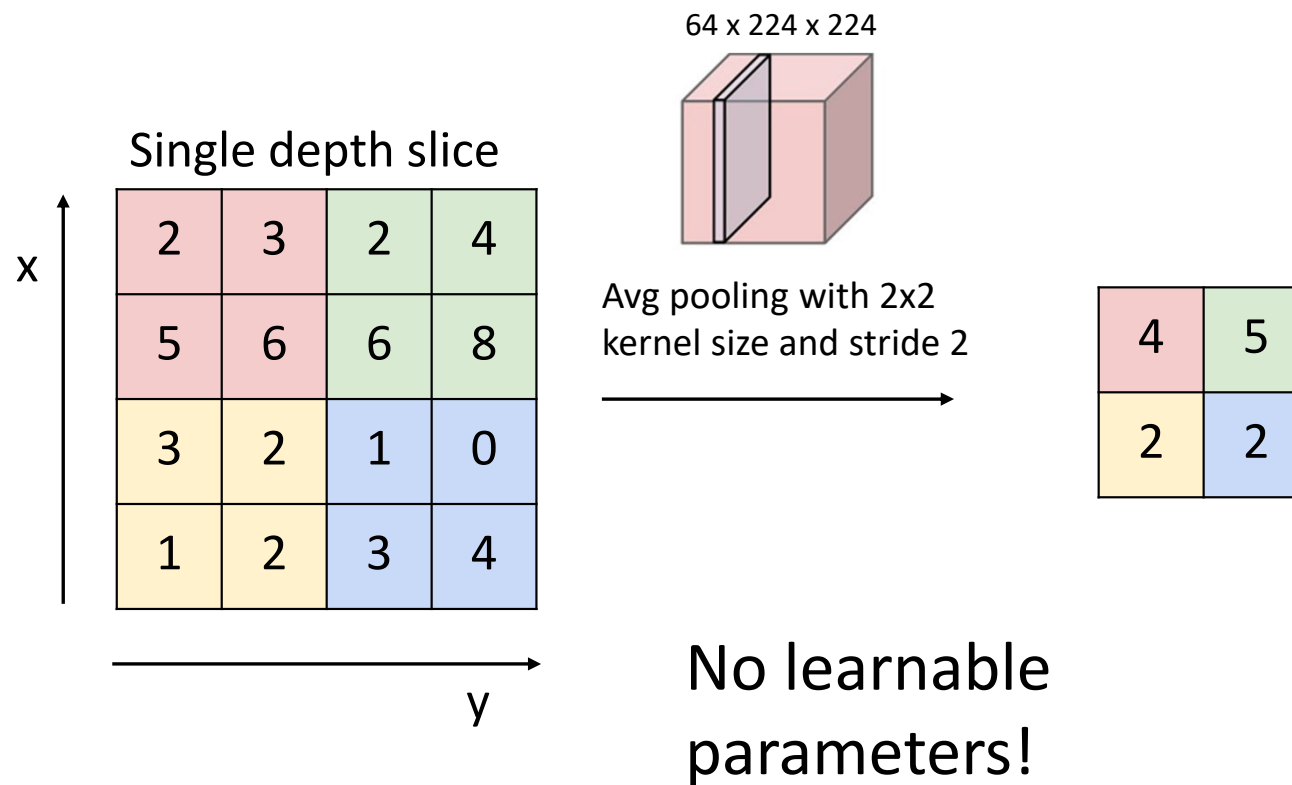
By “pooling” (e.g., taking max) filter responses at different locations we gain robustness to the exact spatial location of features.

2. Max pooling

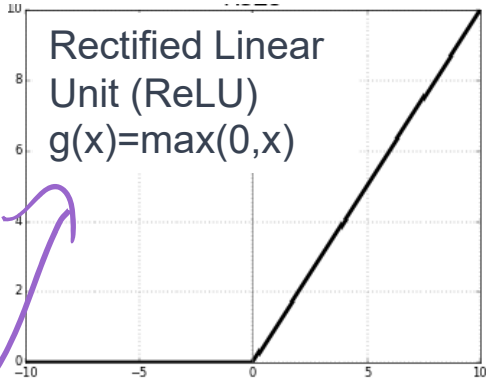
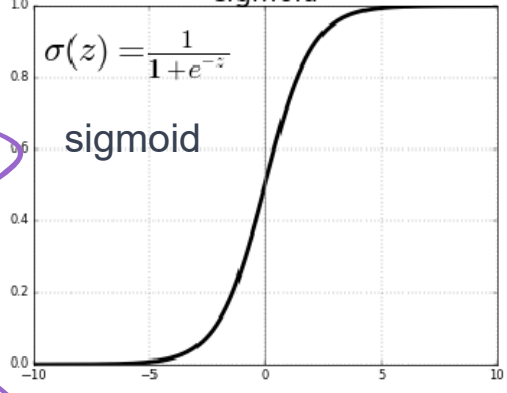
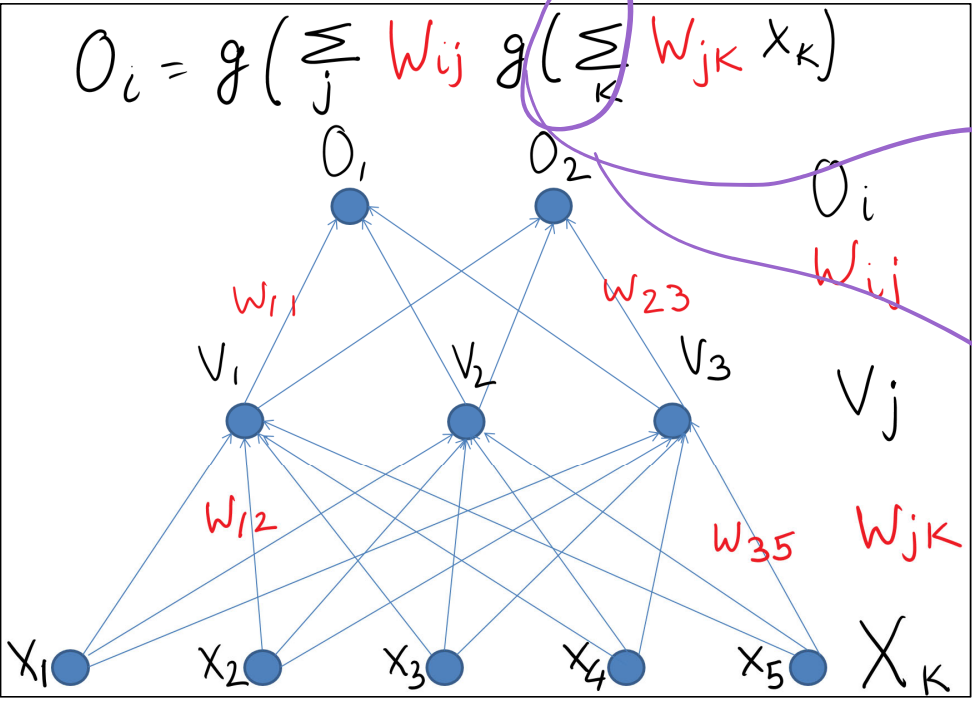


No learnable parameters!

2. Average pooling

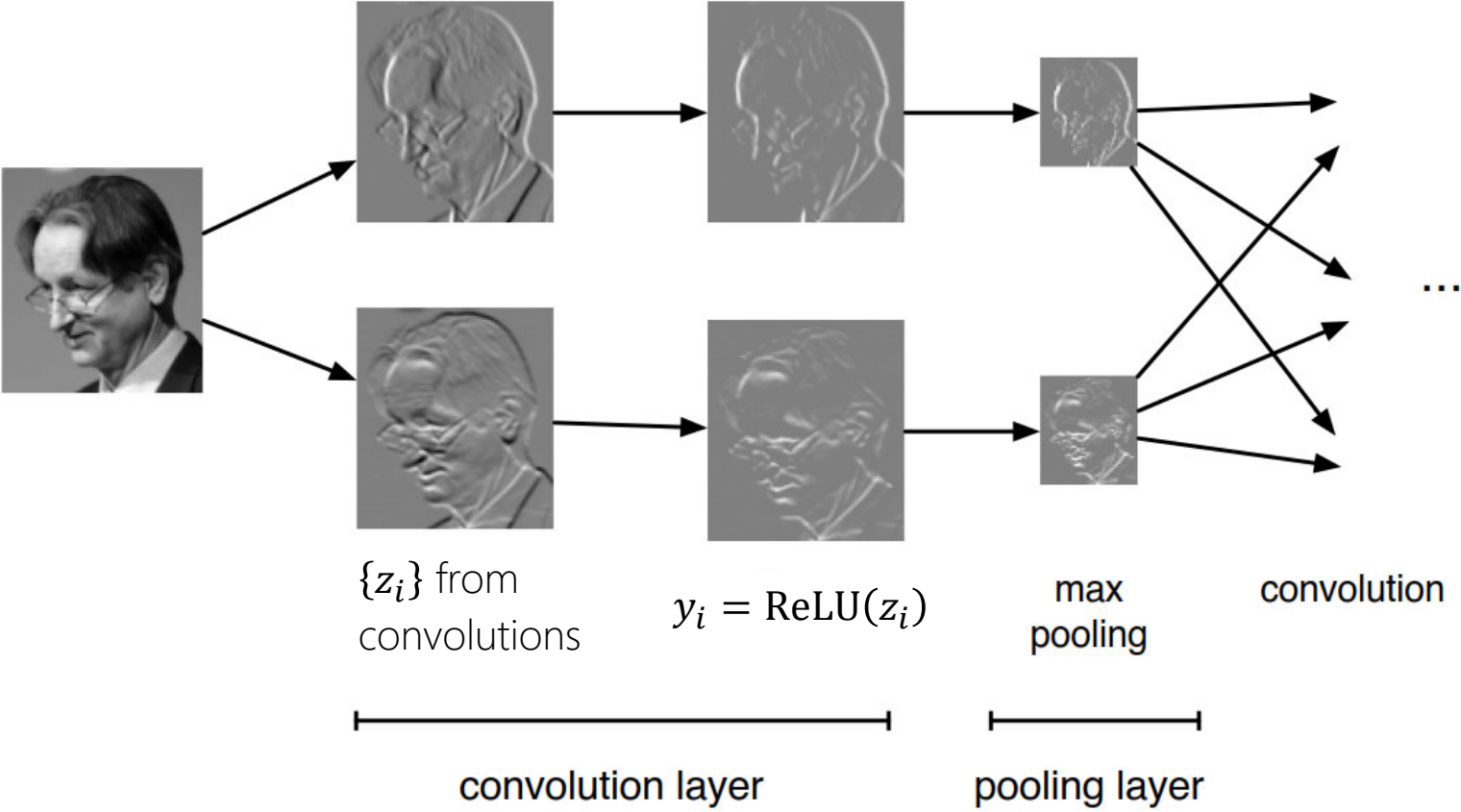


Side note: sigmoid vs ReLU non-linearity in NNs



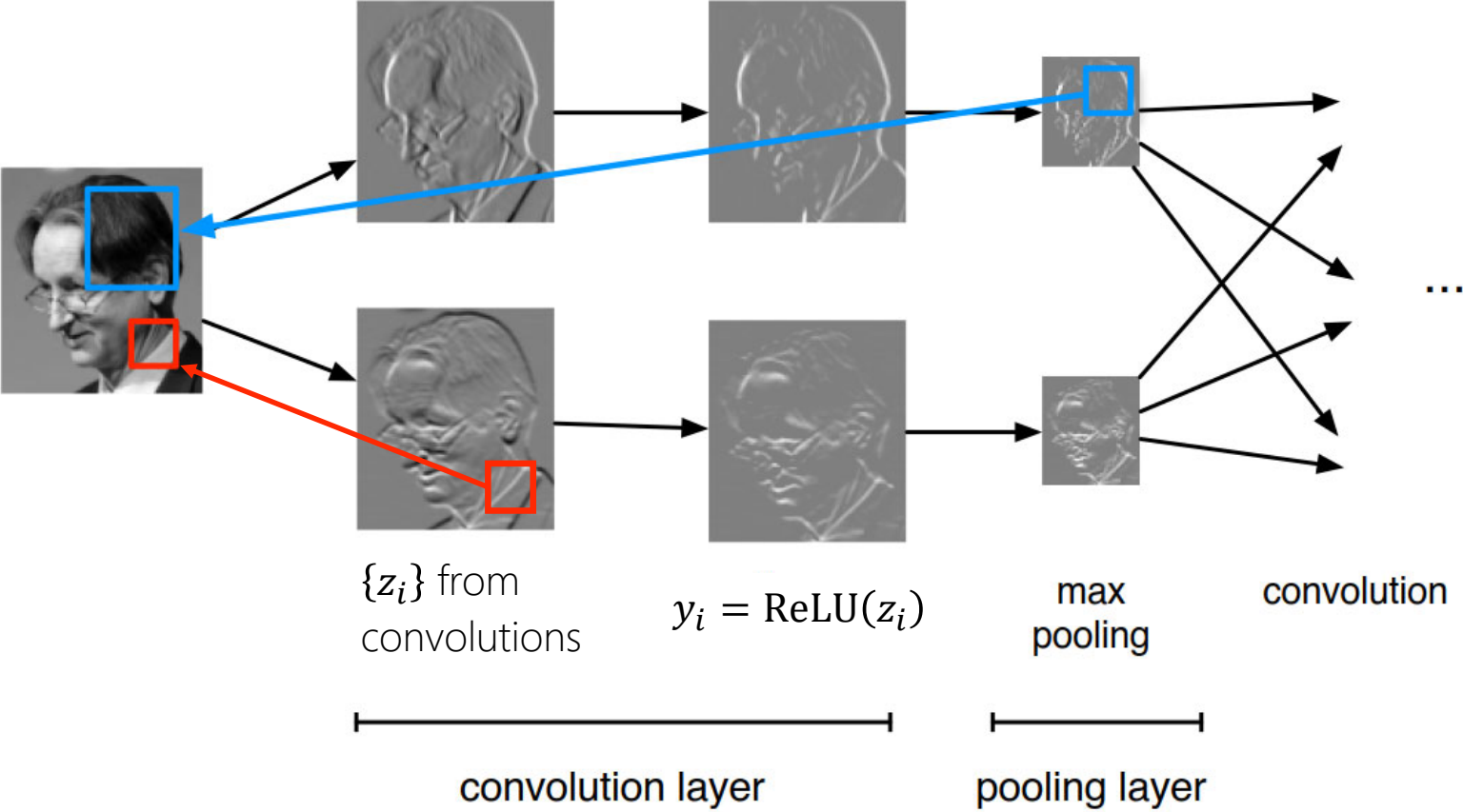
- ReLU:
1. Gradient doesn't die in one direction.
 2. More efficient to compute.
 3. Easier to get exactly zero activations: sparsity.

Putting it altogether! ConvNets: conv + ReLU + pooling

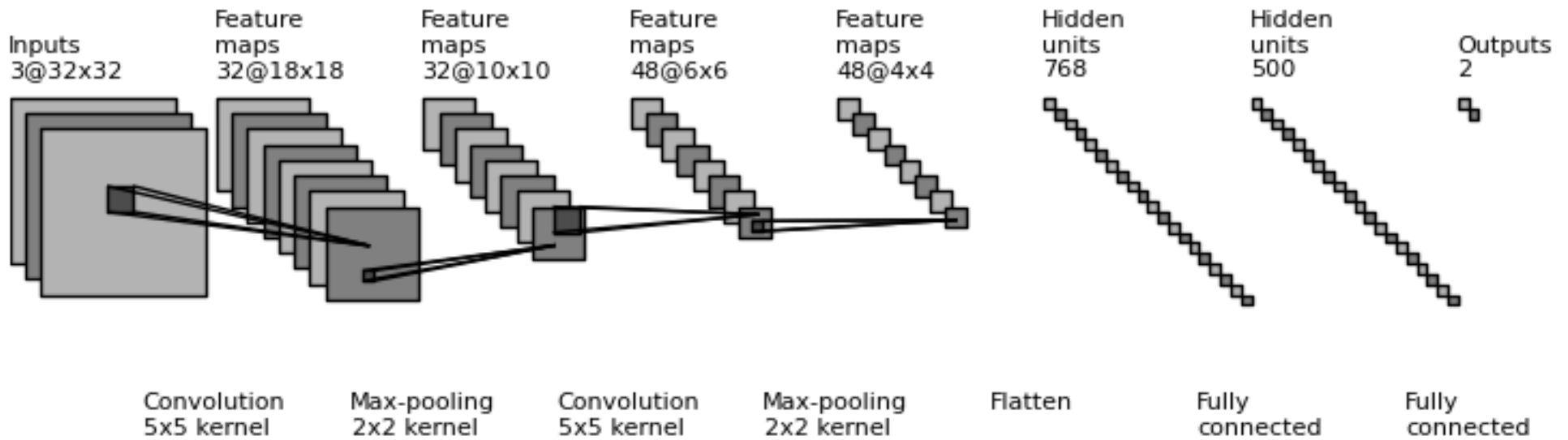


Putting it altogether! ConvNets: conv + ReLU + pooling

Receptive field increases



Example CNN architecture



Training CNNs

Gradient descent with back-propagation algorithm.

1. Goal is still MLE/ maximize cross-entropy.
2. Shared weights (via one convolution filter) → sum over gradient for each use of one filter.
3. Max-pooling → gradient only gets back-propagated through the neuron that “won” the max pool—technically this is a “sub-gradient”.