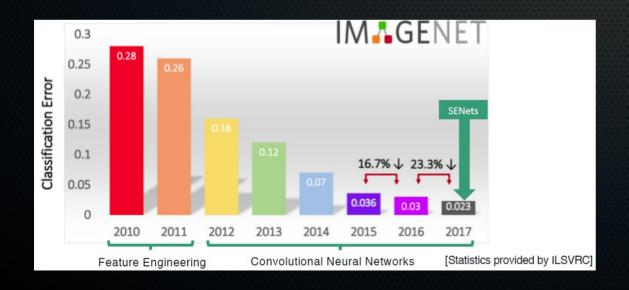
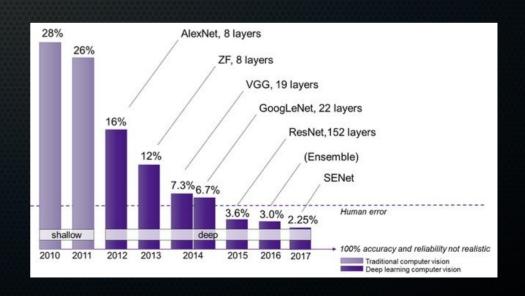
#### Convolutional neural networks

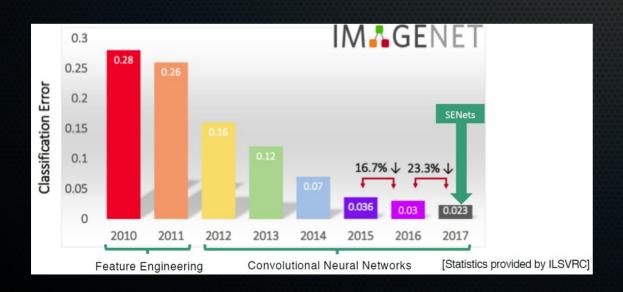
 These dense neural networks are not what made the huge strides in deep learning over the last few years.

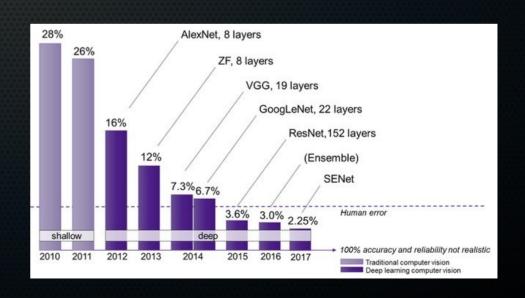




#### Convolutional neural networks

- These dense neural networks are not what made the huge strides in deep learning over the last few years.
- Instead, those are deep convolutional neural networks





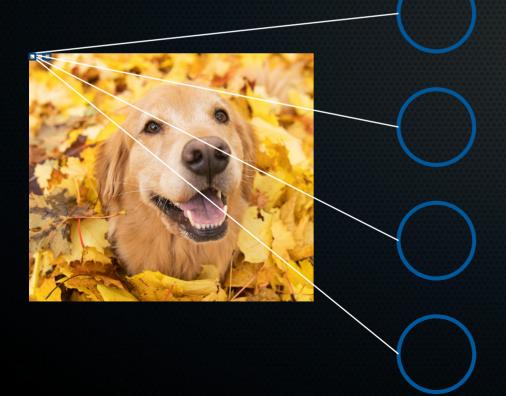
Let's look at an image



Let's look at an image

In a dense architecture, every pixel value is connected to

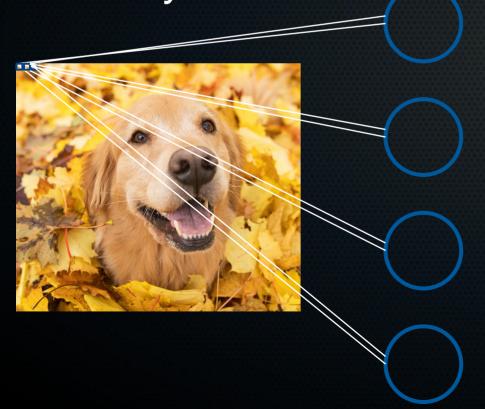
every neuron.



Let's look at an image

In a dense architecture, every pixel value is connected to

every neuron.



Etc.

- Let's look at an image
- In a dense architecture, every pixel value is connected to every neuron.
- This gives problems:
  - You get an *insane* amount of parameters to optimise. 250\*250 pixels \* 20 input neurons = 1,250,020 weights and biases. You can forget about any sort of findable or achievable (global) optimum.
  - There is no locality: if you want your network to know whether or not there is a dog in an image, all these parameters must be optimised so that you can recognise the dog anywhere.



This is madness!



The answer: convolution. Let's look at a 1D example!



When is the heart beating?

The answer: convolution. Let's look at a 1D example!



When is the heart beating?

The answer: convolution. Let's look at a 1D example!



label = 
$$y = [0 \ 0 \ 0 \ 1 \ 1 \ 1 \ 0 \ 0 \ 0 \ 0]$$

Signal can be at different positions in the sequence:



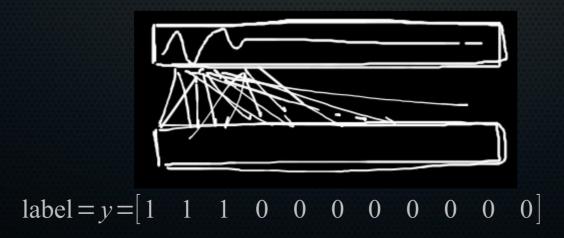




Signal can be at different positions in the sequence:



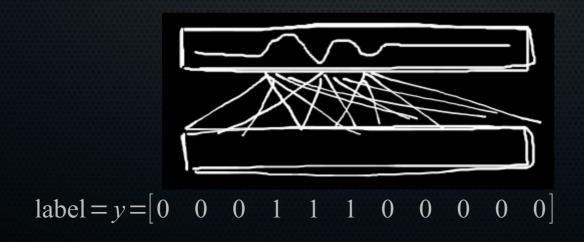
 Dense network needs to optimise such that different weights somehow cause the network to output 1 for different positions of the signal:



Signal can be at different positions in the sequence:



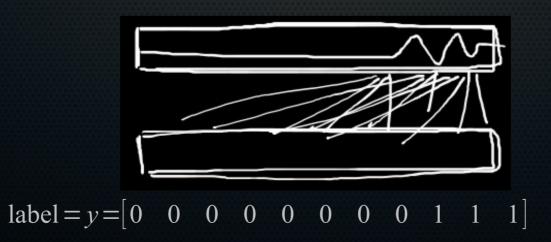
 Dense network needs to optimise such that different weights somehow cause the network to output 1 for different positions of the signal:

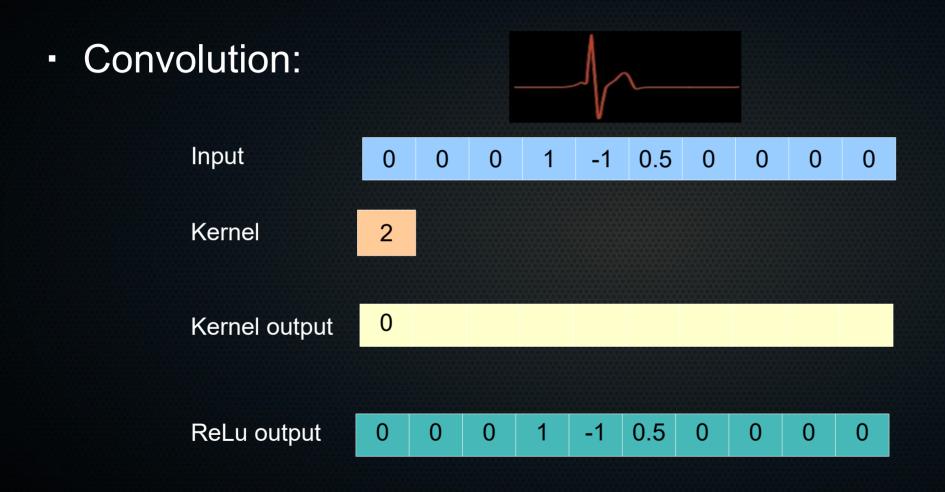


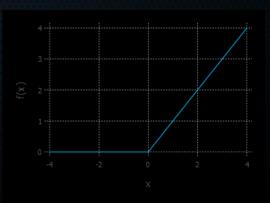
Signal can be at different positions in the sequence:

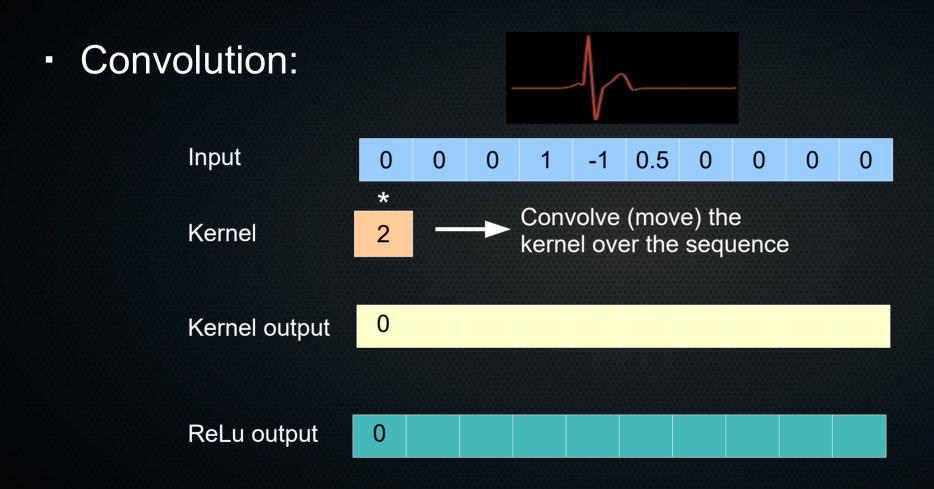


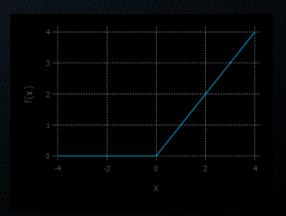
 Dense network needs to optimise such that different weights somehow cause the network to output 1 for different positions of the signal:

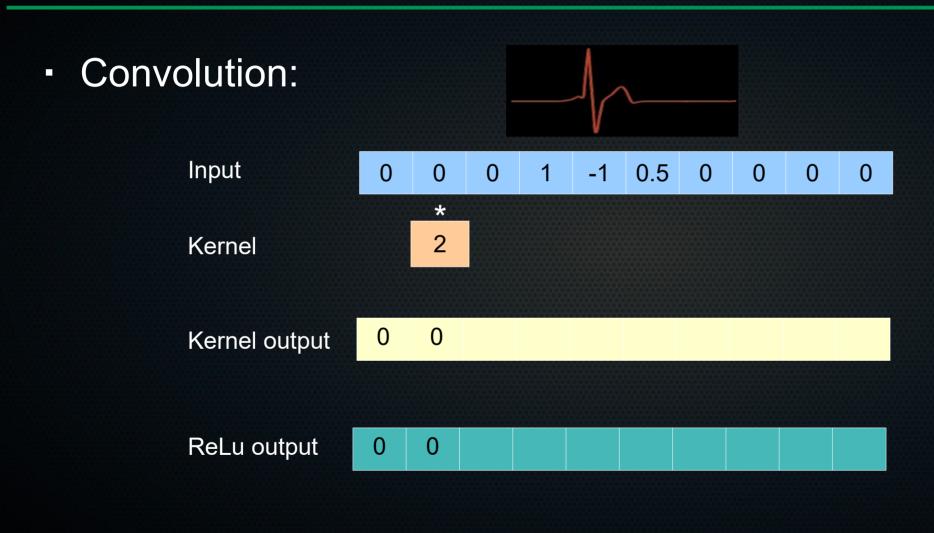


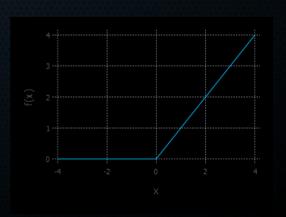


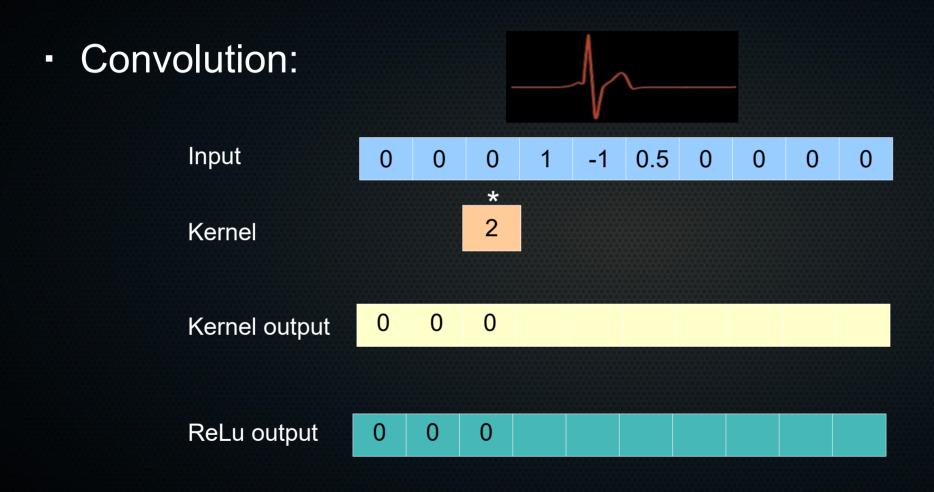


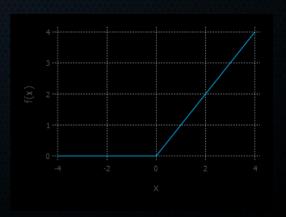


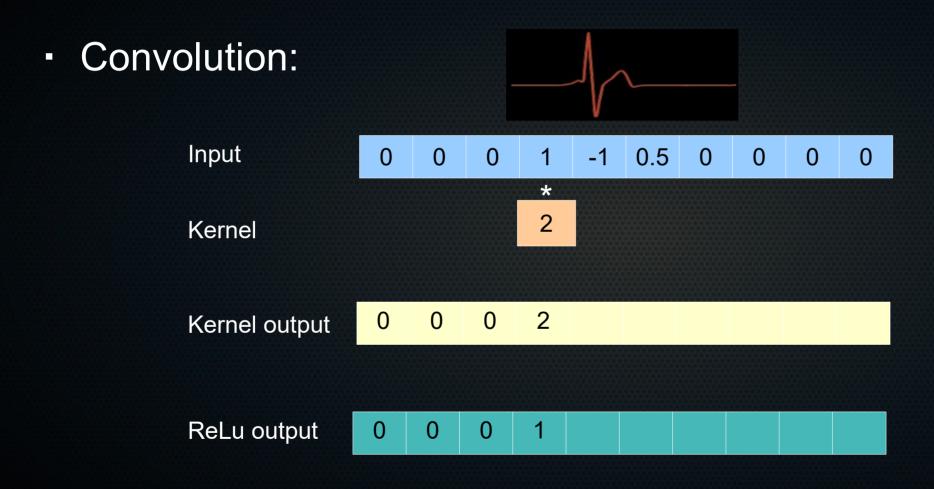


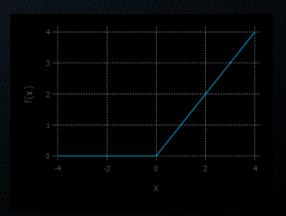


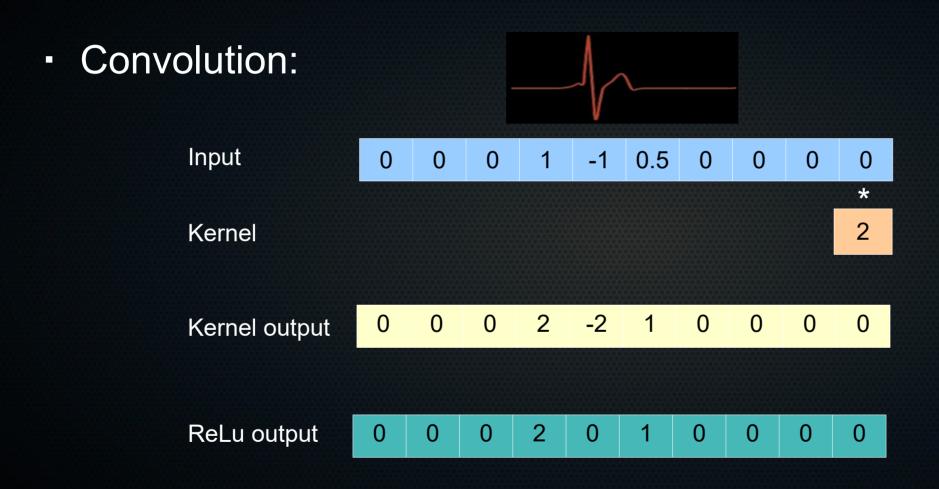


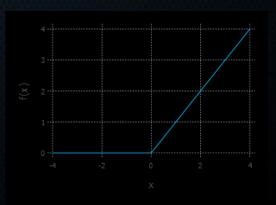


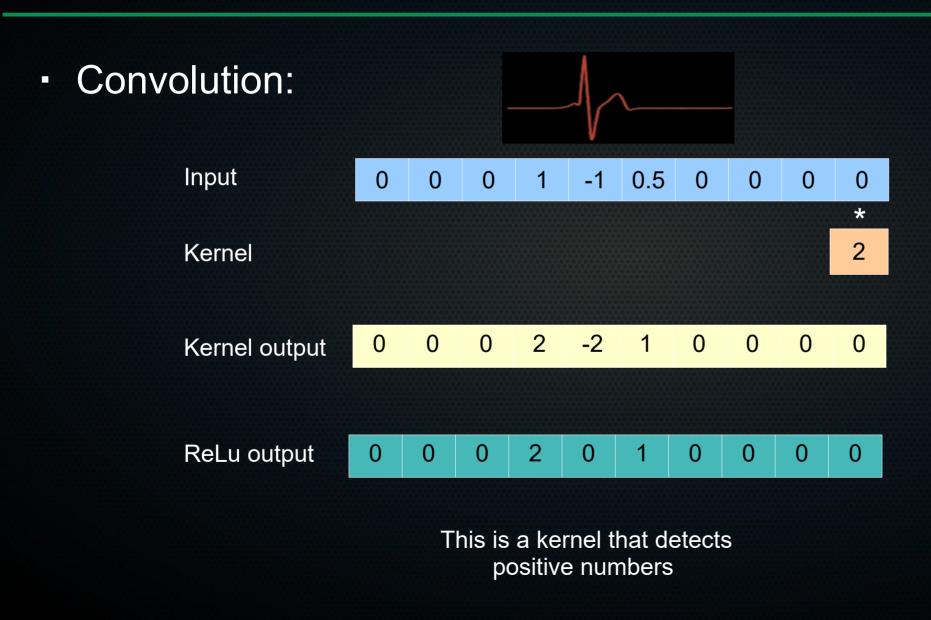


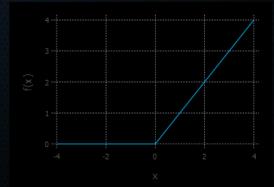


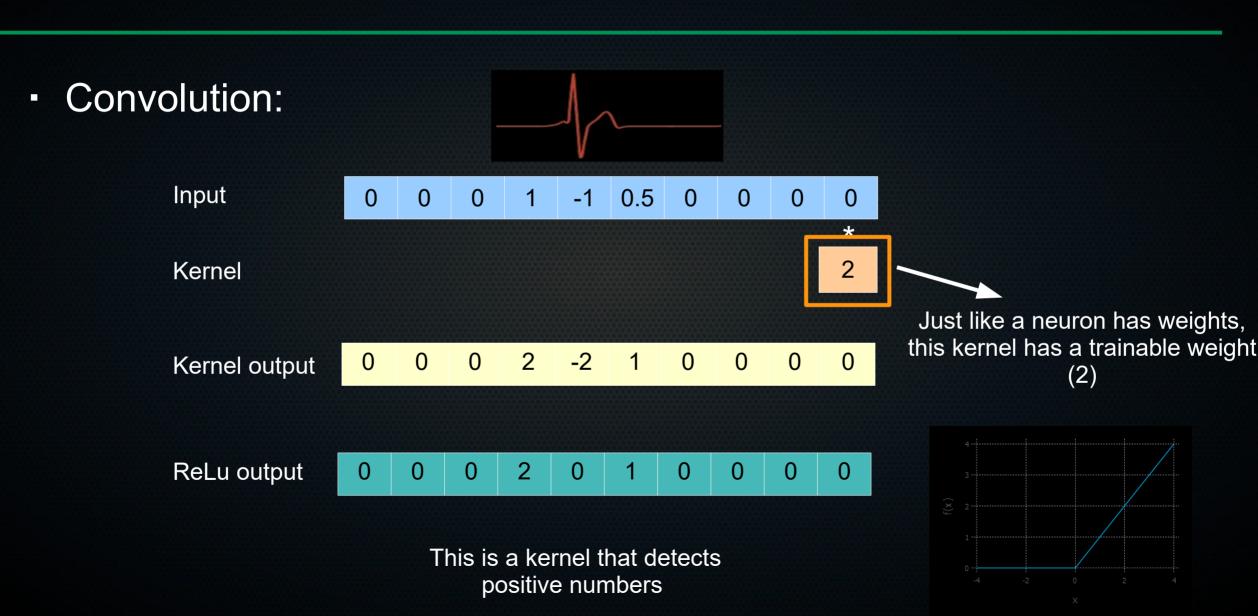


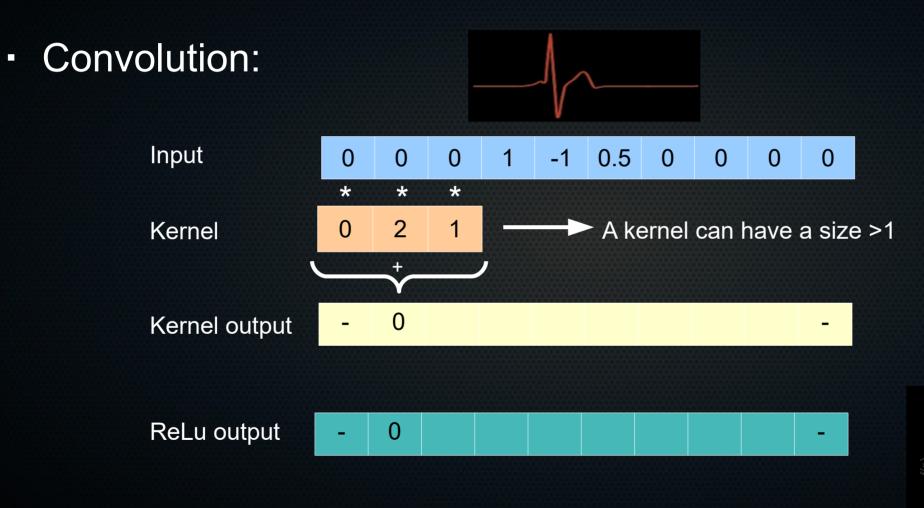


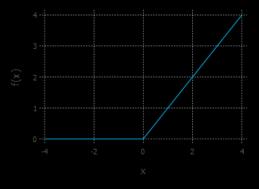


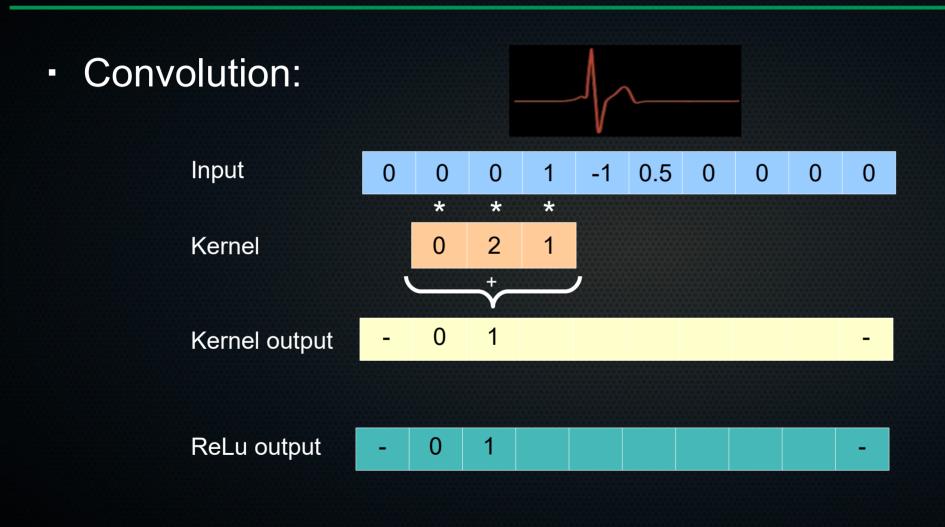


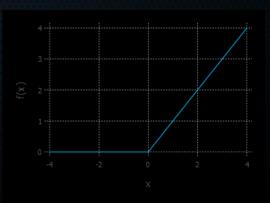


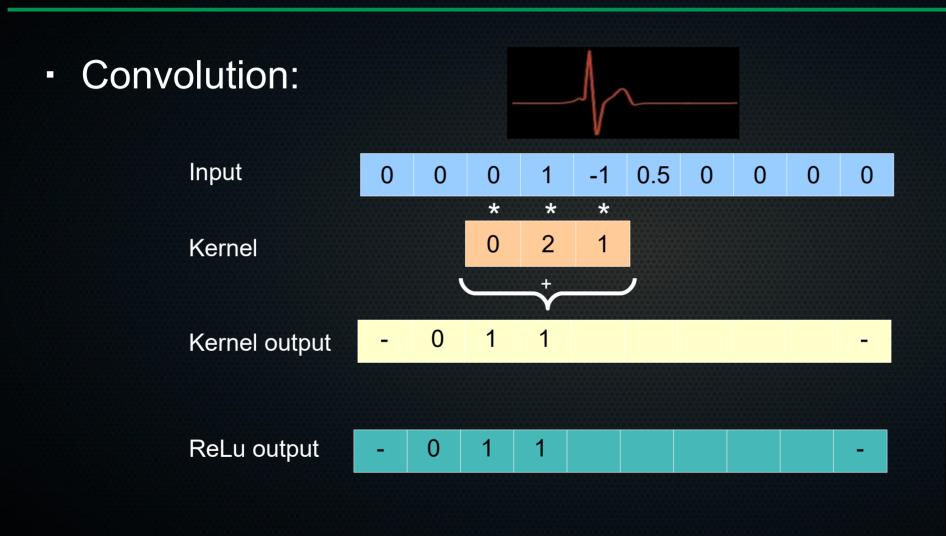


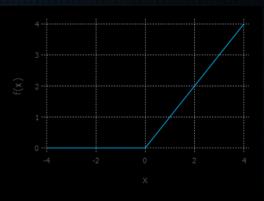


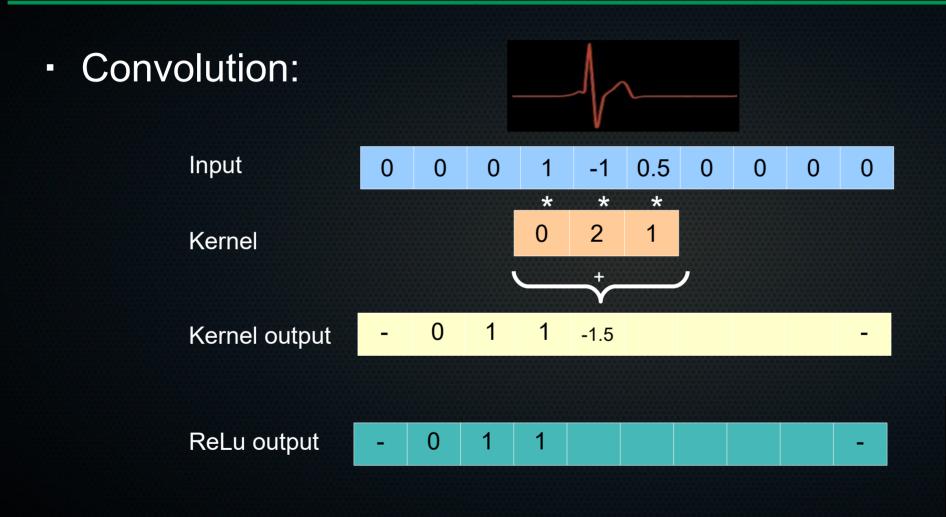


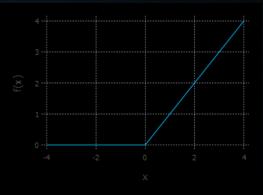


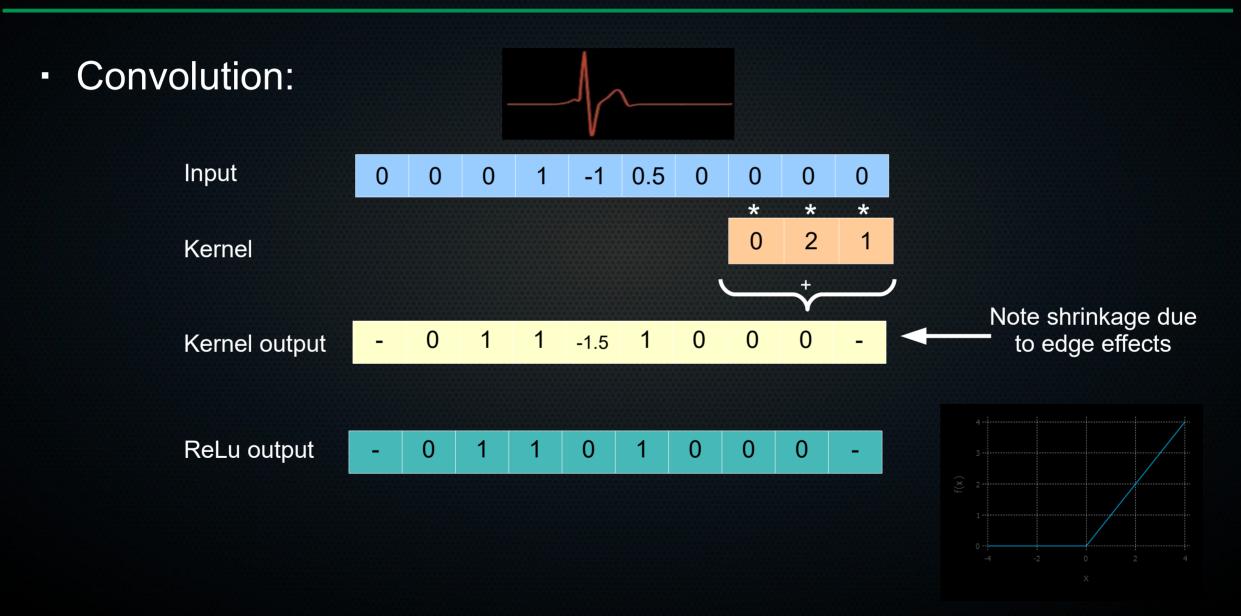


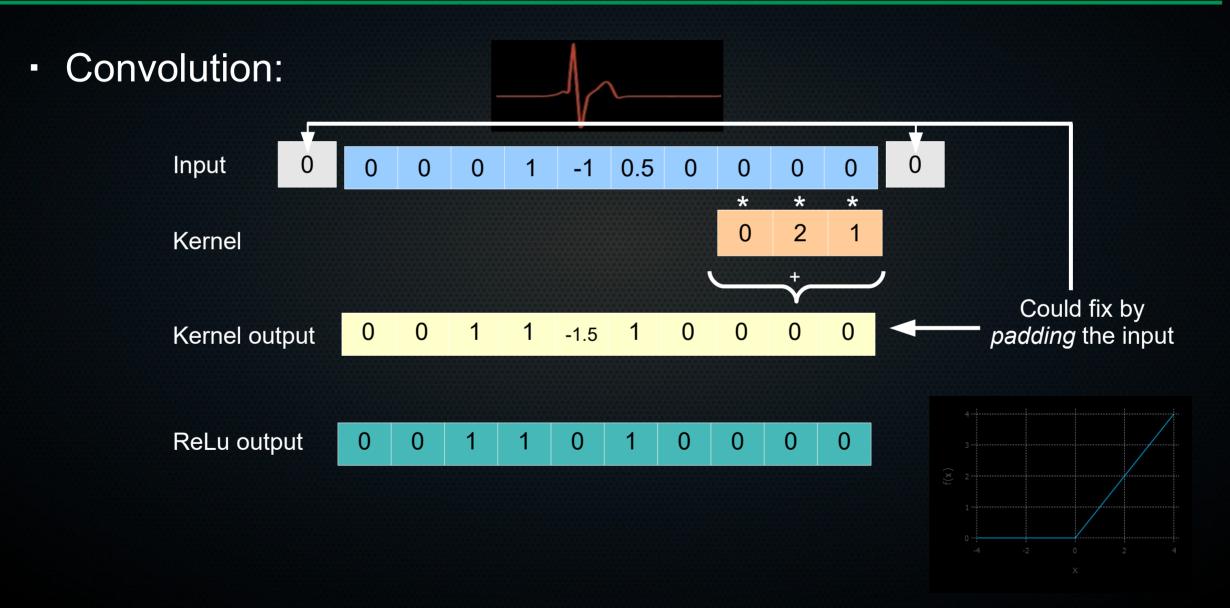


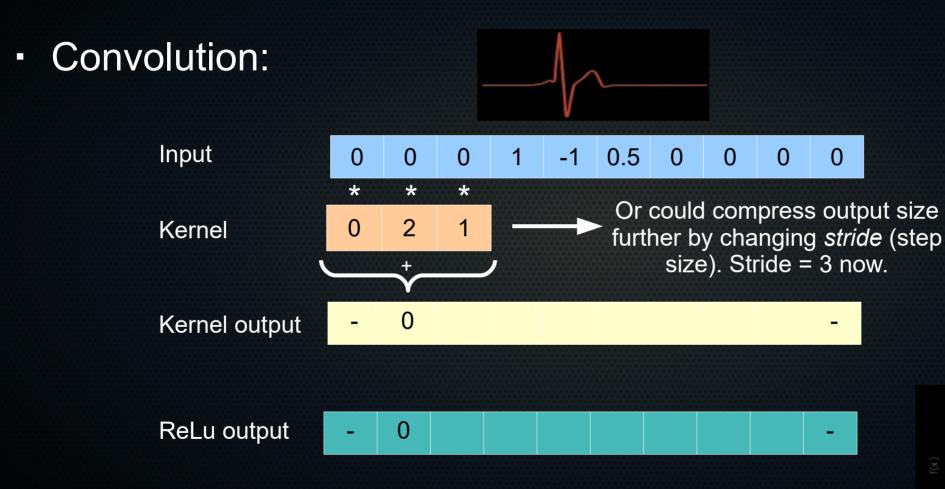


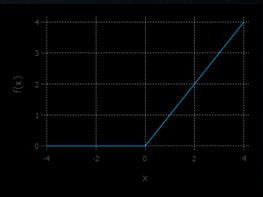


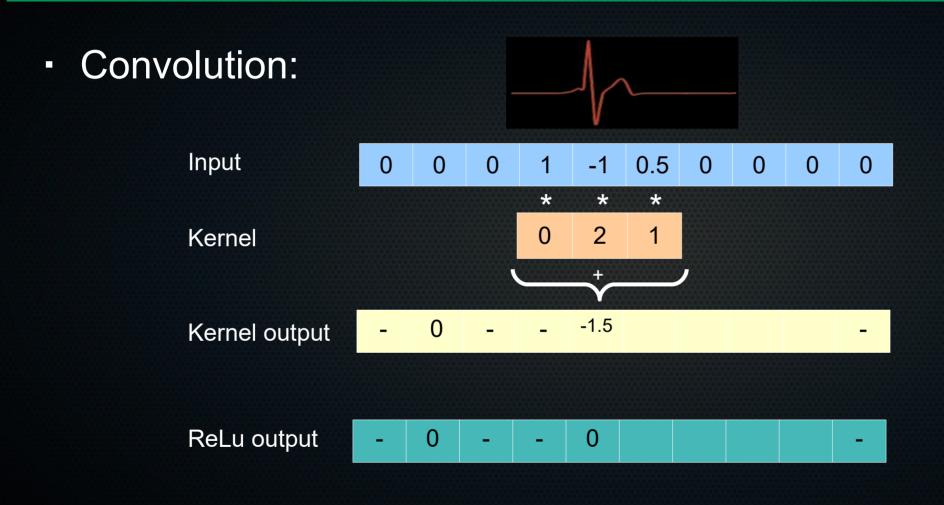


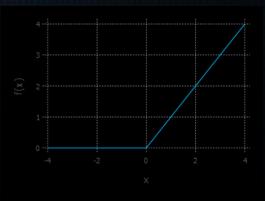


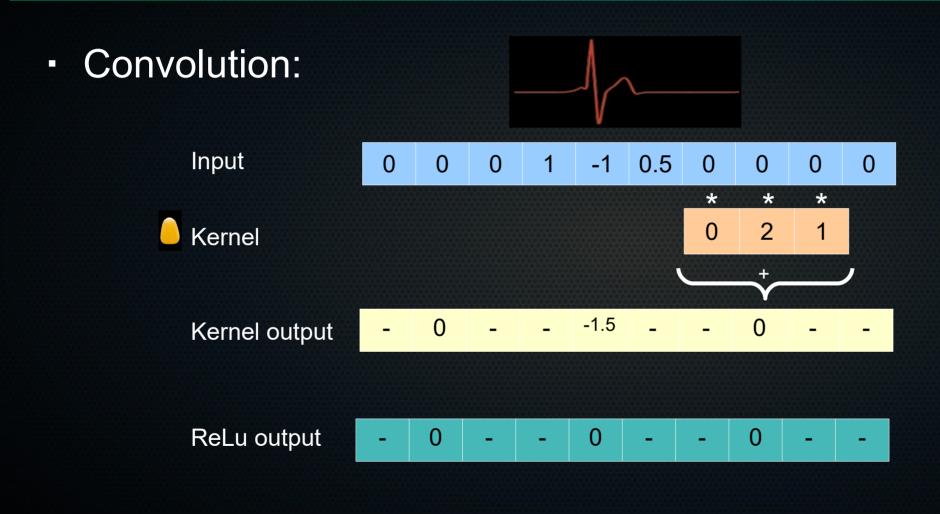


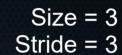


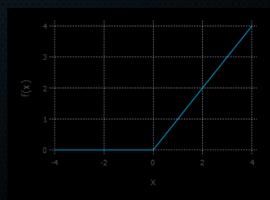


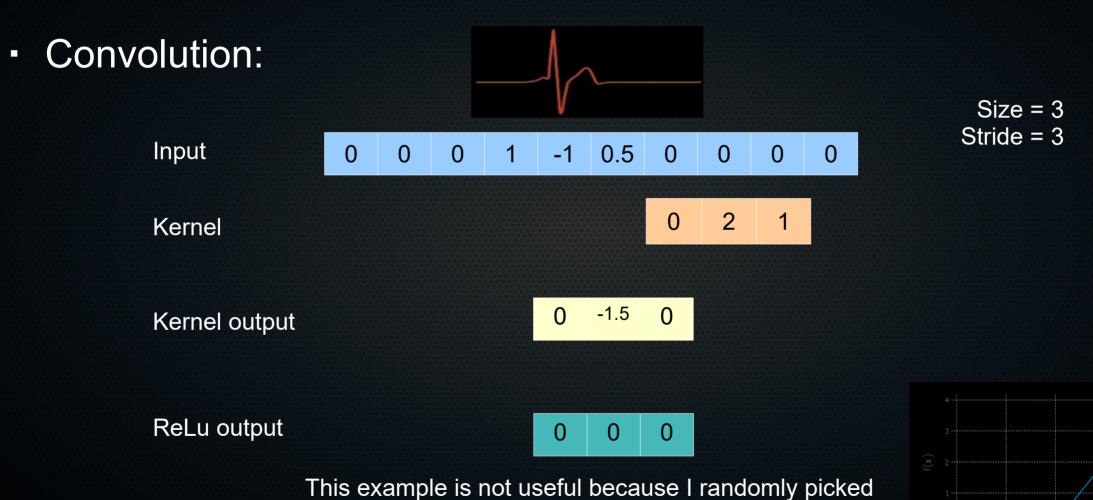




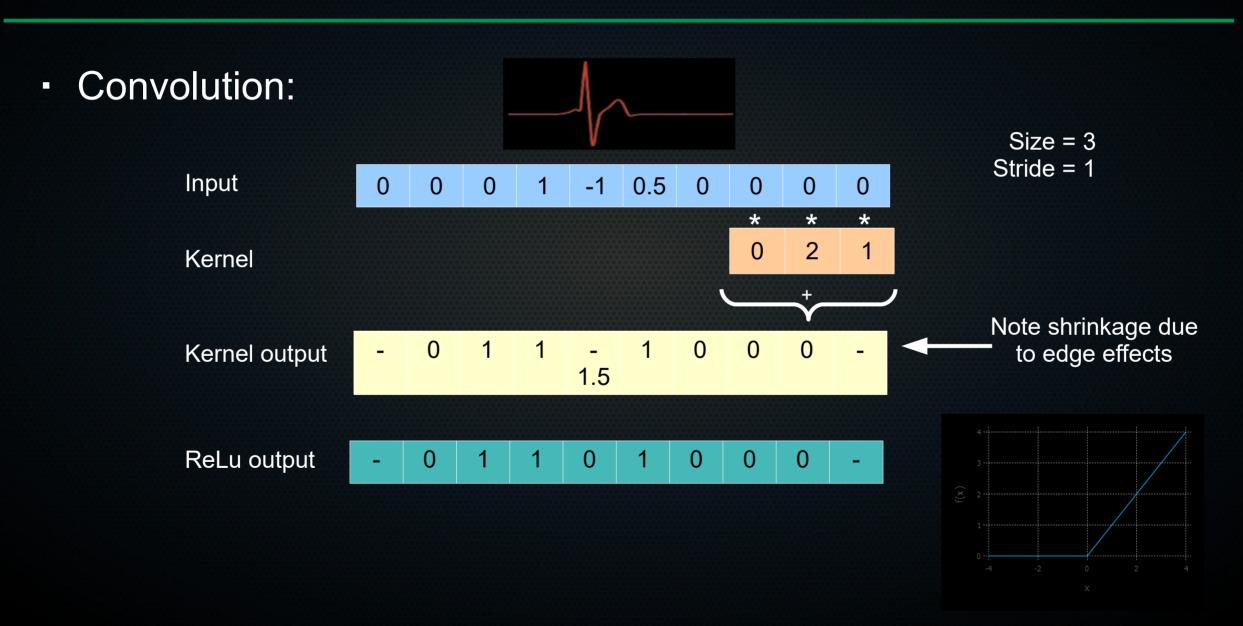


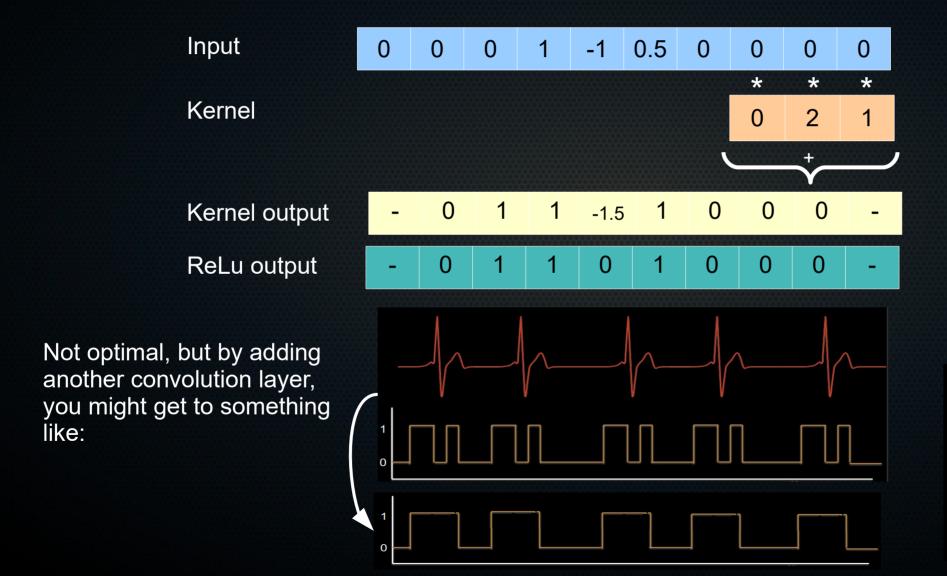




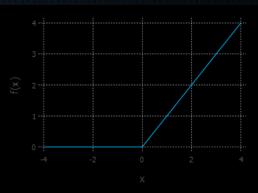


some weights for the kernel. But normally you can train these weights by backpropagation such that the network works well!



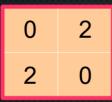


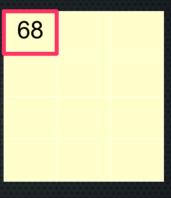
Size = 3 Stride = 1



Size = 2\*2; stride = 1

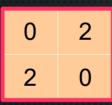
0	22	0	1
12	2	3	23
3	34	26	2
0	22	86	3
4	3	1	4

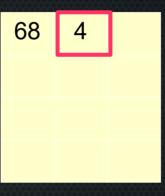




Size = 2\*2; stride = 1

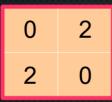
0	22	0	1
12	2	3	23
3	34	26	2
0	22	86	3
4	3	1	4





Size = 2\*2; stride = 1

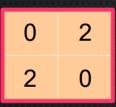
0	22	0	1
12	2	3	23
3	34	26	2
0	22	86	3
4	3	1	4

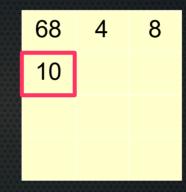




Size = 2\*2; stride = 1

0	22	0	1
12	2	3	23
3	34	26	2
0	22	86	3
4	3	1	4





Etc.

# Another type of convolution: max pooling

 Size = 2\*2; stride = 1; just take the maximum value in the kernel area

0	22	0	1
12	2	3	23
3	34	26	2
0	22	86	3
4	3	1	4

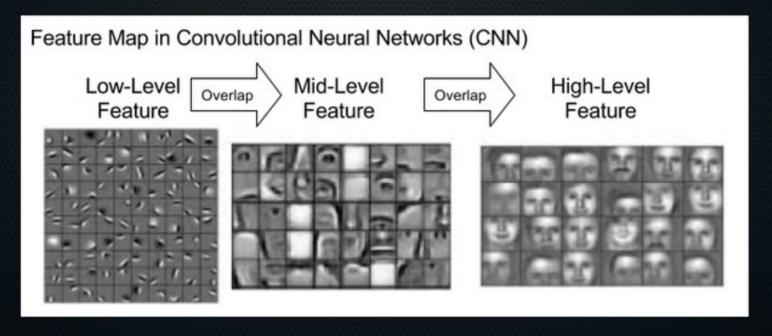
22	22	23
34	26	26
34	86	86
22	86	86

# Another type of convolution: pooling/averaging

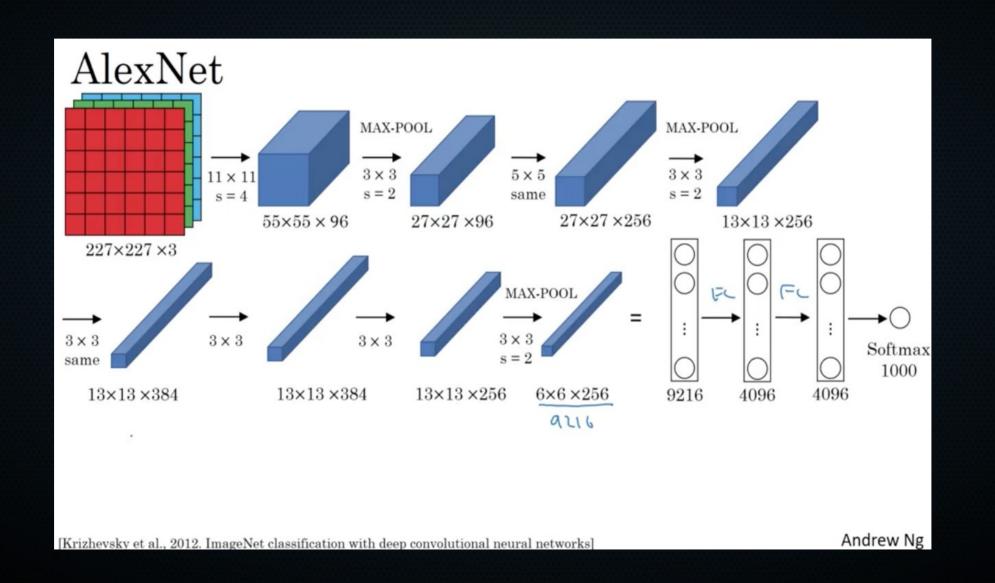


#### Use in face detection

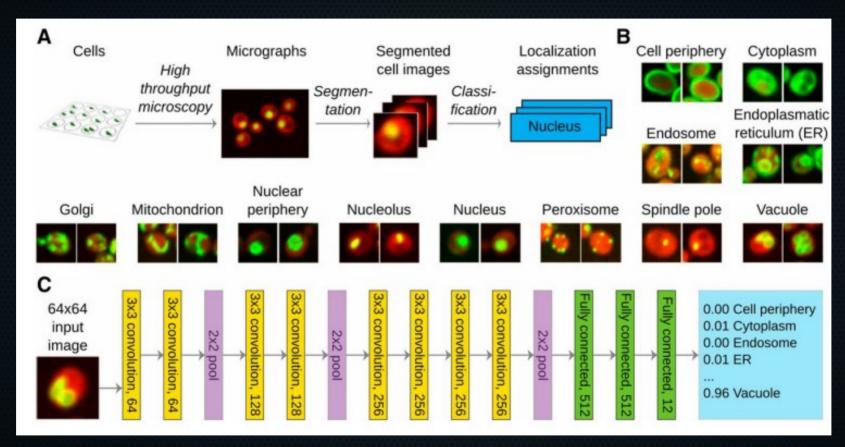
 Since kernels so few parameters: can use many of them per layer → each becomes sensitive to different image features



## Example AlexNet (2012)



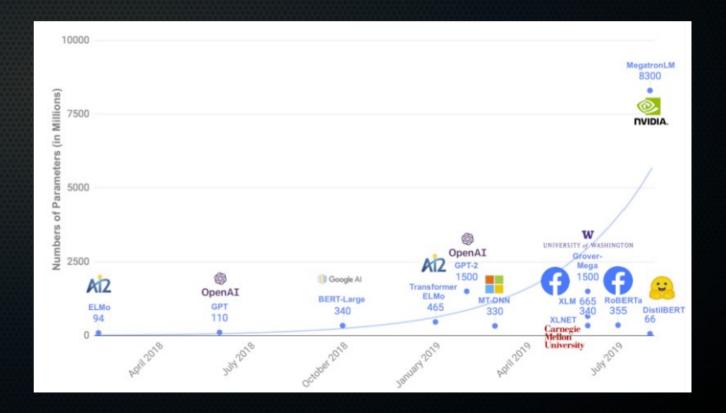
# Biological use



Pärnamaa, T., & Parts, L. (2017). Accurate classification of protein subcellular localization from high-throughput microscopy images using deep learning. G3: Genes, Genomes, Genetics, 7(5), 1385-1392.

## There's a lot more

- Batch normalisation
- Vanishing gradient problem
- Dropout
- Recurrent neural nets



## Implementation

- We are not going to implement convolutional neural networks ourselves: implementing backpropagation properly on a simple dense network is already taxing enough.
- Still, doing that should give you a solid basis for understanding convolutional neural networks, and we'll introduce the Keras library for building (convolutional) neural networks next Monday.

#### Talk 13:15

- Marc Pages Gallego
- Jeroen de Ridder group
- Working with Oxford NanoPore sequencing data
- Talk about ML/deep learning in interpreting sequencing data from nanopores

## Afternoon practical

- Implement backpropagation yourself
- Train a dense neural network on the MNIST dataset