## Deep learning

## 4.4. Convolutions

François Fleuret
https://fleuret.org/dlc/



If they were handled as normal "unstructured" vectors, large-dimension signals such as sound samples or images would require models of intractable size.

For instance a linear layer taking a  $256 \times 256$  RGB image as input, and producing an image of same size would require

$$(256 \times 256 \times 3)^2 \simeq 3.87e + 10$$

parameters, with the corresponding memory footprint (  $\simeq\!150\mbox{Gb}$  !), and excess of capacity.

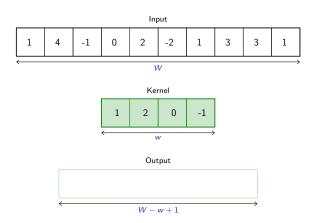
Moreover, this requirement is inconsistent with the intuition that such large signals have some "invariance in translation". A transformation meaningful at a certain location can / should be used everywhere.

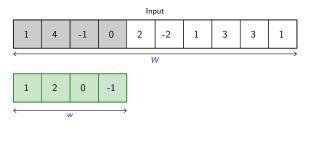
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A convolution layer embodies this idea. It applies the same linear transformation locally, everywhere

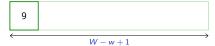
Input

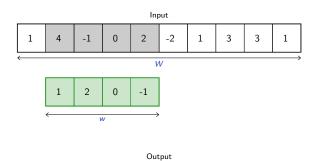
1	4	-1	0	2	-2	1	3	3	1





### Output

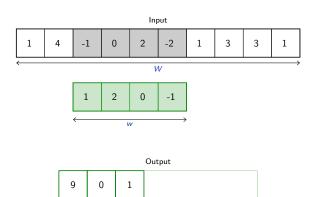




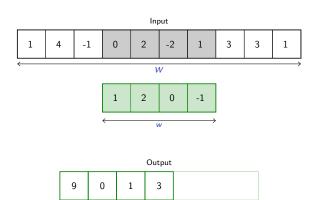
W-w+1

9

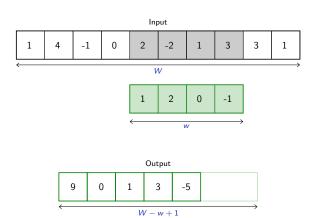
0

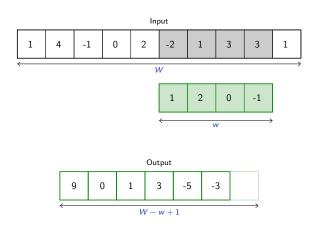


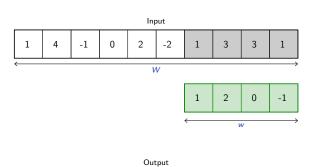
W-w+1

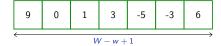


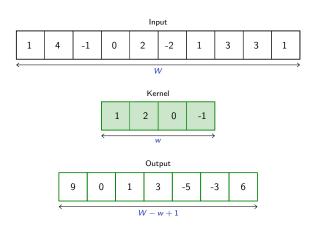
W - w + 1











Formally, in 1d, given

$$x = (x_1, \ldots, x_W)$$

and a "convolution kernel" (or "filter") of width w

$$\textit{u} = (\textit{u}_1, \dots, \textit{u}_w)$$

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the convolution  $x \circledast u$  is a vector of size W - w + 1, with

$$(x \circledast u)_i = \sum_{j=1}^w x_{i-1+j} u_j$$
$$= (x_i, \dots, x_{i+w-1}) \cdot u$$

for instance

$$(1,2,3,4) \circledast (3,2) = (3+4,6+6,9+8) = (7,12,17).$$

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This differs from the usual convolution since the kernel and the signal are both visited in increasing index order.

Convolution can implement in particular differential operators, e.g.

$$(0,0,0,0,1,2,3,4,4,4,4) \otimes (-1,1) = (0,0,0,1,1,1,1,0,0,0).$$

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or crude "template matcher", e.g.



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Its most usual form for "convolutional networks" processes a 3d tensor as input (i.e. a multi-channel 2d signal) to output a 2d tensor. The kernel is not swiped across channels, just across rows and columns.

In this case, if the input tensor is of size  $C \times H \times W$ , and the kernel is  $C \times h \times w$ , the output is  $(H - h + 1) \times (W - w + 1)$ .

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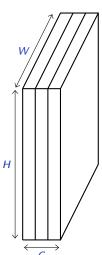
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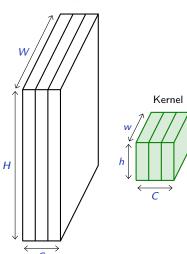
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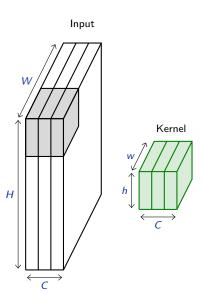
In a standard convolution layer, D such convolutions are combined to generate a  $D \times (H-h+1) \times (W-w+1)$  output.



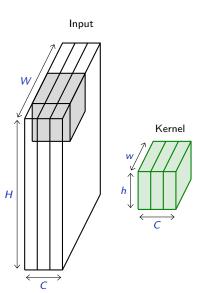


### Input



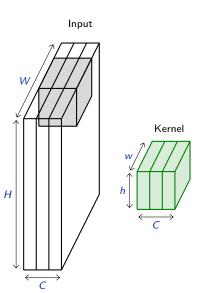




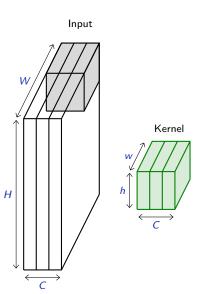




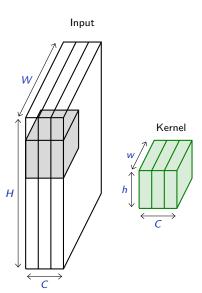


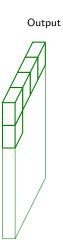


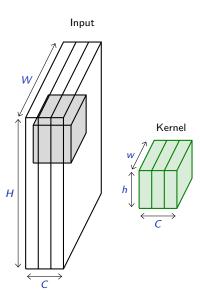


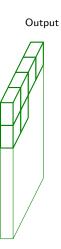


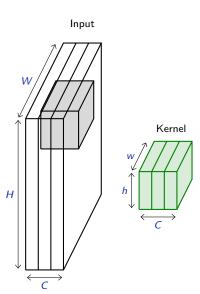










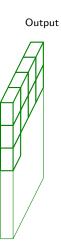


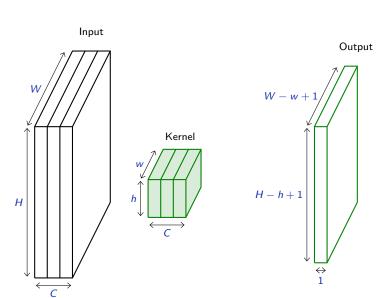


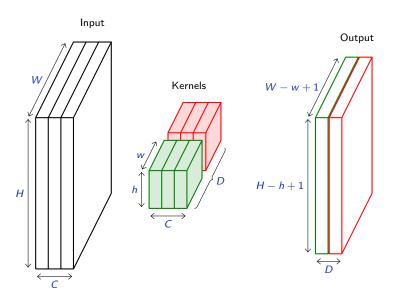
# Input Kernel h Н



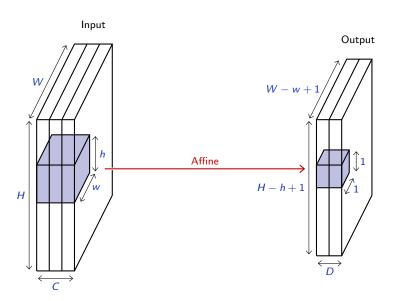
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A 3d convolution can be used if the channel index has some metric meaning, such as time for a series of grayscale video frames. Otherwise swiping across channels makes no sense

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In the context of convolutional networks, a standard linear layer is called a **fully connected layer**, or a **dense layer**, since every input influences every output.

The autograd-compliant function

F.conv2d(input, weight, bias=None, stride=1, padding=0, dilation=1, groups=1)

Implements a 2d convolution, where weight is of dimension  $D \times C \times h \times w$  and contains the kernels, bias is of dimension D, input is of dimension

$$N \times C \times H \times W$$

and the result is of dimension

$$N \times D \times (H-h+1) \times (W-w+1)$$
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```
>>> weight = torch.randn(5, 4, 2, 3)
>>> bias = torch.randn(5)
>>> input = torch.randn(117, 4, 10, 3)
>>> output = F.conv2d(input, weight, bias)
>>> output.size()
torch.Size([117, 5, 9, 1])
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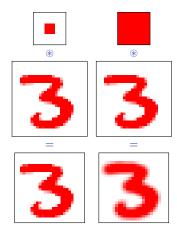
Similar functions implement 1d and 3d convolutions.

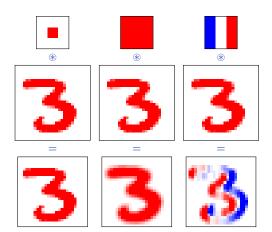
```
x = mnist train.data[12].float().view(1, 1, 28, 28)
weight = torch.emptv(5, 1, 3, 3)
weight[0, 0] = torch.tensor([ [ 0., 0., 0. ],
                             [ 0., 1., 0.],
[ 0., 0., 0.]])
weight[1, 0] = torch.tensor([ [ 1., 1., 1.],
                             [ 1., 1., 1.],
[ 1., 1., 1.])
weight[2, 0] = torch.tensor([ [ -1., 0., 1. ],
                              Γ-1.. O.. 1. ].
                              [-1., 0., 1. ] ])
weight[3, 0] = torch.tensor([ [ -1., -1., -1. ],
                             [ 0., 0., 0.],
[ 1., 1., 1.])
weight[4, 0] = torch.tensor([ [ 0., -1., 0. ],
                              [-1., 4., -1.],
                              [0, -1, 0, 1]
```

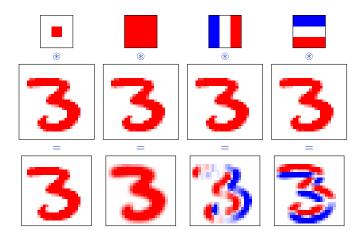
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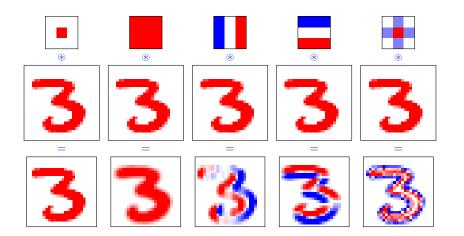
y = F.conv2d(x, weight)











Wraps the convolution into a Module, with the kernels and biases as Parameter properly randomized at creation.

The kernel size is either a pair (h, w) or a single value k interpreted as (k, k).

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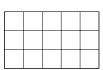
```
>>> f = nn.Conv2d(in_channels = 4, out_channels = 5, kernel_size = (2, 3))
>>> for n, p in f.named_parameters(): print(n, p.size())
...
weight torch.Size([5, 4, 2, 3])
bias torch.Size([5])
>>> x = torch.randn(117, 4, 10, 3)
>>> y = f(x)
>>> y.size()
torch.Size([117, 5, 9, 1])
```

Padding, stride, and dilation

Convolutions have three additional parameters:

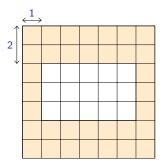
- The padding specifies the size of a zeroed frame added around the input,
- the **stride** specifies a step size when moving the kernel across the signal,
- the dilation modulates the expansion of the filter without adding weights.

Here with  $C \times 3 \times 5$  as input



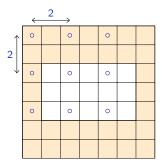
Input

Here with  $C \times 3 \times 5$  as input, a padding of (2,1)

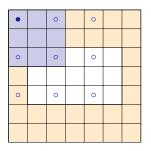


Input

Here with  $C \times 3 \times 5$  as input, a padding of (2,1), a stride of (2,2)

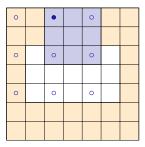


Input



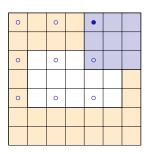
Output

Input



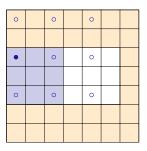


Input



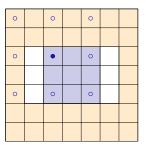


Input



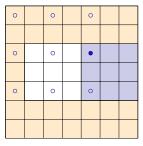
Output

Input



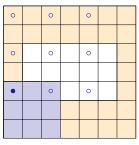
Output

Input



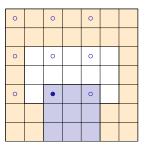


Input





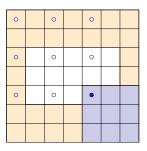
Input



Output

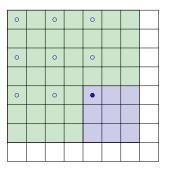
Input

Here with  $C \times 3 \times 5$  as input, a padding of (2,1), a stride of (2,2), and a kernel of size  $C \times 3 \times 3$ , the output is  $1 \times 3 \times 3$ .



Output

Input





A convolution with a stride greater than  $1\ \text{may}$  not cover the input map entirely, hence may ignore some of the input values.

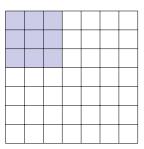
The dilation modulates the expansion of the filter support by adding rows and columns of zeros between coefficients (Yu and Koltun, 2015).

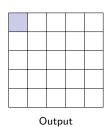
It is 1 for standard convolutions, but can be greater, in which case the resulting operation can be envisioned as a convolution with a regularly sparsified filter.

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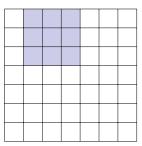
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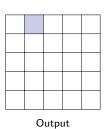
This notion comes from signal processing, where it is referred to as *algorithme à trous*. hence the term sometime used of "convolution à trous".



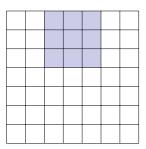


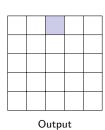
Input



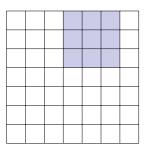


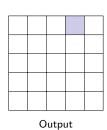
Input



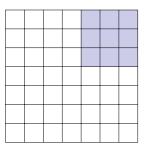


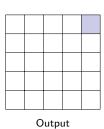
Input



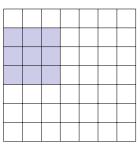


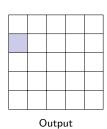
Input



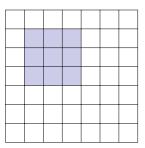


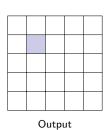
Input



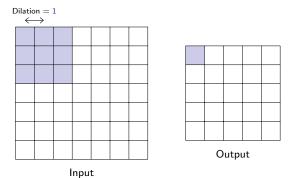


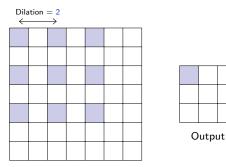
Input

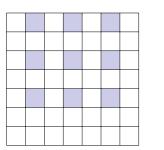




Input

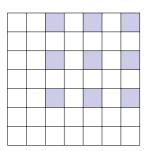






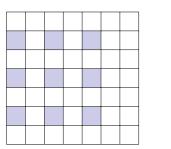


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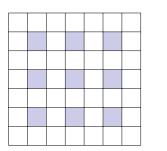


Input



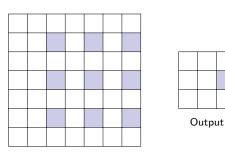
Output

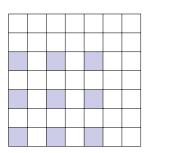
Input





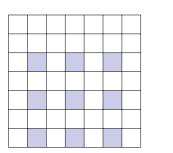
Input





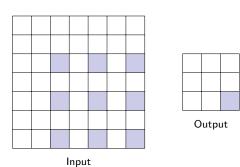
Input

Output



Input

Output



A 1d convolution with a kernel of size k and dilation d can be interpreted as a convolution with a filter of size 1 + (k-1)d with only k non-zero coefficients.

For example with k=3 and d=4, the difference between the input map size and the output map size is 1+(3-1)4-1=8.

```
>>> x = torch.randn(1, 1, 20, 30)
>>> 1 = nn.Conv2d(1, 1, kernel_size = 3, dilation = 4)
>>> 1(x).size()
torch.Size([1, 1, 12, 22])
```

Having a dilation greater than one increases the units' receptive field size without increasing the number of parameters.

Convolutions with stride or dilation strictly greater than one reduce the activation map size, for instance to make a final classification decision.



## References

 $\mathsf{abs}/1511.07122\mathsf{v3},\ 2015.$ 

F. Yu and V. Koltun. Multi-scale context aggregation by dilated convolutions. Col	RR,
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