EE-559 - Deep learning

4.4. Convolutions

François Fleuret

https://fleuret.org/ee559/

Mon Mar 11 08:59:42 UTC 2019





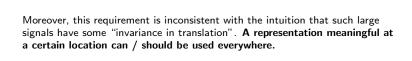
If they were handled as normal "unstructured" vectors, large-dimension signals such as sound samples or images would require models of intractable size.

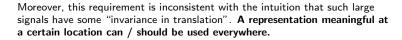
For instance a linear layer taking a 256×256 RGB image as input, and producing an image of same size would require

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

parameters, with the corresponding memory footprint (~150Gb !), and excess of capacity.

1 / 22



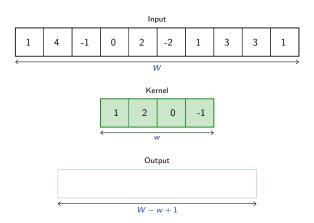


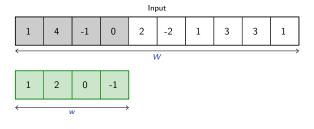
A convolution layer embodies this idea. It applies the same linear transformation locally, everywhere, and preserves the signal structure.

Input

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

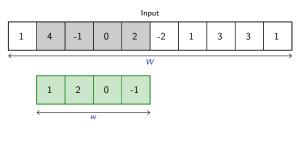
vv





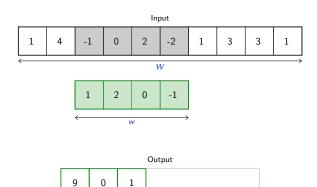
Output



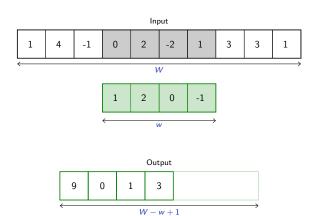


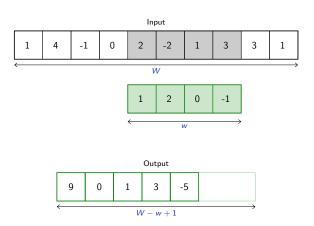
Output

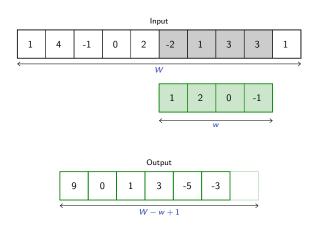


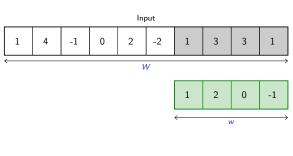


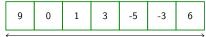
W - w + 1



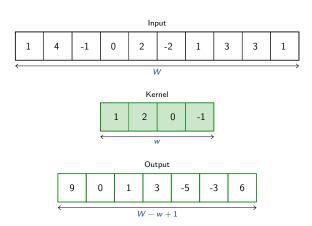








W - w + 1



Formally, in 1d, given

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

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

$$u=(u_1,\ldots,u_w)$$

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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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 \triangle

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

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

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



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



or crude "template matcher", e.g.



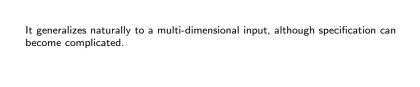
$$(0,0,0,0,1,2,3,4,4,4,4) \otimes (-1,1) = (0,0,0,1,1,1,1,0,0,0).$$



or crude "template matcher", e.g.



Both of these computation examples are indeed "invariant by translation".



It generalizes naturally to a multi-dimensional input, although specification can become complicated.

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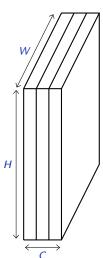
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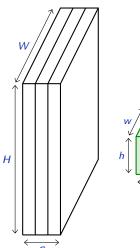
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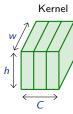
In a standard convolution layer, D such convolutions are combined to generate a $D \times (H-h+1) \times (W-w+1)$ output.

Input

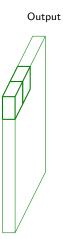


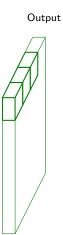
Input

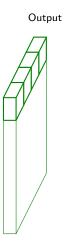


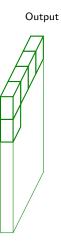


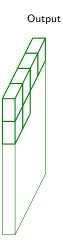


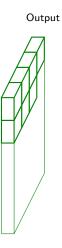




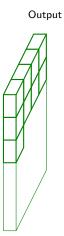


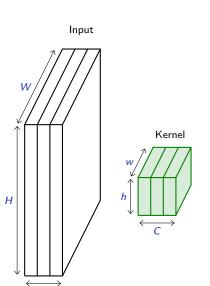


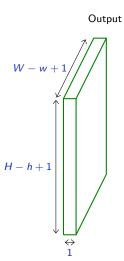


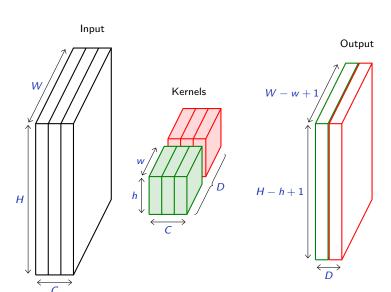














A 1d signal is converted into a 1d signal, a 2d signal into a 2d, and neighboring parts of the input signal influence neighboring parts of the output signal.

Note that a convolution preserves the signal support structure.

A 1d signal is converted into a 1d signal, a 2d signal into a 2d, and neighboring parts of the input signal influence neighboring parts of the output signal.

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.

We usually refer to one of the channels generated by a convolution layer as an activation map.

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The sub-area of an input map that influences a component of the output as the **receptive field** of the latter.

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

Implements a 2d convolution, where weight contains the kernels, and is $D \times C \times h \times w$, 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.empty(5, 4, 2, 3).normal_()
>>> bias = torch.empty(5).normal_()
>>> input = torch.empty(117, 4, 10, 3).normal_()
>>> output = torch.nn.functional.conv2d(input, weight, bias)
>>> output.size()
torch.Size([117, 5, 9, 1])
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```

Similar functions implement 1d and 3d convolutions.

10 / 22

```
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. ],
                              Г-1., 4., -1. ].
                              [0, -1, 0, 1]
y = torch.nn.functional.conv2d(x, weight)
```

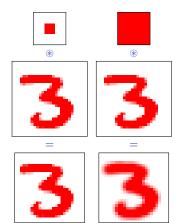


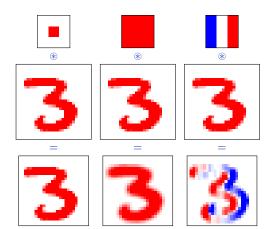
*

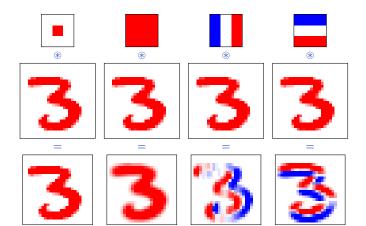


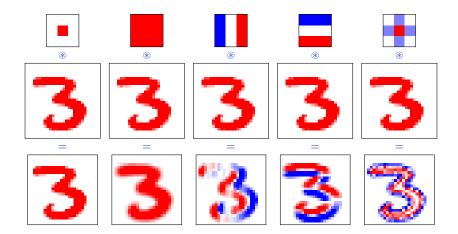
_











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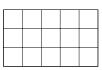
```
>>> f = nn.Conv2d(in_channels = 4, out_channels = 5, kernel_size = (2, 3))
>>> for n, p in f.named_parameters(): print(n, p.size())
...
weight torch.Size([5, 4, 2, 3])
bias torch.Size([5])
>>> x = torch.empty(117, 4, 10, 3).normal_()
>>> y = f(x)
>>> y.size()
torch.Size([117, 5, 9, 1])
```

Padding, stride, and dilation

Convolutions have two additional standard parameters:

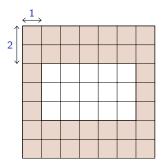
- The padding specifies the size of a zeroed frame added around the input,
- the **stride** specifies a step size when moving the kernel across the signal.

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



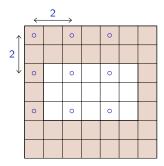
Input

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

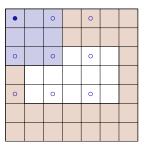


Input

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

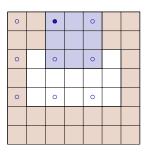


Input



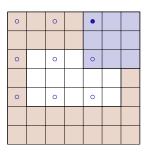


Input



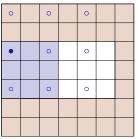


Input

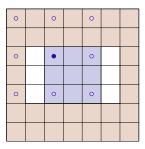




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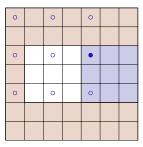






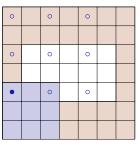
Output

Input



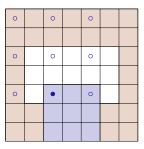


Input



Output

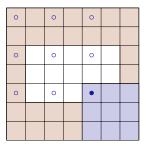
Input



Output

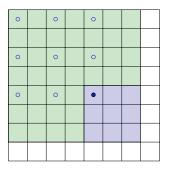
Input

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 ${\bf 1}$ may not cover the input map completely, hence may ignore some of the input values.

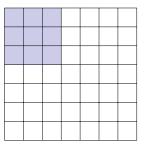
Convolution operations admit one more standard parameter that we have not discussed yet: The dilation, which modulates the expansion of the filter support (Yu and Koltun, 2015).

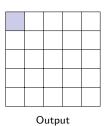
It is 1 for standard convolutions, but can be greater, in which case the resulting operation can be envisioned as a convolution with a regularly sparsified filter.

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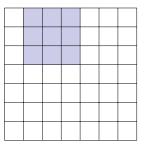
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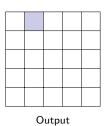
This notion comes from signal processing, where it is referred to as *algorithme à trous*, hence the term sometime used of "convolution à trous".



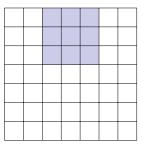


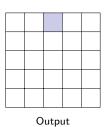
Input



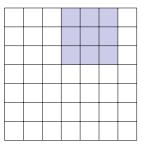


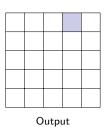
Input



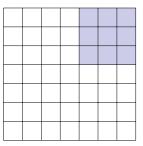


Input



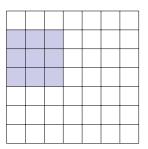


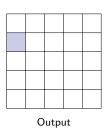
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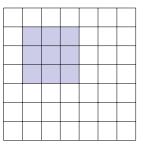
Output

Input



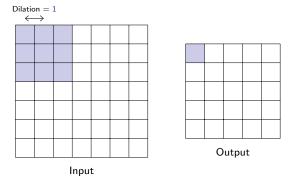


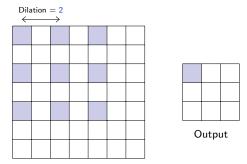
Input

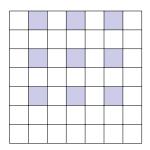


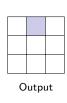
Output

Input

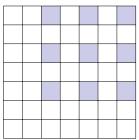






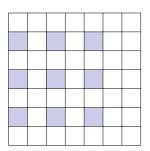


Input



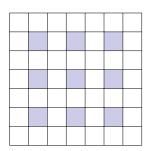


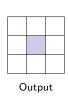
Input



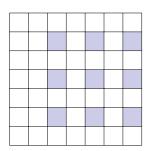


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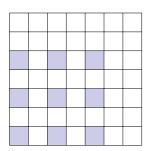


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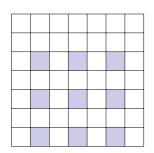


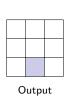
Input



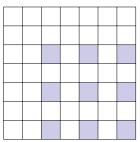


Input





Input





Input

A convolution with a 1d 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 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.empty(1, 1, 20, 30).normal_()
>>> 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.

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Such networks have the advantage of simplicity:

- non-linear operations are only in the activation function,
- joint operations that combine multiple activations to produce one are only in linear layers.

22 / 22



References

abs/1511.07122v3, 2015.

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