Machine Learning & its Application

Convolutional Networks

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Outline

- Introduction
- Convolution
 - Image Processing
 - Convolution in Neural Networks
 - Convolution vs Dense Layers
- Convolutional Neural Networks
 - Residual connections
 - Batch normalization
 - Dropout
- Fully Convolutional Networks
 - Receptive Field
 - Encoder-Decoder Architecture
- Transfer Learning

Introduction

- Dense layers are not suitable for image applications.
- We lose the spatial relation in an image when flattening it.
- We also create a huge vector hard to process.
- Instead, we use convolutional networks.
- We will explore convolution operation and how did it integrate machine learning.
- We will then explore solutions for image-to-image applications.

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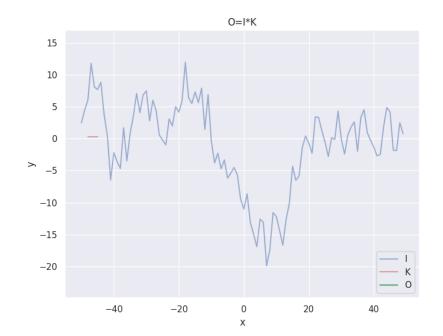
Convolution

- Before talking about convolution we need to talk about images.
- An image is a 2D visual representation of a scene, stored in an array.
- We have different type of images:
 - Color and gray images.
 - Depth images.
 - Thermal images etc..
- We will focus here on color and gray images only.
- Color and gray images are usually stored on 8 bits.

Convolution

- Storing images on 8 bits mean that each pixel can take values in the range [0-255].
- RGB images represent all colors using a combination of red, green and blue.
- Gray images have only a single channel.
- The pixel value represents its intensity:
 - In gray images, the value represents its blackness or whiteness (0 black, 255 white).
 - In color images, the value represent the intensity of each color in each channel.
- We can convert a color image into a gray image.
- We can do a lot of processing on images to extract useful information

- Convolution is a mathematical operation that is widely used in practice.
- We can implement many image processing algorithms with it.
- It is just a moving kernel applied locally. It is defined for 1d and 2d signals.



• Let's take the first image processing application: blurring.





- To blur an image, we can replace each pixel with its neighbors average value.
- This can be implemented by a uniform kernel convolved with the image.

$$K = \begin{bmatrix} \frac{1}{n^2} & \frac{1}{n^2} & \dots & \frac{1}{n^2} \\ \frac{1}{n^2} & \frac{1}{n^2} & \dots & \frac{1}{n^2} \\ \dots & \dots & \dots & \dots \\ \frac{1}{n^2} & \frac{1}{n^2} & \dots & \frac{1}{n^2} \end{bmatrix}; K \in \mathbb{R}^{(n,n)}$$

- Another application is to detect edges in images.
- On a edge, we have a big difference of color intensities. We can compute the absolute difference between consecutive pixels.

absolute difference in x direction



absolute difference in y direction



• We can compute the norm to get a single edge image.

normalized gradient norm

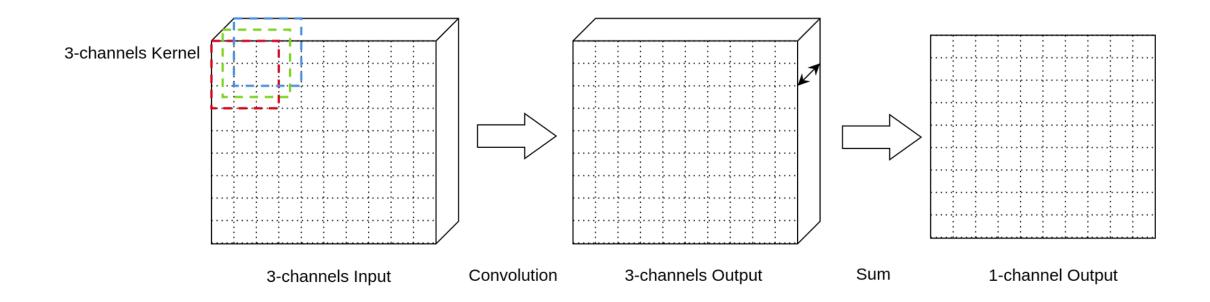


• Interestingly, we can implement the difference in x and y with convolutions.

$$K_x = \begin{bmatrix} -1 & 1 \end{bmatrix}$$
; $K_y = \begin{bmatrix} -1 \\ 1 \end{bmatrix}$

- Convolutional kernels can extract meaningful information from an image.
- Instead of using defined kernels, we want to let the model find the kernel values.
- Convolution layers replace dense layers in deep learning for image applications.

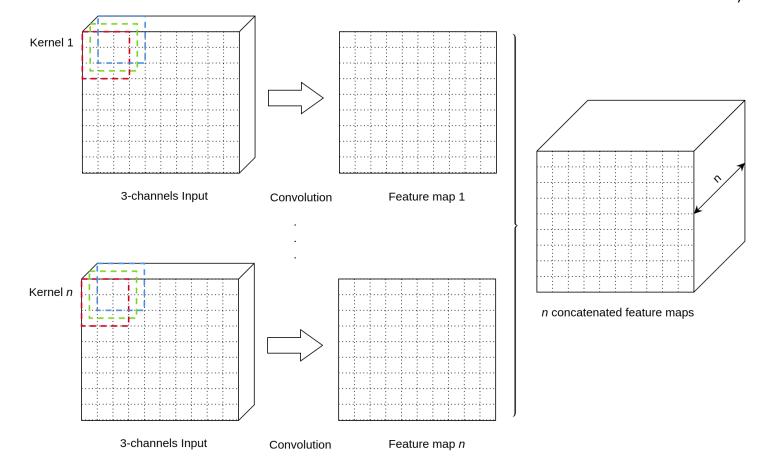
- In neural networks, we learn the values of a convolution kernel.
- In addition, we sum all the channels together after convolving with the kernel.



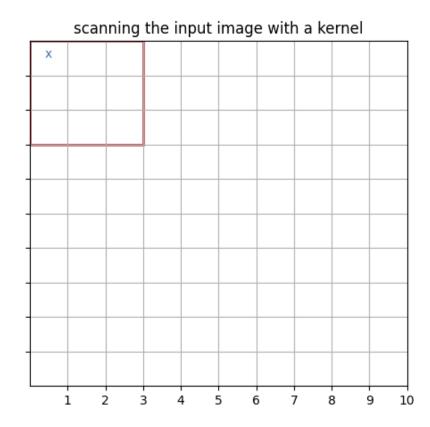
- With dense layers, there is only one parameter to set: the number of units.
- Convolution has multiple parameter to set:
 - Kernel Size.
 - Number of Filters.
 - Padding.
 - Strides.
- We will see what does each one mean.

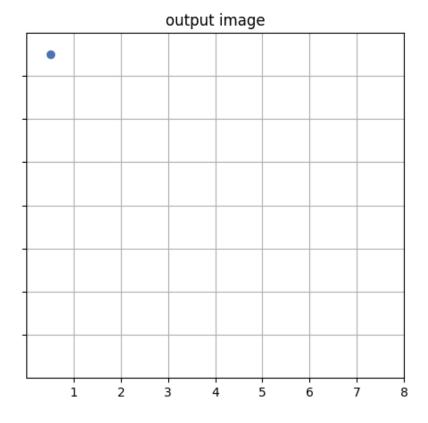
- The kernel size is the spatial size of the learnable kernel (k_x, k_y) .
- However, the total size of the kernel is $k_x^* k_y^* c$, where c is the input channels.
- Number of filters n is the number of kernels to use in a layer.
- Each kernel (or filter) will generate a single "feature" image.
- We concatenate those features by channel to create a feature map.
- A bias term of the same size of the features is used as well.

• The total number of parameters of a convolutional layer is: $k_x^* k_y^* c * n$.

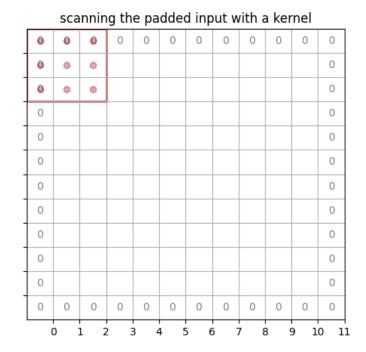


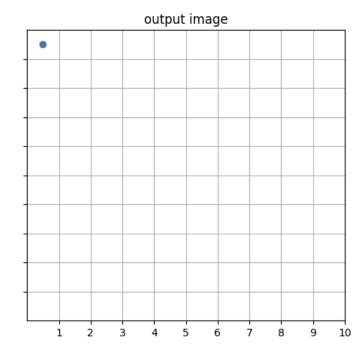
The convolution operation outputs an image smaller than its input.



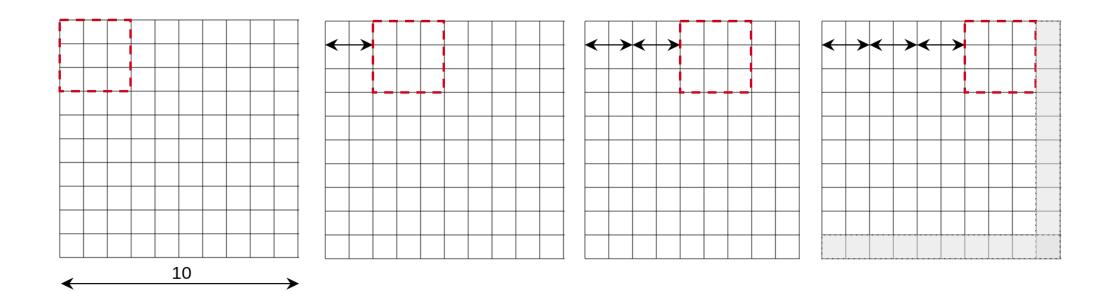


- However, we would to conserve the same output size, thus we pad the input.
- We call this padding "same". If no padding applied, it is called "valid".

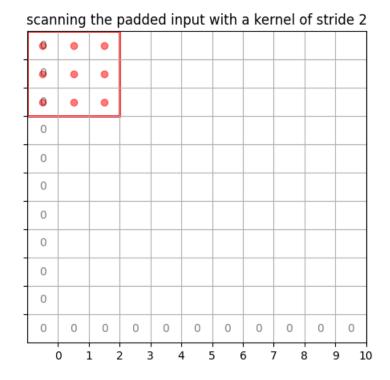


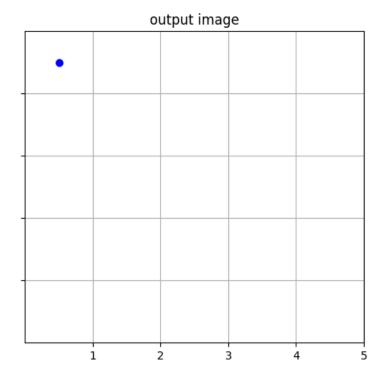


- We presented the case when the kernel move 1 step at a time.
- This step is called the stride, it can be bigger as well.
- What remains from the image is dropped.



- Strides > 1 reduce the output size. We call such convolution strided convolution.
- Using strides = 2 and padding will give an output of half size the input.





Convolution vs Dense Layers

- There are several advantages of using convolution for images:
 - We conserve the image structure and local correlation instead of flattening.
 - We reduce the number of parameters needed by sharing them across the image.
- A convolutional layer is equivalent to a sparse dense layer with shared weights.

$$Image\ I = \begin{bmatrix} I_{11} & I_{12} & \dots & I_{1n} \\ I_{21} & I_{22} & \dots & I_{2n} \\ \dots & \dots & \dots & \dots \\ I_{n1} & I_{n2} & \dots & I_{nn} \end{bmatrix};\ Kernel\ K = \begin{bmatrix} k_1 & k_2 \\ k_3 & k_4 \end{bmatrix};\ O = I*K \ \rightarrow O = \begin{bmatrix} I_{11} & I_{12} & \dots & I_{1n} \\ I_{21} & I_{22} & \dots & I_{2n} \\ \dots & \dots & \dots & \dots \\ I_{n1} & I_{n2} & \dots & I_{nn} \end{bmatrix} * \begin{bmatrix} k_1 & k_2 \\ k_3 & k_4 \end{bmatrix}$$

We can implement a convolution, O = I*K, using a dense layer multiplying the Flattened Image FI with a matrix A, as follows:

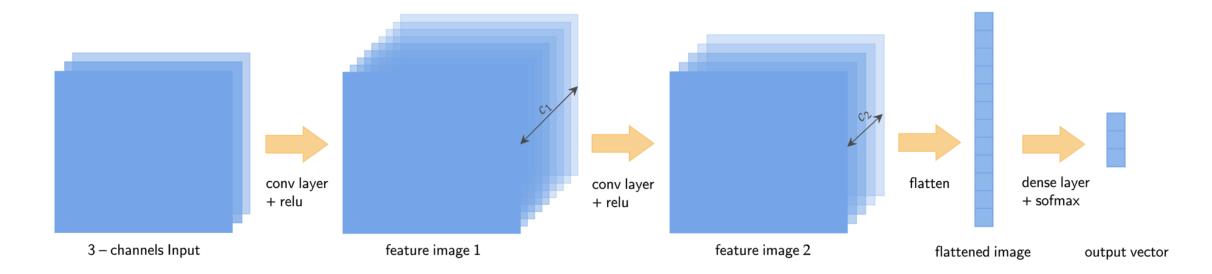
$$FI = \begin{bmatrix} I_{11} & I_{12} & \dots & I_{1n} & I_{21} & I_{22} & \dots & I_{2n} & \dots & I_{n1} & I_{n2} & \dots & I_{nn} \end{bmatrix}; A = \begin{bmatrix} k_1 & 0 & 0 & \dots & 0 \\ k_2 & k_1 & 0 & \dots & 0 \\ 0 & k_2 & k_1 & \dots & \dots \\ 0 & 0 & k_2 & \dots & k_1 \\ \dots & \dots & \dots & \dots & k_2 \\ k_3 & 0 & 0 & \dots & 0 \\ k_4 & k_3 & 0 & \dots & 0 \\ 0 & k_4 & k_3 & \dots & \dots \\ \dots & \dots & k_4 & \dots & k_3 \\ 0 & 0 & 0 & \dots & k_4 \end{bmatrix}$$

Outline

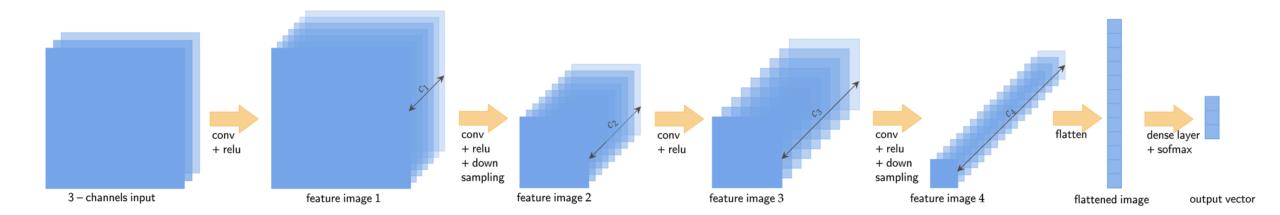
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- We presented how a convolutional layer works. Let's look at convolutional networks.
- Imagine we want to classify images into 3 classes, i.e. multiclass classification.
- We can replace the dense layers in a network with convolutional layers.
- However, we still need at the end a dense layer to apply the linear classifier.
- The convolutional "backbone" is used to extract useful features.
- Then on top we apply one or multiple dense layer.

- We need to flatten the last convolutional layer in order to apply a dense layer.
- However, we get to the same initial problem with a huge dense layer!
- Thus, we should gradually reduce the feature size before flattening.



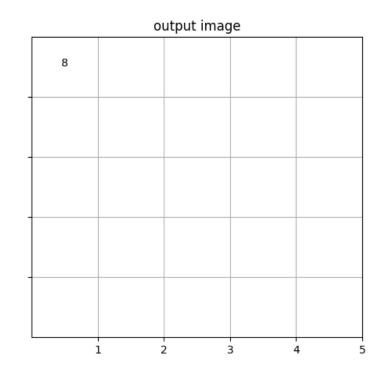
- One way to gradually reduce the size if to use strided convolutions (stride > 1).
- With strides=2, we will reduce at each stage the spatial dimension.
- This will also make also the convolution faster.



- Another way to reduce the spatial dimensions is pooling.
- Pooling compute an average, or max value, in a kernel size. It is not learnable.

pooling								
8	2	1	4	1	9	1	9	1
1	4	4	3	3	3	4	9	6
5	4	4	4	6	9	7	1	3
2	9	3	5	6	7	7	4	9
6	5	6	4	9	7	2	0	1
2	2	2	5	5	3	4	8	9
9	5	0	1	3	5	3	4	7
7	3	4	9	5	4	6	3	2
6	7	4	7	5	2	3	4	6
	1 2	2 3			5 6	5 7	7 8	3 9

nooling



- Downsampling, whether with pooling or striding, is a common pattern.
- It is also related to an important notion called "receptive field" covered later.
- An alternative to flattening is computing an average (or max) value per channel.
- It can help reduce the size of the output vector before the dense layer.
- We can develop our first CNN to classify images.

```
def cnn with pooling(input image, flatten=True):
 input image: a 4D array of type float32 and shape (batch, height, width, channels)
 return: a 2D array of type float32 and shape (batch, 3)
 # Conv2D is a class that implements the convolutional layer
 # filter is the number of filters (or feature maps) to use
  conv 1 = tf.keras.layers.Conv2D(filters=8, kernel size=(3, 3), strides=(1, 1), padding='same', activation='relu')(input image)
  conv 2 = tf.keras.layers.Conv2D(filters=8, kernel size=(3, 3), strides=(1, 1), padding='same', activation='relu')(conv 1)
  downsample 1 = tf.keras.layers.MaxPool2D(pool size=(2, 2), strides=(2, 2), padding='same')(conv 2)
  conv 3 = tf.keras.layers.Conv2D(filters=16, kernel size=(3, 3), strides=(1, 1), padding='same', activation='relu')(downsample 1)
  conv 4 = tf.keras.layers.Conv2D(filters=16, kernel size=(3, 3), strides=(1, 1), padding='same', activation='relu')(conv 3)
  downsample 2 = tf.keras.layers.MaxPool2D(pool size=(2, 2), strides=(2, 2), padding='same')(conv 4)
  conv = tf.keras.layers.Conv2D(filters=32, kernel size=(3, 3), strides=(1, 1), padding='same', activation='relu')(downsample 2)
  conv 6 = tf.keras.layers.Conv2D(filters=32, kernel size=(3, 3), strides=(1, 1), padding='same', activation='relu')(conv 5)
  downsample 3 = tf.keras.layers.MaxPool2D(pool size=(2, 2), strides=(2, 2), padding='same')(conv 6)
  if flatten:
   vector = tf.keras.layers.Flatten()(downsample 3)
  else:
   vector = tf.keras.layers.GlobalAveragePooling2D()(downsample 3)
 dense = tf.keras.layers.Dense(units=3)(vector)
  output = tf.nn.softmax(dense, axis=-1)
  return output
```

• Let's compute the parameters number and output shape with an image of size

(100, 100, 3).

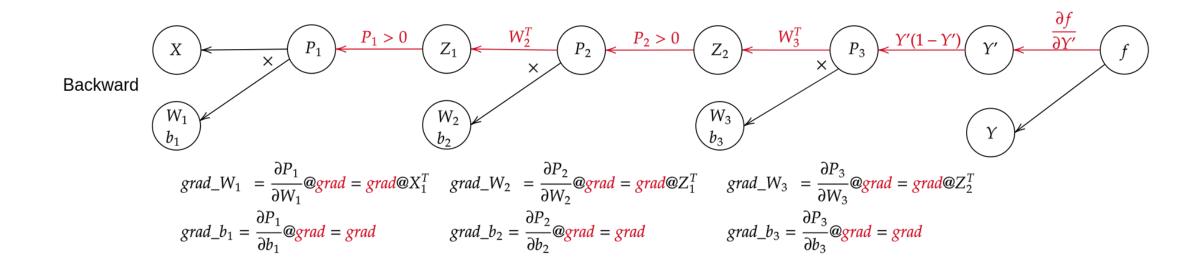
Layer Name	Number of Parameters	Output Shape		
conv_1	3*3*3*8 + 8 = 224	(100, 100, 8)		
conv_2	3*3*8*8 + 8 = 584	(100, 100, 8)		
downsample_1	0	(50, 50, 8)		
conv_3	3*3*8*16 + 16 = 1168	(50, 50, 16)		
conv_4	3*3*16*16 + 16 = 2320	(50, 50, 16)		
downsample_2	0	(25, 25, 16)		
conv_5	3*3*16*32 + 32 = 4640	(25, 25, 32)		
conv_6	3*3*32*32 + 32 = 9248	(25, 25, 32)		
downsample_3	0	(13, 13, 32)		
vector	0	(5408,)		
dense	5408 * 3 = 16227	(3,)		
output	0	(3,)		
Total	34411	-		

- There are 2 things to remark:
 - 1. The convolutional layer parameters depend only the number of channels, not the spatial size.
 - 2. Convolutional layer are not very parameter-intensive, so we can add many of them w/o overfitting.
- The second point implies that we can develop deep networks (a lot of layers).
- However, some problems may arise:
 - Gradient vanishing or explosion: the gradients become too small or too big.
 - Slow convergence and instability: training becomes slow and harder to reach good local minima.
- To solve gradient explosion we can clip it by a maximum value or norm.
- The other problems need more advanced techniques to solve them.

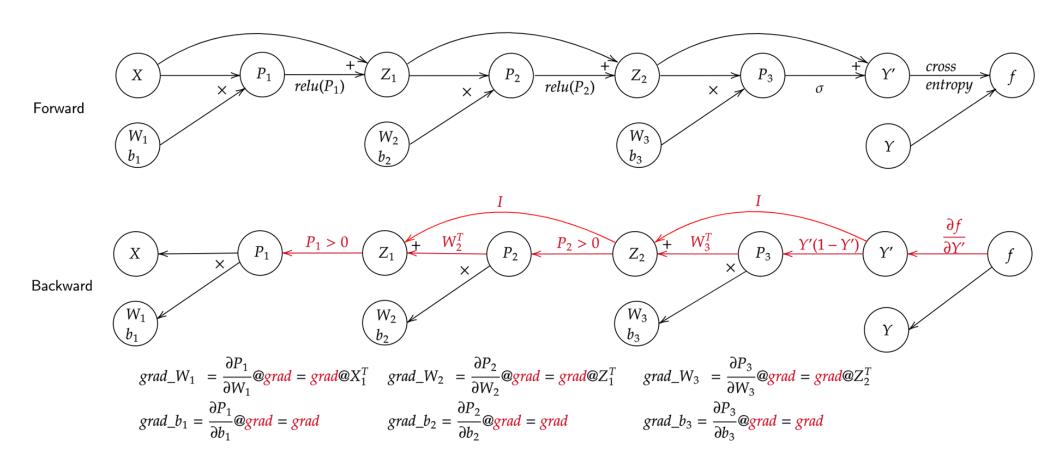
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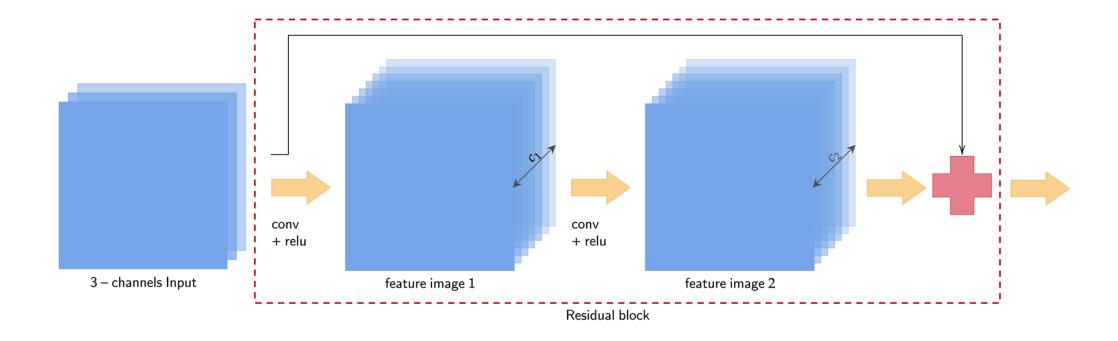
- Residual connections were proposed to solve the problem of vanishing gradients.
- With more depth, gradient we may become too small in early layers, even with ReLU.
- Gradient is accumulated from last to first layer and thus can become too small.



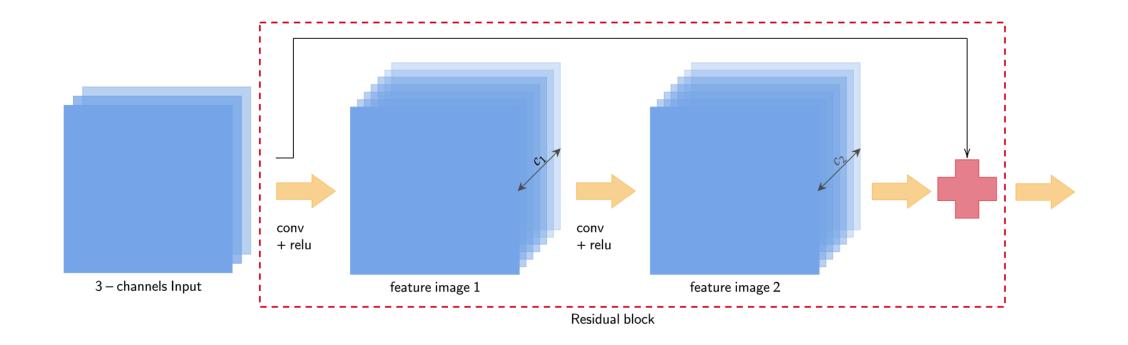
Residual connections ensure a useful gradient at all layers by adding shortcuts.



- It is called residual because output = f(input) + input → f(input) = output input.
- We are learning the difference between output and input.



- We can have multiple convolutions inside a residual block.
- But how can we sum the input with the output if they don't have the same shape?



- The output block has more channel than the input in general (so we can't sum).
- In addition, we may down-sample so the spatial resolution may change as well.
- Thus, instead of directly adding the input, we can add a convolution layer in between.
- We can create a function to create a residual block. It should take:
 - A list of kernel size for each convolution in the block.
 - A list of filters for each convolution in the block.
 - A list of strides for each convolution in the block.

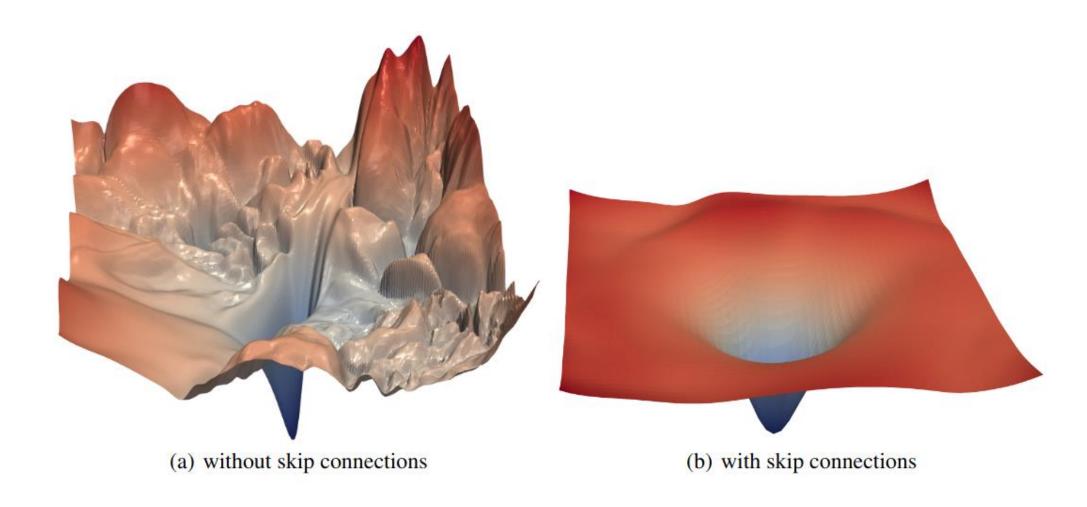
Residual Connections

```
def residual block v1(x, feature maps, strides, kernel size):
  0.00
 x: a 4D array of type float32 and shape (batch, height, width, channels)
  feature maps: list containing the number of feature maps to use
  strides: list containing the strides to use
  kernel size: list containing the kernel size to use
  return: a 4D array of type float32
  residual = x
 # check if the output dimension will be different from the input.
 # we always suppose that the output channels are bigger than the input channels
 if 2 in strides or x.shape[-1] < feature maps[-1]:</pre>
    residual = tf.keras.layers.Conv2D(
           filters=feature maps[-1], kernel size=1,
           strides=max(strides), padding='same', activation='relu')(x)
  conv = x
  for i in range(len(feature maps)):
    conv = tf.keras.layers.Conv2D(
           filters=feature maps[i], kernel size=kernel size[i],
           strides=strides[i], padding='same', activation='relu')(conv)
  return tf.keras.layers.Add()([conv, residual])
```

Residual Connections

- Residual connections solve gradient vanishing problem, and enable deep networks.
- However, deep networks means in general high non-convexity.
- How is it possible to train deep network with gradient descent?
- To understand what happens inside, researcher developed a visualization technique.
- It is called loss landscape visualization and its purpose is to visualize how the loss change w.r.t parameters (in a reduced dimension like 3D).
- As we will see, residual connections make the loss smoother and easier to optimize.

Residual Connections



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- Batch normalization is proposed to stabilize the training and enable faster convergence.
- It enables the use of big learning rate without the risk of divergence and instability.
- Despite it is great success, we still don't have a big understanding why it works.
- It is used with convolutional layers, but also with dense layers as well.
- Every batch normalization layer is composed of 2 steps:
 - Normalization: eliminate the mean and variance of a batch of input.
 - Re-Parameterization: assign new mean and variance that are learnable.

- It is summarized in the following equations.
- Note that it is applied for a batch of images not single one.

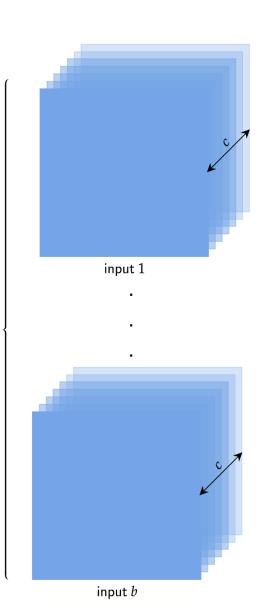
for a given hidden layer, we denote its output as H_B ; H_B of shape (B, ...) is a batch of activation, each activation is denoted H_i of shape (1, ...)

Batch Normalization performs the following:

Normalization :
$$\widehat{H}_B = \frac{H_B - \mu_B}{\sqrt{\sigma_B^2 + \epsilon}}$$
; $\mu_B = \frac{1}{B} \sum_i^B H_i$; $\sigma_B^2 = \frac{1}{B} \sum_i^B (H_i - \mu_B)^2$; μ_B and σ_B has the same shape as H_i ; ϵ small value for stability

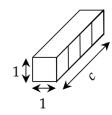
 $\text{Re-Parameterization: } \widetilde{H}_{\text{B}} = \gamma \odot \widehat{H}_{\text{B}} + \beta \text{ ; } \gamma \text{ new learnable standard deviation, } \beta \text{ new learnable mean}$

- For convolutional layers, we compute μ and σ over (batch, height, width).
- But multiple questions arise:
 - How to apply it during inference with a single image?
 - Inference will vary based on the batch size.
- Thus, we use batch norm differently in training and inference.



batch of

b input



size of $\mu/\sigma/\gamma/\beta$ with convolutional layers

- Instead of computing μ and σ for a batch during inference, we will:
 - Estimate a moving mean and variance during training.
 - Apply it in the formula instead of μ and σ .
- Batch normalization doesn't change during training, we just compute an additional quantity.

Batch Normalization during training:

Normalization:
$$\widehat{H}_B = \frac{H_B - \mu_B}{\sqrt{\sigma_B^2 + \epsilon}}$$

Moving averages: $\mu_{moving} = \alpha \mu_{moving} + (1 - \alpha)\mu_B$; $\sigma_{moving} = \alpha \sigma_{moving} + (1 - \alpha)\sigma_B$; $\alpha \in [0, 1)$

$$\operatorname{Re}-\operatorname{Parameterization}:\ \widetilde{H}_{B}=\gamma\odot\widehat{H}_{B}+\beta$$

- Thus, in inference we used the moving μ and σ estimated during training.
- Like this, we can do inference with different batches and have the same results.

Batch Normalization during testing:

Normalization:
$$\widehat{H} = \frac{H - \mu_{moving}}{\sqrt{\sigma_{moving}^2 + \epsilon}}$$

 $\mathsf{Re}-\mathsf{Parameterization}: \ \widetilde{H}_{B}=\gamma \odot \widehat{H}_{B}+\beta$

- Some practical considerations when using batch normalization:
 - Batch norm came most of the time directly after a convolution operation, before activation.
 - If your batch size is too small, batch norm estimates will not be accurate.
 - If you are using multiple GPU, each GPU will have a different batch norm estimates.
 - Think to increase your learning rate with batch normalization (preferably with schedules).
- Other normalization techniques exist like Batch Renormalization and Layer Normalization.

We can write and reuse functions that create convolutional and dense blocks.

```
def conv bn relu(x, feature maps, strides, kernel size):
 x: a 4D array of type float32 and shape (batch, height, width, channels)
  feature maps: (int) number of feature maps to use
  strides: (int) the strides to use
  kernel size: (int) the kernel size to use
  return: a 4D array of type float32
  x = tf.keras.layers.Conv2D(
          filters=feature maps, kernel size=kernel size,
          strides=strides, padding='same')(x)
 x = tf.keras.layers.BatchNormalization()(x)
 x = tf.keras.layers.ReLU()(x)
  return x
def dense bn relu(x, units):
 x: a 2D array of type float32 and shape (batch, features)
 units: (int) number of hidden neurons
  x = tf.keras.layers.Dense(units)(x)
  x = tf.keras.layers.BatchNormalization()(x)
  x = tf.keras.layers.ReLU()(x)
  return x
```

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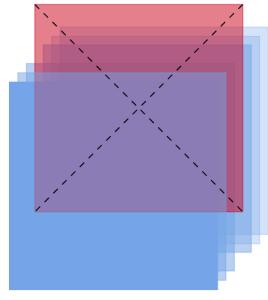
Dropout

- We can use dropout with convolutional layers as well to reduce overfitting.
- However, the standard dropout is not suitable for structured data like image.
- Randomly dropping may hurt spatially correlated information.
- Thus, we replace standard dropout with spatial dropout.
- In such case, we either drop or keep a total feature map (channel).
- This is known as Spatial Dropout 2D. There exist also different forms.

Dropout

We can implement now a residual block with batch norm and dropout.

```
def residual block(x, feature maps, strides, kernel size, drop rate):
 x: a 4D array of type float32 and shape (batch, height, width, channels)
  feature maps: list containing the number of feature maps to use
  strides: list containing the strides to use
  kernel size: list containing the kernel size to use
  drop rate: float, the proportion of feature maps to drop.
  return: a 4D array of type float32
  residual = x
  if 2 in strides or x.shape[-1] < feature maps[-1]:</pre>
    residual = conv bn relu(x, feature maps[-1], strides=max(strides), kernel size=1)
  if drop rate > 0:
   x = tf.keras.layers.SpatialDropout2D(drop rate)(x)
  for i in range(len(feature maps)):
   x = conv bn relu(x, feature maps[i], strides=strides[i], kernel size=kernel size[i])
  return tf.keras.layers.Add()([x, residual])
```



randomly drop a feature map

Outline

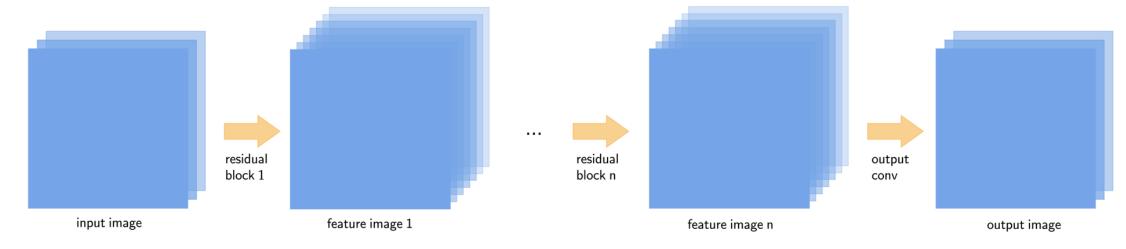
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Fully Convolutional Networks

- CNN can solve image classification and regression problems.
- However, we can imagine many applications that don't fall directly into the
- For instance, we can't solve image-to-image problems with a CNN.
- We need to design a network that takes an image and outputs an image.
- As there is no need for a linear classifier on top, there is no need for a dense layer.
- We can design a fully convolutional network (FCN).

Fully Convolutional Networks

Someone can imagine a network that looks like this:

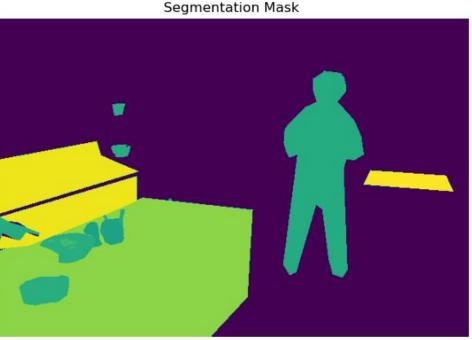


- As there is no need for a dense layer, there is no need for downsampling. True?
- Absolutely not!

Fully Convolutional Networks

- Downsampling reduces the computational time, but this is not the main reason.
- The reason is related to a notion called the network receptive field.
- Let's ask a question: how a local operation like convolution see the whole picture.

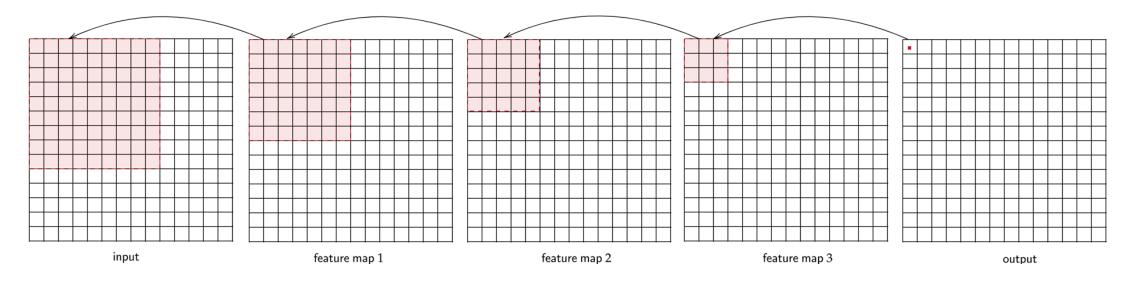




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 - Convolution in Neural Networks
 - Convolution vs Dense Layers
- Convolutional Neural Networks
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 - Batch normalization
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- Fully Convolutional Networks
 - Receptive Field
 - Encoder-Decoder Architecture
- Transfer Learning

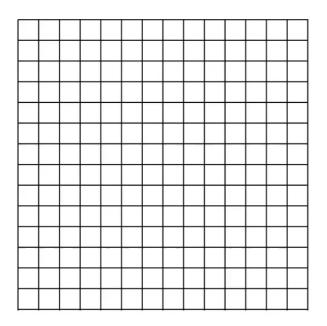
- Receptive field is a notion related to convolutional operations.
- It is the size of a region in the input that contributes to a single pixel in the output.

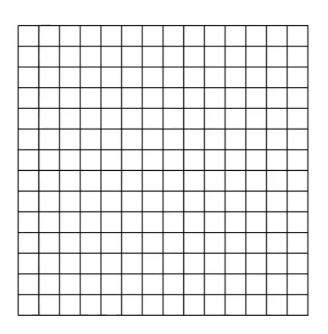


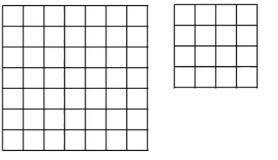
• Each pixel in a layer is connected to a 3x3 pixels in the previous layer.

- To increase the receptive field, we can:
 - Either increase the kernel sizes.
 - Either add more layers.
- Both propositions will increase the number of parameters and operations.
- We would like to have something more efficient.
- Here's come down-sampling into play.

• Down-sampling can exponentially increase the receptive field.







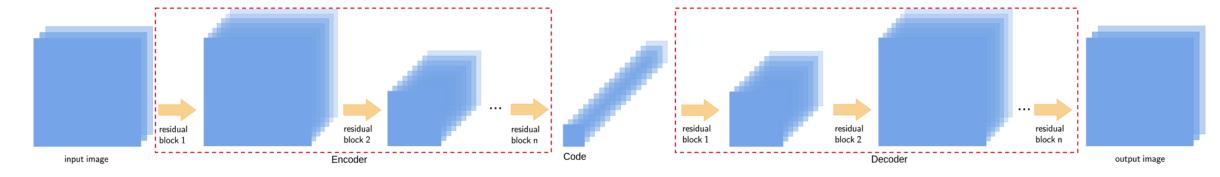


- It is a win-win situation because down-sampling means:
 - Faster training because the feature spatial size is reduced.
 - Bigger receptive field.
- Down-sampling can be implement with pooling or strided convolution.
- The receptive field can be bigger than the input image itself.
- Thus, image-to-image network uses down-sampling to increase their receptive field.
- However, to have an image in the output, we need to upsample. How?

Outline

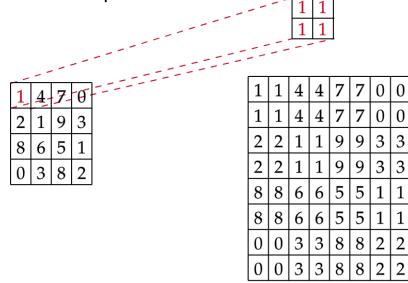
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Encoder-Decoder encodes an image into a "code" and then decode it back.



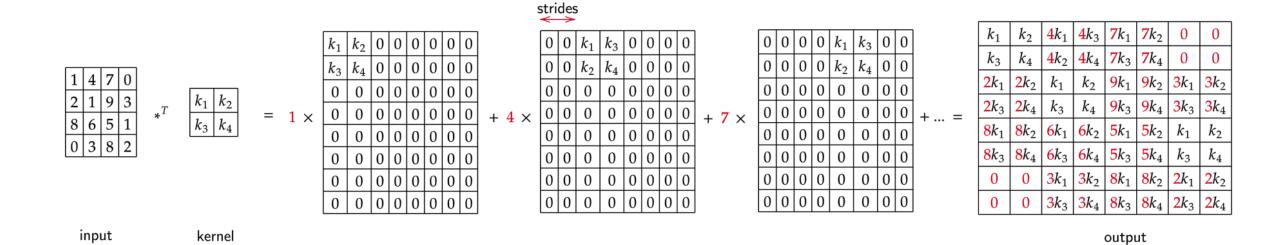
- We know how to downsample, but how to upsample?
- There is 2 ways:
 - Interpolation.
 - Transpose convolution.

- The most used interpolation in deep learning is nearest interpolation.
- To double the image size just repeat the same value in a 2x2 square.



- It is similar to the idea of down-sampling by pooling.
- Transpose convolution is more similar to strided convolution.

- Transpose convolution takes the same parameters as a normal convolution.
- It is usually used only for up-sampling, with a kernel of 2x2 and strides of 2.



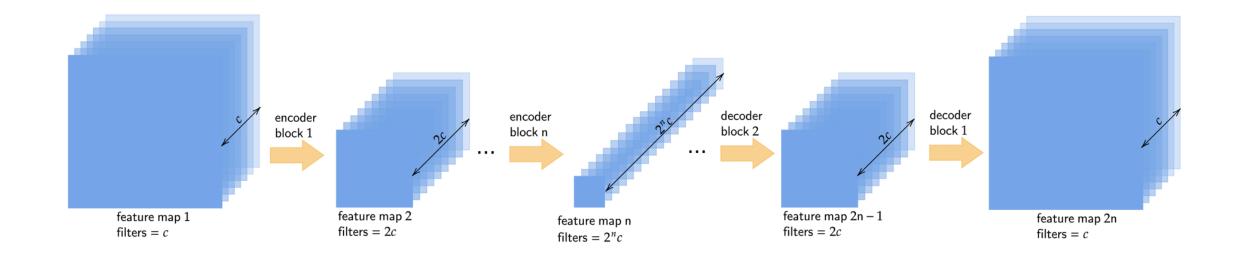
- A encoder residual block is just similar to the residual block of a CNN.
- We can down-sample using strided convolution (learnable) or pooling.

```
def encoder_residual_block(x, feature_maps, kernel_size, drop_rate, learnable_sampling):
    if learnable_sampling:
        x = conv_bn_relu(x, feature_maps[-1], strides=2, kernel_size=1)
    else:
        x = tf.keras.layers.MaxPool2D(padding='same')(x)
    residual = x
    if x.shape[-1] < feature_maps[-1]:
        additional_channels = feature_maps[-1] - x.shape[-1]
        paddings = [[0, 0], [0, 0], [0, 0], [0, additional_channels]]
        residual = tf.pad(residual, paddings)

if drop_rate > 0:
        x = tf.keras.layers.SpatialDropout2D(drop_rate)(x)
    for i in range(len(feature_maps)):
        x = conv_bn_relu(x, feature_maps[i], strides=1, kernel_size=kernel_size[i])
    return tf.keras.layers.Add()([x, residual])
```

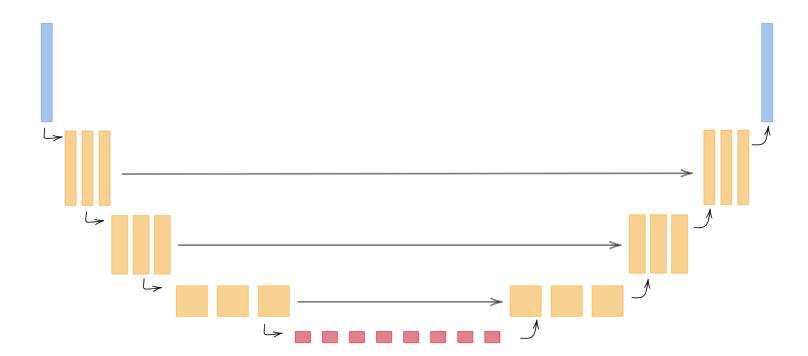
We can implement a decoder residual block similarly to the encoder block.

- Encoder-Decoder is computationally efficient, in addition to its big receptive field.
- A common practice is to double the filters each time we reduce the spatial size.

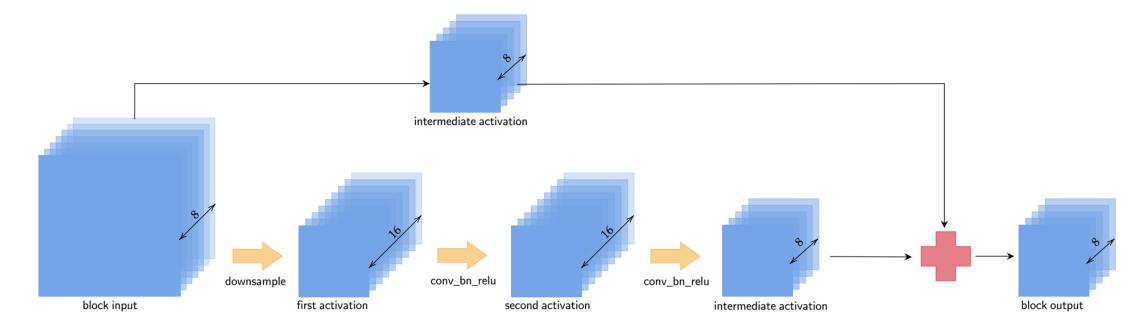


```
input shape = (320, 320, 3)
encoder fmaps = [[8, 8], [16, 16], [32, 32], [64, 64], [128, 128]]
encoder kernels = [[3, 3], [3, 3], [3, 3], [3, 3]]
decoder fmaps = [[128, 128], [64, 64], [32, 32], [16, 16], [8, 8]]
decoder kernels = [[3, 3], [3, 3], [3, 3], [3, 3]]
code fmaps = 256
code kernel = 5
code stride = 1
drop rate = 0
learnable sampling = False
input image = tf.keras.layers.Input(input shape)
encoded = input image
for i in range(len(encoder fmaps)):
 encoded = encoder residual block(encoded, encoder fmaps[i], encoder kernels[i], drop rate, learnable sampling)
code = conv bn relu(encoded, code fmaps, code stride, code kernel)
decoded = code
for i in range(len(encoder fmaps)):
 decoded = decoder residual block(decoded, decoder fmaps[i], decoder kernels[i], drop rate, learnable sampling)
output image = tf.keras.layers.Conv2D(filters=3, kernel size=3, strides=1, padding='same')(decoded)
model = tf.keras.Model(input image, output image)
model.summary()
```

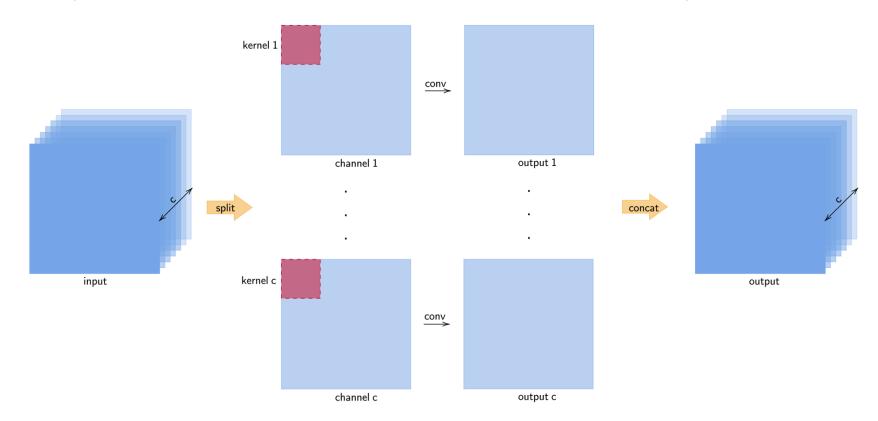
- Another development that improved the architecture performance is UNet.
- The idea is to connect encoder to the decoder to ease the reconstruction step.



- Some tricks were developed for limited resources environment like mobile phones.
- In a network called MobileNet, they introduced "inverted residual block".
- The goal is to reduce the number of parameters of a network.



- A special type of convolution was developed for limited resources environment.
- It is called depthwise convolution and it reduces the size of operations of a model.



- The number of parameters of a depthwise convolution equals $k_x * k_y * c$.
- We can add a multiplier d (how many filters to apply), thus it becomes: $k_x * k_y * c * d$.

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- Convolutional networks learn generic features about images.
- We extract different levels of details in different layers.
- It was shown empirically that early layers learn generic image characteristics (edge ...)
- On the other hand, later layers are more specialized about the task in hand.
- Thus, we can think of using knowledge learned from a dataset into another.
- This is known as transfer learning.

- It is useful in scenarios when we have limited data to learn from.
- In this case, we do the following:
 - We train (or use existing) model on a big and diverse dataset.
 - We finetune the model weights on the dataset of interest in hand.
- Those 2 datasets can be very different in practice.
- Finetuning means updating some or all model layers on the small dataset.
- In the literature, almost all new models are trained on ImageNet for benchmarking.
- ImageNet is a huge dataset for image classification (14M, with 22K classes).

- New developments use self-supervised learning to prepare pretrained models.
- Self-supervised learning aims to train an unsupervised dataset in a supervised manner.
- The idea is to augment each image to create a positive pair.
- All other images are supposed negative pairs.
- This helps to overcome the data scarcity and can enable deep learning for such cases.
- This also help in unsupervised application like clustering.
- We can use those network to output a meaningful representation for an image.

- Another result of this generic learning of convolution is to do multitask learning.
- A multitask network can predict different but related quantities (like age/gender).

