Reference

This is from the coursera deep learning series of Andrew NG.

Convolutional Neural Networks: Step by Step

Implement convolutional (CONV) and pooling (POOL) layers in numpy, including both forward propagation and (optionally) backward propagation.

Notation:

- Superscript [l] denotes an object of the l^{th} layer.
 - Example: $a^{[4]}$ is the 4^{th} layer activation. $W^{[5]}$ and $b^{[5]}$ are the 5^{th} layer parameters.
- Superscript (i) denotes an object from the i^{th} example.
 - Example: $\chi^{(i)}$ is the i^{th} training example input.
- Lowerscript i denotes the i^{th} entry of a vector.
 - Example: $a_i^{[l]}$ denotes the i^{th} entry of the activations in layer l, assuming this is a fully connected (FC) layer.
- n_H , n_W and n_C denote respectively the height, width and number of channels of a given layer. If you want to reference a specific layer l, you can also write $n_H^{[l]}$, $n_W^{[l]}$, $n_C^{[l]}$.
- $n_{H_{prev}}$, $n_{W_{prev}}$ and $n_{C_{prev}}$ denote respectively the height, width and **number of channels of the previous layer**. If referencing a specific layer l, this could also be denoted $n_H^{[l-1]}$, $n_W^{[l-1]}$.

1 - Packages

```
In [1]: import numpy as np
import h5py
import matplotlib.pyplot as plt

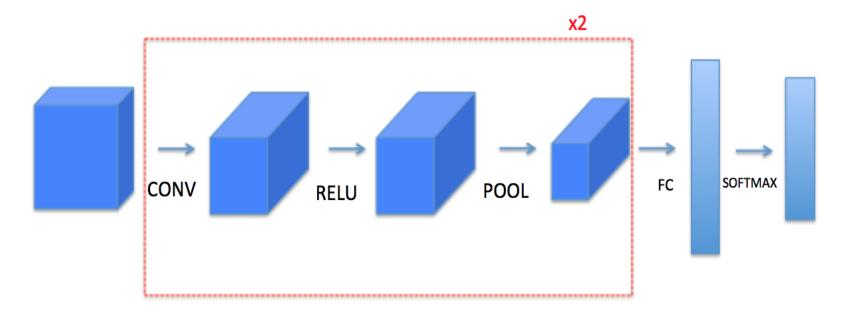
%matplotlib inline
   plt.rcParams['figure.figsize'] = (5.0, 4.0) # set default size of plots
   plt.rcParams['image.interpolation'] = 'nearest'
   plt.rcParams['image.cmap'] = 'gray'

%load_ext autoreload
%autoreload 2

np.random.seed(1)
```

2 - Outline of the Assignment

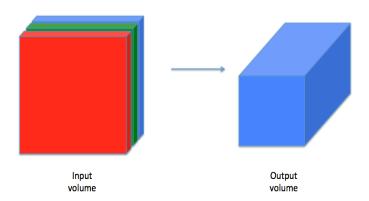
- Convolution functions, including:
 - Zero Padding
 - Convolve window
 - Convolution forward
 - Convolution backward (optional)
- Pooling functions, including:
 - Pooling forward
 - Create mask
 - Distribute value
 - Pooling backward (optional)



Note that for every forward function, there is its corresponding backward equivalent. Hence, at every step of your forward module you will store some parameters in a cache. These parameters are used to compute gradients during backpropagation.

3 - Convolutional Neural Networks

Although programming frameworks make convolutions easy to use, they remain one of the hardest concepts to understand in Deep Learning. A convolution layer transforms an input volume into an output volume of different size, as shown below.



In this part, you will build every step of the convolution layer. You will first implement two helper functions: one for zero padding and the other for computing the convolution function itself.

3.1 - Zero-Padding

Zero-padding adds zeros around the border of an image:

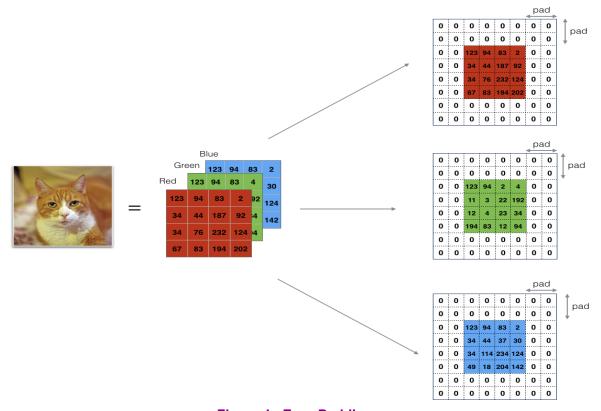


Figure 1 : Zero-Padding Image (3 channels, RGB) with a padding of 2.

The main benefits of padding are the following:

- It allows you to use a CONV layer without necessarily shrinking the height and width of the volumes. This is important for building deeper networks, since otherwise the height/width would shrink as you go to deeper layers. An important special case is the "same" convolution, in which the height/width is exactly preserved after one layer.
- It helps us keep more of the information at the border of an image. Without padding, very few values at the next layer would be affected by pixels as the edges of an image.

Exercise: Implement the following function, which pads all the images of a batch of examples X with zeros. Use np.pad (https://docs.scipy.org/doc/numpy/reference/generated/numpy.pad.html). Note if you want to pad the array "a" of shape (5, 5, 5, 5, 5) with pad = 1 for the 2nd dimension, pad = 3 for the 4th dimension and pad = 0 for the rest, you would do:

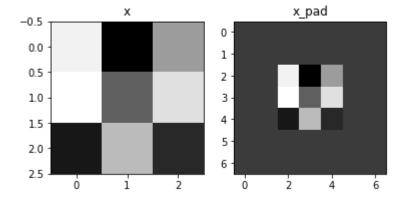
```
a = np.pad(a, ((0,0), (1,1), (0,0), (3,3), (0,0)), 'constant', constant_values = (..,..))
```

X below: -- python numpy array of shape (m, n_H, n_W, n_C) representing a batch of m images with height of n_H, width of n_W and number of channels of n_C

```
In [3]: np.random.seed(1)
        x = np.random.randn(4, 3, 3, 2)
        x pad = zero pad(x, 2)
        print ("x.shape =", x.shape)
        print ("x_pad.shape =", x_pad.shape)
        print ("x[1, 1] =", x[1, 1])
        print ("x_pad[1, 1] =", x_pad[1, 1])
        fig, axarr = plt.subplots(1, 2)
        axarr[0].set title('x')
        axarr[0].imshow(x[0,:,:,0])
        axarr[1].set title('x pad')
        axarr[1].imshow(x_pad[0,:,:,0])
        x.shape = (4, 3, 3, 2)
        x_{pad.shape} = (4, 7, 7, 2)
        x[1, 1] = [[ 0.90085595 -0.68372786]
         [-0.12289023 -0.93576943]
         [-0.26788808 0.53035547]]
        x_pad[1, 1] = [[0. 0.]]
         [0. 0.]
         [0. 0.]
```

Out[3]: <matplotlib.image.AxesImage at 0x2979b2bd828>

[0. 0.] [0. 0.] [0. 0.] [0. 0.]]



3.2 - Single step of convolution

In this part, implement a single step of convolution by applying the filter to a single position of the input. This will be used to build a convolutional unit, which:

- · Takes an input volume
- · Applies a filter at every position of the input
- · Outputs another volume (usually of different size)

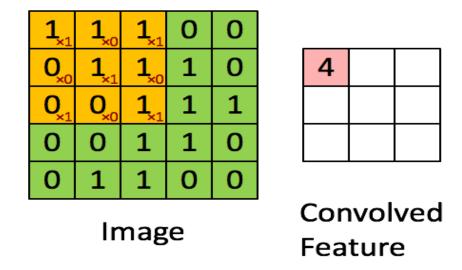


Figure 2: Convolution operation

with a filter of 2x2 and a stride of 1 (stride = amount you move the window each time you slide)

In a computer vision application, each value in the matrix on the left corresponds to a single pixel value, and we convolve a 3x3 filter with the image by multiplying its values element-wise with the original matrix, then summing them up. Here we implement a single step of convolution, corresponding to applying a filter to just one of the positions to get a single real-valued output.

Exercise: Implement conv_single_step(). Hint (https://docs.scipy.org/doc/numpy-1.13.0/reference/generated/numpy.sum.html).

```
In [4]: def conv_single_step(a_slice_prev, W, b):
    """
    Apply one filter defined by parameters W on a single slice (a_slice_prev) of the output activation of the previous layer.

Arguments:
    a_slice_prev -- slice of input data of shape (f, f, n_C_prev)
    W -- Weight parameters contained in a window - matrix of shape (f, f, n_C_prev)
    b -- Bias parameters contained in a window - matrix of shape (1, 1, 1)

Returns:
    Z -- a scalar value, result of convolving the sliding window (W, b) on a slice x of the input data
    """

# Element-wise product between a_slice and W. Add bias.
s = np.multiply(a_slice_prev, W) + b
# Sum over all entries of the volume s
Z = np.sum(s)
    return Z
```

```
In [5]: np.random.seed(1)
    a_slice_prev = np.random.randn(4, 4, 3)
    W = np.random.randn(4, 4, 3)
    b = np.random.randn(1, 1, 1)

Z = conv_single_step(a_slice_prev, W, b)
    print("Z =", Z)
```

Z = -23.16021220252078

3.3 - Convolutional Neural Networks - Forward pass

In the forward pass, you will take many filters and convolve them on the input. Each 'convolution' gives you a 2D matrix output. You will then stack these outputs to get a 3D volume:

0:36 / 0:36

Exercise: Implement the function below to convolve the filters W on an input activation A_prev. This function takes as input A_prev, the activations output by the previous layer (for a batch of m inputs), F filters/weights denoted by W, and a bias vector denoted by b, where each filter has its own (single) bias. Finally you also have access to the hyperparameters dictionary which contains the stride and the padding.

Hint:

1. To select a 2x2 slice at the upper left corner of a matrix "a_prev" (shape (5,5,3)), you would do:

This will be useful when you will define a_slice_prev below, using the start/end indexes you will define.

2. To define a_slice you will need to first define its corners vert_start, vert_end, horiz_start and horiz_end. This figure may be helpful for you to find how each of the corner can be defined using h, w, f and s in the code below.

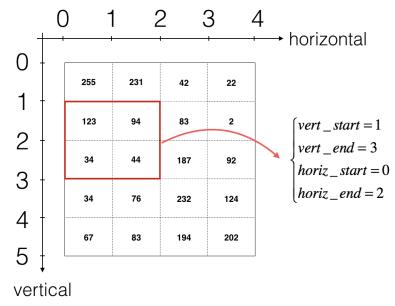


Figure 3: Definition of a slice using vertical and horizontal start/end (with a 2x2 filter)

This figure shows only a single channel.

Reminder: The formulas relating the output shape of the convolution to the input shape is:

$$n_{H} = \lfloor \frac{n_{H_{prev}} - f + 2 \times pad}{stride} \rfloor + 1$$

$$n_{W} = \lfloor \frac{n_{W_{prev}} - f + 2 \times pad}{stride} \rfloor + 1$$

 n_C = number of filters used in the convolution

For this exercise, we won't worry about vectorization, and will just implement everything with for-loops.

```
In [6]: | def conv forward(A prev, W, b, hparameters):
            Implements the forward propagation for a convolution function
            Arguments:
            A prev -- output activations of the previous layer, numpy array of shape (m, n H prev, n W prev, n C prev)
            W -- Weights, numpy array of shape (f, f, n C prev, n C)
            b -- Biases, numpy array of shape (1, 1, 1, n C)
            hparameters -- python dictionary containing "stride" and "pad"
            Returns:
            Z -- conv output, numpy array of shape (m, n H, n W, n C)
            cache -- cache of values needed for the conv backward() function
            # Retrieve dimensions from A prev's shape
            (m, n H prev, n W prev, n C prev) = A prev.shape
            # Retrieve dimensions from W's shape
            (f, f, n C prev, n C) = W.shape
            # Retrieve information from "hparameters"
            stride = hparameters['stride']
            pad = hparameters['pad']
            # Compute the dimensions of the CONV output volume using the formula given above. Hint: use int() to floor.
            n H = int((n H prev - f + 2 * pad) / stride) + 1
            n W = int((n W prev - f + 2 * pad) / stride) + 1
            # Initialize the output volume Z with zeros.
            Z = np.zeros((m, n H, n W, n C))
            # Create A prev pad by padding A prev
            A prev pad = zero pad(A prev, pad)
            for i in range(m):
                                                                # loop over the batch of training examples
                a prev pad = A prev pad[i]
                                                                # Select ith training example's padded activation
                for h in range(n H):
                                                                # loop over vertical axis of the output volume
                    for w in range(n_W):
                                                               # loop over horizontal axis of the output volume
                                                                # loop over channels (= #filters) of the output volume
                        for c in range(n C):
                            vert start = h * stride
                            vert end = vert start + f
```

```
horiz_start = w * stride
horiz_end = horiz_start + f

# Use the corners to define the (3D) slice of a_prev_pad (See Hint above the cell).
a_slice_prev = a_prev_pad[vert_start:vert_end, horiz_start:horiz_end, :]
# Convolve the (3D) slice with the correct filter W and bias b, to get back one output neuror
Z[i, h, w, c] = conv_single_step(a_slice_prev, W[...,c], b[...,c])

# Making sure your output shape is correct
assert(Z.shape == (m, n_H, n_W, n_C))

# Save information in "cache" for the backprop. If using tensorflow, then cache is not necessary.
cache = (A_prev, W, b, hparameters)

return Z, cache
```

```
Z's mean = 0.15585932488906465 cache conv[0][1][2][3] = [-0.20075807   0.18656139   0.41005165]
```

Finally, CONV layer should also contain an activation, in which case we would add the following line of code:

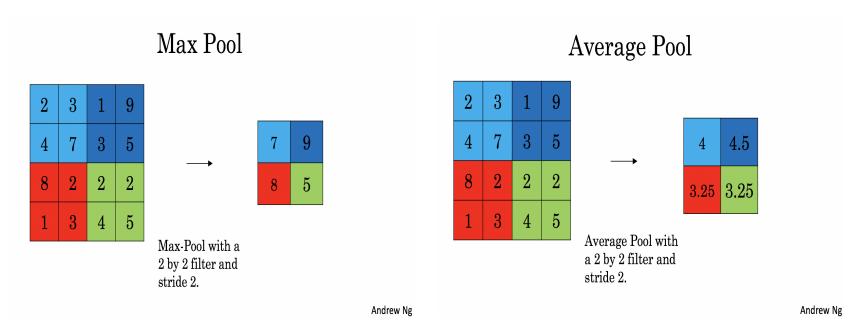
```
# Convolve the window to get back one output neuron
Z[i, h, w, c] = ...
# Apply activation
A[i, h, w, c] = activation(Z[i, h, w, c])
```

You don't need to do it here.

4 - Pooling layer

The pooling (POOL) layer reduces the height and width of the input. It helps reduce computation, as well as helps make feature detectors more invariant to its position in the input. The two types of pooling layers are:

- Max-pooling layer: slides an (f, f) window over the input and stores the max value of the window in the output.
- Average-pooling layer: slides an (f, f) window over the input and stores the average value of the window in the output.



These pooling layers have no parameters for backpropagation to train. However, they have hyperparameters such as the window size f. This specifies the height and width of the fxf window you would compute a max or average over.

4.1 - Forward Pooling

Now, you are going to implement MAX-POOL and AVG-POOL, in the same function.

Exercise: Implement the forward pass of the pooling layer. Follow the hints in the comments below.

Reminder: As there's no padding, the formulas binding the output shape of the pooling to the input shape is:

$$n_{H} = \lfloor \frac{n_{H_{prev}} - f}{stride} \rfloor + 1$$

$$n_{W} = \lfloor \frac{n_{W_{prev}} - f}{stride} \rfloor + 1$$

```
In [8]: | def pool_forward(A_prev, hparameters, mode = "max"):
            Implements the forward pass of the pooling layer
            Arguments:
            A prev -- Input data, numpy array of shape (m, n H prev, n W prev, n C prev)
            hparameters -- python dictionary containing "f" and "stride"
            mode -- the pooling mode you would like to use, defined as a string ("max" or "average")
            Returns:
            A -- output of the pool layer, a numpy array of shape (m, n H, n W, n C)
            cache -- cache used in the backward pass of the pooling layer, contains the input and hparameters
            # Retrieve dimensions from the input shape
            (m, n H prev, n W prev, n C prev) = A prev.shape
            # Retrieve hyperparameters from "hparameters"
            f = hparameters["f"]
            stride = hparameters["stride"]
            # Define the dimensions of the output
            n H = int(1 + (n H prev - f) / stride)
            n W = int(1 + (n W prev - f) / stride)
            n C = n C prev
            # Initialize output matrix A
            A = np.zeros((m, n H, n W, n C))
            for i in range(m):
                                                         # loop over the training examples
                for h in range(n_H):
                                                       # loop on the vertical axis of the output volume
                                                        # loop on the horizontal axis of the output volume
                    for w in range(n W):
                                                         # loop over the channels of the output volume
                        for c in range (n C):
                            # Find the corners of the current "slice" (≈4 lines)
                            vert start = h * stride
                            vert end = vert start + f
                            horiz start = w * stride
                            horiz end = horiz start + f
                            # Use the corners to define the current slice on the ith training example of A prev, channel
```

```
a prev slice = A prev[i, vert start:vert end, horiz start:horiz end, c]
                             # Compute the pooling operation on the slice. Use an if statment to differentiate the modes.
                             if mode == "max":
                                 A[i, h, w, c] = np.max(a_prev_slice)
                             elif mode == "average":
                                 A[i, h, w, c] = np.mean(a prev slice)
            # Store the input and hparameters in "cache" for pool backward()
            cache = (A prev, hparameters)
            # Making sure your output shape is correct
            assert(A.shape == (m, n H, n W, n C))
            return A, cache
In [9]: np.random.seed(1)
        A prev = np.random.randn(2, 4, 4, 3)
        hparameters = {"stride" : 1, "f": 4}
         A, cache = pool forward(A prev, hparameters)
         print("mode = max")
        print("A =", A)
         print()
        A, cache = pool forward(A prev, hparameters, mode = "average")
        print("mode = average")
        print("A =", A)
        mode = max
        A = [[[1.74481176 \ 1.6924546 \ 2.10025514]]]
         [[[1.19891788 1.51981682 2.18557541]]]]
        mode = average
        A = [[[-0.09498456 \quad 0.11180064 \quad -0.14263511]]]
         [[[-0.09525108 0.28325018 0.33035185]]]
```

5 - Backpropagation in convolutional neural networks (OPTIONAL / UNGRADED)

In modern deep learning frameworks, you only have to implement the forward pass, and the framework takes care of the backward pass, so most deep learning engineers don't need to bother with the details of the backward pass. The backward pass for convolutional networks is complicated. If you wish however, you can work through this optional portion of the notebook to get a sense of what backprop in a convolutional network looks like.

When in an earlier course you implemented a simple (fully connected) neural network, you used backpropagation to compute the derivatives with respect to the cost to update the parameters. Similarly, in convolutional neural networks you can to calculate the derivatives with respect to the cost in order to update the parameters. The backprop equations are not trivial and we did not derive them in lecture, but we briefly presented them below.

5.1 - Convolutional layer backward pass

Let's start by implementing the backward pass for a CONV layer.

5.1.1 - Computing dA:

This is the formula for computing dA with respect to the cost for a certain filter W_c and a given training example:

$$dA + = \sum_{h=0}^{n_H} \sum_{w=0}^{n_W} W_c \times dZ_{hw}$$
 (1)

Where W_c is a filter and dZ_{hw} is a scalar corresponding to the gradient of the cost with respect to the output of the conv layer Z at the hth row and wth column (corresponding to the dot product taken at the ith stride left and jth stride down). Note that at each time, we multiply the the same filter W_c by a different dZ when updating dA. We do so mainly because when computing the forward propagation, each filter is dotted and summed by a different a_slice. Therefore when computing the backprop for dA, we are just adding the gradients of all the a slices.

In code, inside the appropriate for-loops, this formula translates into:

5.1.2 - Computing dW:

This is the formula for computing dW_c (dW_c is the derivative of one filter) with respect to the loss:

$$dW_c + = \sum_{h=0}^{n_H} \sum_{w=0}^{n_W} a_{slice} \times dZ_{hw}$$
 (2)

Where a_{slice} corresponds to the slice which was used to generate the acitivation Z_{ij} . Hence, this ends up giving us the gradient for W with respect to that slice. Since it is the same W, we will just add up all such gradients to get dW.

In code, inside the appropriate for-loops, this formula translates into:

$$dW[:,:,:,c] += a slice * dZ[i, h, w, c]$$

5.1.3 - Computing db:

This is the formula for computing db with respect to the cost for a certain filter W_c :

$$db = \sum_{h} \sum_{w} dZ_{hw} \tag{3}$$

As you have previously seen in basic neural networks, db is computed by summing dZ. In this case, you are just summing over all the gradients of the conv output (Z) with respect to the cost.

In code, inside the appropriate for-loops, this formula translates into:

$$db[:,:,:,c] += dZ[i, h, w, c]$$

Exercise: Implement the conv_backward function below. You should sum over all the training examples, filters, heights, and widths. You should then compute the derivatives using formulas 1, 2 and 3 above.

```
In [10]: def conv backward(dZ, cache):
             Implement the backward propagation for a convolution function
             Arguments:
             dZ -- gradient of the cost with respect to the output of the conv layer (Z), numpy array of shape (m, n H, n
             cache -- cache of values needed for the conv backward(), output of conv forward()
             Returns:
             dA prev -- gradient of the cost with respect to the input of the conv layer (A prev),
                        numpy array of shape (m, n_H_prev, n_W_prev, n_C_prev)
             dW -- gradient of the cost with respect to the weights of the conv layer (W)
                   numpy array of shape (f, f, n_C_prev, n_C)
             db -- gradient of the cost with respect to the biases of the conv layer (b)
                   numpy array of shape (1, 1, 1, n C)
             0.00
             # Retrieve information from "cache"
             (A prev, W, b, hparameters) = cache
             # Retrieve dimensions from A prev's shape
             (m, n H prev, n W prev, n C prev) = A prev.shape
             # Retrieve dimensions from W's shape
             (f, f, n C prev, n C) = W.shape
             # Retrieve information from "hparameters"
             stride = hparameters["stride"]
             pad = hparameters["pad"]
             # Retrieve dimensions from dZ's shape
             (m, n H, n W, n C) = dZ.shape
             # Initialize dA prev. dW, db with the correct shapes
             dA_prev = np.zeros((m, n_H_prev, n_W_prev, n_C_prev))
             dW = np.zeros((f, f, n C prev, n C))
             db = np.zeros((1, 1, 1, n C))
             # Pad A prev and dA prev
             A prev pad = zero pad(A prev, pad)
             dA prev pad = zero pad(dA prev, pad)
```

```
for i in range(m):
                                        # loop over the training examples
   # select ith training example from A prev pad and dA prev pad
   a prev pad = A_prev_pad[i]
   da prev pad = dA prev pad[i]
   for h in range(n H):
                                      # loop over vertical axis of the output volume
       for w in range(n W):
                                      # loop over horizontal axis of the output volume
                                        # loop over the channels of the output volume
           for c in range(n C):
               # Find the corners of the current "slice"
               vert start = h * stride
               vert end = vert start + f
               horiz start = w * stride
               horiz end = horiz start + f
               # Use the corners to define the slice from a prev pad
               a slice = a prev pad[vert start:vert end, horiz start:horiz end, :]
               # Update gradients for the window and the filter's parameters using the code formulas given
               da prev pad[vert start:vert end, horiz start:horiz end, :] += W[:,:,:,c] * dZ[i, h, w, c]
               dW[:,:,:,c] += a slice * dZ[i, h, w, c]
               db[:,:,:,c] += dZ[i, h, w, c]
   # Set the ith training example's dA prev to the unpaded da prev pad (Hint: use X[pad:-pad, pad:-pad, :])
   dA prev[i, :, :] = da prev pad[pad:-pad, pad:-pad, :]
# Making sure your output shape is correct
assert(dA prev.shape == (m, n H prev, n W prev, n C prev))
return dA prev, dW, db
```

```
dA_mean = 9.608990675868995
dW_mean = 10.581741275547566
db mean = 76.37106919563735
```

5.2 Pooling layer - backward pass

Next, let's implement the backward pass for the pooling layer, starting with the MAX-POOL layer. Even though a pooling layer has no parameters for backprop to update, you still need to backpropagation the gradient through the pooling layer in order to compute gradients for layers that came before the pooling layer.

5.2.1 Max pooling - backward pass

Before jumping into the backpropagation of the pooling layer, you are going to build a helper function called create mask from window() which does the following:

$$X = \begin{bmatrix} 1 & 3 \\ 4 & 2 \end{bmatrix} \quad \rightarrow \quad M = \begin{bmatrix} 0 & 0 \\ 1 & 0 \end{bmatrix} \tag{4}$$

As you can see, this function creates a "mask" matrix which keeps track of where the maximum of the matrix is. True (1) indicates the position of the maximum in X, the other entries are False (0). You'll see later that the backward pass for average pooling will be similar to this but using a different mask.

Exercise: Implement create_mask_from_window() . This function will be helpful for pooling backward. Hints:

- np.max()() may be helpful. It computes the maximum of an array.
- If you have a matrix X and a scalar x: A = (X == x) will return a matrix A of the same size as X such that:

$$A[i,j] = True if X[i,j] = x$$

 $A[i,j] = False if X[i,j] != x$

• Here, you don't need to consider cases where there are several maxima in a matrix.

```
In [12]: def create_mask_from_window(x):
    """
    Creates a mask from an input matrix x, to identify the max entry of x.

Arguments:
    x -- Array of shape (f, f)

Returns:
    mask -- Array of the same shape as window, contains a True at the position corresponding to the max entry of """

mask = x == np.max(x)
    return mask
```

Why do we keep track of the position of the max? It's because this is the input value that ultimately influenced the output, and therefore the cost. Backprop is computing gradients with respect to the cost, so anything that influences the ultimate cost should have a non-zero gradient. So, backprop will "propagate" the gradient back to this particular input value that had influenced the cost.

5.2.2 - Average pooling - backward pass

In max pooling, for each input window, all the "influence" on the output came from a single input value--the max. In average pooling, every element of the input window has equal influence on the output. So to implement backprop, you will now implement a helper function that reflects this.

For example if we did average pooling in the forward pass using a 2x2 filter, then the mask you'll use for the backward pass will look like:

$$dZ = 1 \quad \rightarrow \quad dZ = \begin{bmatrix} 1/4 & 1/4 \\ 1/4 & 1/4 \end{bmatrix} \tag{5}$$

This implies that each position in the dZ matrix contributes equally to output because in the forward pass, we took an average.

Exercise: Implement the function below to equally distribute a value dz through a matrix of dimension shape. <u>Hint (https://docs.scipy.org/doc/numpy-1.13.0/reference/generated/numpy.ones.html)</u>

```
Distributes the input value in the matrix of dimension shape
             Arguments:
             dz -- input scalar
             shape -- the shape (n H, n W) of the output matrix for which we want to distribute the value of dz
             Returns:
             a -- Array of size (n H, n W) for which we distributed the value of dz
             # Retrieve dimensions from shape (≈1 line)
             (n H, n W) = shape
             # Compute the value to distribute on the matrix (≈1 line)
             average = dz / (n H * n W)
             # Create a matrix where every entry is the "average" value (≈1 line)
             a = np.ones(shape) * average
             return a
In [15]: a = distribute value(2, (2,2))
         print('distributed value =', a)
         distributed value = [[0.5 0.5]
          [0.5 0.5]]
```

5.2.3 Putting it together: Pooling backward

In [14]: | def distribute value(dz, shape):

You now have everything you need to compute backward propagation on a pooling layer.

Exercise: Implement the pool_backward function in both modes ("max" and "average"). You will once again use 4 for-loops (iterating over training examples, height, width, and channels). You should use an if/elif statement to see if the mode is equal to 'max' or 'average' . If it is equal to 'average' you should use the distribute_value() function you implemented above to create a matrix of the same shape as a_slice . Otherwise, the mode is equal to 'max', and you will create a mask with create_mask_from_window() and multiply it by the corresponding value of dZ.

```
In [16]:
         def pool backward(dA, cache, mode = "max"):
             Implements the backward pass of the pooling layer
             Arguments:
             dA -- gradient of cost with respect to the output of the pooling layer, same shape as A
             cache -- cache output from the forward pass of the pooling layer, contains the layer's input and hparameters
             mode -- the pooling mode you would like to use, defined as a string ("max" or "average")
             Returns:
             dA prev -- gradient of cost with respect to the input of the pooling layer, same shape as A prev
             # Retrieve information from cache (≈1 line)
             (A prev, hparameters) = cache
             # Retrieve hyperparameters from "hparameters" (≈2 lines)
             stride = hparameters["stride"]
             f = hparameters["f"]
             # Retrieve dimensions from A prev's shape and dA's shape (≈2 lines)
             m, n H prev, n W prev, n C prev = A prev.shape
             m, n H, n W, n C = dA.shape
             # Initialize dA prev with zeros (≈1 line)
             dA prev = np.zeros(A prev.shape)
             for i in range(m):
                                                      # loop over the training examples
                 # select training example from A prev (≈1 line)
                 a prev = A prev[i]
                 for h in range(n H):
                                                      # loop on the vertical axis
                     for w in range(n W): # loop on the horizontal axis
                         for c in range(n C):
                                                       # loop over the channels (depth)
                             # Find the corners of the current "slice" (≈4 lines)
                             vert start = h
                             vert end = vert start + f
                             horiz start = w
                             horiz end = horiz start + f
                             # Compute the backward propagation in both modes.
                             if mode == "max":
```

```
# Use the corners and "c" to define the current slice from a prev (≈1 line)
                    a prev slice = a prev[vert start:vert end, horiz start:horiz end, c]
                    # Create the mask from a prev slice (≈1 line)
                    mask = create mask from window(a prev slice)
                    # Set dA prev to be dA prev + (the mask multiplied by the correct entry of dA) (≈1 line)
                    dA prev[i, vert start:vert end, horiz start:horiz end, c] += np.multiply(mask, dA[i, h, v
                elif mode == "average":
                    # Get the value a from dA (≈1 line)
                    da = dA[i, h, w, c]
                    # Define the shape of the filter as fxf (≈1 line)
                    shape = (f, f)
                    # Distribute it to get the correct slice of dA prev. i.e. Add the distributed value of do
                    dA prev[i, vert start:vert end, horiz start:horiz end, c] += distribute value(da, shape)
# Making sure your output shape is correct
assert(dA prev.shape == A prev.shape)
return dA prev
```

```
In [17]: np.random.seed(1)
         A prev = np.random.randn(5, 5, 3, 2)
         hparameters = {"stride" : 1, "f": 2}
         A, cache = pool forward(A prev, hparameters)
         dA = np.random.randn(5, 4, 2, 2)
         dA prev = pool backward(dA, cache, mode = "max")
         print("mode = max")
         print('mean of dA = ', np.mean(dA))
         print('dA prev[1,1] = ', dA prev[1,1])
          print()
         dA prev = pool backward(dA, cache, mode = "average")
         print("mode = average")
         print('mean of dA = ', np.mean(dA))
         print('dA prev[1,1] = ', dA prev[1,1])
         mode = max
         mean of dA = 0.14571390272918056
         dA prev[1,1] = [[0.
                                        0.
          [ 5.05844394 -1.68282702]
          [ 0.
                                 ]]
                        0.
```

mode = average

mean of dA = 0.14571390272918056

[1.26461098 -0.25749373] [1.17975636 -0.53624893]]

dA prev[1,1] = $[[0.08485462 \ 0.2787552]$