Optimization Methods

Until now, you've always used Gradient Descent to update the parameters and minimize the cost. In this notebook, you will learn more advanced optimization methods that can speed up learning and perhaps even get you to a better final value for the cost function. Having a good optimization algorithm can be the difference between waiting days vs. just a few hours to get a good result.

Gradient descent goes "downhill" on a cost function J. Think of it as trying to do this:

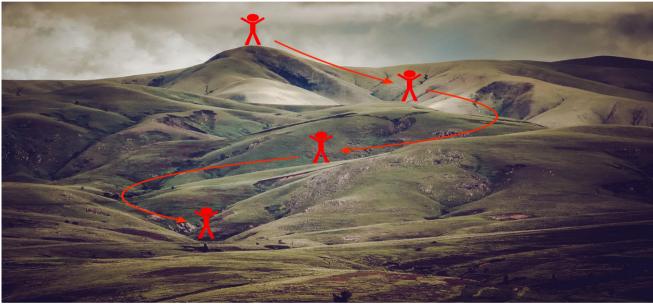


Figure 1: Minimizing the cost is like finding the lowest point in a hilly landscape

At each step of the training, you update your parameters following a certain direction to try to get to the lowest possible point.

Notations: As usual, $\frac{\partial J}{\partial a}=\mathrm{d}a$ for any variable a.

To get started, run the following code to import the libraries you will need.

```
In [1]: import numpy as np
         import matplotlib.pyplot as plt
         import scipy. io
         import math
         import sklearn
         import sklearn.datasets
         from opt utils import load params and grads, initialize parameters, forward propagation, backward propagation
         from opt utils import compute cost, predict, predict dec, plot decision boundary, load dataset
         from testCases import *
         %matplotlib inline
         plt.rcParams['figure.figsize'] = (7.0, 4.0) # set default size of plots
         plt.rcParams['image.interpolation'] = 'nearest'
         plt.rcParams['image.cmap'] = 'gray'
```

1 - Gradient Descent

A simple optimization method in machine learning is gradient descent (GD). When you take gradient steps with respect to all m examples on each step, it is also called Batch Gradient Descent.

Warm-up exercise: Implement the gradient descent update rule. The gradient descent rule is, for $l=1,\ldots,L$:

$$W^{[I]} = W^{[I]} - \alpha \, dW^{[I]} \tag{1}$$

$$b^{[I]} = b^{[I]} - \alpha \, db^{[I]} \tag{2}$$

$$b^{[l]} = b^{[l]} - \alpha \, db^{[l]} \tag{2}$$

where L is the number of layers and α is the learning rate. All parameters should be stored in the parameters dictionary. Note that the iterator 1 starts at 0 in the for loop while the first parameters are $W^{[1]}$ and $b^{[1]}$. You need to shift 1 to 1+1 when coding.

```
In [4]: # GRADED FUNCTION: update parameters with gd
         def update parameters with gd(parameters, grads, learning rate):
             Update parameters using one step of gradient descent
             Arguments:
             parameters -- python dictionary containing your parameters to be updated:
                             parameters ['W' + str(1)] = W1
                             parameters['b' + str(1)] = b1
             grads -- python dictionary containing your gradients to update each parameters:
                             grads['dW' + str(1)] = dW1
                             grads['db' + str(1)] = db1
             learning rate — the learning rate, scalar.
             Returns:
             parameters -- python dictionary containing your updated parameters
             L = len(parameters) // 2 # number of layers in the neural networks
             # Update rule for each parameter
             for 1 in range(L):
                 ### START CODE HERE ### (approx. 2 lines)
                 parameters["W" + str(1+1)] = parameters["W" + str(1+1)] - learning rate * grads["dW" + str(1+1)]
                 parameters["b" + str(1+1)] = parameters["b" + str(1+1)] - learning rate * grads["db" + str(1+1)]
                 ### END CODE HERE ###
             return parameters
```

```
Tn
    [6]: parameters, grads, learning rate = update parameters with gd test case()
           parameters = update parameters with gd(parameters, grads, learning rate)
           print("W1 = " + str(parameters["W1"]))
          print("b1 = " + str(parameters["b1"]))
          print("W2 = " + str(parameters["W2"]))
           print("b2 = " + str(parameters["b2"]))
          W1 = [ [ 1.63535156 -0.62320365 -0.53718766 ]
            \begin{bmatrix} -1.07799357 & 0.85639907 & -2.29470142 \end{bmatrix}
          b1 = [ [ 1.74604067 ]
            [-0.75184921]
          W2 = [ [ 0.32171798 -0.25467393 1.46902454 ] ]
            \begin{bmatrix} -2.05617317 & -0.31554548 & -0.3756023 \end{bmatrix}
            [ 1. 1404819 -1. 09976462 -0. 1612551 ]]
           b2 = \lceil \lceil -0.88020257 \rceil
            [ 0.02561572]
            [ 0.57539477]]
```

Expected Output:

```
W1 [[ 1.63535156 -0.62320365 -0.53718766] [-1.07799357 0.85639907 -2.29470142]]
b1 [[ 1.74604067] [-0.75184921]]
W2 [[ 0.32171798 -0.25467393 1.46902454] [-2.05617317 -0.31554548 -0.3756023 ] [ 1.1404819 -1.09976462 -0.1612551 ]]
b2 [[-0.88020257] [ 0.02561572] [ 0.57539477]]
```

A variant of this is Stochastic Gradient Descent (SGD), which is equivalent to mini-batch gradient descent where each mini-batch has just 1 example. The update rule that you have just implemented does not change. What changes is that you would be computing gradients on just one training example at a time, rather than on the whole training set. The code examples below illustrate the difference between stochastic gradient descent and (batch) gradient descent.

• (Batch) Gradient Descent:

```
X = data_input
Y = labels
parameters = initialize_parameters(layers_dims)
for i in range(0, num_iterations):
    # Forward propagation
    a, caches = forward_propagation(X, parameters)
    # Compute cost.
    cost = compute_cost(a, Y)
    # Backward propagation.
    grads = backward_propagation(a, caches, parameters)
    # Update parameters.
    parameters = update_parameters(parameters, grads)
```

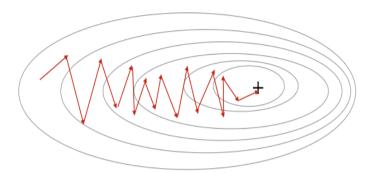
• Stochastic Gradient Descent:

```
X = data_input
Y = labels
parameters = initialize_parameters(layers_dims)
for i in range(0, num_iterations):
    for j in range(0, m):
        # Forward propagation
        a, caches = forward_propagation(X[:, j], parameters)
        # Compute cost
        cost = compute_cost(a, Y[:, j])
        # Backward propagation
        grads = backward_propagation(a, caches, parameters)
        # Update parameters.
        parameters = update_parameters(parameters, grads)
```

In Stochastic Gradient Descent, you use only 1 training example before updating the gradients. When the training set is large, SGD can be faster. But the parameters will "oscillate" toward the minimum rather than converge smoothly. Here is an illustration of this:



Gradient Descent



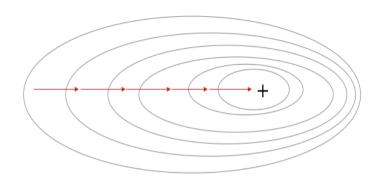


Figure 1: SGD vs GD

"+" denotes a minimum of the cost. SGD leads to many oscillations to reach convergence. But each step is a lot faster to compute for SGD than for GD, as it uses only one training example (vs. the whole batch for GD).

Note also that implementing SGD requires 3 for-loops in total:

- 1. Over the number of iterations
- 2. Over the m training examples
- 3. Over the layers (to update all parameters, from $(W^{[1]},b^{[1]})$ to $(W^{[L]},b^{[L]})$)

In practice, you'll often get faster results if you do not use neither the whole training set, nor only one training example, to perform each update. Minibatch gradient descent uses an intermediate number of examples for each step. With minibatch gradient descent, you loop over the minibatches instead of looping over individual training examples.

Stochastic Gradient Descent

Mini-Batch Gradient Descent

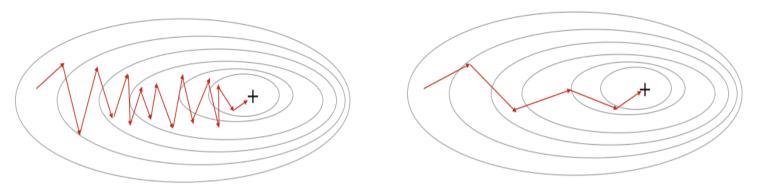


Figure 2: SGD vs Mini-Batch GD

"+" denotes a minimum of the cost. Using mini-batches in your optimization algorithm often leads to faster optimization.

What you should remember:

- The difference between gradient descent, mini-batch gradient descent and stochastic gradient descent is the number of examples you use to perform one update step.
- You have to tune a learning rate hyperparameter α .
- With a well-turned mini-batch size, usually it outperforms either gradient descent or stochastic gradient descent (particularly when the training set is large).

2 - Mini-Batch Gradient descent

Let's learn how to build mini-batches from the training set (X, Y).

There are two steps:

• **Shuffle**: Create a shuffled version of the training set (X, Y) as shown below. Each column of X and Y represents a training example. Note that the random shuffling is done synchronously between X and Y. Such that after the shuffling the i^{th} column of X is the example corresponding to the i^{th} label in Y. The shuffling step ensures that examples will be split randomly into different mini-batches.

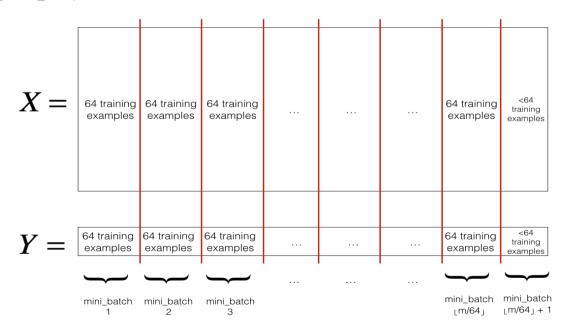
$$X = \begin{pmatrix} x_0^{(1)} & x_0^{(2)} & \dots & x_0^{(m-1)} & x_0^{(m)} \\ x_1^{(1)} & x_1^{(2)} & \dots & x_1^{(m-1)} & x_1^{(m)} \\ \vdots & \vdots & \vdots & \vdots & \vdots \\ x_{12286}^{(1)} & x_{12286}^{(2)} & \dots & x_{12286}^{(m-1)} & x_{12287}^{(m)} \\ x_{12287}^{(1)} & x_{12287}^{(2)} & \dots & x_{122287}^{(m-1)} & x_{12287}^{(m)} \end{pmatrix}$$

$$Y = \begin{pmatrix} y^{(1)} & y^{(2)} & \dots & y^{(m-1)} & y^{(m)} \\ y^{(m)} & y^{(m)} & y^{(m)} & y^{(m)} \end{pmatrix}$$

$$X = \begin{pmatrix} x_0^{(1)} & x_0^{(2)} & \dots & x_0^{(m-1)} & x_0^{(m)} \\ x_1^{(1)} & x_1^{(2)} & \dots & x_1^{(m-1)} & x_1^{(m)} \\ \vdots & \vdots & \vdots & \vdots & \vdots & \vdots \\ x_{12286}^{(1)} & x_{12286}^{(2)} & \dots & x_{12286}^{(m-1)} & x_{12287}^{(m)} \end{pmatrix}$$

$$X = \begin{pmatrix} x_0^{(1)} & x_0^{(2)} & \dots & x_1^{(m-1)} & x_1^{(m)} \\ x_1^{(1)} & x_1^{(2)} & \dots & x_1^{(m-1)} & x_1^{(m)} \\ x_{12287}^{(1)} & x_{12287}^{(2)} & \dots & x_{12287}^{(m-1)} & x_{12287}^{(m)} \end{pmatrix}$$

• Partition: Partition the shuffled (X, Y) into mini-batches of size mini_batch_size (here 64). Note that the number of training examples is not always divisible by mini_batch_size. The last mini batch might be smaller, but you don't need to worry about this. When the final mini-batch is smaller than the full mini batch size, it will look like this:



Exercise: Implement $random_mini_batches$. We coded the shuffling part for you. To help you with the partitioning step, we give you the following code that selects the indexes for the 1^{st} and 2^{nd} mini-batches:

```
first_mini_batch_X = shuffled_X[:, 0 : mini_batch_size]
second_mini_batch_X = shuffled_X[:, mini_batch_size : 2 * mini_batch_size]
...
```

Note that the last mini-batch might end up smaller than mini_batch_size=64. Let $\lfloor s \rfloor$ represents s rounded down to the nearest integer (this is math. floor(s) in Python). If the total number of examples is not a multiple of mini_batch_size=64 then there will be $\lfloor \frac{m}{mini_batch_size} \rfloor$ mini-batches with a full 64 examples, and the number of examples in the final mini-batch will be $(m-mini_batch_size \times \lfloor \frac{m}{mini_batch_size} \rfloor)$.

```
[7]: # GRADED FUNCTION: random mini batches
      def random mini batches(X, Y, mini batch size = 64, seed = 0):
          Creates a list of random minibatches from (X. Y)
          Arguments:
          X — input data, of shape (input size, number of examples)
         Y — true "label" vector (1 for blue dot / 0 for red dot), of shape (1, number of examples)
          mini batch size -- size of the mini-batches, integer
          Returns:
          mini batches -- list of synchronous (mini batch X, mini batch Y)
                                          # To make vour "random" minibatches the same as ours
          np. random. seed (seed)
          m = X. shape[1]
                                           # number of training examples
          mini batches = []
          # Step 1: Shuffle (X, Y)
          permutation = list(np. random. permutation(m))
          shuffled X = X[:, permutation]
          shuffled Y = Y[:, permutation].reshape((1, m))
          # Step 2: Partition (shuffled X, shuffled Y). Minus the end case.
          num complete minibatches = math. floor (m/mini batch size) # number of mini batches of size mini batch size in your partitionning
          for k in range (0, num complete minibatches):
              ### START CODE HERE ### (approx. 2 lines)
              mini batch X = shuffled X[:, k*mini batch size:(k+1)*mini batch size]
              mini batch Y = shuffled Y[:, k*mini batch size:(k+1)*mini batch size]
              ### END CODE HERE ###
              mini batch = (mini batch X, mini batch Y)
              mini batches. append (mini batch)
          # Handling the end case (last mini-batch < mini batch size)
          if m % mini batch size != 0:
              ### START CODE HERE ### (approx. 2 lines)
              mini batch X = shuffled X[:, num complete minibatches*mini batch size:]
              mini batch Y = shuffled Y[:, num complete minibatches*mini batch size:]
              ### END CODE HERE ###
              mini batch = (mini batch X, mini batch Y)
```

```
mini_batches.append(mini_batch)
return mini_batches
```

```
In [8]: X_assess, Y_assess, mini_batch_size = random_mini_batches_test_case()
    mini_batches = random_mini_batches(X_assess, Y_assess, mini_batch_size)

    print ("shape of the 1st mini_batch_X: " + str(mini_batches[0][0].shape))
    print ("shape of the 2nd mini_batch_X: " + str(mini_batches[1][0].shape))
    print ("shape of the 3rd mini_batch_Y: " + str(mini_batches[0][1].shape))
    print ("shape of the 2nd mini_batch_Y: " + str(mini_batches[0][1].shape))
    print ("shape of the 3rd mini_batch_Y: " + str(mini_batches[0][1].shape))
    print ("shape of the 3rd mini_batch_Y: " + str(mini_batches[0][0][0][0][0][0]])

shape of the 1st mini_batch_X: (12288, 64)
    shape of the 2nd mini_batch_X: (12288, 64)
```

Expected Output:

(12288, 64)	shape of the 1st mini_batch_X
(12288, 64)	shape of the 2nd mini_batch_X
(12288, 20)	shape of the 3rd mini_batch_X
(1, 64)	shape of the 1st mini_batch_Y
(1, 64)	shape of the 2nd mini_batch_Y
(1, 20)	shape of the 3rd mini_batch_Y
[0.90085595 -0.7612069 0.2344157]	mini batch sanity check

What you should remember:

- Shuffling and Partitioning are the two steps required to build mini-batches
- Powers of two are often chosen to be the mini-batch size, e.g., 16, 32, 64, 128.

3 - Momentum

Because mini-batch gradient descent makes a parameter update after seeing just a subset of examples, the direction of the update has some variance, and so the path taken by mini-batch gradient descent will "oscillate" toward convergence. Using momentum can reduce these oscillations.

Momentum takes into account the past gradients to smooth out the update. We will store the 'direction' of the previous gradients in the variable v. Formally, this will be the exponentially weighted average of the gradient on previous steps. You can also think of v as the "velocity" of a ball rolling downhill, building up speed (and momentum) according to the direction of the gradient/slope of the hill.

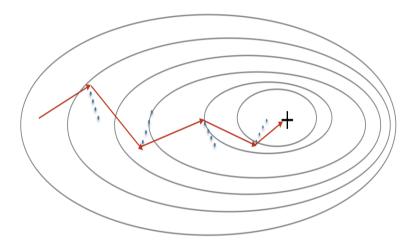


Figure 3: The red arrows shows the direction taken by one step of mini-batch gradient descent with momentum. The blue points show the direction of the gradient (with respect to the current mini-batch) on each step. Rather than just following the gradient, we let the gradient influence v and then take a step in the direction of v.

Exercise: Initialize the velocity. The velocity, v, is a python dictionary that needs to be initialized with arrays of zeros. Its keys are the same as those in the grads dictionary, that is: for l = 1, ..., L:

```
v["dW" + str(1+1)] = \dots #(numpy array of zeros with the same shape as parameters["W" + str(1+1)]) v["db" + str(1+1)] = \dots #(numpy array of zeros with the same shape as parameters["b" + str(1+1)])
```

Note that the iterator I starts at 0 in the for loop while the first parameters are v["dW1"] and v["db1"] (that's a "one" on the superscript). This is why we are shifting I to I+1 in the for loop.

```
In [14]: # GRADED FUNCTION: initialize velocity
          def initialize velocity(parameters):
              Initializes the velocity as a python dictionary with:
                          - keys: "dW1", "db1", ..., "dWL", "dbL"
                          - values: numpy arrays of zeros of the same shape as the corresponding gradients/parameters.
              Arguments:
              parameters -- python dictionary containing your parameters.
                              parameters['W' + str(1)] = W1
                              parameters['b' + str(1)] = b1
              Returns:
              v -- python dictionary containing the current velocity.
                              v['dW' + str(1)] = velocity of dW1
                              v['db' + str(1)] = velocity of db1
              L = len(parameters) // 2 # number of layers in the neural networks
              v = \{\}
               # Initialize velocity
              for 1 in range(L):
                   ### START CODE HERE ### (approx. 2 lines)
                   v["dW" + str(1+1)] = np. zeros((parameters["W" + str(1+1)]. shape))
                   v["db" + str(1+1)] = np. zeros((parameters["b" + str(1+1)]. shape))
                   ### END CODE HERE ###
              return v
```

```
In [15]: parameters = initialize velocity test case()
                v = initialize velocity(parameters)
                print("v[\"dW1\"] = " + str(v["dW1"]))
                print("v[\"db1\"] = " + str(v["db1"]))
                print("v[\"dW2\"] = " + str(v["dW2"]))
                print("v[\"db2\"] = " + str(v["db2"]))
                v\lceil "dW1" \rceil = \lceil \lceil 0. \quad 0. \quad 0. \rceil
                 \begin{bmatrix} 0, & 0, & 0, \end{bmatrix}
                v["db1"] = [[ 0. ]
                 [ 0, ]]
                v\lceil "dW2" \rceil = \lceil \lceil 0. \quad 0. \quad 0. \rceil
                 \begin{bmatrix} 0, & 0, & 0, \end{bmatrix}
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                v["db2"] = [[ 0. ]
                 \begin{bmatrix} 0 \end{bmatrix}
                  \begin{bmatrix} 0. \end{bmatrix}
```

Expected Output:

```
v["dW1"] [[ 0. 0. 0.] [ 0. 0. 0.]]
v["db1"] [[ 0.] [ 0.]]
v["dW2"] [[ 0. 0. 0.] [ 0. 0. 0.] [ 0. 0. 0.]]
v["db2"] [[ 0.] [ 0.] [ 0.]
```

Exercise: Now, implement the parameters update with momentum. The momentum update rule is, for $l=1,\ldots,L$:

$$\begin{cases} v_{dW}[l] = \beta v_{dW}[l] + (1 - \beta)dW^{[l]} \\ W^{[l]} = W^{[l]} - \alpha v_{dW}[l] \end{cases}$$
(3)

$$\begin{cases} v_{db}[l] = \beta v_{db}[l] + (1 - \beta)db^{[l]} \\ b^{[l]} = b^{[l]} - \alpha v_{db}[l] \end{cases}$$
(4)

where L is the number of layers, β is the momentum and α is the learning rate. All parameters should be stored in the parameters dictionary. Note that the iterator 1 starts at 0 in the for loop while the first parameters are $W^{[1]}$ and $b^{[1]}$ (that's a "one" on the superscript). So you will need to shift 1 to 1+1 when coding.

```
In [16]: # GRADED FUNCTION: update parameters with momentum
           def update parameters with momentum (parameters, grads, v, beta, learning rate):
               Update parameters using Momentum
               Arguments:
               parameters -- python dictionary containing your parameters:
                               parameters ['W' + str(1)] = W1
                               parameters ['b' + str(1)] = b1
               grads -- python dictionary containing your gradients for each parameters:
                               grads['dW' + str(1)] = dW1
                               grads['db' + str(1)] = db1
              v -- python dictionary containing the current velocity:
                               v['dW' + str(1)] = \dots
                               v['db' + str(1)] = \dots
              beta — the momentum hyperparameter, scalar
               learning rate — the learning rate, scalar
               Returns:
               parameters -- python dictionary containing your updated parameters
               v -- python dictionary containing your updated velocities
              L = len(parameters) // 2 # number of layers in the neural networks
               # Momentum update for each parameter
               for 1 in range(L):
                   ### START CODE HERE ### (approx. 4 lines)
                   # compute velocities
                   v["dW" + str(1+1)] = beta * v["dW" + str(1+1)] + (1-beta) * grads['dW' + str(1+1)]
                   v["db" + str(1+1)] = beta * v["db" + str(1+1)] + (1-beta) * grads['db' + str(1+1)]
                   # update parameters
                   parameters ["W" + str(1+1)] = parameters ["W" + str(1+1)] - learning rate * v ["dW" + str(1+1)]
                   parameters \lceil "b" + str(1+1) \rceil = parameters \lceil "b" + str(1+1) \rceil - 1earning rate * v \lceil "db" + str(1+1) \rceil
                   ### END CODE HERE ###
               return parameters, v
```

```
In [17]: parameters, grads, v = update parameters with momentum test case()
           parameters, v = update parameters with momentum (parameters, grads, v, beta = 0.9, learning rate = 0.01)
           print("W1 = " + str(parameters["W1"]))
           print("b1 = " + str(parameters["b1"]))
           print("W2 = " + str(parameters["W2"]))
           print("b2 = " + str(parameters["b2"]))
           print("v[\"dW1\"] = " + str(v["dW1"]))
           print("v[\'"db1\'"] = " + str(v["db1"]))
           print("v[\"dW2\"] = " + str(v["dW2"]))
           print("v[\"db2\"] = " + str(v["db2"]))
           W1 = [ [ 1.62544598 -0.61290114 -0.52907334 ]
            \begin{bmatrix} -1.07347112 & 0.86450677 & -2.30085497 \end{bmatrix}
           b1 = [ [ 1.74493465 ]
            [-0.76027113]
           W2 = \begin{bmatrix} 0.31930698 & -0.24990073 & 1.4627996 \end{bmatrix}
            [-2.05974396 -0.32173003 -0.38320915]
            [ 1.13444069 -1.0998786 -0.1713109 ]]
           b2 = [-0.87809283]
            [ 0.04055394]
            [ 0.58207317]]
           v["dW1"] = [[-0.11006192 \ 0.11447237 \ 0.09015907]
             \begin{bmatrix} 0.05024943 & 0.09008559 & -0.06837279 \end{bmatrix}
           v["db1"] = [[-0.01228902]
             [-0.09357694]
           v["dW2"] = [[-0.02678881 \ 0.05303555 \ -0.06916608]
            [-0.03967535 -0.06871727 -0.08452056]
            [-0.06712461 -0.00126646 -0.11173103]
           v["db2"] = [[ 0.02344157]
             [ 0.16598022]
             [ 0.07420442]]
```

Expected Output:

W1 [[1.62544598 -0.61290114 -0.52907334] [-1.07347112 0.86450677 -2.30085497]]
b1 [[1.74493465] [-0.76027113]]
W2 [[0.31930698 -0.24990073 1.4627996] [-2.05974396 -0.32173003 -0.38320915] [1.13444069 -1.0998786 -0.1713109]

b2	[[-0.87809283] [0.04055394] [0.58207317]]
v["dW1"]	[[-0.11006192 0.11447237 0.09015907] [0.05024943 0.09008559 -0.06837279]]
v["db1"]	[[-0.01228902] [-0.09357694]]
v["dW2"]	[[-0.02678881 0.05303555 -0.06916608] [-0.03967535 -0.06871727 -0.08452056] [-0.06712461 -0.00126646 -0.11173103]]
v["db2"]	[[0.02344157] [0.16598022] [0.07420442]]

Note that:

- The velocity is initialized with zeros. So the algorithm will take a few iterations to "build up" velocity and start to take bigger steps.
- If $\beta = 0$, then this just becomes standard gradient descent without momentum.

How do you choose β ?

- The larger the momentum β is, the smoother the update because the more we take the past gradients into account. But if β is too big, it could also smooth out the updates too much.
- Common values for β range from 0.8 to 0.999. If you don't feel inclined to tune this, $\beta = 0.9$ is often a reasonable default.
- Tuning the optimal β for your model might need trying several values to see what works best in term of reducing the value of the cost function J.

What you should remember:

- Momentum takes past gradients into account to smooth out the steps of gradient descent. It can be applied with batch gradient descent, minibatch gradient descent or stochastic gradient descent.
- You have to tune a momentum hyperparameter β and a learning rate α .

4 - Adam

Adam is one of the most effective optimization algorithms for training neural networks. It combines ideas from RMSProp (described in lecture) and Momentum.

How does Adam work?

- 1. It calculates an exponentially weighted average of past gradients, and stores it in variables v (before bias correction) and $v^{corrected}$ (with bias correction).
- 2. It calculates an exponentially weighted average of the squares of the past gradients, and stores it in variables s (before bias correction) and s (with bias correction).
- 3. It updates parameters in a direction based on combining information from "1" and "2".

The update rule is, for $l = 1, \dots, L$:

$$\begin{cases} v_{dW}[l] = \beta_1 v_{dW}[l] + (1 - \beta_1) \frac{\partial \mathcal{J}}{\partial W^{[l]}} \\ v_{dW}^{corrected} = \frac{v_{dW}[l]}{1 - (\beta_1)^l} \\ s_{dW}[l] = \beta_2 s_{dW}[l] + (1 - \beta_2) (\frac{\partial \mathcal{J}}{\partial W^{[l]}})^2 \\ s_{dW}^{corrected} = \frac{s_{dW}[l]}{1 - (\beta_1)^l} \\ W^{[l]} = W^{[l]} - \alpha \frac{v_{dW}^{corrected}}{\sqrt{s_{dW}^{corrected}}} \end{cases}$$

where:

- t counts the number of steps taken of Adam
- L is the number of layers
- β_1 and β_2 are hyperparameters that control the two exponentially weighted averages.
- α is the learning rate
- $m{\cdot}$ $\ensuremath{arepsilon}$ is a very small number to avoid dividing by zero

As usual, we will store all parameters in the parameters dictionary

Exercise: Initialize the Adam variables v, s which keep track of the past information.

Instruction: The variables v, s are python dictionaries that need to be initialized with arrays of zeros. Their keys are the same as for grads, that is: for $l = 1, \ldots, L$:

```
v["dW" + str(1+1)] = \dots \#(numpy \ array \ of \ zeros \ with \ the \ same \ shape \ as \ parameters["W" + str(1+1)]) v["db" + str(1+1)] = \dots \#(numpy \ array \ of \ zeros \ with \ the \ same \ shape \ as \ parameters["b" + str(1+1)]) s["dW" + str(1+1)] = \dots \#(numpy \ array \ of \ zeros \ with \ the \ same \ shape \ as \ parameters["W" + str(1+1)]) s["db" + str(1+1)] = \dots \#(numpy \ array \ of \ zeros \ with \ the \ same \ shape \ as \ parameters["b" + str(1+1)])
```

```
In [18]: # GRADED FUNCTION: initialize adam
           def initialize adam(parameters) :
               Initializes v and s as two python dictionaries with:
                           - keys: "dW1", "db1", ..., "dWL", "dbL"
                           - values: numpy arrays of zeros of the same shape as the corresponding gradients/parameters.
               Arguments:
               parameters — python dictionary containing your parameters.
                               parameters["W" + str(1)] = W1
                               parameters["b" + str(1)] = b1
               Returns:
               v -- python dictionary that will contain the exponentially weighted average of the gradient.
                               v["dW" + str(1)] = \dots
                               v\lceil "db" + str(1) \rceil = \dots
               s -- python dictionary that will contain the exponentially weighted average of the squared gradient.
                               s["dW" + str(1)] = \dots
                               s["db" + str(1)] = \dots
               L = len(parameters) // 2 # number of layers in the neural networks
               v = \{\}
               S = \{\}
               # Initialize v, s. Input: "parameters". Outputs: "v, s".
               for 1 in range(L):
               ### START CODE HERE ### (approx. 4 lines)
                   v["dW" + str(1+1)] = np. zeros(parameters["W" + str(1+1)]. shape)
                   v["db" + str(1+1)] = np. zeros(parameters["b" + str(1+1)]. shape)
                   s["dW" + str(1+1)] = np. zeros(parameters["W" + str(1+1)]. shape)
                   s["db" + str(1+1)] = np. zeros(parameters["b" + str(1+1)]. shape)
               ### END CODE HERE ###
               return v, s
```

```
In [19]: parameters = initialize adam test case()
                 v, s = initialize adam(parameters)
                 print("v[\"dW1\"] = " + str(v["dW1"]))
                 print("v[\'"db1\'"] = " + str(v["db1"]))
                 print("v[\"dW2\"] = " + str(v["dW2"]))
                 print("v[\'"db2\'"] = " + str(v["db2"]))
                 print("s[\"dW1\"] = " + str(s["dW1"]))
                 print("s[\''db1\''] = " + str(s["db1"]))
                 print("s[\"dW2\"] = " + str(s["dW2"]))
                 print("s[\"db2\"] = " + str(s["db2"]))
                v\lceil "dW1" \rceil = \lceil \lceil 0. \quad 0. \quad 0. \rceil
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                v["db1"] = [[ 0. ]
                  [ 0, ]]
                v\lceil "dW2" \rceil = \lceil \lceil 0. \quad 0. \quad 0. \rceil
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                v["db2"] = [[ 0. ]
                 \begin{bmatrix} 0. \end{bmatrix}
                  \lceil 0. \rceil \rceil
                s\lceil "dW1" \rceil = \lceil \lceil 0. \quad 0. \quad 0. \rceil
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                s["db1"] = [[ 0. ]
                 \begin{bmatrix} 0. \end{bmatrix}
                s["dW2"] = [[ 0.  0.  0. ]
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                 \begin{bmatrix} 0. & 0. & 0. \end{bmatrix}
                s["db2"] = [[ 0. ]
                 \lceil 0. \rceil
                  \lceil 0. \rceil \rceil
```

Expected Output:

v["dW1"] [[0. 0. 0.] [0. 0. 0.]] v["db1"] [[0.] [0.]] v["dW2"] [[0. 0. 0.] [0. 0. 0.] [0. 0. 0.]

v["db2"]	[[0.] [0.] [0.]]
s["dW1"]	[[0. 0. 0.] [0. 0. 0.]]
s["db1"]	[[0.] [0.]]
s["dW2"]	[[0. 0. 0.] [0. 0. 0.] [0. 0. 0.]]
s["db2"]	[[0.] [0.] [0.]]

Exercise: Now, implement the parameters update with Adam. Recall the general update rule is, for $l = 1, \dots, L$:

$$\begin{cases} v_W[l] = \beta_1 v_W[l] + (1 - \beta_1) \frac{\partial J}{\partial W^{[l]}} \\ v_W^{corrected} = \frac{v_W[l]}{1 - (\beta_1)^l} \\ s_W[l] = \beta_2 s_W[l] + (1 - \beta_2) (\frac{\partial J}{\partial W^{[l]}})^2 \\ s_W^{corrected} = \frac{s_W[l]}{1 - (\beta_2)^l} \\ W^{[l]} = W^{[l]} - \alpha \frac{v_W^{corrected}}{\sqrt{s_W^{corrected}}} \end{cases}$$

Note that the iterator 1 starts at 0 in the for loop while the first parameters are $W^{[1]}$ and $b^{[1]}$. You need to shift 1 to 1+1 when coding.

```
In [32]: # GRADED FUNCTION: update parameters with adam
          def update parameters with adam(parameters, grads, v, s, t, learning rate = 0.01,
                                           beta1 = 0.9, beta2 = 0.999, ensilon = 1e-8):
              Update parameters using Adam
              Arguments:
              parameters -- python dictionary containing your parameters:
                              parameters ['W' + str(1)] = W1
                              parameters ['b' + str(1)] = b1
              grads -- python dictionary containing your gradients for each parameters:
                              grads['dW' + str(1)] = dW1
                              grads['db' + str(1)] = db1
              v -- Adam variable, moving average of the first gradient, python dictionary
              s -- Adam variable, moving average of the squared gradient, python dictionary
              learning rate — the learning rate, scalar.
              betal -- Exponential decay hyperparameter for the first moment estimates
              beta2 -- Exponential decay hyperparameter for the second moment estimates
              epsilon -- hyperparameter preventing division by zero in Adam updates
              Returns:
              parameters -- python dictionary containing your updated parameters
              v -- Adam variable, moving average of the first gradient, python dictionary
              s -- Adam variable, moving average of the squared gradient, python dictionary
              L = len(parameters) // 2
                                                        # number of layers in the neural networks
              v corrected = {}
                                                        # Initializing first moment estimate, python dictionary
              s corrected = {}
                                                        # Initializing second moment estimate, python dictionary
               # Perform Adam update on all parameters
              for 1 in range(L):
                  # Moving average of the gradients. Inputs: "v, grads, beta1". Output: "v".
                  ### START CODE HERE ### (approx. 2 lines)
                  v["dW" + str(1+1)] = beta1 * v["dW" + str(1+1)] + (1-beta1) * grads['dW' + str(1+1)]
                  v["db" + str(1+1)] = beta1 * v["db" + str(1+1)] + (1-beta1) * grads['db' + str(1+1)]
                  ### END CODE HERE ###
                  # Compute bias-corrected first moment estimate. Inputs: "v, beta1, t". Output: "v corrected".
                  ### START CODE HERE ### (approx. 2 lines)
```

```
v = v = v'' dW'' + str(1+1) = v''' dW'' + str(1+1) = v'''' dW'' + str(1+1) = v''' dW'' + str(1+1) = v'''' dW'' + str(1+1) = v''' dW''' + str(1+1) = v''' dW'' + str(1
                 v \operatorname{corrected}["db" + \operatorname{str}(1+1)] = v["db" + \operatorname{str}(1+1)] / (1-\operatorname{beta1})
                  ### END CODE HERE ###
                 # Moving average of the squared gradients. Inputs: "s, grads, beta2". Output: "s".
                 ### START CODE HERE ### (approx. 2 lines)
                 s["dW" + str(1+1)] = beta2 * s["dW" + str(1+1)] + (1-beta2) * np. power(grads['dW' + str(1+1)], 2)
                 s["db" + str(1+1)] = beta2 * s["db" + str(1+1)] + (1-beta2) * np. power(grads['db' + str(1+1)], 2)
                  ### END CODE HERE ###
                 # Compute bias-corrected second raw moment estimate. Inputs: "s, beta2, t". Output: "s corrected".
                 ### START CODE HERE ### (approx. 2 lines)
                 s corrected \lceil "dW" + str(1+1) \rceil = s \lceil "dW" + str(1+1) \rceil / (1-beta2)
                 s \ corrected["db" + str(1+1)] = s["db" + str(1+1)] / (1-beta2)
                  ### END CODE HERE ###
                 # Update parameters. Inputs: "parameters, learning rate, v corrected, s corrected, epsilon". Output: "parameters".
                 ### START CODE HERE ### (approx. 2 lines)
                 parameters["W" + str(1+1)] = parameters["W" + str(1+1)] - learning rate * v corrected["dW" + str(1+1)] / np. sqrt(s corrected[
                 parameters["b" + str(1+1)] = parameters["b" + str(1+1)] - learning rate * v corrected["db" + str(1+1)] / np. sqrt(s corrected["db" + str(1+1)])
                 ### END CODE HERE ###
return parameters, v, s
```

https://hub.coursera-notebooks.org/user/tvdrtwvoywmtxjsfxnxsbg/notebooks/week6/Optimization%20methods.ipynb

```
In [33]: parameters, grads, v, s = update parameters with adam test case()
           parameters, v, s = update parameters with adam(parameters, grads, v, s, t = 2)
           print("W1 = " + str(parameters["W1"]))
           print("b1 = " + str(parameters["b1"]))
           print("W2 = " + str(parameters["W2"]))
           print("b2 = " + str(parameters["b2"]))
           print("v[\"dW1\"] = " + str(v["dW1"]))
           print("v[\'"db1\'"] = " + str(v["db1"]))
           print("v[\"dW2\"] = " + str(v["dW2"]))
           print("v[\'"db2\'"] = " + str(v["db2"]))
           print("s[\"dW1\"] = " + str(s["dW1"]))
           print("s[\"db1\"] = " + str(s["db1"]))
           print("s[\"dW2\"] = " + str(s["dW2"]))
           print("s[\''db2\''] = " + str(s["db2"]))
           W1 = \begin{bmatrix} 1.63434536 & -0.62175641 & -0.53817175 \end{bmatrix}
            \begin{bmatrix} -1.08296862 & 0.85540763 & -2.2915387 \end{bmatrix}
           b1 = [ [ 1.75481176 ]
            [-0.7512069]]
           W2 = \begin{bmatrix} 0.3290391 & -0.25937038 & 1.47210794 \end{bmatrix}
            \begin{bmatrix} -2.05014071 & -0.3124172 & -0.37405435 \end{bmatrix}
            [ 1.14376944 -1.08989158 -0.16242821]]
           b2 = [-0.88785842]
            [ 0.03221375]
            [0.57281521]
           v\lceil "dW1" \rceil = \lceil \lceil -0.11006192 \quad 0.11447237 \quad 0.09015907 \rceil
            v["db1"] = [[-0.01228902]]
            [-0.09357694]
           v["dW2"] = [[-0.02678881 \ 0.05303555 \ -0.06916608]
            [-0.03967535 -0.06871727 -0.08452056]
            [-0.06712461 -0.00126646 -0.11173103]
           v["db2"] = [[ 0.02344157]
            [ 0.16598022]
            [0.07420442]
           s["dW1"] = [[ 0.00121136   0.00131039   0.00081287]
            s["db1"] = [[ 1.51020075e-05]]
            [ 8.75664434e-04]]
           s["dW2"] = [[7.17640232e-05] 2.81276921e-04] 4.78394595e-04]
```

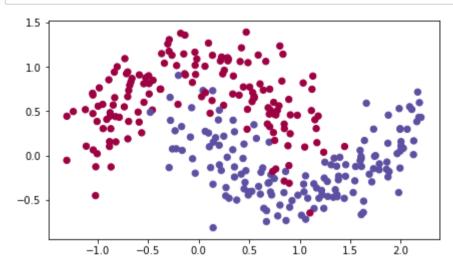
Expected Output:

W1	[[1.63178673 -0.61919778 -0.53561312] [-1.08040999 0.85796626 -2.29409733]]
b1	[[1.75225313] [-0.75376553]]
W2	[[0.32648046 -0.25681174 1.46954931] [-2.05269934 -0.31497584 -0.37661299] [1.14121081 -1.09245036 -0.16498684]]
b2	[[-0.88529978] [0.03477238] [0.57537385]]
v["dW1"]	[[-0.11006192 0.11447237 0.09015907] [0.05024943 0.09008559 -0.06837279]]
v["db1"]	[[-0.01228902] [-0.09357694]]
v["dW2"]	[[-0.02678881 0.05303555 -0.06916608] [-0.03967535 -0.06871727 -0.08452056] [-0.06712461 -0.00126646 -0.11173103]]
v["db2"]	[[0.02344157] [0.16598022] [0.07420442]]
s["dW1"]	[[0.00121136 0.00131039 0.00081287] [0.0002525 0.00081154 0.00046748]]
s["db1"]	[[1.51020075e-05] [8.75664434e-04]]
s["dW2"]	[[7.17640232e-05 2.81276921e-04 4.78394595e-04] [1.57413361e-04 4.72206320e-04 7.14372576e-04] [4.50571368e-04 1.60392066e-07 1.24838242e-03]]
s["db2"]	[[5.49507194e-05] [2.75494327e-03] [5.50629536e-04]]

You now have three working optimization algorithms (mini-batch gradient descent, Momentum, Adam). Let's implement a model with each of these optimizers and observe the difference.

5 - Model with different optimization algorithms

Lets use the following "moons" dataset to test the different optimization methods. (The dataset is named "moons" because the data from each of the two classes looks a bit like a crescent-shaped moon.)



We have already implemented a 3-layer neural network. You will train it with:

- Mini-batch **Gradient Descent**: it will call your function:
 - update_parameters_with_gd()
- Mini-batch **Momentum**: it will call your functions:
 - initialize_velocity() and update_parameters_with_momentum()
- Mini-batch Adam: it will call your functions:
 - initialize_adam() and update_parameters_with_adam()

```
In [36]: def model(X, Y, layers dims, optimizer, learning rate = 0.0007, mini batch size = 64, beta = 0.9,
                    beta1 = 0.9, beta2 = 0.999, epsilon = 1e-8, num epochs = 10000, print cost = True):
              3-layer neural network model which can be run in different optimizer modes.
              Arguments:
              X — input data, of shape (2, number of examples)
              Y -- true "label" vector (1 for blue dot / 0 for red dot), of shape (1, number of examples)
              layers dims -- python list, containing the size of each layer
              learning rate — the learning rate, scalar.
              mini batch size -- the size of a mini batch
              beta -- Momentum hyperparameter
              betal -- Exponential decay hyperparameter for the past gradients estimates
              beta2 -- Exponential decay hyperparameter for the past squared gradients estimates
               epsilon -- hyperparameter preventing division by zero in Adam updates
              num epochs — number of epochs
              print cost -- True to print the cost every 1000 epochs
              Returns:
              parameters -- python dictionary containing your updated parameters
              L = len(layers dims)
                                               # number of layers in the neural networks
              costs = []
                                                # to keep track of the cost
              t = 0
                                                # initializing the counter required for Adam update
                                                # For grading purposes, so that your "random" minibatches are the same as ours
              seed = 10
               # Initialize parameters
              parameters = initialize parameters(layers dims)
               # Initialize the optimizer
              if optimizer == "gd":
                  pass # no initialization required for gradient descent
              elif optimizer == "momentum":
                  v = initialize velocity(parameters)
              elif optimizer == "adam":
                  v, s = initialize adam(parameters)
               # Optimization loop
              for i in range (num epochs):
```

```
# Define the random minibatches. We increment the seed to reshuffle differently the dataset after each epoch
    seed = seed + 1
    minibatches = random mini batches(X, Y, mini batch size, seed)
    for minibatch in minibatches:
        # Select a minibatch
        (minibatch X, minibatch Y) = minibatch
        # Forward propagation
        a3. caches = forward propagation(minibatch X, parameters)
        # Compute cost
        cost = compute cost(a3, minibatch Y)
        # Backward propagation
        grads = backward propagation (minibatch X, minibatch Y, caches)
        # Update parameters
        if optimizer == "gd":
            parameters = update parameters with gd(parameters, grads, learning rate)
        elif optimizer == "momentum":
            parameters, v = update parameters with momentum (parameters, grads, v, beta, learning rate)
        elif optimizer == "adam":
            t = t + 1 # Adam counter
            parameters, v, s = update parameters with adam(parameters, grads, v, s,
                                                            t, learning rate, betal, beta2, epsilon)
    # Print the cost every 1000 epoch
    if print cost and i % 1000 == 0:
        print ("Cost after epoch %i: %f" %(i, cost))
    if print cost and i % 100 == 0:
        costs. append (cost)
# plot the cost
plt.plot(costs)
plt. vlabel ('cost')
plt. xlabel ('epochs (per 100)')
plt.title("Learning rate = " + str(learning rate))
plt.show()
return parameters
```

You will now run this 3 layer neural network with each of the 3 optimization methods.

5.1 - Mini-batch Gradient descent

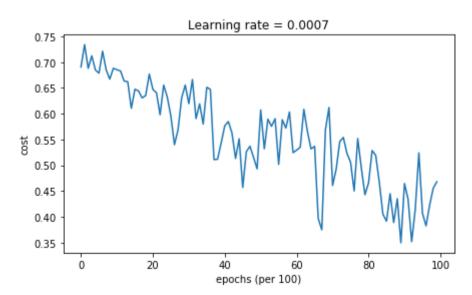
Run the following code to see how the model does with mini-batch gradient descent.

```
In [37]: # train 3-layer model
layers_dims = [train_X.shape[0], 5, 2, 1]
parameters = model(train_X, train_Y, layers_dims, optimizer = "gd")

# Predict
predictions = predict(train_X, train_Y, parameters)

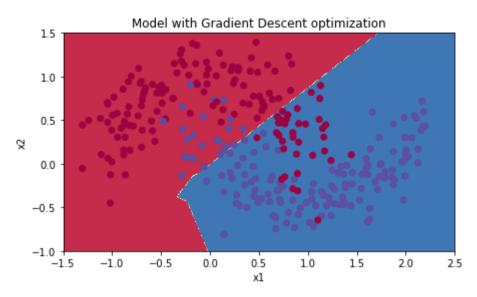
# Plot decision boundary
plt.title("Model with Gradient Descent optimization")
axes = plt.gca()
axes.set_xlim([-1.5, 2.5])
axes.set_ylim([-1, 1.5])
plot_decision_boundary(lambda x: predict_dec(parameters, x.T), train_X, train_Y)
```

Cost after epoch 0: 0.690736 Cost after epoch 1000: 0.685273 Cost after epoch 2000: 0.647072 Cost after epoch 3000: 0.619525 Cost after epoch 4000: 0.576584 Cost after epoch 5000: 0.607243 Cost after epoch 6000: 0.529403 Cost after epoch 7000: 0.460768 Cost after epoch 8000: 0.465586 Cost after epoch 9000: 0.464518



Accuracy: 0.796666666667

2018/1/16



5.2 - Mini-batch gradient descent with momentum

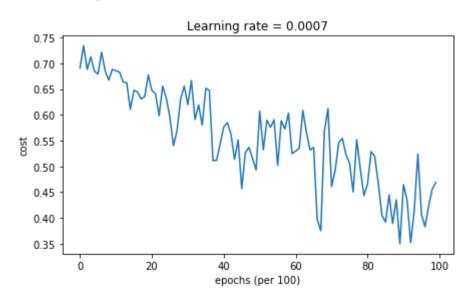
Run the following code to see how the model does with momentum. Because this example is relatively simple, the gains from using momentum are small; but for more complex problems you might see bigger gains.

```
In [43]: # train 3-layer model
layers_dims = [train_X.shape[0], 5, 2, 1]
parameters = model(train_X, train_Y, layers_dims, beta = 0.9, optimizer = "momentum")

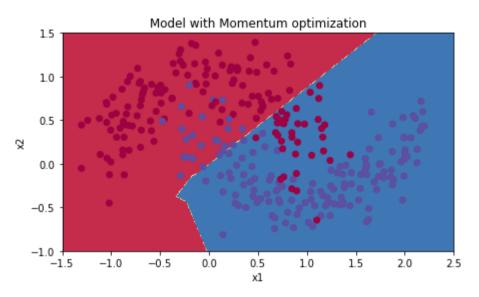
# Predict
predictions = predict(train_X, train_Y, parameters)

# Plot decision boundary
plt.title("Model with Momentum optimization")
axes = plt.gca()
axes.set_xlim([-1.5, 2.5])
axes.set_ylim([-1, 1.5])
plot_decision_boundary(lambda x: predict_dec(parameters, x.T), train_X, train_Y)
```

Cost after epoch 0: 0.690741
Cost after epoch 1000: 0.685341
Cost after epoch 2000: 0.647145
Cost after epoch 3000: 0.619594
Cost after epoch 4000: 0.576665
Cost after epoch 5000: 0.607324
Cost after epoch 6000: 0.529476
Cost after epoch 7000: 0.460936
Cost after epoch 8000: 0.465780
Cost after epoch 9000: 0.464740



Accuracy: 0.796666666667



5.3 - Mini-batch with Adam mode

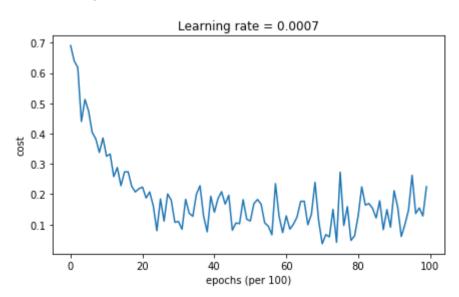
Run the following code to see how the model does with Adam.

```
In [41]: # train 3-layer model
layers_dims = [train_X.shape[0], 5, 2, 1]
parameters = model(train_X, train_Y, layers_dims, optimizer = "adam")

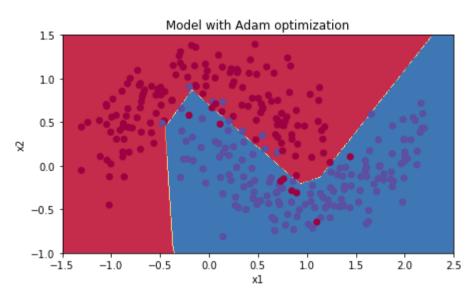
# Predict
predictions = predict(train_X, train_Y, parameters)

# Plot decision boundary
plt.title("Model with Adam optimization")
axes = plt.gca()
axes.set_ylim([-1, 5, 2, 5])
axes.set_ylim([-1, 1, 5])
plot_decision_boundary(lambda x: predict_dec(parameters, x.T), train_X, train_Y)
```

Cost after epoch 0: 0.690468
Cost after epoch 1000: 0.325335
Cost after epoch 2000: 0.223432
Cost after epoch 3000: 0.109831
Cost after epoch 4000: 0.140488
Cost after epoch 5000: 0.111569
Cost after epoch 6000: 0.128532
Cost after epoch 7000: 0.036301
Cost after epoch 8000: 0.128247
Cost after epoch 9000: 0.211582



Accuracy: 0.943333333333



5.4 - Summary

cost shape	accuracy	optimization method
oscillations	79.7%	Gradient descent
oscillations	79.7%	Momentum
smoother	94%	Adam

Momentum usually helps, but given the small learning rate and the simplistic dataset, its impact is almost negligeable. Also, the huge oscillations you see in the cost come from the fact that some minibatches are more difficult thans others for the optimization algorithm.

Adam on the other hand, clearly outperforms mini-batch gradient descent and Momentum. If you run the model for more epochs on this simple dataset, all three methods will lead to very good results. However, you've seen that Adam converges a lot faster.

Some advantages of Adam include:

- Relatively low memory requirements (though higher than gradient descent and gradient descent with momentum)
- Usually works well even with little tuning of hyperparameters (except α)

References:

• Adam paper: https://arxiv.org/pdf/1412.6980.pdf (https://arxiv.org/pdf/1412.6980.pdf)