Instructions

Welcome to the first assignment of the hyper parameters tuning specialization. It is very important that you regularize your model properly because it could dramatically improve your results.

Chart

Description automatically generated

By completing this assignment you will:

- Understand that different regularization methods that could help your model.

- Implement dropout and see it work on data.

- Recognize that a model without regularization gives you a better accuracy on the training set but nor necessarily on the test set.

- Understand that you could use both dropout and regularization on your model.

This assignment prepares you well for the upcoming assignment. Take your time to complete it and make sure you get the expected outputs when working through the different exercises. In some code blocks, you will find a "#GRADED FUNCTION: functionName" comment. Please do not modify it. After you are done, submit your work and check your results. You need to score 80% to pass. Good luck :) !

**Regularization**

Welcome to the second assignment of this week. Deep Learning models have so much flexibility and capacity that **overfitting can be a serious problem**, if the training dataset is not big enough. Sure it does well on the training set, but the learned network **doesn't generalize to new examples** that it has never seen!

**You will learn to:** Use regularization in your deep learning models.

Let's first import the packages you are going to use.

**Updates to Assignment**

**If you were working on a previous version**

* The current notebook filename is version "2a".
* You can find your work in the file directory as version "2".
* To see the file directory, click on the Coursera logo at the top left of the notebook.

**List of Updates**

* Clarified explanation of 'keep\_prob' in the text description.
* Fixed a comment so that keep\_prob and 1-keep\_prob add up to 100%
* Updated print statements and 'expected output' for easier visual comparisons.

In [1]:

*# import packages*

**import** numpy **as** np

**import** matplotlib.pyplot **as** plt

**from** reg\_utils **import** sigmoid, relu, plot\_decision\_boundary, initialize\_parameters, load\_2D\_dataset, predict\_dec

**from** reg\_utils **import** compute\_cost, predict, forward\_propagation, backward\_propagation, update\_parameters

**import** sklearn

**import** sklearn.datasets

**import** scipy.io

**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'

**Problem Statement**: You have just been hired as an AI expert by the French Football Corporation. They would like you to recommend positions where France's goal keeper should kick the ball so that the French team's players can then hit it with their head.

**Figure 1** : **Football field**  
The goal keeper kicks the ball in the air, the players of each team are fighting to hit the ball with their head

They give you the following 2D dataset from France's past 10 games.

In [2]:

train\_X, train\_Y, test\_X, test\_Y = load\_2D\_dataset()

Chart, scatter chart

Description automatically generated

Each dot corresponds to a position on the football field where a football player has hit the ball with his/her head after the French goal keeper has shot the ball from the left side of the football field.

* If the dot is blue, it means the French player managed to hit the ball with his/her head
* If the dot is red, it means the other team's player hit the ball with their head

**Your goal**: Use a deep learning model to find the positions on the field where the goalkeeper should kick the ball.

**Analysis of the dataset**: This dataset is a little noisy, but it looks like a diagonal line separating the upper left half (blue) from the lower right half (red) would work well.

You will first try a non-regularized model. Then you'll learn how to regularize it and decide which model you will choose to solve the French Football Corporation's problem.

**1 - Non-regularized model**

You will use the following neural network (already implemented for you below). This model can be used:

* in *regularization mode* -- by setting the lambd input to a non-zero value. We use "lambd" instead of "lambda" because "lambda" is a reserved keyword in Python.
* in *dropout mode* -- by setting the keep\_prob to a value less than one

You will first try the model without any regularization. Then, you will implement:

* *L2 regularization* -- functions: "compute\_cost\_with\_regularization()" and "backward\_propagation\_with\_regularization()"
* *Dropout* -- functions: "forward\_propagation\_with\_dropout()" and "backward\_propagation\_with\_dropout()"

In each part, you will run this model with the correct inputs so that it calls the functions you've implemented. Take a look at the code below to familiarize yourself with the model.

In [3]:

**def** model(X, Y, learning\_rate = 0.3, num\_iterations = 30000, print\_cost = **True**, lambd = 0, keep\_prob = 1):

"""

Implements a three-layer neural network: LINEAR->RELU->LINEAR->RELU->LINEAR->SIGMOID.

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 (output size, number of examples)

learning\_rate -- learning rate of the optimization

num\_iterations -- number of iterations of the optimization loop

print\_cost -- If True, print the cost every 10000 iterations

lambd -- regularization hyperparameter, scalar

keep\_prob - probability of keeping a neuron active during drop-out, scalar.

Returns:

parameters -- parameters learned by the model. They can then be used to predict.

"""

grads = {}

costs = [] *# to keep track of the cost*

m = X.shape[1] *# number of examples*

layers\_dims = [X.shape[0], 20, 3, 1]

*# Initialize parameters dictionary.*

parameters = initialize\_parameters(layers\_dims)

​

*# Loop (gradient descent)*

​

**for** i **in** range(0, num\_iterations):

​

*# Forward propagation: LINEAR -> RELU -> LINEAR -> RELU -> LINEAR -> SIGMOID.*

**if** keep\_prob == 1:

a3, cache = forward\_propagation(X, parameters)

**elif** keep\_prob **<** 1:

a3, cache = forward\_propagation\_with\_dropout(X, parameters, keep\_prob)

*# Cost function*

**if** lambd == 0:

cost = compute\_cost(a3, Y)

**else**:

cost = compute\_cost\_with\_regularization(a3, Y, parameters, lambd)

*# Backward propagation.*

**assert**(lambd==0 **or** keep\_prob==1) *# it is possible to use both L2 regularization and dropout,*

*# but this assignment will only explore one at a time*

**if** lambd == 0 **and** keep\_prob == 1:

grads = backward\_propagation(X, Y, cache)

**elif** lambd **!**= 0:

grads = backward\_propagation\_with\_regularization(X, Y, cache, lambd)

**elif** keep\_prob **<** 1:

grads = backward\_propagation\_with\_dropout(X, Y, cache, keep\_prob)

*# Update parameters.*

parameters = update\_parameters(parameters, grads, learning\_rate)

*# Print the loss every 10000 iterations*

**if** print\_cost **and** i **%** 10000 == 0:

print("Cost after iteration {}: {}".format(i, cost))

**if** print\_cost **and** i **%** 1000 == 0:

costs.append(cost)

*# plot the cost*

plt.plot(costs)

plt.ylabel('cost')

plt.xlabel('iterations (x1,000)')

plt.title("Learning rate =" **+** str(learning\_rate))

plt.show()

**return** parameters

Let's train the model without any regularization, and observe the accuracy on the train/test sets.

In [4]:

parameters = model(train\_X, train\_Y)

print ("On the training set:")

predictions\_train = predict(train\_X, train\_Y, parameters)

print ("On the test set:")

predictions\_test = predict(test\_X, test\_Y, parameters)

Cost after iteration 0: 0.6557412523481002

Cost after iteration 10000: 0.16329987525724213

Cost after iteration 20000: 0.13851642423253263

Shape

Description automatically generated

On the training set:

Accuracy: 0.947867298578

On the test set:

Accuracy: 0.915

The train accuracy is 94.8% while the test accuracy is 91.5%. This is the **baseline model** (you will observe the impact of regularization on this model). Run the following code to plot the decision boundary of your model.

In [5]:

plt.title("Model without regularization")

axes = plt.gca()

axes.set\_xlim([**-**0.75,0.40])

axes.set\_ylim([**-**0.75,0.65])

plot\_decision\_boundary(**lambda** x: predict\_dec(parameters, x.T), train\_X, train\_Y)

Map

Description automatically generated

The non-regularized model is obviously overfitting the training set. It is fitting the noisy points! Lets now look at two techniques to reduce overfitting.

**2 - L2 Regularization**

The standard way to avoid overfitting is called **L2 regularization**. It consists of appropriately modifying your cost function, from:

J=−1m∑i=1m(y(i)log(a[L](i))+(1−y(i))log(1−a[L](i)))(1)(1)J=−1m∑i=1m(y(i)log⁡(a[L](i))+(1−y(i))log⁡(1−a[L](i)))

To:

Jregularized=−1m∑i=1m(y(i)log(a[L](i))+(1−y(i))log(1−a[L](i)))cross-entropy cost+1mλ2∑l∑k∑jW[l]2k,jL2 regularization cost(2)(2)Jregularized=−1m∑i=1m(y(i)log⁡(a[L](i))+(1−y(i))log⁡(1−a[L](i)))⏟cross-entropy cost+1mλ2∑l∑k∑jWk,j[l]2⏟L2 regularization cost

Let's modify your cost and observe the consequences.

**Exercise**: Implement compute\_cost\_with\_regularization() which computes the cost given by formula (2). To calculate ∑k∑jW[l]2k,j∑k∑jWk,j[l]2 , use :

np.sum(np.square(Wl))

Note that you have to do this for W[1]W[1], W[2]W[2] and W[3]W[3], then sum the three terms and multiply by 1mλ21mλ2.

In [6]:

*# GRADED FUNCTION: compute\_cost\_with\_regularization*

​

**def** compute\_cost\_with\_regularization(A3, Y, parameters, lambd):

"""

Implement the cost function with L2 regularization. See formula (2) above.

Arguments:

A3 -- post-activation, output of forward propagation, of shape (output size, number of examples)

Y -- "true" labels vector, of shape (output size, number of examples)

parameters -- python dictionary containing parameters of the model

Returns:

cost - value of the regularized loss function (formula (2))

"""

m = Y.shape[1]

W1 = parameters["W1"]

W2 = parameters["W2"]

W3 = parameters["W3"]

cross\_entropy\_cost = compute\_cost(A3, Y) *# This gives you the cross-entropy part of the cost*

*### START CODE HERE ### (approx. 1 line)*

L2\_regularization\_cost = (1 **/** m) **\*** (lambd **/** 2) **\*** (np.sum(np.square(W1)) **+** np.sum(np.square(W2)) **+** np.sum(np.square(W3)))

*### END CODER HERE ###*

cost = cross\_entropy\_cost **+** L2\_regularization\_cost

**return** cost

In [7]:

A3, Y\_assess, parameters = compute\_cost\_with\_regularization\_test\_case()

​

print("cost = " **+** str(compute\_cost\_with\_regularization(A3, Y\_assess, parameters, lambd = 0.1)))

cost = 1.78648594516

**Expected Output**:

|  |  |
| --- | --- |
| **cost** | 1.78648594516 |

Of course, because you changed the cost, you have to change backward propagation as well! All the gradients have to be computed with respect to this new cost.

**Exercise**: Implement the changes needed in backward propagation to take into account regularization. The changes only concern dW1, dW2 and dW3. For each, you have to add the regularization term's gradient (ddW(12λmW2)=λmWddW(12λmW2)=λmW).

In [8]:

*# GRADED FUNCTION: backward\_propagation\_with\_regularization*

​

**def** backward\_propagation\_with\_regularization(X, Y, cache, lambd):

"""

Implements the backward propagation of our baseline model to which we added an L2 regularization.

Arguments:

X -- input dataset, of shape (input size, number of examples)

Y -- "true" labels vector, of shape (output size, number of examples)

cache -- cache output from forward\_propagation()

lambd -- regularization hyperparameter, scalar

Returns:

gradients -- A dictionary with the gradients with respect to each parameter, activation and pre-activation variables

"""

m = X.shape[1]

(Z1, A1, W1, b1, Z2, A2, W2, b2, Z3, A3, W3, b3) = cache

dZ3 = A3 **-** Y

*### START CODE HERE ### (approx. 1 line)*

dW3 = 1.**/**m **\*** np.dot(dZ3, A2.T) **+** (lambd **/** m) **\*** W3

*### END CODE HERE ###*

db3 = 1.**/**m **\*** np.sum(dZ3, axis=1, keepdims = **True**)

dA2 = np.dot(W3.T, dZ3)

dZ2 = np.multiply(dA2, np.int64(A2 **>** 0))

*### START CODE HERE ### (approx. 1 line)*

dW2 = 1.**/**m **\*** np.dot(dZ2, A1.T) **+** (lambd **/** m) **\*** W2

*### END CODE HERE ###*

db2 = 1.**/**m **\*** np.sum(dZ2, axis=1, keepdims = **True**)

dA1 = np.dot(W2.T, dZ2)

dZ1 = np.multiply(dA1, np.int64(A1 **>** 0))

*### START CODE HERE ### (approx. 1 line)*

dW1 = 1.**/**m **\*** np.dot(dZ1, X.T) **+** (lambd **/** m) **\*** W1

*### END CODE HERE ###*

db1 = 1.**/**m **\*** np.sum(dZ1, axis=1, keepdims = **True**)

gradients = {"dZ3": dZ3, "dW3": dW3, "db3": db3,"dA2": dA2,

"dZ2": dZ2, "dW2": dW2, "db2": db2, "dA1": dA1,

"dZ1": dZ1, "dW1": dW1, "db1": db1}

**return** gradients

In [9]:

X\_assess, Y\_assess, cache = backward\_propagation\_with\_regularization\_test\_case()

​

grads = backward\_propagation\_with\_regularization(X\_assess, Y\_assess, cache, lambd = 0.7)

print ("dW1 = \n"**+** str(grads["dW1"]))

print ("dW2 = \n"**+** str(grads["dW2"]))

print ("dW3 = \n"**+** str(grads["dW3"]))

dW1 =

[[-0.25604646 0.12298827 -0.28297129]

[-0.17706303 0.34536094 -0.4410571 ]]

dW2 =

[[ 0.79276486 0.85133918]

[-0.0957219 -0.01720463]

[-0.13100772 -0.03750433]]

dW3 =

[[-1.77691347 -0.11832879 -0.09397446]]

**Expected Output**:

dW1 =

[[-0.25604646 0.12298827 -0.28297129]

[-0.17706303 0.34536094 -0.4410571 ]]

dW2 =

[[ 0.79276486 0.85133918]

[-0.0957219 -0.01720463]

[-0.13100772 -0.03750433]]

dW3 =

[[-1.77691347 -0.11832879 -0.09397446]]

Let's now run the model with L2 regularization (λ=0.7)(λ=0.7). The model() function will call:

* compute\_cost\_with\_regularization instead of compute\_cost
* backward\_propagation\_with\_regularization instead of backward\_propagation

In [10]:

parameters = model(train\_X, train\_Y, lambd = 0.7)

print ("On the train set:")

predictions\_train = predict(train\_X, train\_Y, parameters)

print ("On the test set:")

predictions\_test = predict(test\_X, test\_Y, parameters)

Cost after iteration 0: 0.6974484493131264

Cost after iteration 10000: 0.2684918873282239

Cost after iteration 20000: 0.26809163371273015

Shape

Description automatically generated

On the train set:

Accuracy: 0.938388625592

On the test set:

Accuracy: 0.93

Congrats, the test set accuracy increased to 93%. You have saved the French football team!

You are not overfitting the training data anymore. Let's plot the decision boundary.

In [11]:

plt.title("Model with L2-regularization")

axes = plt.gca()

axes.set\_xlim([**-**0.75,0.40])

axes.set\_ylim([**-**0.75,0.65])

plot\_decision\_boundary(**lambda** x: predict\_dec(parameters, x.T), train\_X, train\_Y)

Chart

Description automatically generated

**Observations**:

* The value of λλ is a hyperparameter that you can tune using a dev set.
* L2 regularization makes your decision boundary smoother. If λλ is too large, it is also possible to "oversmooth", resulting in a model with high bias.

**What is L2-regularization actually doing?**:

L2-regularization relies on the assumption that a model with small weights is simpler than a model with large weights. Thus, by penalizing the square values of the weights in the cost function you drive all the weights to smaller values. It becomes too costly for the cost to have large weights! This leads to a smoother model in which the output changes more slowly as the input changes.

**What you should remember** -- the implications of L2-regularization on:

* The cost computation:
  + A regularization term is added to the cost
* The backpropagation function:
  + There are extra terms in the gradients with respect to weight matrices
* Weights end up smaller ("weight decay"):
  + Weights are pushed to smaller values.

**3 - Dropout**

Finally, **dropout** is a widely used regularization technique that is specific to deep learning. **It randomly shuts down some neurons in each iteration.** Watch these two videos to see what this means!

Figure 2 : Drop-out on the second hidden layer.  
At each iteration, you shut down (= set to zero) each neuron of a layer with probability 1−keep\_prob1−keep\_prob or keep it with probability keep\_probkeep\_prob (50% here). The dropped neurons don't contribute to the training in both the forward and backward propagations of the iteration.

Figure 3 : Drop-out on the first and third hidden layers.  
1st1st layer: we shut down on average 40% of the neurons. 3rd3rd layer: we shut down on average 20% of the neurons.

When you shut some neurons down, you actually modify your model. The idea behind drop-out is that at each iteration, you train a different model that uses only a subset of your neurons. With dropout, your neurons thus become less sensitive to the activation of one other specific neuron, because that other neuron might be shut down at any time.

**3.1 - Forward propagation with dropout**

**Exercise**: Implement the forward propagation with dropout. You are using a 3 layer neural network, and will add dropout to the first and second hidden layers. We will not apply dropout to the input layer or output layer.

**Instructions**: You would like to shut down some neurons in the first and second layers. To do that, you are going to carry out 4 Steps:

1. In lecture, we dicussed creating a variable d[1]d[1] with the same shape as a[1]a[1] using np.random.rand() to randomly get numbers between 0 and 1. Here, you will use a vectorized implementation, so create a random matrix D[1]=[d[1](1)d[1](2)...d[1](m)]D[1]=[d[1](1)d[1](2)...d[1](m)] of the same dimension as A[1]A[1].
2. Set each entry of D[1]D[1] to be 1 with probability (keep\_prob), and 0 otherwise.

**Hint:** Let's say that keep\_prob = 0.8, which means that we want to keep about 80% of the neurons and drop out about 20% of them. We want to generate a vector that has 1's and 0's, where about 80% of them are 1 and about 20% are 0. This python statement:  
X = (X < keep\_prob).astype(int)

is conceptually the same as this if-else statement (for the simple case of a one-dimensional array) :

for i,v in enumerate(x):

if v < keep\_prob:

x[i] = 1

else: # v >= keep\_prob

x[i] = 0

Note that the X = (X < keep\_prob).astype(int) works with multi-dimensional arrays, and the resulting output preserves the dimensions of the input array.

Also note that without using .astype(int), the result is an array of booleans True and False, which Python automatically converts to 1 and 0 if we multiply it with numbers. (However, it's better practice to convert data into the data type that we intend, so try using .astype(int).)

1. Set A[1]A[1] to A[1]∗D[1]A[1]∗D[1]. (You are shutting down some neurons). You can think of D[1]D[1] as a mask, so that when it is multiplied with another matrix, it shuts down some of the values.
2. Divide A[1]A[1] by keep\_prob. By doing this you are assuring that the result of the cost will still have the same expected value as without drop-out. (This technique is also called inverted dropout.)

In [14]:

*# GRADED FUNCTION: forward\_propagation\_with\_dropout*

​

**def** forward\_propagation\_with\_dropout(X, parameters, keep\_prob = 0.5):

"""

Implements the forward propagation: LINEAR -> RELU + DROPOUT -> LINEAR -> RELU + DROPOUT -> LINEAR -> SIGMOID.

Arguments:

X -- input dataset, of shape (2, number of examples)

parameters -- python dictionary containing your parameters "W1", "b1", "W2", "b2", "W3", "b3":

W1 -- weight matrix of shape (20, 2)

b1 -- bias vector of shape (20, 1)

W2 -- weight matrix of shape (3, 20)

b2 -- bias vector of shape (3, 1)

W3 -- weight matrix of shape (1, 3)

b3 -- bias vector of shape (1, 1)

keep\_prob - probability of keeping a neuron active during drop-out, scalar

Returns:

A3 -- last activation value, output of the forward propagation, of shape (1,1)

cache -- tuple, information stored for computing the backward propagation

"""

np.random.seed(1)

*# retrieve parameters*

W1 = parameters["W1"]

b1 = parameters["b1"]

W2 = parameters["W2"]

b2 = parameters["b2"]

W3 = parameters["W3"]

b3 = parameters["b3"]

*# LINEAR -> RELU -> LINEAR -> RELU -> LINEAR -> SIGMOID*

Z1 = np.dot(W1, X) **+** b1

A1 = relu(Z1)

*### START CODE HERE ### (approx. 4 lines) # Steps 1-4 below correspond to the Steps 1-4 described above.*

D1 = np.random.rand(A1.shape[0], A1.shape[1]) *# Step 1: initialize matrix D1 = np.random.rand(..., ...)*

D1 = D1 **<** keep\_prob *# Step 2: convert entries of D1 to 0 or 1 (using keep\_prob as the threshold)*

A1 = A1 **\*** D1 *# Step 3: shut down some neurons of A1*

A1 = A1 **/** keep\_prob *# Step 4: scale the value of neurons that haven't been shut down*

*### END CODE HERE ###*

Z2 = np.dot(W2, A1) **+** b2

A2 = relu(Z2)

*### START CODE HERE ### (approx. 4 lines)*

D2 = np.random.rand(A2.shape[0], A2.shape[1]) *# Step 1: initialize matrix D2 = np.random.rand(..., ...)*

D2 = D2 **<** keep\_prob *# Step 2: convert entries of D2 to 0 or 1 (using keep\_prob as the threshold)*

A2 = A2 **\*** D2 *# Step 3: shut down some neurons of A2*

A2 = A2 **/** keep\_prob *# Step 4: scale the value of neurons that haven't been shut down*

*### END CODE HERE ###*

Z3 = np.dot(W3, A2) **+** b3

A3 = sigmoid(Z3)

cache = (Z1, D1, A1, W1, b1, Z2, D2, A2, W2, b2, Z3, A3, W3, b3)

**return** A3, cache

In [15]:

X\_assess, parameters = forward\_propagation\_with\_dropout\_test\_case()

​

A3, cache = forward\_propagation\_with\_dropout(X\_assess, parameters, keep\_prob = 0.7)

print ("A3 = " **+** str(A3))

A3 = [[ 0.36974721 0.00305176 0.04565099 0.49683389 0.36974721]]

**Expected Output**:

|  |  |
| --- | --- |
| **A3** | [[ 0.36974721 0.00305176 0.04565099 0.49683389 0.36974721]] |

**3.2 - Backward propagation with dropout**

**Exercise**: Implement the backward propagation with dropout. As before, you are training a 3 layer network. Add dropout to the first and second hidden layers, using the masks D[1]D[1] and D[2]D[2] stored in the cache.

**Instruction**: Backpropagation with dropout is actually quite easy. You will have to carry out 2 Steps:

1. You had previously shut down some neurons during forward propagation, by applying a mask D[1]D[1] to A1. In backpropagation, you will have to shut down the same neurons, by reapplying the same mask D[1]D[1] to dA1.
2. During forward propagation, you had divided A1 by keep\_prob. In backpropagation, you'll therefore have to divide dA1 by keep\_prob again (the calculus interpretation is that if A[1]A[1] is scaled by keep\_prob, then its derivative dA[1]dA[1] is also scaled by the same keep\_prob).

In [17]:

*# GRADED FUNCTION: backward\_propagation\_with\_dropout*

​

**def** backward\_propagation\_with\_dropout(X, Y, cache, keep\_prob):

"""

Implements the backward propagation of our baseline model to which we added dropout.

Arguments:

X -- input dataset, of shape (2, number of examples)

Y -- "true" labels vector, of shape (output size, number of examples)

cache -- cache output from forward\_propagation\_with\_dropout()

keep\_prob - probability of keeping a neuron active during drop-out, scalar

Returns:

gradients -- A dictionary with the gradients with respect to each parameter, activation and pre-activation variables

"""

m = X.shape[1]

(Z1, D1, A1, W1, b1, Z2, D2, A2, W2, b2, Z3, A3, W3, b3) = cache

dZ3 = A3 **-** Y

dW3 = 1.**/**m **\*** np.dot(dZ3, A2.T)

db3 = 1.**/**m **\*** np.sum(dZ3, axis=1, keepdims = **True**)

dA2 = np.dot(W3.T, dZ3)

*### START CODE HERE ### (≈ 2 lines of code)*

dA2 = dA2 **\*** D2 *# Step 1: Apply mask D2 to shut down the same neurons as during the forward propagation*

dA2 = dA2 **/** keep\_prob *# Step 2: Scale the value of neurons that haven't been shut down*

*### END CODE HERE ###*

dZ2 = np.multiply(dA2, np.int64(A2 **>** 0))

dW2 = 1.**/**m **\*** np.dot(dZ2, A1.T)

db2 = 1.**/**m **\*** np.sum(dZ2, axis=1, keepdims = **True**)

dA1 = np.dot(W2.T, dZ2)

*### START CODE HERE ### (≈ 2 lines of code)*

dA1 = dA1 **\*** D1 *# Step 1: Apply mask D1 to shut down the same neurons as during the forward propagation*

dA1 = dA1 **/** keep\_prob *# Step 2: Scale the value of neurons that haven't been shut down*

*### END CODE HERE ###*

dZ1 = np.multiply(dA1, np.int64(A1 **>** 0))

dW1 = 1.**/**m **\*** np.dot(dZ1, X.T)

db1 = 1.**/**m **\*** np.sum(dZ1, axis=1, keepdims = **True**)

gradients = {"dZ3": dZ3, "dW3": dW3, "db3": db3,"dA2": dA2,

"dZ2": dZ2, "dW2": dW2, "db2": db2, "dA1": dA1,

"dZ1": dZ1, "dW1": dW1, "db1": db1}

**return** gradients

In [18]:

X\_assess, Y\_assess, cache = backward\_propagation\_with\_dropout\_test\_case()

​

gradients = backward\_propagation\_with\_dropout(X\_assess, Y\_assess, cache, keep\_prob = 0.8)

​

print ("dA1 = \n" **+** str(gradients["dA1"]))

print ("dA2 = \n" **+** str(gradients["dA2"]))

dA1 =

[[ 0.36544439 0. -0.00188233 0. -0.17408748]

[ 0.65515713 0. -0.00337459 0. -0. ]]

dA2 =

[[ 0.58180856 0. -0.00299679 0. -0.27715731]

[ 0. 0.53159854 -0. 0.53159854 -0.34089673]

[ 0. 0. -0.00292733 0. -0. ]]

**Expected Output**:

dA1 =

[[ 0.36544439 0. -0.00188233 0. -0.17408748]

[ 0.65515713 0. -0.00337459 0. -0. ]]

dA2 =

[[ 0.58180856 0. -0.00299679 0. -0.27715731]

[ 0. 0.53159854 -0. 0.53159854 -0.34089673]

[ 0. 0. -0.00292733 0. -0. ]]

Let's now run the model with dropout (keep\_prob = 0.86). It means at every iteration you shut down each neurons of layer 1 and 2 with 14% probability. The function model() will now call:

* forward\_propagation\_with\_dropout instead of forward\_propagation.
* backward\_propagation\_with\_dropout instead of backward\_propagation.

In [19]:

parameters = model(train\_X, train\_Y, keep\_prob = 0.86, learning\_rate = 0.3)

​

print ("On the train set:")

predictions\_train = predict(train\_X, train\_Y, parameters)

print ("On the test set:")

predictions\_test = predict(test\_X, test\_Y, parameters)

Cost after iteration 0: 0.6543912405149825

/home/jovyan/work/week5/Regularization/reg\_utils.py:236: RuntimeWarning: divide by zero encountered in log

logprobs = np.multiply(-np.log(a3),Y) + np.multiply(-np.log(1 - a3), 1 - Y)

/home/jovyan/work/week5/Regularization/reg\_utils.py:236: RuntimeWarning: invalid value encountered in multiply

logprobs = np.multiply(-np.log(a3),Y) + np.multiply(-np.log(1 - a3), 1 - Y)

Cost after iteration 10000: 0.06101698657490562

Cost after iteration 20000: 0.060582435798513114

Shape, rectangle

Description automatically generated

On the train set:

Accuracy: 0.928909952607

On the test set:

Accuracy: 0.95

Dropout works great! The test accuracy has increased again (to 95%)! Your model is not overfitting the training set and does a great job on the test set. The French football team will be forever grateful to you!

Run the code below to plot the decision boundary.

In [20]:

plt.title("Model with dropout")

axes = plt.gca()

axes.set\_xlim([**-**0.75,0.40])

axes.set\_ylim([**-**0.75,0.65])

plot\_decision\_boundary(**lambda** x: predict\_dec(parameters, x.T), train\_X, train\_Y)

Chart

Description automatically generated

**Note**:

* A **common mistake** when using dropout is to use it both in training and testing. You should use dropout (randomly eliminate nodes) only in training.
* Deep learning frameworks like [tensorflow](https://www.tensorflow.org/api_docs/python/tf/nn/dropout), [PaddlePaddle](http://doc.paddlepaddle.org/release_doc/0.9.0/doc/ui/api/trainer_config_helpers/attrs.html), [keras](https://keras.io/layers/core/#dropout) or [caffe](http://caffe.berkeleyvision.org/tutorial/layers/dropout.html) come with a dropout layer implementation. Don't stress - you will soon learn some of these frameworks.

**What you should remember about dropout:**

* Dropout is a regularization technique.
* You only use dropout during training. Don't use dropout (randomly eliminate nodes) during test time.
* Apply dropout both during forward and backward propagation.
* During training time, divide each dropout layer by keep\_prob to keep the same expected value for the activations. For example, if keep\_prob is 0.5, then we will on average shut down half the nodes, so the output will be scaled by 0.5 since only the remaining half are contributing to the solution. Dividing by 0.5 is equivalent to multiplying by 2. Hence, the output now has the same expected value. You can check that this works even when keep\_prob is other values than 0.5.

**4 - Conclusions**

**Here are the results of our three models**:

|  |  |  |
| --- | --- | --- |
| **model** | **train accuracy** | **test accuracy** |
| 3-layer NN without regularization | 95% | 91.5% |
| 3-layer NN with L2-regularization | 94% | 93% |
| 3-layer NN with dropout | 93% | 95% |

Note that regularization hurts training set performance! This is because it limits the ability of the network to overfit to the training set. But since it ultimately gives better test accuracy, it is helping your system.

Congratulations for finishing this assignment! And also for revolutionizing French football. :-)

**What we want you to remember from this notebook**:

* Regularization will help you reduce overfitting.
* Regularization will drive your weights to lower values.
* L2 regularization and Dropout are two very effective regularization techniques.