Machine learning Exercise sheet 6 Optimization

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# Convexity of functions

## Problem (1):

a) 
$$h(x) = g_2(g_1(x))$$
  
 $h'(x) = g_2' [g_1(x)] \cdot g_1'(x)$   
 $h''(x) = g_2'' [g_1(x)] [g_1'(x)]^2 + g_2' [g_1(x)] \cdot g''(x)$   
 $0 > 0 > 0$   
We know that  $g_1(x) \Rightarrow convex$  and  $g_2(x) \Rightarrow convex$   
then:  $g_1''(x) > 0$   
 $g_2''(x) > 0$   
 $g_2''(x) = g_2'' [g_1(x)] \cdot g_1'(x)$ 

In order h(x) to be convex, then h"(x) >,0, but we don't know what 92' [91(x)] is. It can be both positive and negative. So the function is not longer.

b) 
$$h(x) = 92(91(x))$$
  
 $h'(x) = 92'[91(x)] \cdot 91(x)$   
 $h''(x) = 92'[91(x)] \cdot [91(x)] + 92'[91(x)] \cdot 9''(x)$   
 $h''(x) = 92'[91(x)] \cdot [91(x)] + 92'[91(x)] \cdot 9''(x)$   
 $91/92 \quad convex \rightarrow (92(x))'' > 0$   
 $91/(x) > 0$   
 $91/(x) > 0$   
 $92 \quad non \ decreasing \rightarrow 92'(x) > 0$ 

Therefore h(x) convex.

C) Pick any 
$$x,y \in dom(h)$$
,  $\lambda \in [0,1]$ .  
Then,  
 $h(\lambda x + (1-\lambda)y) = gi(\lambda x + (1-\lambda)y)$   
for some  $j \in \{1,...,m\}$   
 $gj(onvex \langle \lambda gj(x) + (1-\lambda)gj(y)$ 

λ max(g1, , , gn(x))+(1-2)max (91(y)...9;(y)

Problem (Problem) Problem (2)

a) 
$$\frac{df}{dx_1} = x_1 + 2$$
,  $\frac{df}{dx_2} = 2x_2 + 1$ 

$$\nabla f(X_1, X_2) = \begin{bmatrix} X_1 + 2 \\ 2X_2 + 1 \end{bmatrix}$$

$$\nabla f = 0 \iff \boxed{X_1 = -2} \quad \boxed{X_2 = -\frac{1}{2}}$$

$$Hf(X_1, X_2) = \begin{bmatrix} 1 & 0 \\ 0 & 2 \end{bmatrix}$$

To determine, whether a matrix is positive definite, we first compute it's determinant.

$$det(H) = 2-0 = 2 > 0$$
 (1)

The product of the eigenvalues is positive, which means they have same sign.

We check the tr(H) which is the sum of eigenvalues in order to determine the sign.  $tr(H) = 1+2 = 3(2) \Longrightarrow both eigenvalues positive$ 

Therefore H > positive definite and (X1, X2)=(-2, 3

b) 
$$(X_{1}^{n}) = (X_{1}^{n-1}) - \alpha \nabla f(X_{1}^{n-1}, X_{2}^{n-1})^{2}$$

our original function value is:
$$f(o, o) = \cos(\sin(\pi)) = 0.9999.$$

$$\nabla f(X_{1}, X_{2}) = \begin{bmatrix} X_{1} + 2 \\ 2X_{2} + 1 \end{bmatrix}$$

$$\nabla f(o, o) = \begin{bmatrix} 2 \\ 1 \end{bmatrix}$$

Step 1
$$(X_{1}^{n}) = \begin{bmatrix} 0 \\ 0 \end{bmatrix} - \begin{bmatrix} 2 \\ 1 \end{bmatrix} = \begin{bmatrix} -2 \\ -1 \end{bmatrix}$$

$$f(X_{1}^{n}, X_{2}^{n}) = -1,000000146$$

Step 2
$$\nabla f(X_{1}^{n}, X_{2}^{n}) = \begin{bmatrix} 0 \\ -1 \end{bmatrix} = \begin{bmatrix} -2 \\ 0 \end{bmatrix}$$

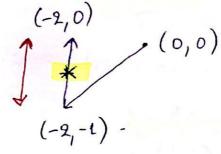
$$f(X_{1}^{n}, X_{2}^{n}) = -1,000000146$$

Minimum value for  $(X_{1}^{n}, X_{2}^{n}) = -1,000000146$ 

 $f(X_1, X_2) = -1.750000146$ 

c) if we draw the isolines we see the following behaviour:

# : MINIMUM



Gradient descent will not converge to the minimum X\*. As we see above, we overshoot and then oscilate between the points (-2,-1) and (-2,0). In order to overcome this problem we need smaller step, which means smaller learning rate. We could also have adaptive learning rate where at the beginning we do larger steps and later on we do smaller in order not to overshoot the minimum.

# Problem (4):

a) The shaded region is not convex because we cannot pass between the points (3.5, L) and (6, 3.5) without leaving the shaded region.

b) In order to find maximizer x\* of of over the shaded region S we will split the mon-convex set into maker ones which are the following ones.

4 triangles

L square in the middle.

Afterwards, we will find the maximum value in each one of the convex sets and then pick the largest one.

We start working on the right triangle:

Maximum over a convex function on a convex set is obtained on a vertex.

Therefore, we have to check the vertexes in the right triangle. The vertexes are the following

- . (6,3.5)
- · (H.5,3)
- . (4.5,4)

$$f(6,3.5) = e^{6+3.5} - 5 \log(3.5) = 13357,00649$$

$$f(6,3.5) = e^{4.5+3} - 5 \log(3) = 1805,0322114$$

$$f(4.5,3) = e^{4.5+3} - 5 \log(3) = 4911,75854$$

$$f(4.5,4) = e^{4.5+4} - 5 \log(4) = 4911,75854$$

We do the same process for the rest convex sets and we choose the maximum value out of all of them.

c) in order to find the minimum we can follow the same process as before but instead we will transform the function to concave (-f(xi,x) and find minimum in the vertices.

## exercise\_06\_optimization

November 24, 2019

### 1 Programming assignment 3: Optimization - Logistic Regression

```
[1]: import numpy as np
  import matplotlib.pyplot as plt
  %matplotlib inline

from sklearn.datasets import load_breast_cancer
  from sklearn.model_selection import train_test_split
  from sklearn.metrics import accuracy_score, f1_score
```

#### 1.1 Your task

In this notebook code skeleton for performing logistic regression with gradient descent is given. Your task is to complete the functions where required. You are only allowed to use built-in Python functions, as well as any numpy functions. No other libraries / imports are allowed.

For numerical reasons, we actually minimize the following loss function

$$\mathcal{L}(\mathbf{w}) = \frac{1}{N} NLL(\mathbf{w}) + \frac{1}{2} \lambda ||\mathbf{w}||_2^2$$

where  $NLL(\mathbf{w})$  is the negative log-likelihood function, as defined in the lecture (see Eq. 33).

#### 1.2 Exporting the results to PDF

Once you complete the assignments, export the entire notebook as PDF and attach it to your homework solutions. The best way of doing that is 1. Run all the cells of the notebook. 2. Export/download the notebook as PDF (File -> Download as -> PDF via LaTeX (.pdf)). 3. Concatenate your solutions for other tasks with the output of Step 2. On a Linux machine you can simply use pdfunite, there are similar tools for other platforms too. You can only upload a single PDF file to Moodle.

Make sure you are using nbconvert Version 5.5 or later by running jupyter nbconvert --version. Older versions clip lines that exceed page width, which makes your code harder to grade.

#### 1.3 Load and preprocess the data

In this assignment we will work with the UCI ML Breast Cancer Wisconsin (Diagnostic) dataset https://goo.gl/U2Uwz2.

Features are computed from a digitized image of a fine needle aspirate (FNA) of a breast mass. They describe characteristics of the cell nuclei present in the image. There are 212 malignant examples and 357 benign examples.

```
[2]: X, y = load_breast_cancer(return_X_y=True)

# Add a vector of ones to the data matrix to absorb the bias term
X = np.hstack([np.ones([X.shape[0], 1]), X])

# Set the random seed so that we have reproducible experiments
np.random.seed(123)

# Split into train and test
test_size = 0.3
X_train, X_test, y_train, y_test = train_test_split(X, y, test_size=test_size)
```

#### 1.4 Task 1: Implement the sigmoid function

```
[4]: def sigmoid(t):
    """
    Applies the sigmoid function elementwise to the input data.

Parameters
-----
t: array, arbitrary shape
    Input data.

Returns
-----
t_sigmoid: array, arbitrary shape.
    Data after applying the sigmoid function.
"""

t_sigmoid = 1/ (1+np.exp(-t))
return t_sigmoid
```

#### 1.5 Task 2: Implement the negative log likelihood

As defined in Eq. 33

```
[7]: def negative_log_likelihood(X, y, w):
    """

Negative Log Likelihood of the Logistic Regression.
```

#### 1.5.1 Computing the loss function $\mathcal{L}(\mathbf{w})$ (nothing to do here)

```
[8]: def compute_loss(X, y, w, lmbda):
         Negative Log Likelihood of the Logistic Regression.
         Parameters
         X : array, shape [N, D]
             (Augmented) feature matrix.
         y : array, shape [N]
             Classification targets.
         w : array, shape [D]
             Regression coefficients (w[0] is the bias term).
         lmbda : float
             L2 regularization strength.
         Returns
         _____
         loss : float
             Loss of the regularized logistic regression model.
         # The bias term w[0] is not regularized by convention
         return negative_log_likelihood(X, y, w) / len(y) + lmbda * 0.5 * np.linalg.
      \rightarrownorm(w[1:])**2
```

#### 1.6 Task 3: Implement the gradient $\nabla_{\mathbf{w}} \mathcal{L}(\mathbf{w})$

Make sure that you compute the gradient of the loss function  $\mathcal{L}(\mathbf{w})$  (not simply the NLL!)

```
[9]: def get_gradient(X, y, w, mini_batch_indices, lmbda):
         Calculates the gradient (full or mini-batch) of the negative log likelilhood_
      \hookrightarrow w.r.t. w.
         Parameters
         X: array, shape [N, D]
             (Augmented) feature matrix.
         y : array, shape [N]
             Classification targets.
         w : array, shape [D]
             Regression coefficients (w[0] is the bias term).
         mini_batch_indices: array, shape [mini_batch_size]
              The indices of the data points to be included in the (stochastic)_{\sqcup}
      \rightarrow calculation of the gradient.
              This includes the full batch gradient as well, if mini_batch_indices =_ 
      \hookrightarrow np.arange(n\_train).
          lmbda: float
              Regularization strentgh. lmbda = 0 means having no regularization.
         Returns
         dw : array, shape [D]
             Gradient w.r.t. w.
         X = X[mini_batch_indices]
         y = y[mini_batch_indices]
         return (1/len(X)) * X.T.dot(sigmoid(w.dot(X.T)) - y) - lmbda * w
```

#### 1.6.1 Train the logistic regression model (nothing to do here)

```
The learning rate to use when updating the parameters w.
   mini_batch_size: int
       The number of examples in each mini-batch.
       If mini_batch_size=n_train we perform full batch gradient descent.
   lmbda: float
       Regularization strentgh. lmbda = 0 means having no regularization.
   verbose : bool
       Whether to print the loss during optimization.
   Returns
   w : array, shape [D]
      Optimal regression coefficients (w[0] is the bias term).
   trace: list
      Trace of the loss function after each step of gradient descent.
  trace = [] # saves the value of loss every 50 iterations to be able to plotu
\rightarrow it later
  n_train = X.shape[0] # number of training instances
  w = np.zeros(X.shape[1]) # initialize the parameters to zeros
   # run gradient descent for a given number of steps
  for step in range(num_steps):
       permuted_idx = np.random.permutation(n_train) # shuffle the data
       # go over each mini-batch and update the paramters
       # if mini_batch_size = n_train we perform full batch GD and this loop_l
→runs only once
       for idx in range(0, n_train, mini_batch_size):
           # get the random indices to be included in the mini batch
           mini_batch_indices = permuted_idx[idx:idx+mini_batch_size]
           gradient = get_gradient(X, y, w, mini_batch_indices, lmbda)
           # update the parameters
           w = w - learning_rate * gradient
       # calculate and save the current loss value every 50 iterations
       if step \% 50 == 0:
           loss = compute_loss(X, y, w, lmbda)
           trace.append(loss)
           # print loss to monitor the progress
           if verbose:
               print('Step {0}, loss = {1:.4f}'.format(step, loss))
   return w, trace
```

#### 1.7 Task 4: Implement the function to obtain the predictions

#### 1.7.1 Full batch gradient descent

[12]: # Change this to True if you want to see loss values over iterations.

```
mini_batch_size=n_train,
lmbda=0.1,
verbose=verbose)
```

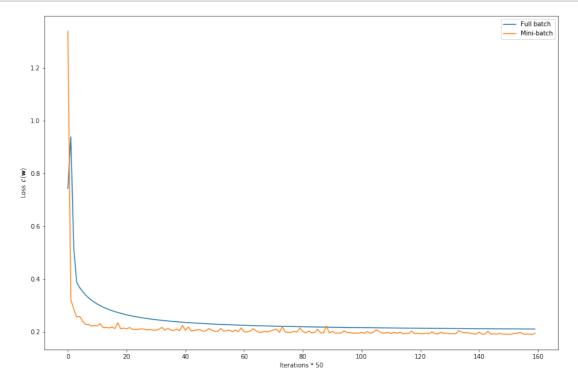
learning\_rate=1e-5,

Our reference solution produces, but don't worry if yours is not exactly the same.

```
Full batch: accuracy: 0.9240, f1_score: 0.9384 Mini-batch: accuracy: 0.9415, f1_score: 0.9533
```

Full batch: accuracy: 0.9240, f1\_score: 0.9384 Mini-batch: accuracy: 0.9415, f1\_score: 0.9533

```
[16]: plt.figure(figsize=[15, 10])
   plt.plot(trace_full, label='Full batch')
   plt.plot(trace_minibatch, label='Mini-batch')
   plt.xlabel('Iterations * 50')
   plt.ylabel('Loss $\mathcal{L}(\mathbf{w})$')
   plt.legend()
   plt.show()
```



[]: