STATS 542: Homework 5

Diego Kleiman (diegoek2)

Due: Tuesday 11:59 PM CT, Mar 9th

About HW5

We utilize the coordinate descent algorithm introduced in the class with the one variable Lasso algorithm from the last homework to complete the entire Lasso solution. This involves two steps: in the first step, we solve the solution for a fixed λ value, while in the second step, we consider a grid of λ value and solve it using the path-wise coordinate descent.

Question 1 [40 Points] Lasso solution for fixed λ

For this question, you cannot use functions from any additional library in your algorithm. Following HW4, we use the this version of the objective function:

$$\arg\min_{\beta} \frac{1}{2n} \|\mathbf{y} - \mathbf{X}\boldsymbol{\beta}\|^2 + \lambda \|\boldsymbol{\beta}\|_1$$

The following data is used to fit this model. Note that the MASS package can only be used to generate multivariate normal data. You can consider using similar functions in Python if needed.

```
library(MASS)
  set.seed(10)
  n = 100
  p = 200

# generate data
V = matrix(0.3, p, p)
  diag(V) = 1

X_org = as.matrix(mvrnorm(n, mu = rep(0, p), Sigma = V))
  true_b = c(runif(10, -1, 1), rep(0, p-10))
  y_org = X_org %*% true_b + rnorm(n)

# pre-scale and center X and y

X = scale(X_org)*sqrt(n/(n-1))
  y = scale(y_org)*sqrt(n/(n-1))
  lambda = 0.3
```

We will use the pre-scale and centered data x and y for this question, hence no intercept is needed. Write a Lasso algorithm function myLasso(x, y, lambda, tol, maxitr) which will output a vector of β values **without** the intercept. You need to consider the following while completing this question:

- · Do not use functions from any additional library
- Start with a vector $\beta = 0$
- Use the soft-threshold function you developed in HW4.
- Use the efficient **r** update algorithm we introduced during the lecture.
- Run your coordinate descent algorithm for a maximum of $\mathtt{maxitr} = 100$ iterations (while each iteration will loop through all variables). However, stop your algorithm if the β value of the current iteration is sufficiently similar to the previous one, i.e., $\|\beta^{(k)} \beta^{(k-1)}\|^2 \le \text{tol}$. Set tol = 1e-7.
- After running the algorithm, print out the first 10 variables.
- Finally, check and compare your answer to the glmnet package using the following code:

```
library(glmnet)
   glmnetfit = glmnet(X, y, lambda = 0.3, intercept = FALSE)
   glmnetfit$beta[1:10]

In [1]: import numpy as np
   from scipy.stats import multivariate_normal # Python equivalent of mvrnorm
   from sklearn.preprocessing import scale

In [2]: seed = 1

In [3]: np.random.seed(seed)
```

```
In [4]: n = 100
p = 200

V = np.ones((p, p))*0.3
np.fill_diagonal(V, 1)
X_org = np.asarray(multivariate_normal(mean=np.zeros(p), cov=V, seed=seed).rvs(n))
true_b = np.append(np.random.uniform(low=-1, high=1, size=10), np.zeros(p-10))
y_org = np.matmul(X_org, true_b) + np.random.normal(n)

# Python adaptation note: the function sklearn.preprocessing.scale does NOT divide by (n-1) to comput
e the standard
# deviation. Instead it uses n. For this reason, we do not need to multiply by sqrt(n/(n-1)) (if we d
id, we wouldn't
# get Xj^T*Xj=n).

X = scale(X_org)
y = scale(Y_org)
1 = 0.3 # lambda is a reserved keyword in Python
```

```
In [5]: | def myLasso(X, y, 1, tol=1e-7, maxitr=100):
            Fit Lasso regression parameters (no intercept).
            This function uses the coordinate descent algorithm to find the parameters.
            Arguments
            X: np.ndarray of shape (n, p). Design matrix.
            y: np.nndarray of shape (n). Outcomes.
            1: float. Lambda (one-norm penalty consitant for Lasso).
            tol: float. Default 1e-7. Stop coordinate descent when ||beta^k - beta^(k-1)||^2 \le tol.
            maxitr: int. Default 100. Stop coordinate descent after maxitr iterations (if tolerance
                criterion has not been met already).
            Returns
            ____
            b lasso: np.ndarray of shape (p). Lasso fit parameters.
            (n, p) = X.shape
            b lasso = np.zeros(p) # Start with vector of zeros
            b prev = np.ones(p) # To compare succesive iterations of coordinate descent
            itr = 0
            while (np.linalq.norm(b lasso - b prev)**2 > tol) and (itr < maxitr):
                b prev = b lasso.copy()
                itr += 1
                for j in range(p): # Update one parameter at a time (Gauss-Seidel style)
                    X j = X[:, j]
                    if (i = 0):
                        b minus j = np.delete(b lasso, j, axis=0) # Remove jth parameter
                        X minus j = np.delete(X, j, axis=1) # Remove jth column
                        r = y - np.matmul(X minus j, b minus j) # r is initialized for first j
                    else:
                        # In future iterations, we update r by including the effect of the previous variable
         on the outcome
                        # and excluding the effect of the variable we're currently updating
                        X_j_prev = X[:, j-1]
                        b j prev = b lasso[j-1]
```

```
b_j_cur = b_lasso[j]
                r = r prev - X j prev*b j prev + X j*b j cur
            b_ols_j = np.asarray([np.matmul(X_j.T, r)/n]) # OLS parameter
            b_lasso[j] = soft_th(b_ols_j, 1) # Apply regularization (update jth parameter from beta^k
to beta^(k+1)
            r prev = r.copy() # Store r for efficient update step
    return b_lasso
# Soft threshold function from HW4
def soft_th(b, 1):
    Return the Lasso regression fit parameters from the corresponding OLS parameters (b) and penalty
 constant (1).
    1 1 1
    assert(1 > 0) # lambda must be greater than zero
    size = b.shape[0]
    b_lasso = np.empty(size)
    for j in range(size):
        if b[j] > 1:
            b_{asso[j]} = b[j] - 1
       elif np.abs(b[j]) < 1:
            b lasso[j] = 0
        elif b[j] < -1:
            b_{asso[j]} = b[j] + 1
    return b_lasso
```

```
In [6]: b_lasso = myLasso(X, y, 1)
```

In [7]: from sklearn.linear_model import Lasso # Python equivalent of glmnet
model = Lasso(alpha=1, fit_intercept=False, max_iter=100, tol=1e-7, random_state=seed).fit(X, y)
b_lasso_library = model.coef_

Conclusion: the fitting parameters from my function are virtually identical to the ones from the sklearn module.

Question 2 [40 Points] Path-wise Coordinate Descent

Let's modify our Lasso code to perform path-wise coordinate descent. The idea is simple: we will solve the solution on a grid of λ values, starting from the largest one. After obtaining the optimal β for a given λ , we simply use this solution as the initial value (instead of all zero) for the next (smaller) λ . This is referred to as a warm start in optimization problems. For more details, please watch the lecture video. We will consider the following grid of λ , with the glmnet solution of the first 10 variables plotted.

```
glmnetfit = glmnet(X, y, intercept = FALSE)
  lambda_all = glmnetfit$lambda

matplot(t(glmnetfit$beta[1:10, ]), type = "l", xlab = "Lambda Index", ylab = "Estimated Beta")
```

You need to add an additional input argument lambda_all to your Lasso function. After finishing your algorithm, output a matrix that records all the fitted parameters on your λ grid.

- Provide a plot same as the above glmnet solution plot of the first 10 variables.
- Which two variables entering (start to have nonzero values) the model first?
- What is the maximum discrepancy between your solution and glmnet?

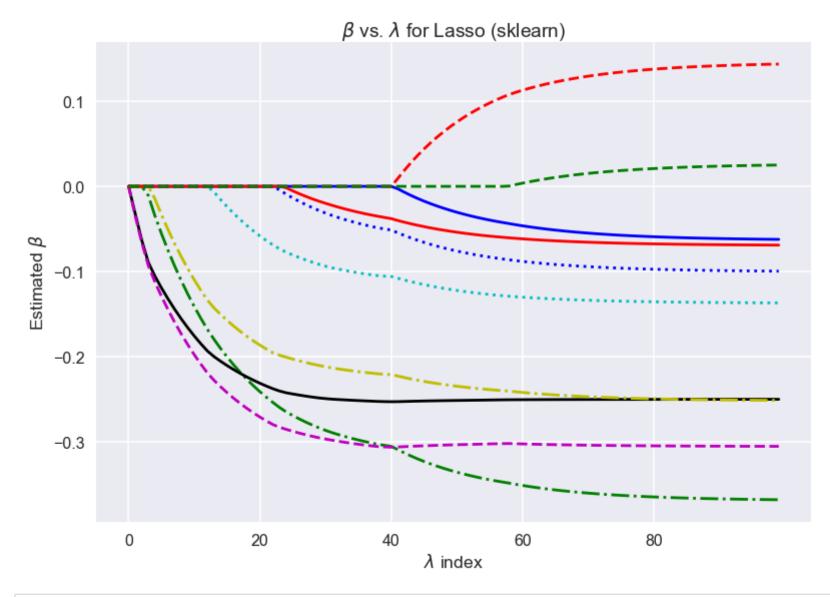
```
In [10]: def myLasso grid(X, y, lambda all, tol=1e-7, maxitr=100):
             Perform Lasso path coordinate descent through all lambda values in lambda all.
             Arguments
             X: np.ndarray of shape (n, p). Design matrix.
             y: np.ndarray of shape (n). Outcomes.
             lambda all: np.ndarray of shape (n lambda). Vector of successively smaller one-norm
                 penalty constants for Lasso.
             tol: float. Default 1e-7. Stop coordinate descent when ||beta^k - beta^(k-1)||^2 \le tol.
             maxitr: int. Default 100. Stop coordinate descent after maxitr iterations (if tolerance
                 criterion has not been met already).
             Returns
             ____
             b lasso: np.ndarray of shape (p, n lambda). Lasso fit parameters for each lambda value.
             (n, p) = X.shape
             b = np.zeros(p) # Start with zeros
             n lambda = lambda all.shape[0]
             b lasso = np.empty((p, n lambda)) # Output matrix
             for i, l in enumerate(lambda all):
                 b = myLasso grid helper(X, y, l, b, tol, maxitr)
                 b lasso[:, i] = b
             return b lasso
         def myLasso grid helper(X, y, l, b, tol, maxitr):
             Fit Lasso regression parameters (no intercept). Helper function to explore lambda grid.
             This function uses the coordinate descent algorithm to find the parameters.
             Arguments
             X: np.ndarray of shape (n, p). Design matrix.
             y: np.ndarray of shape (n). Outcomes.
             1: float. Lambda (one-norm penalty constant for Lasso).
             b: np.ndarray of shape (p). Initial parameters.
             tol: float. Default 1e-7. Stop coordinate descent when ||beta^k - beta^(k-1)||^2 \le tol.
```

```
maxitr: int. Default 100. Stop coordinate descent after maxitr iterations (if tolerance
        criterion has not been met already).
   Returns
    ____
   b lasso: np.ndarray of shape (p). Lasso fit parameters.
   (n, p) = X.shape
   b lasso = b # Warm start
   b prev = np.ones(p) # To compare succesive iterations of coordinate descent
   itr = 0
   while (np.linalg.norm(b_lasso - b_prev)**2 > tol) and (itr < maxitr):</pre>
        b prev = b lasso.copy()
       itr += 1
       for j in range(p): # Update one parameter at a time (Gauss-Seidel style)
           X_j = X[:, j]
           if (j == 0):
                b_minus_j = np.delete(b_lasso, j, axis=0) # Remove jth parameter
                X_minus_j = np.delete(X, j, axis=1) # Remove jth column
               r = y - np.matmul(X minus j, b minus j) # r is initialized for first j
           else:
                # In future iterations, we update r by including the effect of the previous variable
on the outcome
                # and excluding the effect of the variable we're currently updating
                X_j prev = X[:, j-1]
                b_j prev = b_lasso[j-1]
                b_j_cur = b_lasso[j]
               r = r prev - X j prev*b j prev + X j*b j cur
           b ols j = np.asarray([np.matmul(X j.T, r)/n]) # OLS parameter
           b lasso[j] = soft th(b ols j, l) # Apply regularization (update jth parameter from beta^k
to beta^(k+1)
           r prev = r.copy() # Store r for efficient update step
   return b lasso
```

```
In [11]: from itertools import cycle
    from matplotlib import pyplot as plt
    plt.style.use('seaborn-poster')
    plt.style.use('seaborn-darkgrid')
```

```
In [12]: colors = cycle(['b', 'r', 'g', 'c', 'k', 'm', 'y'])
    styles = cycle(['-', '--', '--', ':'])
    neg_log_lambda_all = -np.log(lambda_all)
    for coef, c, s in zip(beta_library[:10, :], colors, styles):
        11 = plt.plot(neg_log_lambda_all, coef, s, c=c)

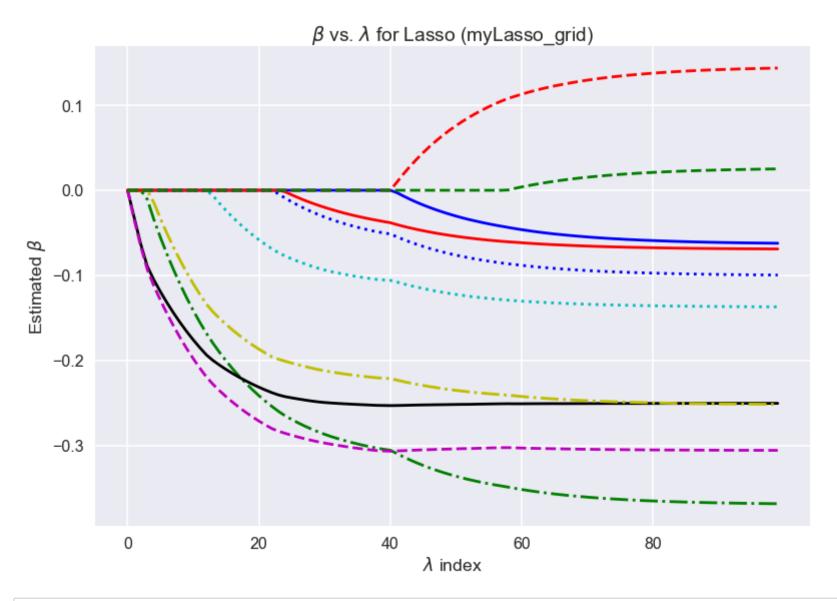
    xticks = neg_log_lambda_all.copy()
    xticks.sort()
    step = 20
    plt.xticks(xticks[::step], labels=np.arange(0, 100)[::step])
    plt.xlabel("$\lambda$ index")
    plt.ylabel(r'Estimated $\beta$')
    plt.title(r'$\beta$ vs. $\lambda$ for Lasso (sklearn)')
    plt.show()
    plt.close()
```



In [13]: my_beta_lasso = myLasso_grid(X, y, lambda_all) # Prints some debugging values

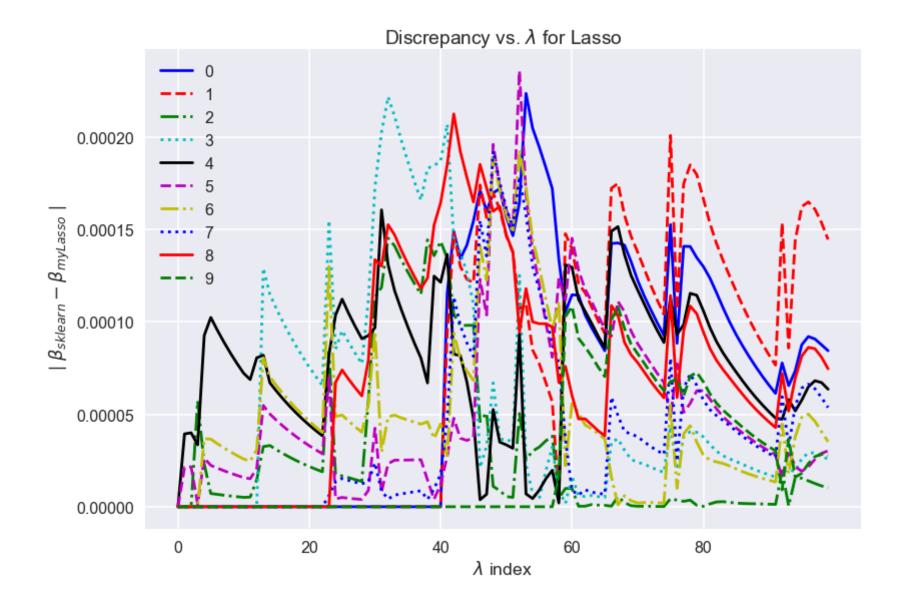
```
In [14]: colors = cycle(['b', 'r', 'g', 'c', 'k', 'm', 'y'])
    styles = cycle(['-', '--', '--', ':'])
    neg_log_lambda_all = -np.log(lambda_all)
    for coef, c, s in zip(my_beta_lasso[:10, :], colors, styles):
        l1 = plt.plot(neg_log_lambda_all, coef, s, c=c)

    xticks = neg_log_lambda_all.copy()
    xticks.sort()
    step = 20
    plt.xticks(xticks[::step], labels=np.arange(0, 100)[::step])
    plt.xlabel("$\lambda$ index")
    plt.ylabel(r'Estimated $\beta$')
    plt.title(r'$\beta$ vs. $\lambda$ for Lasso (myLasso_grid)')
    # plt.legend(np.arange(10), loc='upper left')
    plt.show()
    plt.close()
```



First two variables to enter model according to my implementation: [4 5]
First two variables to enter model according to sklearn: [4 5]

```
In [16]: # Max discrepancy
         discrepancies = np.abs(my beta lasso - beta library)
         colors = cycle(['b', 'r', 'g', 'c', 'k', 'm', 'y'])
         styles = cycle(['-', '--', '--', ':'])
         neg log lambda all = -np.log(lambda all)
         for coef, c, s in zip(discrepancies[:10, :], colors, styles):
             11 = plt.plot(neg log lambda all, coef, s, c=c)
         xticks = neg log lambda all.copy()
         xticks.sort()
         step = 20
         plt.xticks(xticks[::step], labels=np.arange(0, 100)[::step])
         plt.xlabel("$\lambda$ index")
         plt.ylabel(r'$ \mid \beta {sklearn} - \beta {myLasso} \mid$')
         plt.title(r'Discrepancy vs. $\lambda$ for Lasso')
         plt.legend(np.arange(10))
         plt.show()
         plt.close()
```



```
In [17]: # Max discrepancy
  val = np.max(np.abs(my_beta_lasso[:10, :] - beta_library[:10, :]))
  var_index, l_index = np.where(np.abs(my_beta_lasso - beta_library) == val)

  print("Max discrepancy value:", val)
  print("Occuring for lambda value:", lambda_all[l_index[0]])
  print("Occuring for variable:", var_index[0])

Max discrepancy value: 0.00023564668637593433
  Occuring for lambda value: 0.020393229658758557
  Occuring for variable: 5
```

Conclusions:

We can see form the plot that my function and the library show excellent agreement.

The two variables that enter first into the model (that is, the two variables that are non-zero for a small lambda) are variables 5 and 6 (indices 4 and 5).

The maximum discrepancy occurs for the lambda value of 0.020393229658758557 and it is for the variable 6 (index 5).

Question 3 [20 Points] Recovering the Original Scale

The formula provided in HW4 can also be used when there are multiple variables.

$$\frac{Y - \bar{Y}}{\operatorname{sd}_{y}} = \sum_{j=1}^{p} \frac{X_{j} - \bar{X}_{j}}{\operatorname{sd}_{j}} \gamma_{j}$$

$$Y = \bar{Y} - \sum_{j=1}^{p} \bar{X}_{j} \frac{\operatorname{sd}_{y} \cdot \gamma_{j}}{\operatorname{sd}_{j}} + \sum_{j=1}^{p} X_{j} \underbrace{\frac{\operatorname{sd}_{y} \cdot \gamma_{j}}{\operatorname{sd}_{j}}}_{\beta_{j}},$$

Use this formula to recover the original scale of the β , including the intercept term β_0 .

- Use the following code of glmnet to obtain a solution path.
- After recovering your β values, produce a plot of your solution path.
- What is the maximum discrepancy between your solution and glmnet?
- [Bonus 5 Points] If we do not specify lambda in the following glmnet() function, the package will pick a different grid, which lead to a different set of solution. Explain how the lambda values are picked in this case. What is the largest lambda being considered? and why we don't need to consider a larger lambda value? Consider reading the following paper (section 2.5) and the documentation of the glmnet() function at the CRAN website. However, please note that the descriptions from these two sources are slightly different, with similar ideas.
 - Friedman, Jerome, Trevor Hastie, and Rob Tibshirani. "Regularization paths for generalized linear models via coordinate descent." Journal of statistical software 33, no. 1 (2010): 1.

```
glmnetfit2 = glmnet(X_org, y_org, lambda = lambda_all*sd(y_org)*sqrt(n/(n-1)))
  lassobeta2 = coef(glmnetfit2)[2:11, ]
  matplot(t(as.matrix(coef(glmnetfit2)[2:11, ])), type = "l", xlab = "Lambda Index", ylab = "Estimated Beta")
```

```
In [18]: from glmnet import ElasticNet # Using glmnet because sklearn fucntion behaves differently

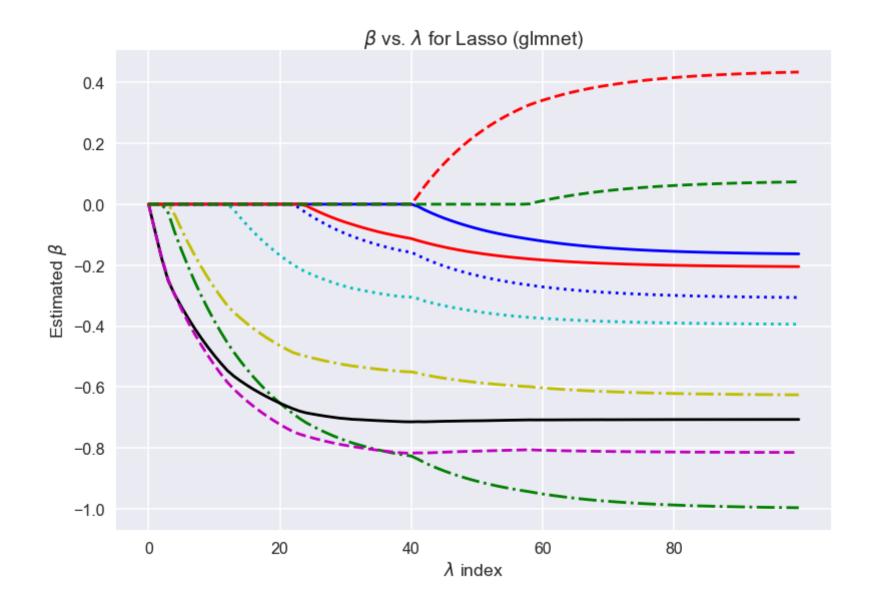
# Python adaptation note: the function np.std does NOT divide by (n-1) to compute the standard

# deviation. Instead, it divides by n. For this reason, we do not need to multiply np.std(y_org)

# by sqrt(n/(n-1)).

model2 = ElasticNet(lambda_path=lambda_all*np.std(y_org)).fit(X_org, y_org)

lassobeta2 = model2.coef_path_
```

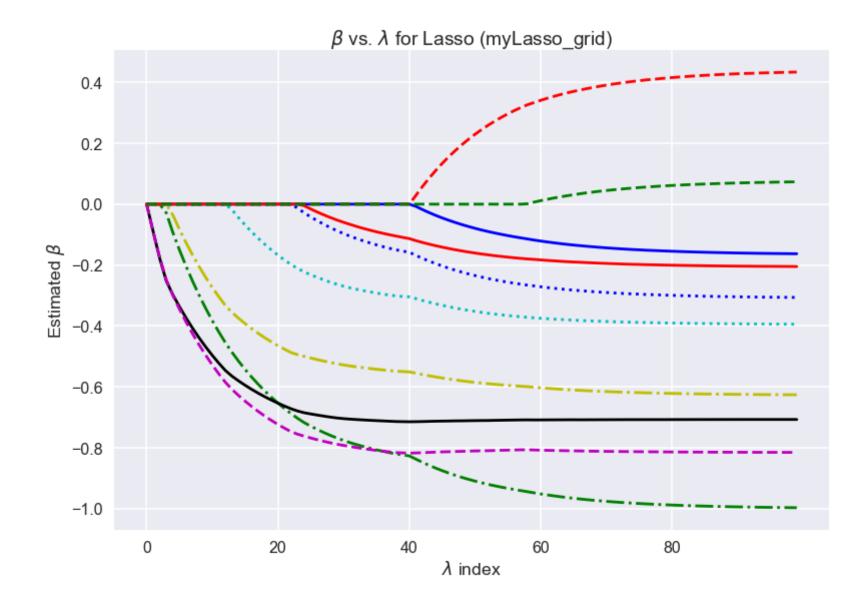


```
In [20]: # Scaling function from HW4 --> Used to get mu_j, sd_j, mu_y, sd_y to recover scale
          def scale data(X, Y):
              Standardize data and predictions (center on mean = 0 and scale to stadard deviation = 1).
              Returns stadardized data (X std, Y std) and necessary parameters (mu j, sd j, mu y, sd y)
              to recover the predictions in the original scale.
              mu j = np.mean(X, axis=0)
              sd j = np.std(X, axis=0)
              mu y = np.mean(Y)
              sd y = np.std(Y)
              X \text{ std} = (X - \text{mu j}) / \text{sd j}
              Y \text{ std} = (Y - \text{mu } y) / \text{sd } y
              return X std, Y std, mu_j, sd_j, mu_y, sd_y
In [21]: _, _, mu j, sd j, mu y, sd y = scale data(X org, y org)
         my beta 0 = mu y - np.sum(mu j*sd y/sd j)
         my beta j = my beta lasso*sd y/sd j[:, None]
In [22]: my beta 0
```

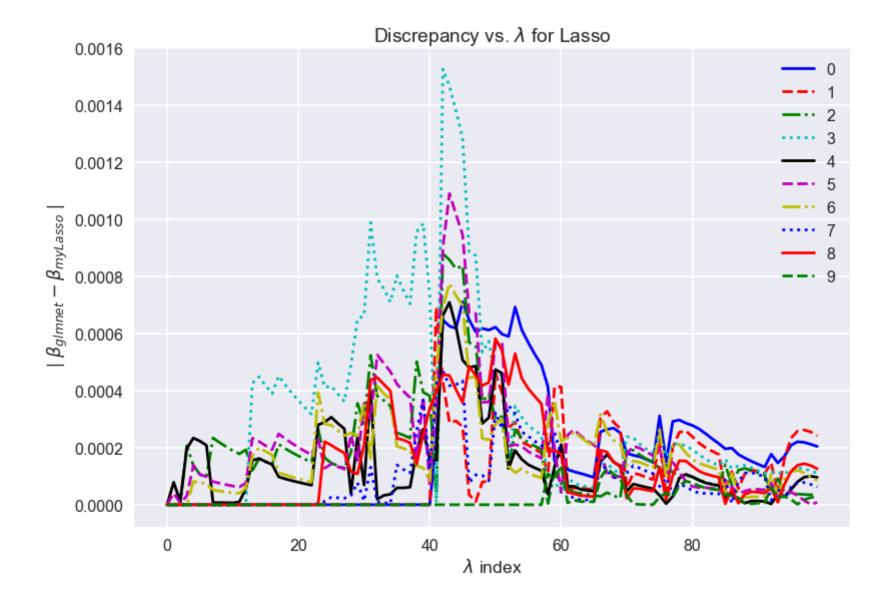
Out[22]: 74.2261988308727

```
In [23]: colors = cycle(['b', 'r', 'g', 'c', 'k', 'm', 'y'])
    styles = cycle(['-', '--', '--', ':'])
    neg_log_lambda_all = -np.log(lambda_all*np.std(y_org))
    for coef, c, s in zip(my_beta_j[:10, :], colors, styles):
        l1 = plt.plot(neg_log_lambda_all, coef, s, c=c)

    xticks = neg_log_lambda_all.copy()
    xticks.sort()
    step = 20
    plt.xticks(xticks[::step], labels=np.arange(0, 100)[::step])
    plt.xlabel("$\lambda\sindex")
    plt.ylabel(r'Estimated \stacks')
    plt.title(r'\s\beta\stacks vs. \stacks')
    plt.show()
    plt.close()
```



```
In [24]: # Max discrepancy
         discrepancies = np.abs(my beta j - lassobeta2)
         colors = cycle(['b', 'r', 'g', 'c', 'k', 'm', 'y'])
         styles = cycle(['-', '--', '-.', ':'])
         neg log lambda all = -np.log(lambda all*np.std(y org))
         for coef, c, s in zip(discrepancies[:10, :], colors, styles):
             11 = plt.plot(neg log lambda all, coef, s, c=c)
         xticks = neg log lambda all.copy()
         xticks.sort()
         step = 20
         plt.xticks(xticks[::step], labels=np.arange(0, 100)[::step])
         plt.xlabel("$\lambda$ index")
         plt.ylabel(r'$ \mid \beta {glmnet} - \beta {myLasso} \mid$')
         plt.title(r'Discrepancy vs. $\lambda$ for Lasso')
         plt.legend(np.arange(10))
         plt.show()
         plt.close()
```



```
In [25]: # Max discrepancy
  val = np.max(np.abs(my_beta_j[:10, :] - lassobeta2[:10, :]))
  var_index, l_index = np.where(np.abs(my_beta_j - lassobeta2) == val)

print("Max discrepancy value:", val)
  print("Occuring for lambda value:", lambda_all[l_index[0]])
  print("Occuring for variable:", var_index[0])

Max discrepancy value: 0.0015254544176474538
  Occuring for lambda value: 0.04097475005926602
  Occuring for variable: 3
```

Conclusion: we can see that the plots look very similar to those in Q2, except that the parameters are in the original scale of the data.

The maximum discrepancy occurs for lambda value 0.04097475005926602 and corresponds to variable 4 (index 3).

[Bonus]

The lambda values are picked in the following way:

1. The maximum lambda value is selected such that all parameters are zero (it is not necessary to try with larger lambda values because all the parameters would still be zero). In mathematical terms, this means that we can pick our largest lambda using the formula:

$$N\alpha\lambda_{max} = max_l(|\langle X_l^T, y \rangle|)$$

- 1. The minimum value is selected based on some constant ratio ϵ . I.e., $\lambda_{min} = \epsilon \lambda_{max}$
- 2. The rest of the values on the grid are selected on a logarithmic scale (they are ordered in decreasing values).