SDS 385 Ex 07: ADMM for LASSO

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ADMM for LASSO

Algorithm Details

I used the ADMM algorithm as detailed in Section 6.4 of Boyd, "Distributed Optimization and Statistical Learning via the Alternating Direction Method of Multipliers", 2011.

The setting for this algorithm as applied to lasso is below.

Objective Function:

minimize
$$\frac{1}{2}||X\beta - y||_2^2 + \lambda||x||_1$$
 with $\lambda > 0$ as the l1 regularization penalty parameter.

ADMM Form of Problem:

minimize
$$f(x) + g(x)$$

subject to $x - z = 0$
where:

$$f(x) = \frac{1}{2}||X\beta - y||_2^2$$

$$g(x) = \lambda||x||_1$$

ADMM Steps:

$$\beta^{k+1} = (X^T X + \rho I)^{-1} (X^T y + \rho (z^k - u_k))$$

$$z^{k+1} = S_{\lambda/\rho} (\beta^{k+1} + u^k)$$

$$u^{k+1} = u^k + \beta^{k+1} - z^{k+1}$$

where $S_{\lambda\rho}(a) = sign(a)(|a| - \lambda/\rho)_+$ and u^k is the augmented lagrangian at the k^{th} iteration.

Implementation Notes:

- Since the step size is fixed, to increase efficiency, I cached the matrix inverse $(X^TX + \rho I)^{-1}$ before beginning iterations.
- Boyd notes that when not using fixed step size, it is best to still cache this inverse, and only update it when the ρ step size is updated.

My Implementation

My ADMM implementation for LASSO converged in 230 iterations. This was considerably faster than my proximal gradient descent (7687 iterations) and my accelerated proximal gradient descent (1853 iterations). I used $\lambda = .01$ for the l1 lasso penalty and $\rho = .01$ for the step size. I used the convergence criteria of absolute change in the lasso objective function < tol, where tol = 1E - 10.

Note that the lambda and rho were the same values used for exercise 6, for consistency. In practice, I would select an optimal lambda value via cross-validation.

The plot of the lasso objective function is as follows.

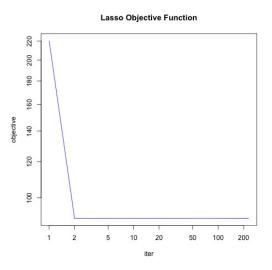


Figure 1: LASSO objective using ADMM($\hat{\alpha}$)

The estimated beta coefficients obtained using admm are compared with the coefficients from glmnet below. Most appear to be of reasonably similar sign and magnitude.

```
glmnet
                               admm
                         0.03007395
         0.0084994686
age
         -0.1274718132 -0.16390462
sex
                         0.28268400
         0.3053578438
bmi
         0.1928804642
                         0.21180397
map
         -0.0387865853 -0.10040001
tc
ldl
         -0.1635236545 -0.10958174
hdl
tch
                         0.05421951
ltg
         0.3230766076
                         0.42695740
glu
         0.0337240035
                         0.03968565
age.2
         0.0351045119
                         0.04230375
bmi.2
         0.0238787313
                         0.02913876
                        -0.00496873
map.2
tc.2
                         3.32992803
```

```
ld1.2
                       1.78879470
hdl.2
                       0.86057254
tch.2
                       0.45016330
ltg.2
                       0.71081295
glu.2 0.0636440798
                       0.07115055
age.sex 0.0961729059
                       0.09425725
age.bmi .
                      -0.00974232
age.map 0.0145699540 0.01097956
                      -0.10307316
age.tc
age.ldl -0.0389574124 -0.03850246
age.hdl 0.0258647293 0.13295890
age.tch .
                       0.11473838
age.ltg 0.0407916229 0.07678043
 age.glu 0.0184222925 0.03852093
sex.bmi 0.0270461479
                       0.04070618
sex.map 0.0387600107 0.05455798
                       0.25201571
sex.tc .
sex.ldl -0.0181954985 -0.20580754
sex.hdl 0.0429324533 -0.06940488
sex.tch .
                     -0.07673176
sex.ltg .
                     -0.07203117
sex.glu .
                       0.02857881
bmi.map 0.0837040826 0.09607870
bmi.tc -0.0002777938 -0.22702825
bmi.ldl .
                     0.18330034
bmi.hdl .
                       0.08852731
bmi.tch .
                    -0.01969956
bmi.ltg .
                     0.07900017
bmi.glu .
                       0.01519782
map.tc
         0.0083855329 0.28434387
map.ldl .
                      -0.19175559
map.hdl 0.0233788843 -0.11053647
map.tch .
                     -0.03567046
                      -0.09184804
map.ltg
map.glu -0.0362793960 -0.08424349
tc.ldl .
                      -4.61510706
tc.hdl 0.0072487602 -1.93720146
tc.tch -0.0477242186 -1.10984540
tc.ltg -0.0098056312 -2.26554521
              -0.08709231
tc.glu .
ldl.hdl .
                      1.26150669
ldl.tch .
                       0.52479375
ldl.ltg 0.0742412609 1.70305929
ldl.glu 0.0067025050 0.03256134
hdl.tch -0.0522309970 0.62815286
hdl.ltg .
                       0.84045023
hdl.glu .
                       0.12679103
tch.ltg -0.0607125224 0.15900559
tch.glu 0.0214432611
                       0.14614305
```

SDS 383C - Stats Modeling : SDS 385 Ex 07: ADMM for LASSO

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65 | ltg.glu . 0.04353096

Appendix: R Code

```
### SDS 385 - Exercises 07 - ADMM for LASSO.
   ### ADMM = Alternating Direction Method of Multipliers
   #Jennifer Starling
  #18 October 2016
   rm(list=ls()) #Clean workspace.
   library(glmnet)
  library(Matrix)
   #Read in Diabetes.csv data.
   X <- read.csv(file='/Users/jennstarling/UTAustin/2016_Fall_SDS 385_Big_Data/
      Exercise 05 R Code/DiabetesX.csv',header=T)
   y <- read.csv(file='/Users/jennstarling/UTAustin/2016_Fall_SDS 385_Big_Data/
      Exercise 05 R Code/DiabetesY.csv',header=F)
   #Scale X and y.
   X = scale(X)
   y = scale(y)
   #LASSO objective function:
   #Inputs:
   \# X = X matrix (scaled)
     y = response data (scaled)
      lambda = a chosen lambda value
     beta = a vector of beta coefficients.
   #Output:
      Value of the LASSO objective function at specified inputs.
   fx <- function(X,y,lambda,beta){</pre>
      #f = (1/nrow(X)) * (t(y - X %*% beta) %*% (y - X %*% beta))
30
      f = (1/2) * (t(y - X %*% beta) %*% (y - X %*% beta))
      g = lambda * sum(abs(beta))
      obj = (f+g)
      return(as.numeric(obj))
  }
35
   #Proximal L1 Operator function: (soft thresholding operator)
   #Inputs:
   \# x = vector of values.
      lambda = the scaling factor of the 11 norm.
     t = the step size.
   #Output:
  # Value of the soft-thresholding proximal operator.
   prox_l1 <- function(x,gamma,tau=1) {</pre>
      thresh <- gamma*tau
      prox = rep(0, length(x))
       idx.1 = which(x < -thresh)
       idx.2 = which(x > thresh)
       idx.3 = which(abs(x) \le thresh)
```

```
if (length(idx.1) > 0) prox[idx.1] = x[idx.1] + thresh
55
       if (length(idx.2) > 0) prox[idx.2] = x[idx.2] - thresh
       if (length(idx.3) > 0) prox[idx.3] = 0
       return(prox)
   }
    #ADMM for Lasso:
   #Inputs:
      X = design matrix
       y = response vector
      rho = step size
      maxiter = maximum iterations
       tol = tolerance for convergence
       lambda = 11 norm penalty constant.
   #Output:
       List including estimated beta values and objective function.
   admmLasso = function(X,Y,rho=.01,lambda=.1,maxiter=1000,tol=1E-10){
       i=0
                            #Initialize iterator.
75
       converged <- 0
                            #Indicator for whether convergence met.
       #1. Initialize matrix to hold beta vector for each iteration.
       betas <- matrix(0, nrow=maxiter, ncol=ncol(X))</pre>
80
       #2. Initialize values for objective function.
       obj <- rep(0,maxiter)</pre>
                               #Initialize vector to hold loglikelihood fctn.
       obj[1] <- fx(X,y,lambda,betas[1,])</pre>
       #3. Cache matrix inverse, since using fixed step size for each iter.
       inv_cache = solve(t(X) %*% X + rho*diag(1,ncol(betas)))
       #4. Initialize values.
       betas[1,] <- rep(0,ncol(X)) #Initialize beta vector to 0 to start.
       u = rep(0, ncol(betas)) #Initialize the lagrangian to all zeros.
       z = rep(0, ncol(betas)) #Initialize z vector to all zeros.
       #4. ADMM looping.
       for (i in 2:maxiter){
            #Update betas.
            betas[i,] = inv_cache %*% (t(X) %*% y + rho * (z-u) )
            #Update z.
100
           z = prox_l1(betas[i,] + u,lambda/rho)
            #Update u (lagrangian).
           u = u + betas[i,] - z
            #Update objective function.
            obj[i] = fx(X,y,lambda=lambda,beta=betas[i,])
            #Convergence check.
            #Check if convergence met: If yes, exit loop.
110
            if (abs(obj[i]-obj[i-1])/abs(obj[i-1]+1E-3) < tol ){
                converged=1;
                break;
```

```
}
       }
115
       #Return function values.
       return(list(obj=obj, betas=betas, beta_hat=betas[i,], converged=converged,
           iter=i))
120
   #Run admm for lasso.
   output <- admmLasso(X,y,rho=.01,lambda=.01,maxiter=1000,tol=1E-14)
   #Iterations to convergence:
   print(output$iter)
   print(output$converged)
   #Plot objective function.
   jpeg(file='/Users/jennstarling/UTAustin/2016_Fall_SDS 385_Big_Data/Exercise 07
       LaTeX Files/admm_objective.jpg')
   plot(1:output$iter,output$obj[1:output$iter],type='l',col='blue',log='xy',
       main='Lasso Objective Function',xlab='iter',ylab='objective')
   dev.off()
   #Compare results to glmnet:
   myLasso <- glmnet(X,y,family='gaussian',alpha=1,lambda=.01) #Fit lasso glmnet
   beta_glmnet <- myLasso$beta</pre>
                                                                 #Save glmnet betas.
   cbind(glmnet=beta_glmnet,admm=round(output$beta_hat,8))
                                                                 #Output comparison
```