# proximal gradient method

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function to calculate negative likelihood with lasso penalty

```
nll_lasso = function(beta, x, y, lambda)
{
   xbeta = x %*% beta
   res = y - xbeta
   nll = crossprod(res)/2 + lambda * sum( abs(beta) )
   return (nll)
}
```

function to calculate negative likelihood

```
nll = function(beta, x, y)
{
   xbeta = x %*% beta
   res = y - xbeta
   nll = crossprod(res)/2

   return (nll)
}
```

function to calculate gradient

```
grad = function(beta, x, y)
{
   xbeta = x %*% beta
   res = y - xbeta
   grad = -t(x) %*% res

   return (grad)
}
```

function to calculate soft threshold

```
soft_threshold = function(y,lambda)
{
  z = abs(y) - lambda
  theta = sign(y)*z*(z > 0)
  return (theta) # this is max(z, 0) element wise
}
```

#### proximal gradient algorithm

```
prox_grad = function(x, y, beta0, step=0.01, lambda, iter_max=10000, precision=1e-10)
 p = length(beta0)
  beta = matrix(0, nrow = iter_max, ncol = p)
  beta[1,] = beta0
  nll_track = rep(0, iter_max)
  nll_track[1] = nll(beta0, x, y)
  nll_track_lasso = rep(0, iter_max)
  nll_track_lasso[1] = nll_lasso(beta0, x, y, lambda)
  delta = 1
  iter = 1
  while ( (delta >= precision) && (iter <= iter_max) )</pre>
   u = beta[iter,] - step*grad(beta[iter,], x, y)
   beta[iter+1,] = soft_threshold(u, lambda*step)
   nll_track_lasso[iter+1] = nll_lasso(beta[iter+1,], x, y, lambda)
   nll_track[iter+1] = nll(beta[iter+1,], x, y)
   delta = abs(nll_track_lasso[iter+1] - nll_track_lasso[iter])/nll_track_lasso[iter]
   iter = iter + 1
 return (list(iter, nll_track_lasso, nll_track, beta))
```

### accelerated proximal gradient algorithm

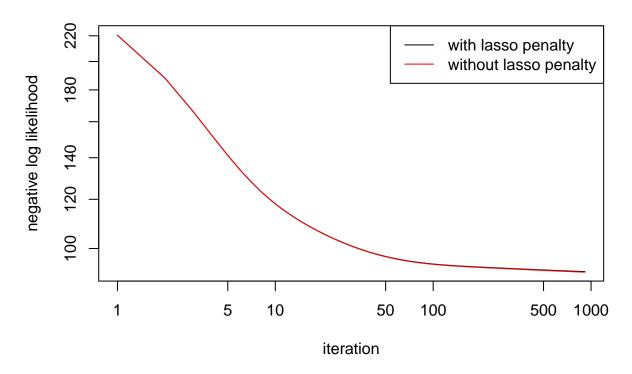
```
acc_prox_grad = function(x, y, beta0, step=0.01, s=2, lambda, iter_max=10000, precision=1e-10)
 p = length(beta0)
 beta = matrix(0, nrow = iter_max, ncol = p)
 beta[1,] = beta0
 nll_track = rep(0, iter_max)
 nll_track[1] = nll(beta0, x, y)
  nll_track_lasso = rep(0, iter_max)
 nll_track_lasso[1] = nll_lasso(beta0, x, y, lambda)
  delta = 1
  iter = 1
  z = beta0
  while ( (delta >= precision) && (iter < iter_max) )</pre>
   u = z - step*grad(z, x, y)
   beta[iter+1,] = soft_threshold(u, lambda*step)
   s_{update} = (1 + sqrt(1 + 4*s^2))/2
   z = beta[iter+1,] + (s - 1)/s\_update * (beta[iter+1,] - beta[iter,])
   \#z = beta[iter,] + (s - 1)/s\_update * (beta[iter,] - beta[iter+1,])
   nll track lasso[iter+1] = nll lasso(beta[iter+1,], x, y, lambda)
   nll_track[iter+1] = nll(beta[iter+1,], x, y)
```

```
delta = abs(nll_track_lasso[iter+1] - nll_track_lasso[iter])/nll_track_lasso[iter]
  iter = iter + 1
    s = s_update
}
#nll_track_lasso[1:100]
return (list(iter, nll_track_lasso, nll_track, beta))
}
```

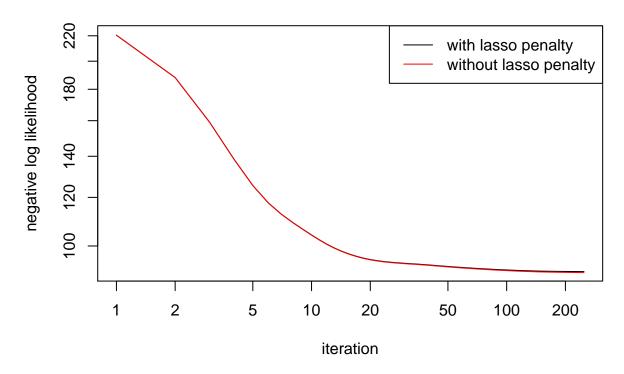
run and compare beta with glmnet output

```
diabetesX = read.csv("C:/Users/schen/Dropbox/toChensu/Stats/2016Fall/Big Data/Assignment5/diabetesX.csv
diabetesY = read.csv("C:/Users/schen/Dropbox/toChensu/Stats/2016Fall/Big Data/Assignment5/diabetesY.csv
X = as.matrix(diabetesX)
Y = as.matrix(diabetesY)
Xs = scale(X)
Ys = scale(Y)
beta0 = rep(0, ncol(Xs))
### use cv.glmnet to choose optimal lambda ###
library(glmnet)
## Loading required package: Matrix
## Loading required package: foreach
## Loaded glmnet 2.0-5
cv.fit = cv.glmnet(x = Xs, y = Ys)
cv.lambda = cv.fit$lambda.min
fit.cv.lambda = glmnet(x = Xs, y = Ys, lambda = cv.lambda)
glmnet_beta = fit.cv.lambda$beta
result = prox_grad(x=Xs, y=Ys, beta0, step=0.0001, lambda=cv.lambda, iter_max=10000, precision=1e-5)
plot_iter = result[[1]]
result_nll_lasso = result[[2]][1:plot_iter]
result_nll = result[[3]][1:plot_iter]
result_beta = result[[4]][plot_iter,]
plot(x = 1:plot_iter, y = result_nll_lasso, type = "l",
     xlab = "iteration", ylab = "negative log likelihood",
     log = "xy", main = "convergence of proximal gradient")
lines(x = 1:plot_iter, y = result_nll, col = "red")
legend("topright", c("with lasso penalty", "without lasso penalty"),
       col = c("black", "red"), lty = c(1,1))
```

### convergence of proximal gradient

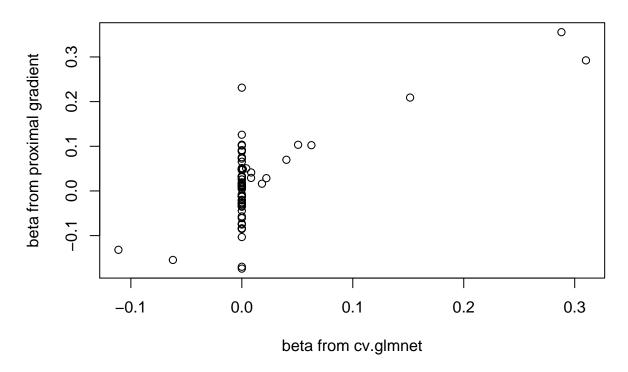


## convergence of accelerated proximal gradient



```
plot(x = glmnet_beta, y = result_beta, xlab = "beta from cv.glmnet",
    ylab = "beta from proximal gradient",
    main = "compare results of proximal gradient with cv.glmnet")
```

## compare results of proximal gradient with cv.glmnet



```
plot(x = glmnet_beta, y = result_beta_acc, xlab = "beta from cv.glmnet",
    ylab = "beta from accelated proximal gradient",
    main = "compare results of accelated proximal gradient with cv.glmnet")
```

## compare results of accelated proximal gradient with cv.glmnet

