## STAT230 HW 2 University of California, Berkeley

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1

```
load('family.rda')
attach(family)
```

```
intercept = rep(1,length(weight))
X = cbind(intercept,height, bmi)
betahat = solve(crossprod(X,X), t(X) %*% weight)
betahat
##
## intercept -310.346943
## height 4.675473
## bmi
              6.308645
residuals = weight - X %*% betahat
residuals
##
                 [,1]
   [1,] -0.676730632
   [2,] 0.474100055
## [3,] -0.264728345
## [4,] -1.371495045
## [5,] 2.176912620
## [6,] -0.219261947
## [7,] -0.413102763
```

```
## [8,] -0.007851258

## [9,] -0.800784395

## [10,] 3.819187100

## [11,] -1.272889595

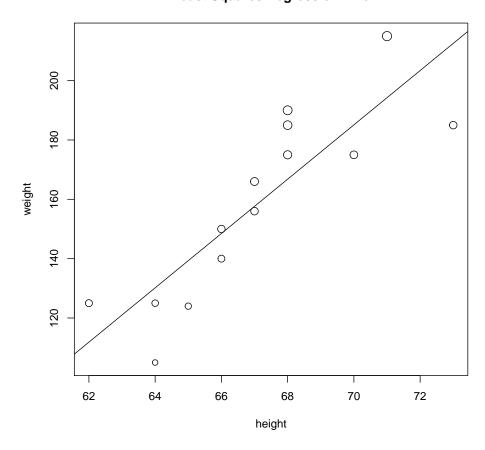
## [12,] -1.086205171

## [13,] -1.289915027

## [14,] 0.932764403
```

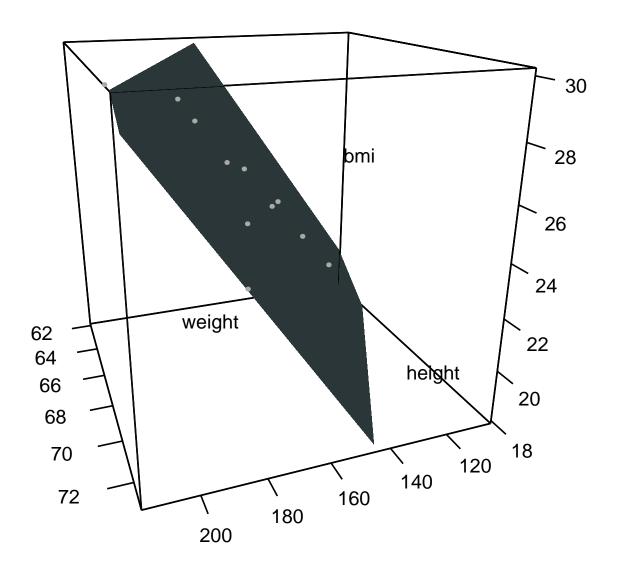
2

## **Least Squares Regression Line**



bmi is increasing with both weight and height.

3



```
M = X[,1:2]
N = X[,3]
Y = weight
# step 1
gamma_1 = solve(crossprod(M,M), crossprod(M,Y))
gamma_1
##
                  [,1]
## intercept -455.6660
## height
                9.1537
f = Y - M %*% gamma_1
# step 2
gamma_2 = solve(crossprod(M,M), crossprod(M,N))
gamma_2
##
## intercept -23.0349135
## height
               0.7098557
g = N - M \%*\% gamma_2
# step 3
gamma_3 = sum(f*g)/sum(g*g)
gamma_3
## [1] 6.308645
e = f-g*gamma_3
# step 4
betahatBis = c(gamma_1-gamma_2*gamma_3,gamma_3)
betahatBis
## [1] -310.346943   4.675473   6.308645
```

```
# Check validity
abs(betahat-betahatBis)

## [,1]
## intercept 6.082246e-12
## height 1.803002e-13
## bmi 2.522427e-13
```