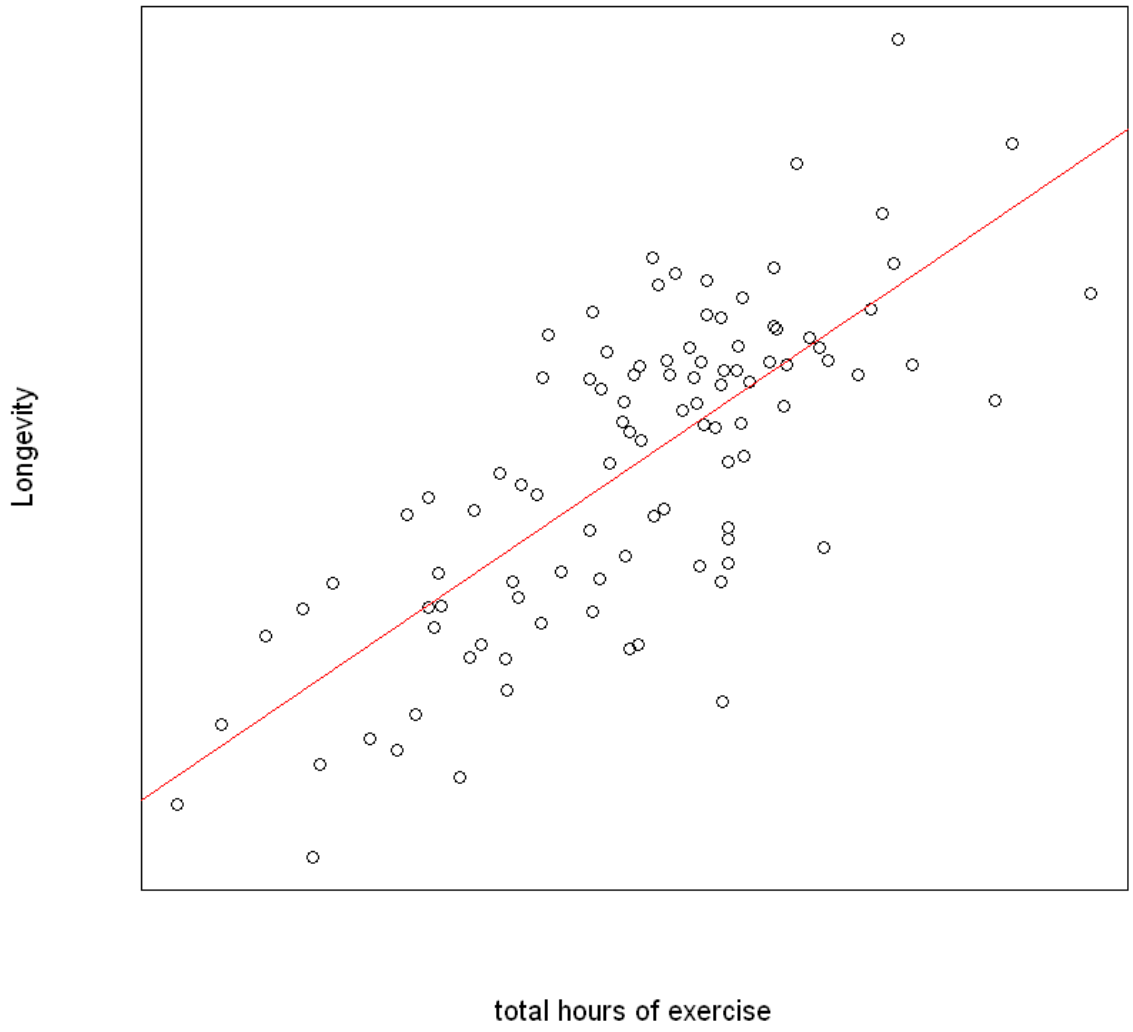
Answer #1

In [81]:
```
#x = total hours of exercise
#y = Longevity
x<- rnorm(100,1,5)
y<- 10+0.2*x+rnorm(100)
data.set<-data.frame(x,y)
plot(x, y, axes=FALSE, frame.plot=TRUE, xlab= "total hours of exercise", ylab=
"Longevity")
abline(lm(y ~ x, data=data.set), col="red")
a<-lm(y ~ x)
summary(a)
par(new=TRUE)
x[100]= - 200
y[100]= 90
data.set2<-data.frame(x,y)
b<-lm(y ~ x)
summary(b)
plot(x,y,xlim= c(-200, -20),ylim= c(1:90),xlab= "total hours of exercise", yla
b="Longevity", col= "blue")
abline(lm(y ~ x, data=data.set2), col="blue",new=TRUE)
```

```
Call:
lm(formula = y ~ x)

Residuals:
    Min      1Q  Median      3Q     Max
-2.8008 -0.7138  0.1434  0.7661  2.3287

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 10.06632    0.10174   98.94   <2e-16 ***
x            0.20700    0.01884   10.99   <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 1.001 on 98 degrees of freedom
Multiple R-squared:  0.552,     Adjusted R-squared:  0.5474
F-statistic: 120.7 on 1 and 98 DF,  p-value: < 2.2e-16
```

```
Call:
lm(formula = y ~ x)

Residuals:
    Min      1Q  Median      3Q     Max
-8.3413 -2.0335  0.2979  2.1149  7.9181

Coefficients:
            Estimate Std. Error t value Pr(>|t|)
(Intercept) 10.68686    0.32899   32.48   <2e-16 ***
x           -0.35698    0.01588  -22.48   <2e-16 ***
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

Residual standard error: 3.286 on 98 degrees of freedom
Multiple R-squared:  0.8376,    Adjusted R-squared:  0.8359
F-statistic: 505.3 on 1 and 98 DF,  p-value: < 2.2e-16
```

```
Error in plot.window(...): invalid 'ylim' value
Traceback:

1. plot(x, y, xlim = c(-200, -20), ylim = c(1:90), xlab = "total hours of exe
rcise",
 .     ylab = "Longevity", col = "blue")
2. plot.default(x, y, xlim = c(-200, -20), ylim = c(1:90), xlab = "total hour
s of exercise",
 .     ylab = "Longevity", col = "blue")
3. localWindow(xlim, ylim, log, asp, ...)
4. plot.window(...)
```



Answer #2

In [83]:
```
#ANSWER 2
library(Matching)
library(arm)
library()
data("lalonde")
newdata=lalonde[which(lalonde$treat==0),]

lmlalonde<-glm(newdata$re78 ~ newdata$age+newdata$educ+newdata$re74+newdata$re
75+newdata$educ:newdata$re74+newdata$educ:newdata$re75+newdata$age:newdata$re7
4+newdata$age:newdata$re75+newdata$re74:newdata$re75)
summary(lmlalonde)
#lmlalonde$coefficients
Simulation <- sim(lmlalonde, n.sims = 10000)
Simulation@coef[1,5]


storage.vector <- rep(NA,10000)

m_educ=median(newdata$educ)
m_re74=median(newdata$re74)
m_re75=median(newdata$re75)


q90.edu<-quantile(newdata$educ,.90)
q90.re74<-quantile(newdata$re74,.90)
q90.re75<-quantile(newdata$re75,.90)

summary(newdata$age)

#quantile(simulation@coef[,1], probs = c(0.025, 0.975))

storagedf<- matrix(NA, nrow = 10000, ncol= length(newdata$age))

for(i in 1:10000){

    predictedYs <- Simulation@coef[i,1] * 1 +
      Simulation@coef[i,2] * newdata$age +
      Simulation@coef[i,3] * m_educ +
      Simulation@coef[i,4] * m_re74 +
      Simulation@coef[i,5] * m_re75 +
      Simulation@coef[i,6] * m_educ*m_re74 +
      Simulation@coef[i,7] * m_educ*m_re75 +
      Simulation@coef[i,8] * newdata$age*m_re74 +
      Simulation@coef[i,9] * newdata$age*m_re75 +
      Simulation@coef[i,10] * m_re74*m_re75 +
      rnorm(1,0,Simulation@sigma[i])


    # put preductedYs into a matrix
      storagedf[i,] = predictedYs


      #* c(1,newdata$age,m_educ,m_re74,m_re75,m_educ*m_re74,m_educ*m_re75,newd
ata$age*m_re74,newdata$age*m_re75, m_re74*m_re75))
```

```
    #storagedf[i,age-16]<-exp(beta)/(1+exp(beta))
}

#confidence.intervals <- quantile(storagedf, probs = c(0.025, 0.975))
head(storagedf)

lowbounds.medians = rep(NA, length(newdata$age))
upperbounds.medians = rep(NA, length(newdata$age))

for (i in 1:ncol(storagedf)) {
  lowbounds.medians[i] = quantile(storagedf[,i],0.025)
  upperbounds.medians[i] = quantile(storagedf[,i],0.975)

}


plot(newdata$age, lowbounds.medians, ylim = c(-8000,18000), col= "blue",, xlab
 = "lower and upper bounds for median")
points(newdata$age, upperbounds.medians, col="red")

#Simulation@sigma

storagedf_2<- matrix(NA, nrow = 10000, ncol= length(newdata$age))

for(i in 1:10000){

  predictedYs_q<- Simulation@coef[i,1] * 1 +
    Simulation@coef[i,2] * newdata$age +
    Simulation@coef[i,3] * q90.edu +
    Simulation@coef[i,4] * q90.re74 +
    Simulation@coef[i,5] * q90.re75 +
    Simulation@coef[i,6] * q90.edu*q90.re74 +
    Simulation@coef[i,7] * q90.edu*q90.re75 +
    Simulation@coef[i,8] * newdata$age*q90.re74 +
    Simulation@coef[i,9] * newdata$age*q90.re75 +
    Simulation@coef[i,10] * q90.re74*q90.re75 +
    rnorm(1,0,Simulation@sigma[i])


  # put preductedYs into a matrix
  storagedf_2[i,] = predictedYs_q



}

head(storagedf_2)

lowbounds.q90 = rep(NA, length(newdata$age))
upperbounds.q90 = rep(NA, length(newdata$age))

for (i in 1:ncol(storagedf)) {
  lowbounds.q90[i] = quantile(storagedf_2[,i],0.025)
  upperbounds.q90[i] = quantile(storagedf_2[,i],0.975)

}
```

```
plot(newdata$age, lowbounds.q90, ylim = c(-8000,18000), col= "blue")
points(newdata$age, upperbounds.q90, col="red")


Summary(lmlalonde)
```

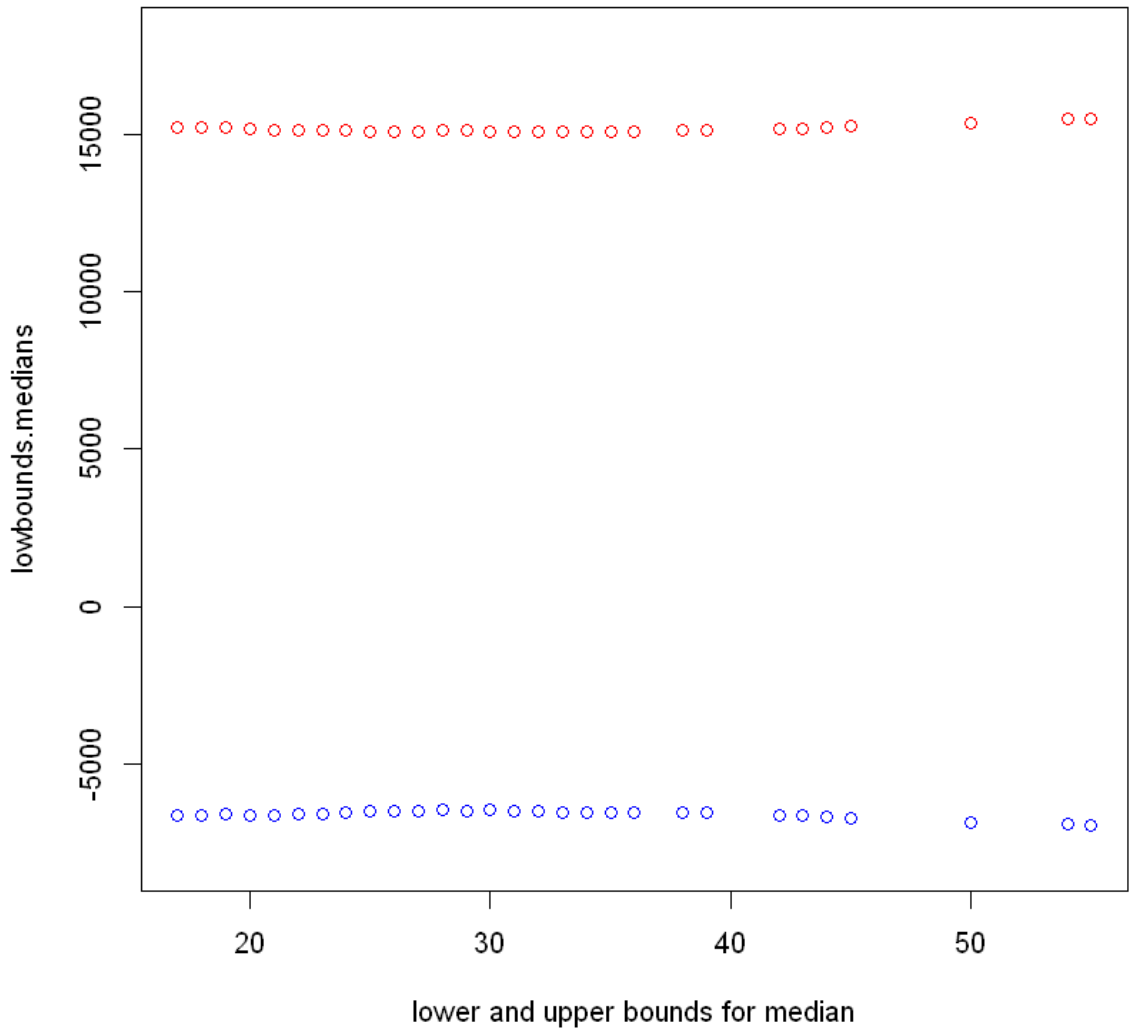
```
Call:
glm(formula = newdata$re78 ~ newdata$age + newdata$educ + newdata$re74 +
    newdata$re75 + newdata$educ:newdata$re74 + newdata$educ:newdata$re75 +
    newdata$age:newdata$re74 + newdata$age:newdata$re75 + newdata$re74:newdat
a$re75)

Deviance Residuals:
   Min      1Q  Median      3Q     Max
 -7264   -4148   -1590    3014   33846

Coefficients:
                              Estimate Std. Error t value Pr(>|t|)
(Intercept)                  3.686e+03  2.630e+03   1.401    0.162
newdata$age                  2.216e+00  5.206e+01   0.043    0.966
newdata$educ                 3.907e+01  2.292e+02   0.170    0.865
newdata$re74                -1.552e-02  4.790e-01  -0.032    0.974
newdata$re75                 7.845e-01  1.878e+00   0.418    0.677
newdata$educ:newdata$re74    3.441e-02  6.671e-02   0.516    0.606
newdata$educ:newdata$re75   -7.204e-02  1.341e-01  -0.537    0.592
newdata$age:newdata$re74    -5.705e-03  2.783e-02  -0.205    0.838
newdata$age:newdata$re75     9.309e-03  4.212e-02   0.221    0.825
newdata$re74:newdata$re75   -2.294e-05  1.413e-05  -1.623    0.106

(Dispersion parameter for gaussian family taken to be 30371884)

    Null deviance: 7788768802  on 259  degrees of freedom
Residual deviance: 7592971078  on 250  degrees of freedom
AIC: 5229.2

Number of Fisher Scoring iterations: 2
```

newdata$re75: -0.894239894276869

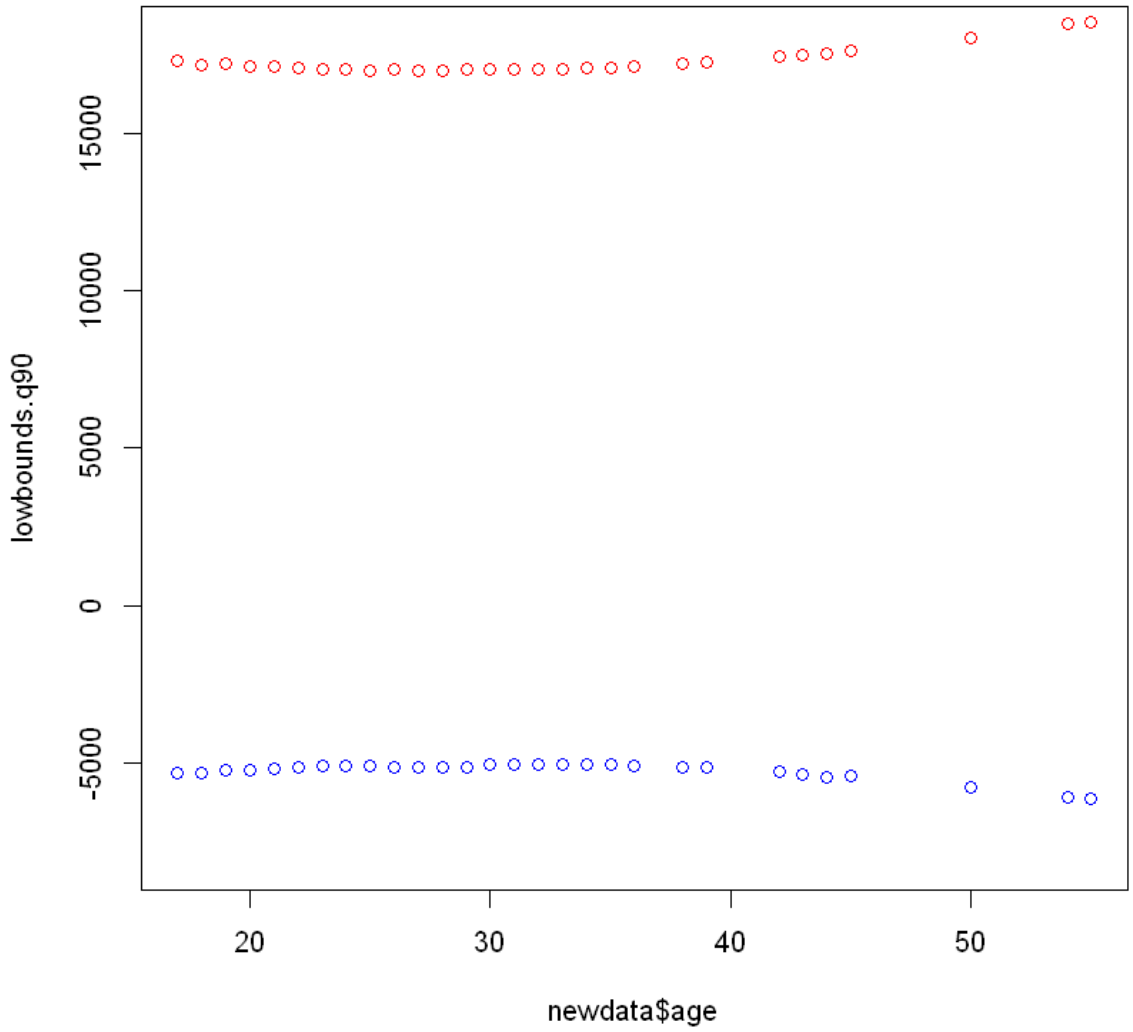
```
   Min. 1st Qu.  Median    Mean 3rd Qu.    Max.
  17.00   19.00   24.00   25.05   28.00   55.00
```

3227.1640	3226.77450	3227.2938	3227.8131	3224.3077	3227.8131	3227.0342	3225.735
4331.6234	4186.19845	4380.0984	4573.9984	3265.1735	4573.9984	4283.1484	3798.398
4696.1660	4673.31004	4703.7846	4734.2592	4528.5559	4734.2592	4688.5473	4612.360
5164.6430	4981.00153	5225.8568	5470.7121	3817.9390	5470.7121	5103.4292	4491.291
308.9804	434.20347	267.2393	100.2752	1227.2830	100.2752	350.7214	768.1317
181.2045	85.73219	213.0286	340.3251	-518.9259	340.3251	149.3804	-168.8607

-5562.0480	-5356.840	-5630.4505	-5904.0605	-4057.193	-5904.0605	-5493.645	-4809.
3999.3481	3875.109	4040.7611	4206.4133	3088.261	4206.4133	3957.935	3543.8
9152.2489	9021.080	9195.9721	9370.8646	8190.340	9370.8646	9108.526	8671.2
10267.3443	10236.244	10277.7110	10319.1780	10039.276	10319.1780	10256.978	10153.
5747.4440	6012.843	5658.9777	5305.1127	7693.702	5305.1127	5835.910	6720.5
944.3398	1575.825	733.8446	-108.1361	5575.234	-108.1361	1154.835	3259.7



```
Error in (function (classes, fdef, mtable) : unable to find an inherited meth
od for function 'Summary' for signature '"glm"'
Traceback:

1. Summary(lmlalonde)
2. (function (classes, fdef, mtable)
 . {
 .     methods <- .findInheritedMethods(classes, fdef, mtable)
 .     if (length(methods) == 1L)
 .         return(methods[[1L]])
 .     else if (length(methods) == 0L) {
 .         cnames <- paste0("\"", vapply(classes, as.character,
 .             ""), "\"", collapse = ", ")
 .         stop(gettextf("unable to find an inherited method for function %s
for signature %s",
 .             sQuote(fdef@generic), sQuote(cnames)), domain = NA)
 .     }
 .     else stop("Internal error in finding inherited methods; didn't return
a unique method",
 .         domain = NA)
 . })(list("glm"), new("groupGenericFunction", .Data = function (x,
 .     ..., na.rm = FALSE)
 . standardGeneric("Summary"), groupMembers = list("max", "min",
 .     "range", "prod", "sum", "any", "all"), generic = structure("Summary",
package = "base"),
 .     package = "base", group = list(), valueClass = character(0),
 .     signature = c("x", "na.rm"), default = NULL, skeleton = (function (x,
 .         ..., na.rm = FALSE)
 .     stop("invalid call in method dispatch to 'Summary' (no default metho
d)",
 .         domain = NA))(x, ..., na.rm = na.rm)), <environment>)
3. stop(gettextf("unable to find an inherited method for function %s for sign
ature %s",
 .     sQuote(fdef@generic), sQuote(cnames)), domain = NA)
```



In [85]:
```
head(data.frame(upperbounds.medians,lowbounds.medians,lowbounds.q90,upperbounds.q90,m_educ,m_re74,m_re75,q90.edu,q90.re74,q90.re75,newdata$age))
```

```
Warning message in data.frame(upperbounds.medians, lowbounds.medians, lowbounds.q90, :
"row names were found from a short variable and have been discarded"
```

upperbounds.medians	lowbounds.medians	lowbounds.q90	upperbounds.q90	m_educ
15133.82	-6570.202	-5108.423	17034.12	10
15086.65	-6504.220	-5117.491	17023.95	10
15151.71	-6580.343	-5117.607	17086.50	10
15230.41	-6634.446	-5316.342	17185.75	10
15269.59	-6707.853	-5428.120	17614.52	10
15230.41	-6634.446	-5316.342	17185.75	10

Answer #3

In [36]:
```
library(foreign)
getwd()
setwd("C:/Users/Hanna/Downloads")
mydata<-read.dta("nsw.dta")
cof_storage = rep(1:10000)
for (i in 1:10000) { boot.sample = sample(1:nrow(mydata), nrow(mydata), replac
e= TRUE)
 newset<-mydata[boot.sample,]
 mb<-lm(re78~ treat,data = newset)
 cof_storage[i]<-mb$coef[2]
  }
quantile(cof_storage, c(0.25,0.975))
manual<-quantile(cof_storage, c(0.25,0.975))
#analytical
ma<-lm(mydata$re78~ mydata$treat)
confint(ma)
analytical<-confint(ma)[2,]

table(analytical,manual)
```

'C:/Users/Hanna/Downloads'

25%	546.244009746679
97.5%	1864.21311818525

	2.5 %	97.5 %
(Intercept)	4495.60595	5684.491
mydata$treat	-40.52635	1813.134

```
                  manual
analytical          546.244009746679 1864.21311818525
  -40.5263533904598                1                0
  1813.13379829327                 0                1
```

Answer #4

In [31]:
```
preds <- cof_storage[c(1:722)] #taking the first 722 values for the example
actual <- mydata$re78
rss <- sum((preds - actual) ^ 2)
tss <- sum((actual - mean(actual)) ^ 2)
rsq <- 1 - rss/tss
print(rsq)
#shows how much better the predictive model is to the mean model.
```

```
[1] -0.5435433
```

Answer #5

In [37]:
```
mydata<- mydata[, -1]
glm.fit= glm(treat~age+education+black+hispanic+married+nodegree+re75, data= m
ydata, family= "binomial")
summary(glm.fit)

Probability_distribution_for_treatment_group<- predict (glm.fit,type = "respon
se")[which(predict(glm.fit, mydata,type="response")>0.5)]
hist(Probability_distribution_for_treatment_group,col="red")
predict (glm.fit ,mydata,type="response")
predicted_Y<- rep(0,722)
predicted_Y[which(predict(glm.fit,type = "response")>0.5)] <- 1

Probability_distribution_for_control_group<- predict (glm.fit,type = "respons
e")[which(predict(glm.fit, mydata,type="response")<0.5)]
hist(Probability_distribution_for_control_group, col = "blue")
length(which(predict(glm.fit,type = "response")>0.5))
```

```
Call:
glm(formula = treat ~ age + education + black + hispanic + married +
    nodegree + re75, family = "binomial", data = mydata)

Deviance Residuals:
    Min       1Q   Median       3Q      Max
-1.2318  -0.9981  -0.9696   1.3521   1.4851

Coefficients:
              Estimate Std. Error z value Pr(>|z|)
(Intercept)  4.343e-01  8.599e-01   0.505   0.6135
age         -1.624e-03  1.204e-02  -0.135   0.8927
education   -2.378e-02  5.966e-02  -0.399   0.6901
black       -9.743e-02  2.635e-01  -0.370   0.7116
hispanic    -2.525e-01  3.481e-01  -0.725   0.4682
married      9.292e-02  2.158e-01   0.431   0.6668
nodegree    -5.292e-01  2.443e-01  -2.167   0.0303 *
re75        -3.121e-06  1.550e-05  -0.201   0.8404
---
Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

(Dispersion parameter for binomial family taken to be 1)

    Null deviance: 978.09  on 721  degrees of freedom
Residual deviance: 970.25  on 714  degrees of freedom
AIC: 986.25

Number of Fisher Scoring iterations: 4
```

1	0.396225912847882
2	0.354962802210249
3	0.500673520701647
4	0.378035995583314
5	0.392626208225703
6	0.391216275670585
7	0.503516107943042
8	0.376128249440221
9	0.480148483656082
10	0.546905278532309
11	0.392377497458912
12	0.498382302003517
13	0.398451628952347
14	0.40584047870905
15	0.404557184787712
16	0.386721670740282
17	0.495945832527907
18	0.385181816524545
19	0.496612651796742
20	0.502297873385584
21	0.379564918924397
22	0.508671660273811
23	0.408357413918736
24	0.37918246254408
25	0.370005674809203
26	0.387106989806774
27	0.399338860495208
28	0.378800156250085
29	0.347691721170464
30	0.387492450218655
31	0.37918246254408
32	0.387492450218655
33	0.398446089491543
34	0.409625190185038
35	0.400898406747964
36	0.504734300753949
37	0.502703955317149
38	0.483913861664219
39	0.412912352185081
40	0.526720914379688
41	0.40987497784081
42	0.384797210626191
43	0.504328242014872
44	0.415834519861758
45	0.398841034609666
46	0.383648092348568
47	0.380713184415002
48	0.400898406747964

49 0.398841034609666
50 0.393152323082491
51 0.412912352185081
52 0.503516107943042
53 0.394564710868809
54 0.398559850069125
55 0.38147943751925
56 0.502703955317149
57 0.398949291347399
58 0.387492450218655
59 0.385428859429113
60 0.435062960585381
61 0.507138053937616
62 0.403154495296983
63 0.398841034609666
64 0.394176742740828
65 0.381170063653447
66 0.380576137675207
67 0.378035995583314
68 0.417109143989018
69 0.388897535409484
70 0.378035995583314
71 0.425844514424865
72 0.42763540390194
73 0.495133713108439
74 0.389283646309638
75 0.488232167083217
76 0.409906379711107
77 0.502703955317149
78 0.483653371781889
79 0.386482381315026
80 0.401982479502198
81 0.503922177566035
82 0.376712778780428
83 0.391603213461201
84 0.373083853171877
85 0.379947524974568
86 0.502703955317149
87 0.410882206992114
88 0.406702851978354
89 0.379637965884288
90 0.505546398960552
91 0.52176914303544
92 0.495945832527907
93 0.394952811910835
94 0.373844020453997
95 0.360052764879864
96 0.378418000458156
97 0.403401081302816

98	0.528655045742064
99	0.39572941096051
100	0.398559850069125
101	0.387492450218655
102	0.396077230921326
103	0.488659185791586
104	0.378800156250085
105	0.504328242014872
106	0.371565401639847
107	0.38643655600967
108	0.527035716640512
109	0.386515642186538
110	0.392764842787508
111	0.391783245230585
112	0.390320791324527
113	0.399465138172406
114	0.495800514497468
115	0.384412748644843
116	0.351622643808709
117	0.384797210626191
118	0.38402843100667
119	0.504439755372049
120	0.40605548281577
121	0.378800156250085
122	0.441510138748802
123	0.522572992845292
124	0.38147943751925
125	0.381096236968731
126	0.383122834074522
127	0.495945832527907
128	0.399923603049733
129	0.340721754360203
130	0.393152323082491
131	0.376890890600614
132	0.387376208379314
133	0.545453003371793
134	0.392873074107087
135	0.505107477013104
136	0.343532007761267
137	0.375489900772844
138	0.341750430400933
139	0.497535337087372
140	0.386159266943549
141	0.378176829797095
142	0.368163240823253
143	0.415397085976937
144	0.349176407606222
145	0.351434150382385
146	0.380954335952557

147 0.347393590228248
148 0.466406372769029
149 0.379501389467873
150 0.391499125052317
151 0.380201059481912
152 0.386588990899109
153 0.386857967276357
154 0.501448345785281
155 0.384774987592466
156 0.503829077034124
157 0.397554644781705
158 0.391808596641207
159 0.386092375389039
160 0.385704790934062
161 0.355225992727685
162 0.349819744030351
163 0.379678144041376
164 0.398514843846204
165 0.384346261835187
166 0.353738614310888
167 0.381275846273798
168 0.399335946053566
169 0.401360282380325
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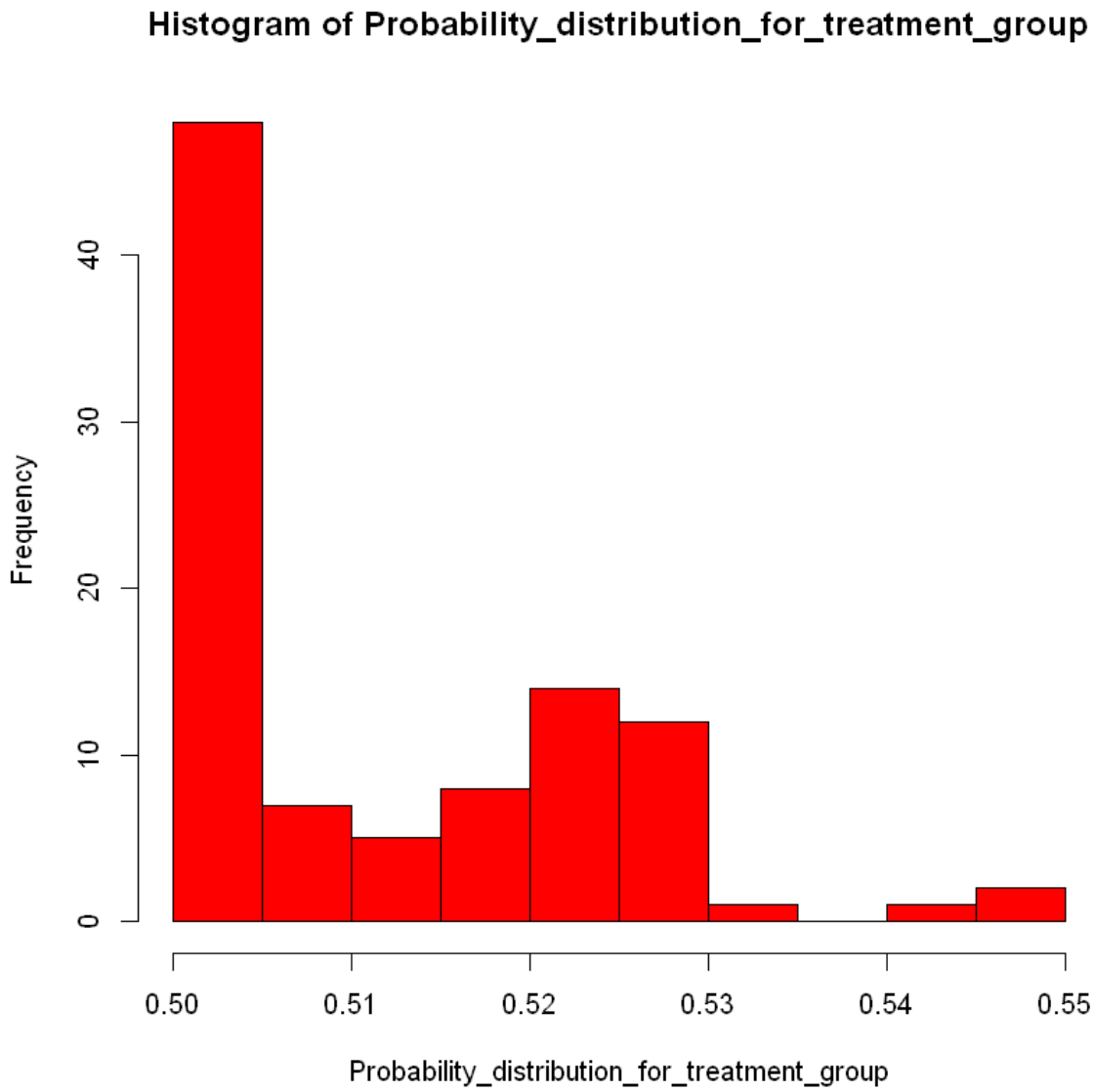
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0.54
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Probability_distribution_for_treatment_group

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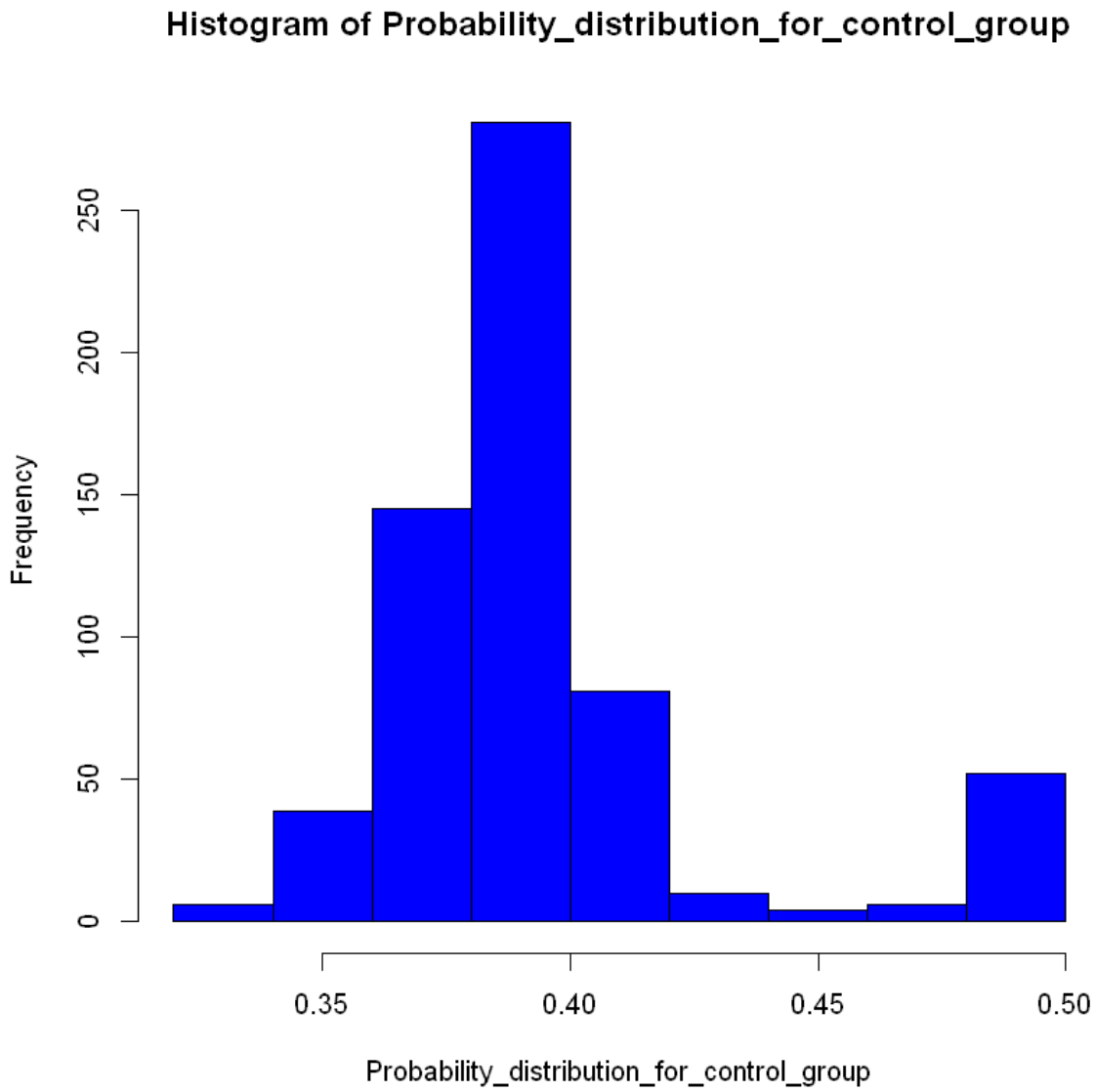
Histogram of Probability_distribution_for_control_group
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200
250
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0.45
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