HW1

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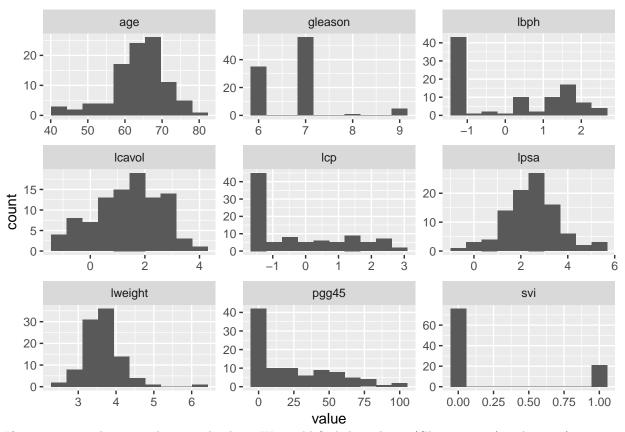
2/1/2020

Problem 1

From the numerical and graphical summary, we can see that lbph (log(benign prostatic hyperplasia amount)), lcp (log(capsular penetration)) and pgg45 (percentage Gleason scores 4 or 5) are quite skewed to the right. Most of men have small value for those three variables. svi is binary which takes value between 0 and 1. gleason is also categorical variable which takes value among 6,7,8 and 9.

summary(prostate,digits=3)

```
##
        lcavol
                         lweight
                                                           1bph
                                           age
##
                              :2.37
    Min.
           :-1.347
                      Min.
                                      Min.
                                              :41.0
                                                      Min.
                                                              :-1.39
##
    1st Qu.: 0.513
                      1st Qu.:3.38
                                      1st Qu.:60.0
                                                      1st Qu.:-1.39
    Median : 1.447
                      Median:3.62
                                      Median:65.0
                                                      Median: 0.30
##
##
    Mean
           : 1.350
                      Mean
                              :3.65
                                      Mean
                                              :63.9
                                                      Mean
                                                              : 0.10
                                      3rd Qu.:68.0
##
    3rd Qu.: 2.127
                      3rd Qu.:3.88
                                                      3rd Qu.: 1.56
                                              :79.0
                                                              : 2.33
##
    Max.
           : 3.821
                      Max.
                              :6.11
                                      Max.
                                                      Max.
##
         svi
                          lcp
                                          gleason
                                                           pgg45
##
    Min.
           :0.000
                             :-1.386
                                       Min.
                                               :6.00
                                                               : 0.0
                     Min.
                                                       Min.
##
    1st Qu.:0.000
                     1st Qu.:-1.386
                                       1st Qu.:6.00
                                                       1st Qu.: 0.0
                                       Median:7.00
                                                       Median: 15.0
##
    Median :0.000
                     Median :-0.799
                             :-0.179
                                                               : 24.4
##
    Mean
           :0.216
                     Mean
                                       Mean
                                               :6.75
                                                       Mean
##
    3rd Qu.:0.000
                     3rd Qu.: 1.179
                                       3rd Qu.:7.00
                                                       3rd Qu.: 40.0
##
    Max.
           :1.000
                     Max.
                            : 2.904
                                       Max.
                                               :9.00
                                                       Max.
                                                               :100.0
##
         lpsa
##
    Min.
           :-0.431
##
    1st Qu.: 1.732
##
    Median : 2.592
##
    Mean
           : 2.478
##
    3rd Qu.: 3.056
    Max.
           : 5.583
ggplot(gather(prostate), aes(value)) +
    geom_histogram(bins=10) +
    facet_wrap(~key, scales = 'free')
```



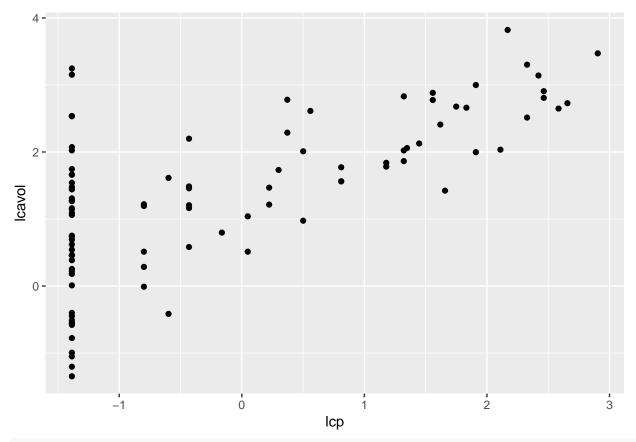
If we run a correlation analysis on the data. We could find that gleason (Gleason score) and pgg45 (percentage Gleason scores 4 or 5) are strongly correlated with value of 0.752. It is not surprising since both of them are related to Gleason score.

round(cor(prostate),3)

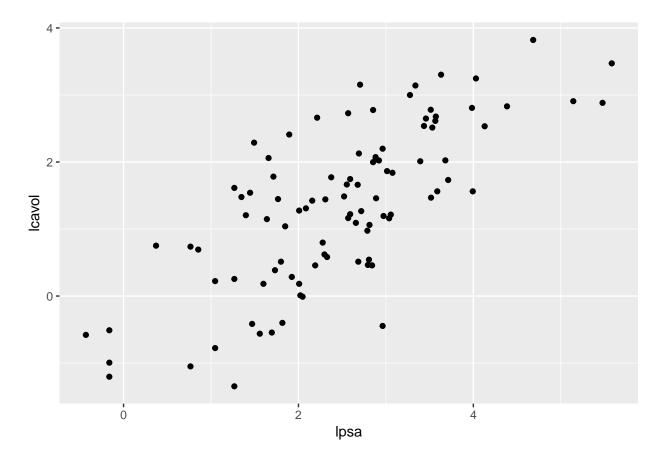
```
##
           lcavol lweight
                             age
                                   lbph
                                            svi
                                                   1cp gleason pgg45
## lcavol
            1.000
                     0.194 0.225
                                  0.027
                                          0.539
                                                 0.675
                                                          0.432 0.434 0.734
## lweight
            0.194
                     1.000 0.308
                                  0.435
                                          0.109
                                                 0.100
                                                         -0.001 0.051 0.354
## age
            0.225
                     0.308 1.000
                                  0.350
                                          0.118
                                                 0.128
                                                          0.269 0.276 0.170
  lbph
            0.027
                     0.435 0.350
                                   1.000 -0.086 -0.007
                                                          0.078 0.078 0.180
##
##
  svi
            0.539
                     0.109 0.118
                                 -0.086
                                          1.000
                                                 0.673
                                                          0.320 0.458 0.566
            0.675
##
   lcp
                     0.100 0.128
                                 -0.007
                                          0.673
                                                 1.000
                                                          0.515 0.632 0.549
  gleason
##
            0.432
                    -0.001 0.269
                                  0.078
                                         0.320
                                                 0.515
                                                          1.000 0.752 0.369
## pgg45
            0.434
                     0.051 0.276
                                  0.078
                                          0.458
                                                 0.632
                                                          0.752 1.000 0.422
            0.734
                                         0.566
                                                 0.549
                                                          0.369 0.422 1.000
## lpsa
                     0.354 0.170 0.180
```

If we assume that lcavol (log(cancer volume)) is the dependent variable. We could find that lcp (log(capsular penetration)) and lpsa (log(prostate specific antigen)) are highly correlated to lcavol. Scatter plots of those two are shown below and it is clear that they both are positively correlated to lcavol.

```
ggplot(prostate,aes(x=lcp,y=lcavol))+geom_point()
```



ggplot(prostate,aes(x=lpsa,y=lcavol))+geom_point()

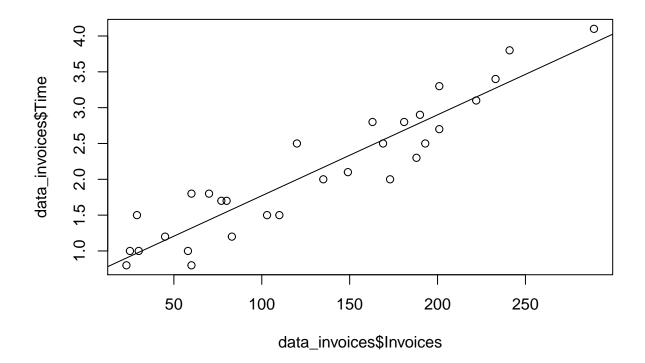


Problem 3

3(a)

Linear regression model fits the data well since most of the points are near the regression line. The processing time increases as number of invoices increases. There is no obvious outlier.

```
data_invoices = read.csv('invoices.txt',sep='\t')
plot(data_invoices$Invoices,data_invoices$Time)
lm_invoices = lm(Time~Invoices,data_invoices)
abline(lm_invoices)
```



3(b)

The 95% confidence interval for the start-up time is (0.391,0.892)

predict(lm_invoices, newdata=data.frame(Invoices=0), interval='confidence')

fit lwr upr ## 1 0.6417099 0.3912496 0.8921701

3(c)

Since p-value=0.1257 > 0.05, we do not have enough evidence to reject the null hypothesis that the average processing time for an additional invoice is 0.01 hours.

```
mycoef = summary(lm_invoices)$coefficients
2*pt((0.01-mycoef[2,1])/mycoef[2,2], 28)
```

[1] 0.1257402

3(d)

The point estimate is 2.109 and 95% prediction interval is (1.986, 2.232) for the time taken to process 130 invoices.

```
predict(lm_invoices, newdata=data.frame(Invoices=130), interval='confidence')
```

```
## fit lwr upr
## 1 2.109624 1.986293 2.232954
```

Problem 5

5(a)

R-squared is 0.2792 which means that 27.92% of the variance in the data can be explained by the linear regression model. Adjusted R-squared is 0.2341 which means that 23.41% of the variance in the data can be explained by the linear regression model. R-squared always increases as number of predictors increases, adjusted R-squared penalizes additional predictors that are not helpful to explain the variance in data. Relatively low adjusted R-squared means that the linear regression model does not fit data well.

```
data_indicators = read.csv('indicators.txt',sep='\t')
lm_indicators = lm(PriceChange~LoanPaymentsOverdue,data=data_indicators)
summary(lm_indicators)
##
## Call:
## lm(formula = PriceChange ~ LoanPaymentsOverdue, data = data_indicators)
##
## Residuals:
##
       Min
                1Q Median
                                3Q
                                       Max
## -4.6541 -3.3419 -0.6944 2.5288 6.9163
##
## Coefficients:
##
                       Estimate Std. Error t value Pr(>|t|)
                         4.5145
                                    3.3240
                                             1.358
                                                     0.1933
## (Intercept)
                        -2.2485
                                    0.9033
                                                     0.0242 *
## LoanPaymentsOverdue
                                           -2.489
## ---
## Signif. codes: 0 '***' 0.001 '**' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 3.954 on 16 degrees of freedom
## Multiple R-squared: 0.2792, Adjusted R-squared: 0.2341
## F-statistic: 6.196 on 1 and 16 DF, p-value: 0.02419
```

The 95% confidence interval for the slope is (-4.163,-0.333). Since the confidence interval is negative, there is evidence of a significant negative linear association.

```
confint(lm_indicators)
```

```
## 2.5 % 97.5 %
## (Intercept) -2.532112 11.5611000
## LoanPaymentsOverdue -4.163454 -0.3335853
```

5(c)

5(b)

E(Y|X=4) = -4.479 with 95% confidence interval (-6.648, -2.310). Since the confidence interval does not include 0%, it is not a feasible value.

```
predict(lm indicators,newdata=data.frame(LoanPaymentsOverdue=4),interval='confidence')
```

```
## fit lwr upr
## 1 -4.479585 -6.648849 -2.310322
```