Babies and ANOVA

January 7, 2022

```
[1]: babies<-read.csv("babies.csv", header=TRUE, sep = ";")
head(babies)</pre>
```

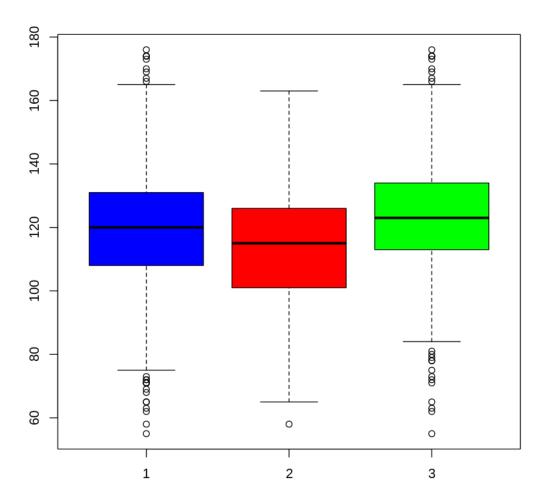
```
smoke
                          bwt
                                   gestation
                                              parity
                                                                height
                                                                         weight
                                                       age
                          <int>
                                   <int>
                                               <int>
                                                        <int>
                                                                 <int>
                                                                          <int>
                                                                                   <int>
                          120
                                   284
                                                       27
                                                                62
                                                                          100
                                               0
                                                                                   0
                          113
                                   282
                                                                          135
                                                                                   0
                                               0
                                                       33
                                                                64
A data.frame: 6 \times 7
                          128
                                   279
                                                       28
                                               0
                                                                64
                                                                          115
                          123
                                   999
                                               0
                                                       36
                                                                                   0
                                                                 69
                                                                          190
                          108
                                   282
                                               0
                                                       23
                                                                 67
                                                                          125
                                                                                   1
                          136
                                   286
                                                       25
                                                                62
                                                                          93
```

- [2]: babies<-babies[(babies\$weight<999 & babies\$gestation<999 & babies\$height<99 & babies\$age< 99),]
- [3]: smoking<-babies[babies\$smoke==1,]
 nonsmoking<-babies[babies\$smoke==0,]
 head(smoking)
 head(nonsmoking)</pre>

		bwt <int></int>	gestation <int></int>	parity <int></int>	age <int></int>	height <int></int>	weight <int></int>	$\begin{array}{c} {\rm smoke} \\ {\rm } \end{array}$
A data.frame: 6×7	3	128	279	0	28	64	115	1
	5	108	282	0	23	67	125	1
	10	143	299	0	30	66	136	1
	12	144	282	0	32	64	124	1
	13	141	279	0	23	63	128	1
	14	110	281	0	36	61	99	1
		bwt	gestation	parity	age	height	weight	smoke

			0		0			
		<int $>$						
A data.frame: 6×7	1	120	284	0	27	62	100	0
	2	113	282	0	33	64	135	0
	6	136	286	0	25	62	93	0
	7	138	244	0	33	62	178	0
	8	132	245	0	23	65	140	0
	9	120	289	0	25	62	125	0

[4]: boxplot(babies\$bwt, smoking\$bwt, nonsmoking\$bwt, col=c("blue", "red", "green"))



[5]: summary(babies\$bwt) summary(smoking\$bwt) summary(nonsmoking\$bwt)

Min. 1st Qu. Median Mean 3rd Qu. Max. 55.0 108.0 120.0 119.5 131.0 176.0 Min. 1st Qu. Median Mean 3rd Qu. Max. 58.0 101.0 115.0 113.8 126.0 163.0 Min. 1st Qu. Median Mean 3rd Qu. Max. 55.0 113.0 123.0 123.1 134.0 176.0

```
[6]: fit<-aov(bwt ~ smoke, data=babies)
      summary(fit)
                   Df Sum Sq Mean Sq F value
                                                Pr(>F)
                        3835
                                3835
                                       11.48 0.000728 ***
     smoke
                 1182 395024
                                 334
     Residuals
     Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1
[13]: avg<-mean(babies$bwt)
      mus<-mean(smoking$bwt)</pre>
      muns<-mean(nonsmoking$bwt)</pre>
      mu<-(mus+muns)/2
      alphas<-(mus-muns)/2
      alphans<-(muns-mus)/2
      cat("avg = " ,avg, "\n")
      cat("mus = " ,mus, "\n")
      cat("muns = ", muns, "\n")
      cat("mu = ", mu, "\n")
      cat("alphas = ", alphas, "\n")
      cat("alphans = ", alphans, "\n")
     avg = 119.5236
     mus = 113.8192
     muns = 123.0853
     mu = 118.4522
     alphas = -4.633071
     alphans = 4.633071
 []:
```