

*Data Analysis Tools and
Practice(Using R)*

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R语言实战



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- US baby names数据集：
- 美国1880年到2008年Top1000的男婴和女婴的名字

- 258000条记录

- year

- name

- sex

- percent

year	name	percent	sex
1880	John	0.081541	boy
1880	William	0.080511	boy
1880	James	0.050057	boy
1880	Charles	0.045167	boy
1880	George	0.043292	boy
1880	Frank	0.027380	boy
1880	Joseph	0.022229	boy
1880	Thomas	0.021401	boy

- 见： bnames.csv

```
> head(bnames,15)
  year   name percent sex
1 1880    John 0.081541 boy
2 1880  William 0.080511 boy
3 1880    James 0.050057 boy
4 1880  Charles 0.045167 boy
5 1880   George 0.043292 boy
6 1880    Frank 0.027380 boy
7 1880  Joseph 0.022229 boy
8 1880  Thomas 0.021401 boy
9 1880   Henry 0.020641 boy
10 1880  Robert 0.020404 boy
11 1880  Edward 0.019965 boy
12 1880   Harry 0.018175 boy
13 1880  Walter 0.014822 boy
14 1880  Arthur 0.013504 boy
15 1880    Fred 0.013251 boy
```

- * 该数据集中每年有多少记录
- * 数据集中男孩和女孩各自排名
- * 男孩名和女孩名的Top100
- * Top100中男孩名和女孩名的所占比例
- * 画图显示每一年男孩名和女孩名在Top100的比例
- * 哪些名字仅仅在一年中使用，哪些名字每一年都使用
- * 显示每个名字的平均百分比
- * 那个名字被使用的时间最长

dplyr

方差分析

方差分析定义

- 方差分析 (analysis of variance, ANOVA) 是分析各个自变量对因变量影响的一种方法。
- 这里的自变量就是定性变量的因子及可能出现的称为协变量 (covariate) 的定量变量。
- 分析结果是由一个方差分析表表示的
- 原理为：把因变量的值随着自变量的不同取值而得到的变化进行分解，使得每一个自变量都有一份贡献，最后剩下无法用已知的原因解释的则看成随机误差的贡献。
- 然后用各自变量的贡献和随机误差的贡献进行比较 (F检验)，以判断该自变量的不同水平是否对因变量的变化有显著贡献。输出就是F-值和检验的一些p-值。

一个例子

表9-1 单因素组间方差分析

治疗方案	
CBT	EMDR
s1	s6
s2	s7
s3	s8
s4	s9
s5	s10

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表9-2 单因素组内方差分析

患 者	时 间	
	5周	6个月
s1		
s2		
s3		
s4		
s5		
s6		
s7		
s8		
s9		
s10		

一个例子

表9-3 含组间和组内因子的双因素方差分析

		患 者	时 间	
			5周	6个月
疗法	CBT	s1 s2 s3 s4 s5		
	EMDR	s6		
		s7		
		s8		
		s9		
	EMDR	s10		

协方差分析

多元方差分析

aov()

- **aov(formula, data = dataframe)**

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表9-4 R表达式中的特殊符号

符 号	用 法
~	分隔符号，左边为响应变量，右边为解释变量。例如，用A、B和C预测y，代码为 $y \sim A + B + C$
+	分隔解释变量
:	表示变量的交互项。例如，用A、B和A与B的交互项来预测y，代码为 $y \sim A + B + A:B$
*	表示所有可能交互项。代码 $y \sim A * B * C$ 可展开为 $y \sim A + B + C + A:B + A:C + B:C + A:B:C$
^	表示交互项达到某个次数。代码 $y \sim (A + B + C)^2$ 可展开为 $y \sim A + B + C + A:B + A:C + B:C$
.	表示包含除因变量外的所有变量。例如，若一个数据框包含变量y、A、B和C，代码 $y \sim .$ 可展开为 $y \sim A + B + C$

表9-5 常见研究设计的表达式

设 计	表 达 式
单因素ANOVA	$y \sim A$
含单个协变量的单因素ANCOVA	$y \sim x + A$
双因素ANOVA	$y \sim A * B$
含两个协变量的双因素ANCOVA	$y \sim x1 + x2 + A*B$
随机化区组	$y \sim B + A$ (B是区组因子)
单因素组内ANOVA	$y \sim A + Error(Subject/A)$
含单个组内因子(w) 和单个组间因子(b) 的重复测量ANOVA	$y \sim B * W + Error(Subject/W)$

单因素方差分析

```
> table(trt)
trt
 1time 2times 4times drugD drugE
      10     10     10    10    10
> aggregate(response, by = list(trt), FUN = mean)
  Group.1
1  1time 5.78197
2  2times 9.22497
3  4times 12.37478
4  drugD 15.36117
5  drugE 20.94752
> aggregate(response, by = list(trt), FUN = sd)
  Group.1
1  1time 2.878113
2  2times 3.483054
3  4times 2.923119
4  drugD 3.454636
5  drugE 3.345003
```

```
> library(multcomp)
> attach(cholesterol)
```

```
> cholesterol
   trt response
1  1time  3.8612
2  1time 10.3868
3  1time  5.9059
4  1time  3.0609
5  1time  7.7204
6  1time  2.7139
7  1time  4.9243
8  1time  2.3039
9  1time  7.5301
10 1time  9.4123
11 2times 10.3993
12 2times  8.6027
13 2times 13.6320
14 2times  3.5054
15 2times  7.7703
16 2times  8.6266
17 2times  9.2274
18 2times  6.3159
19 2times 15.8258
20 2times  8.3443
21 4times 13.9621
-- -- -- -- --
```

单因素方差分析表

表 7.3: 单因素方差分析表

方差来源	自由度	平方和	均方	F 比	p 值
因素 A	$r - 1$	S_A	$MS_A = \frac{S_A}{r-1}$	$F = \frac{MS_A}{MS_E}$	p
误差	$n - r$	S_E	$MS_E = \frac{S_E}{n-r}$		
总和	$n - 1$	S_T			

```

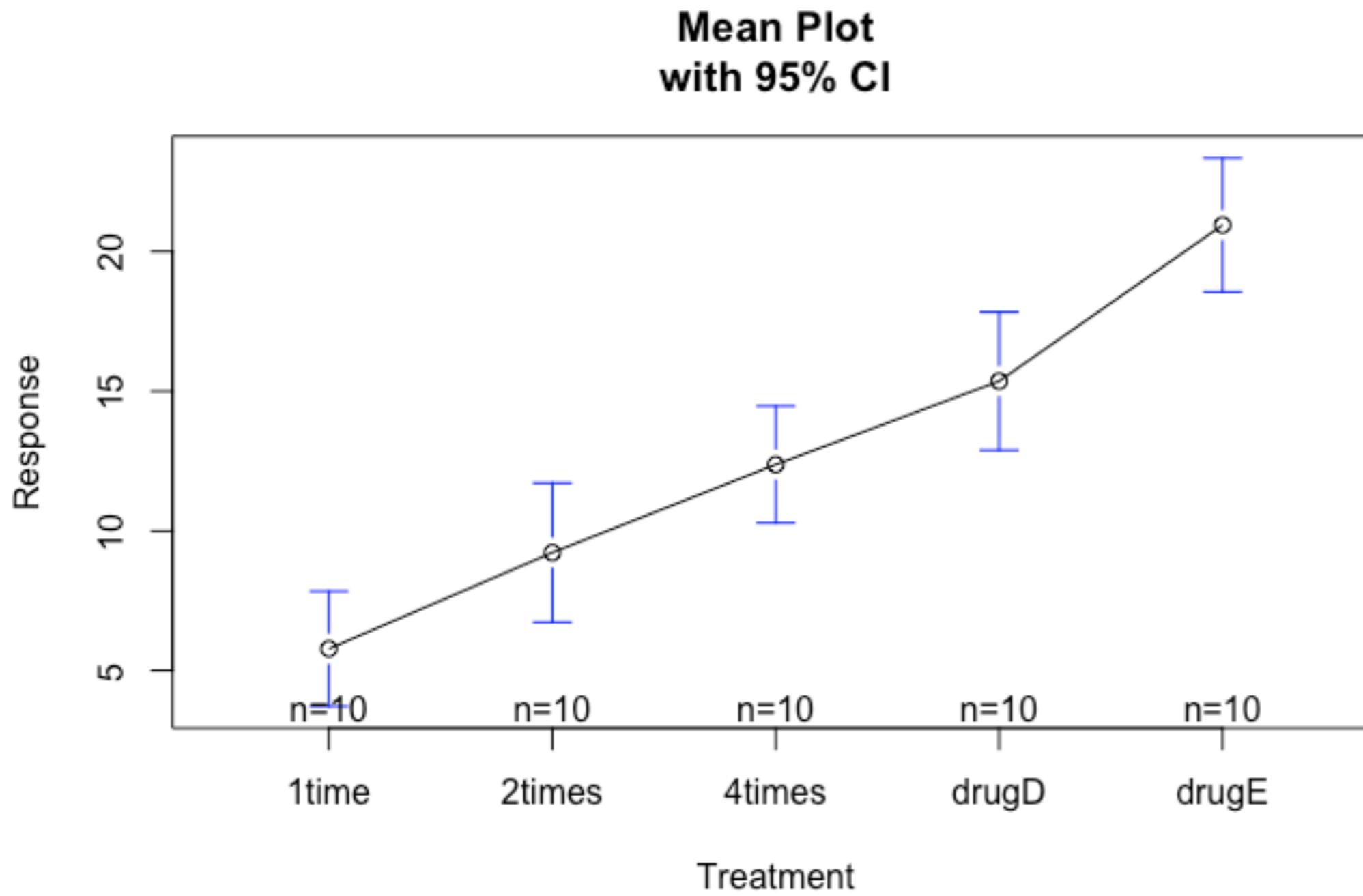
> fit <- aov(response ~ trt)
> summary(fit)

      Df Sum Sq Mean Sq F value    Pr(>F)
trt      4 1351.4   337.8   32.43 9.82e-13 ***
Residuals 45  468.8    10.4
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

```

单因素方差分析例子

```
> library(gplots)
> plotmeans(response ~ trt, xlab = "Treatment", ylab = "Response",
+     main = "Mean Plot\\nwith 95% CI")
|
```



多重比较

```
> TukeyHSD(fit)
Tukey multiple comparisons of means
95% family-wise confidence level
```

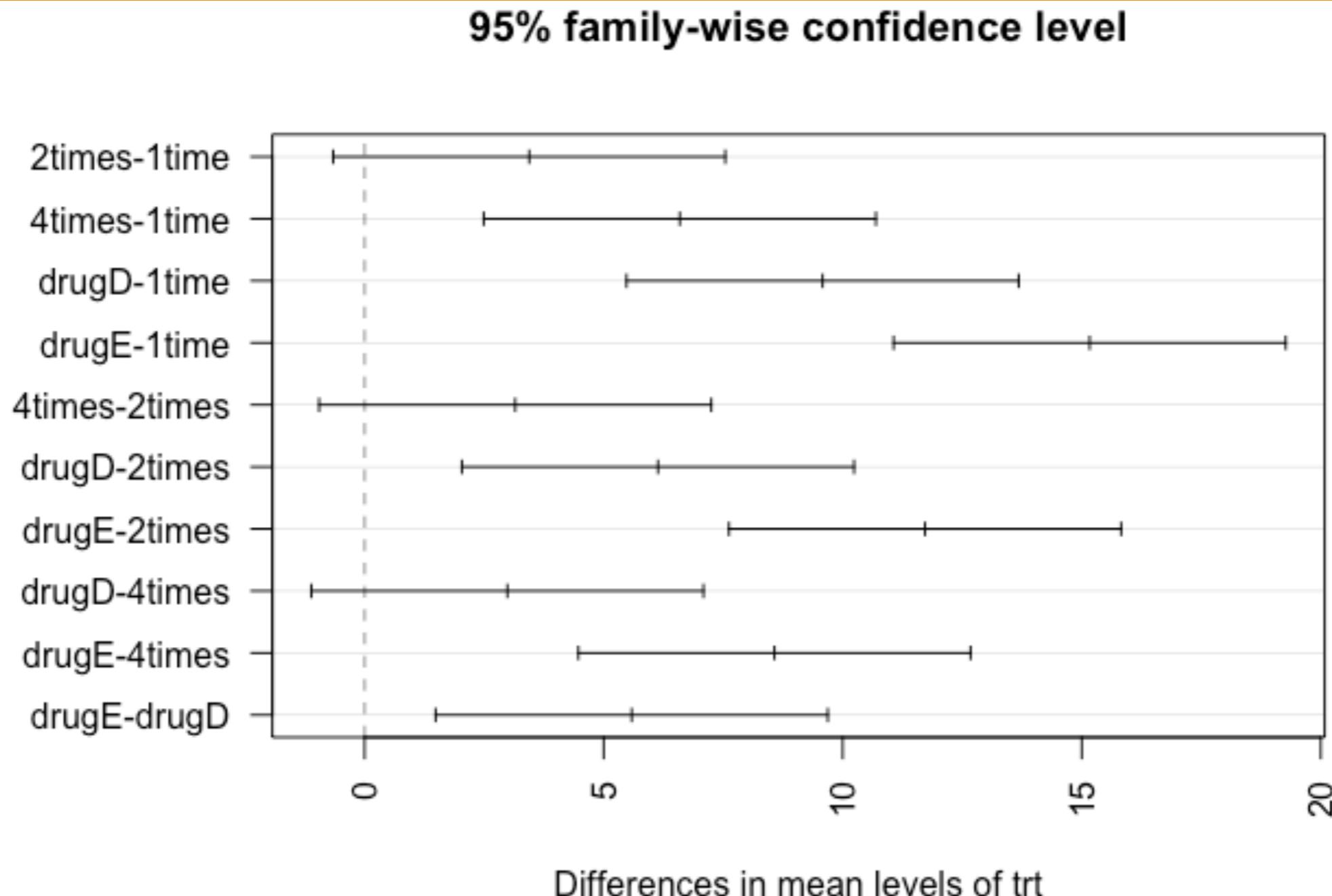
```
Fit: aov(formula = response ~ trt)
```

```
$trt
```

	diff	lwr	upr	p adj
2times-1time	3.44300	-0.6582817	7.544282	0.1380949
4times-1time	6.59281	2.4915283	10.694092	0.0003542
drugD-1time	9.57920	5.4779183	13.680482	0.0000003
drugE-1time	15.16555	11.0642683	19.266832	0.0000000
4times-2times	3.14981	-0.9514717	7.251092	0.2050382
drugD-2times	6.13620	2.0349183	10.237482	0.0009611
drugE-2times	11.72255	7.6212683	15.823832	0.0000000
drugD-4times	2.98639	-1.1148917	7.087672	0.2512446
drugE-4times	8.57274	4.4714583	12.674022	0.0000037
drugE-drugD	5.58635	1.4850683	9.687632	0.0030633

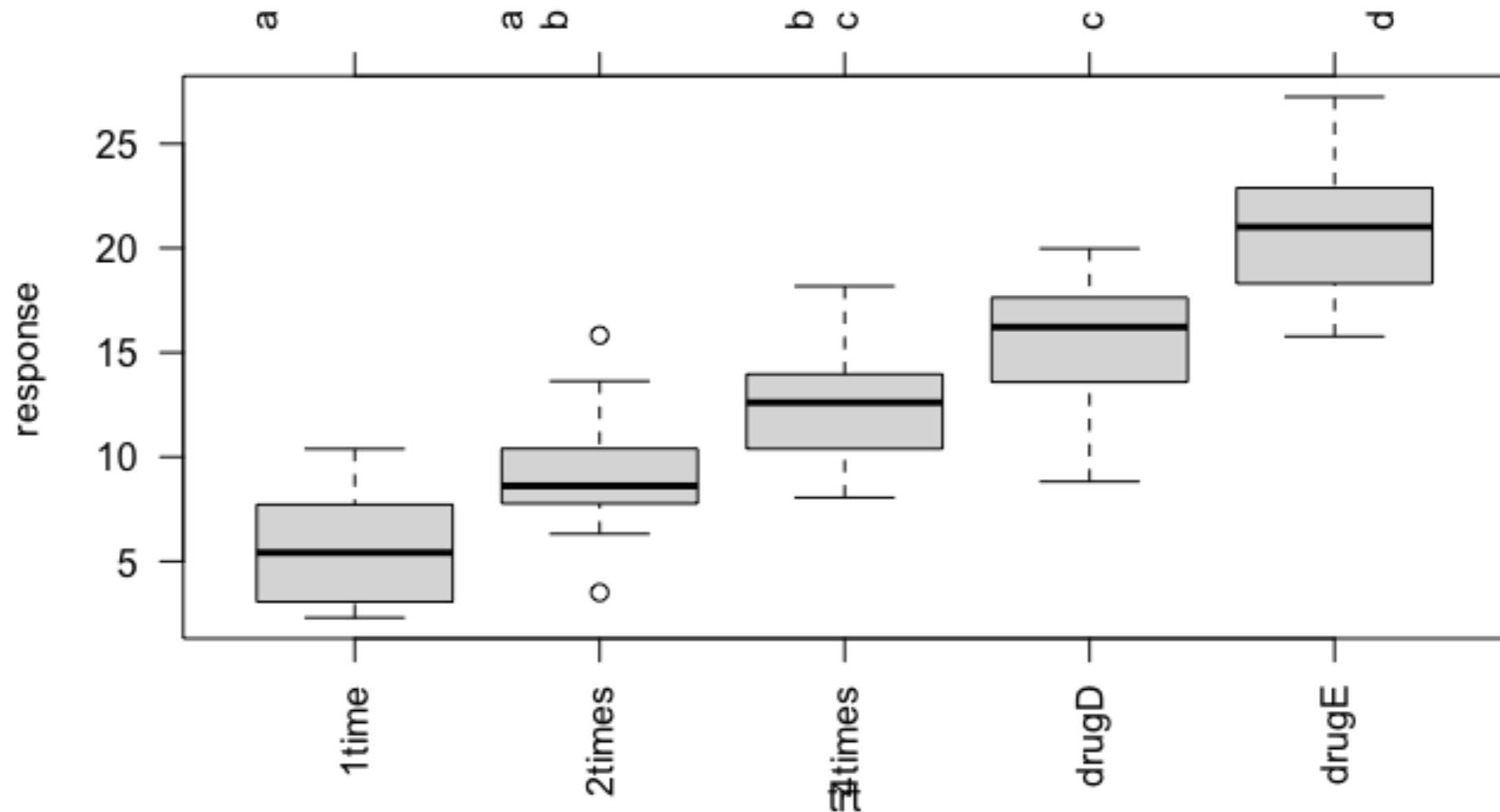
多重比较

```
> par(las = 2)
> par(mar = c(5, 8, 4, 2))
> plot(TukeyHSD(fit))
> par(opar)
```



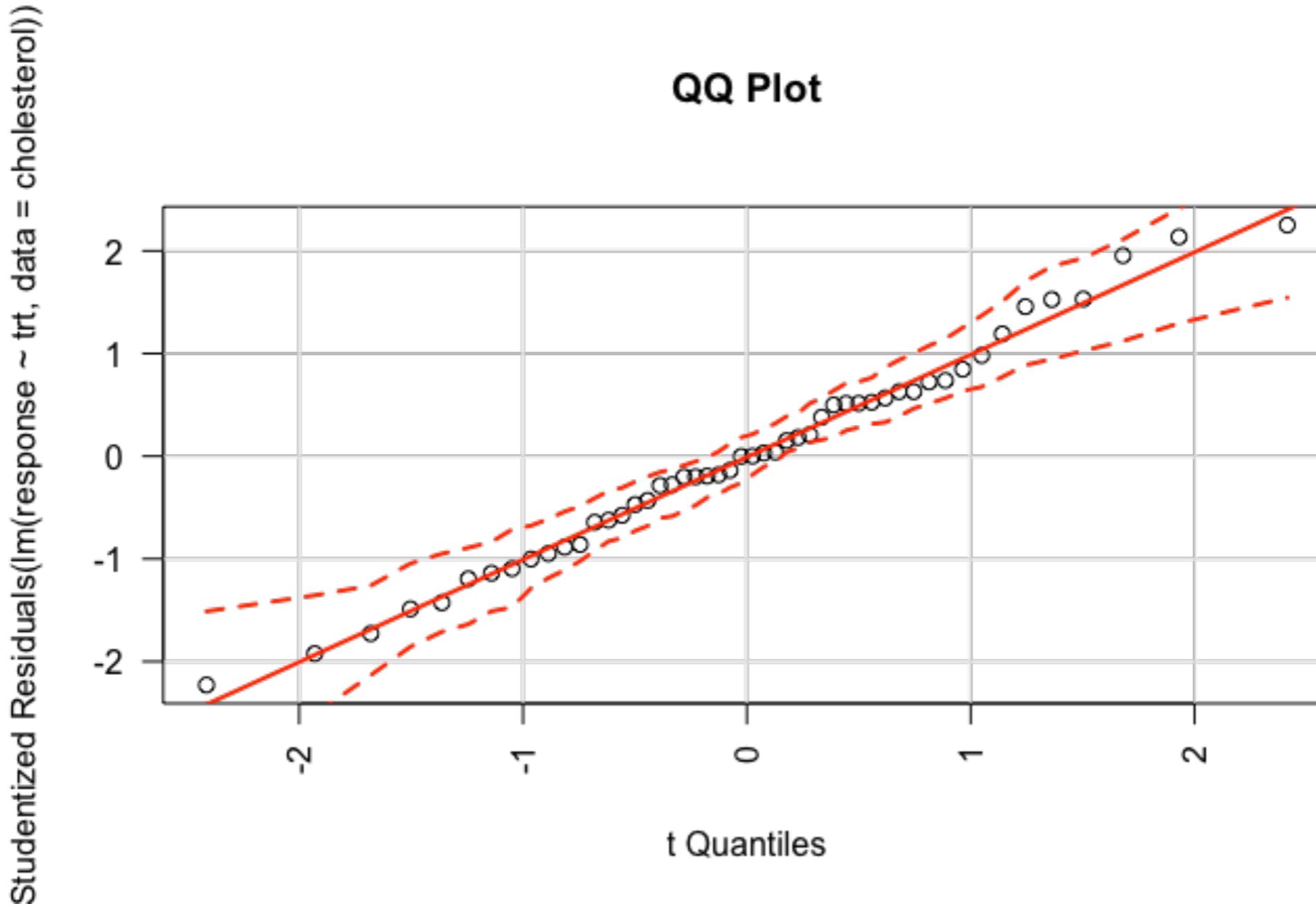
多重比较

```
library(multcomp)
par(mar = c(5, 4, 6, 2))
tuk <- glht(fit, linfct = mcp(trt = "Tukey"))
plot(cld(tuk, level = 0.05), col = "lightgrey")
par(opar)
```



正态假设检验

```
library(car)  
qqPlot(lm(response ~ trt, data = cholesterol), simulate = TRUE,  
       main = "QQ Plot", labels = FALSE)
```



单因素协方差分析

```
> data(litter, package = "multcomp")
> attach(litter)
> table(dose)
dose
  0   5   50  500
20  19   18   17
> aggregate(weight, by = list(dose), FUN = mean)
  Group.1      x
1       0 32.30850
2       5 29.30842
3      50 29.86611
4     500 29.64647
> fit <- aov(weight ~ gesttime + dose)
> summary(fit)

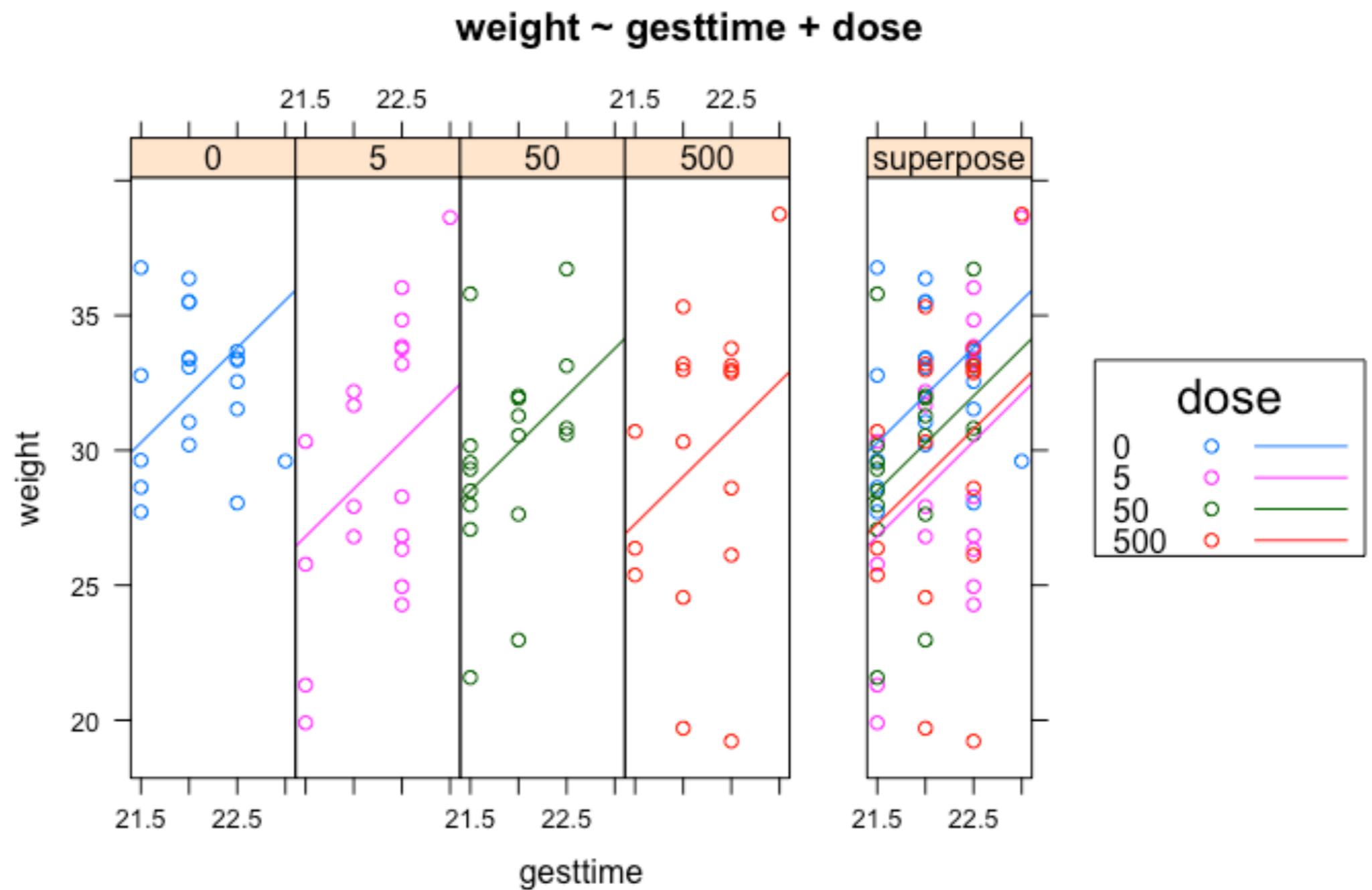
             Df Sum Sq Mean Sq F value    Pr(>F)
gesttime     1 134.3  134.30   8.049 0.00597 **
dose         3 137.1   45.71   2.739 0.04988 *
Residuals   69 1151.3   16.69

---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1
```

> litter				
	dose	weight	gesttime	number
1	0	28.05	22.5	15
2	0	33.33	22.5	14
3	0	36.37	22.0	14
4	0	35.52	22.0	13
5	0	36.77	21.5	15
6	0	29.60	23.0	5
7	0	27.72	21.5	16
8	0	33.67	22.5	15
9	0	32.55	22.5	14
10	0	32.78	21.5	15
11	0	31.05	22.0	12
12	0	33.40	22.5	15
13	0	30.20	22.0	16
14	0	28.63	21.5	7
15	0	33.38	22.0	15
16	0	33.43	22.0	13
17	0	29.63	21.5	14
18	0	33.08	22.0	15
19	0	31.53	22.5	16
20	0	35.48	22.0	9

单因素协方差分析

```
library(HH)  
ancova(weight ~ gesttime + dose, data = litter)
```



双因素方差分析

```
> table(supp, dose)
  dose
supp 0.5 1 2
  OJ 10 10 10
  VC 10 10 10
> aggregate(len, by = list(supp, dose), FUN = mean)
  Group.1 Group.2      x
1      OJ     0.5 13.23
2      VC     0.5  7.98
3      OJ     1.0 22.70
4      VC     1.0 16.77
5      OJ     2.0 26.06
6      VC     2.0 26.14
> aggregate(len, by = list(supp, dose), FUN = sd)
  Group.1 Group.2      x
1      OJ     0.5 4.459709
2      VC     0.5 2.746634
3      OJ     1.0 3.910953
4      VC     1.0 2.515309
5      OJ     2.0 2.655058
6      VC     2.0 4.797731
...   ...  ...  ...  ...  ...  ...
```

双因素方差分析表

表 7.12: 双因素方差分析表

方差来源	自由度	平方和	均方	F 比	p 值
因素 A	$r - 1$	S_A	$MS_A = \frac{S_A}{r-1}$	$F_A = \frac{MS_A}{MS_E}$	p_A
因素 B	$s - 1$	S_B	$MS_B = \frac{S_B}{s-1}$	$F_B = \frac{MS_B}{MS_E}$	p_B
误差	$(r - 1)(s - 1)$	S_E	$MS_E = \frac{S_E}{(r-1)(s-1)}$		
总和	$rs - 1$	S_T			

```

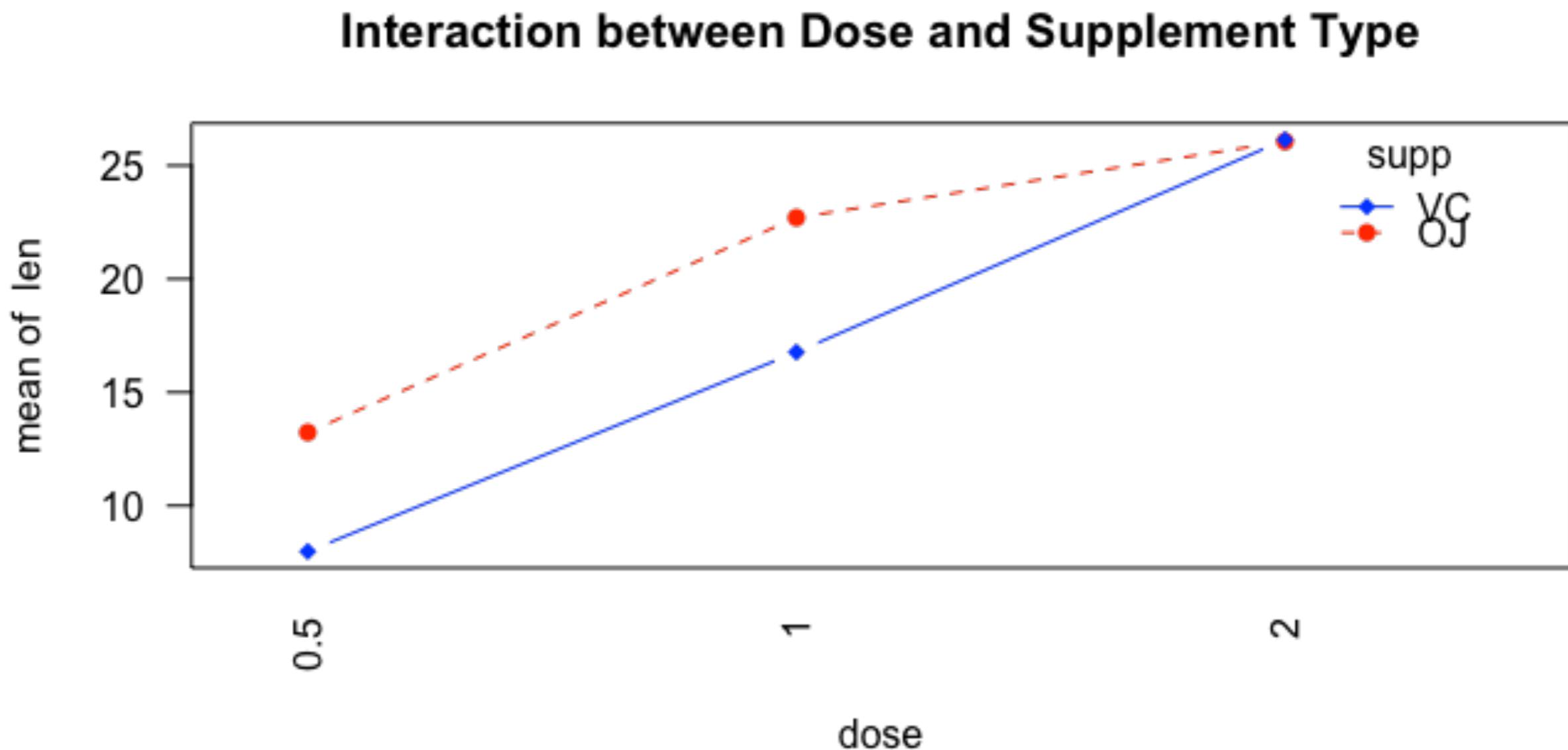
> fit <- aov(len ~ supp * dose)
> summary(fit)

  Df Sum Sq Mean Sq F value    Pr(>F)
supp      1  205.4   205.4  12.317 0.000894 ***
dose      1 2224.3  2224.3 133.415 < 2e-16 ***
supp:dose 1    88.9    88.9   5.333 0.024631 *
Residuals 56   933.6   16.7
---
Signif. codes:  0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

```

双因素方差分析例子

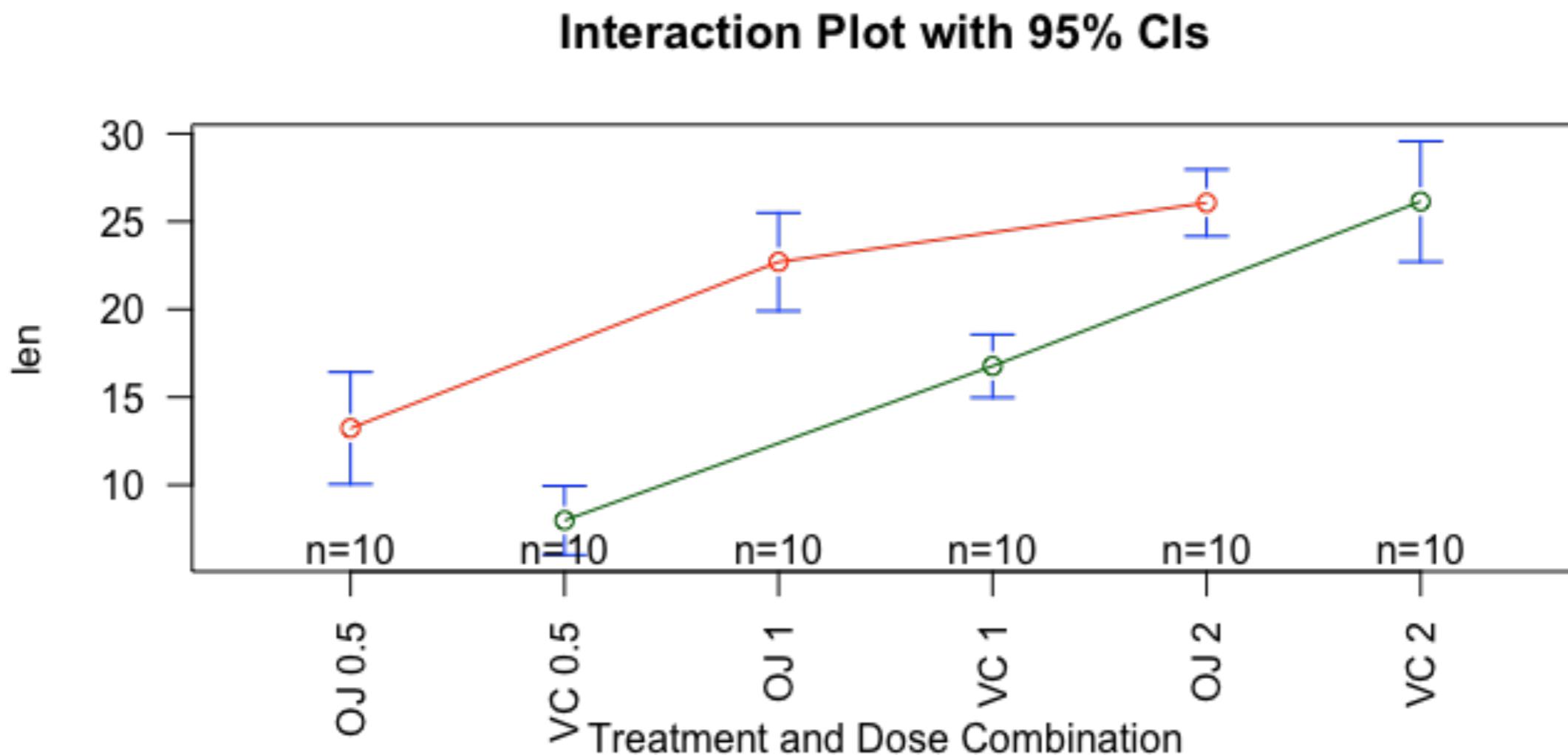
```
interaction.plot(dose, supp, len,  
    type = "b", col = c("red", "blue"), pch = c(16, 18),  
    main = "Interaction between Dose and Supplement Type")
```



双因素方差分析例子

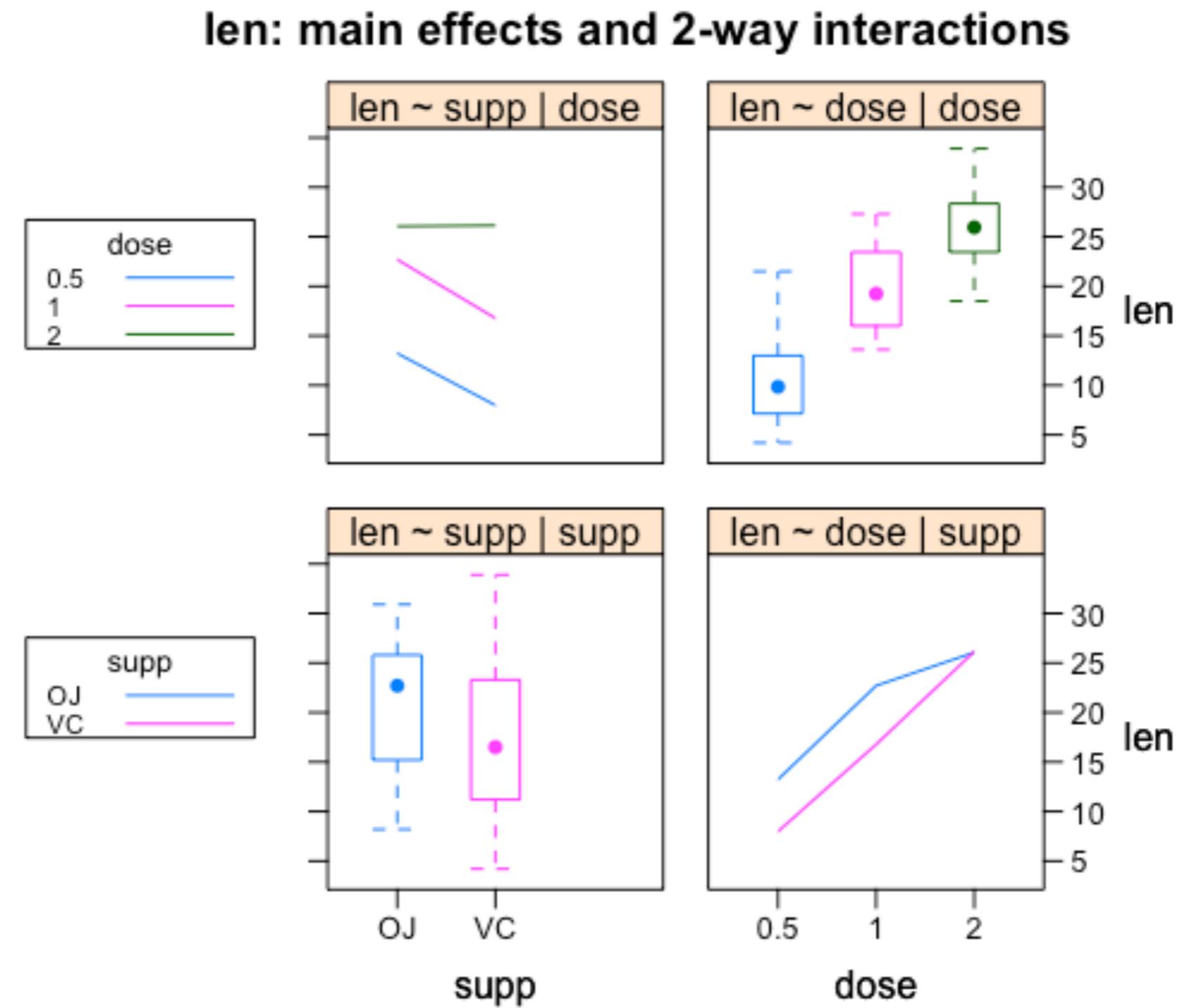
```
library(gplots)
plotmeans(len ~ interaction(supp, dose, sep = " "),
            connect = list(c(1, 3, 5), c(2, 4, 6)), col = c("red", "darkgreen"),
            main = "Interaction Plot with 95% CIs",
            xlab = "Treatment and Dose Combination")
```

交互效应



双因素方差分析例子

```
library(HH)  
interaction2wt(len ~ supp * dose)
```



重复测量方差分析

```
w1b1 <- subset(CO2, Treatment == "chilled")
fit <- aov(uptake ~ (conc * Type) + Error(Plant/(conc)),
w1b1)
summary(fit)
```

> summary(fit)

Error: Plant

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Type	1	2667.2	2667.2	60.41	0.00148 **
Residuals	4	176.6	44.1		

Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

Error: Plant:conc

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
conc	1	888.6	888.6	215.46	0.000125 ***
conc:Type	1	239.2	239.2	58.01	0.001595 **
Residuals	4	16.5	4.1		

Signif. codes: 0 ‘***’ 0.001 ‘**’ 0.01 ‘*’ 0.05 ‘.’ 0.1 ‘ ’ 1

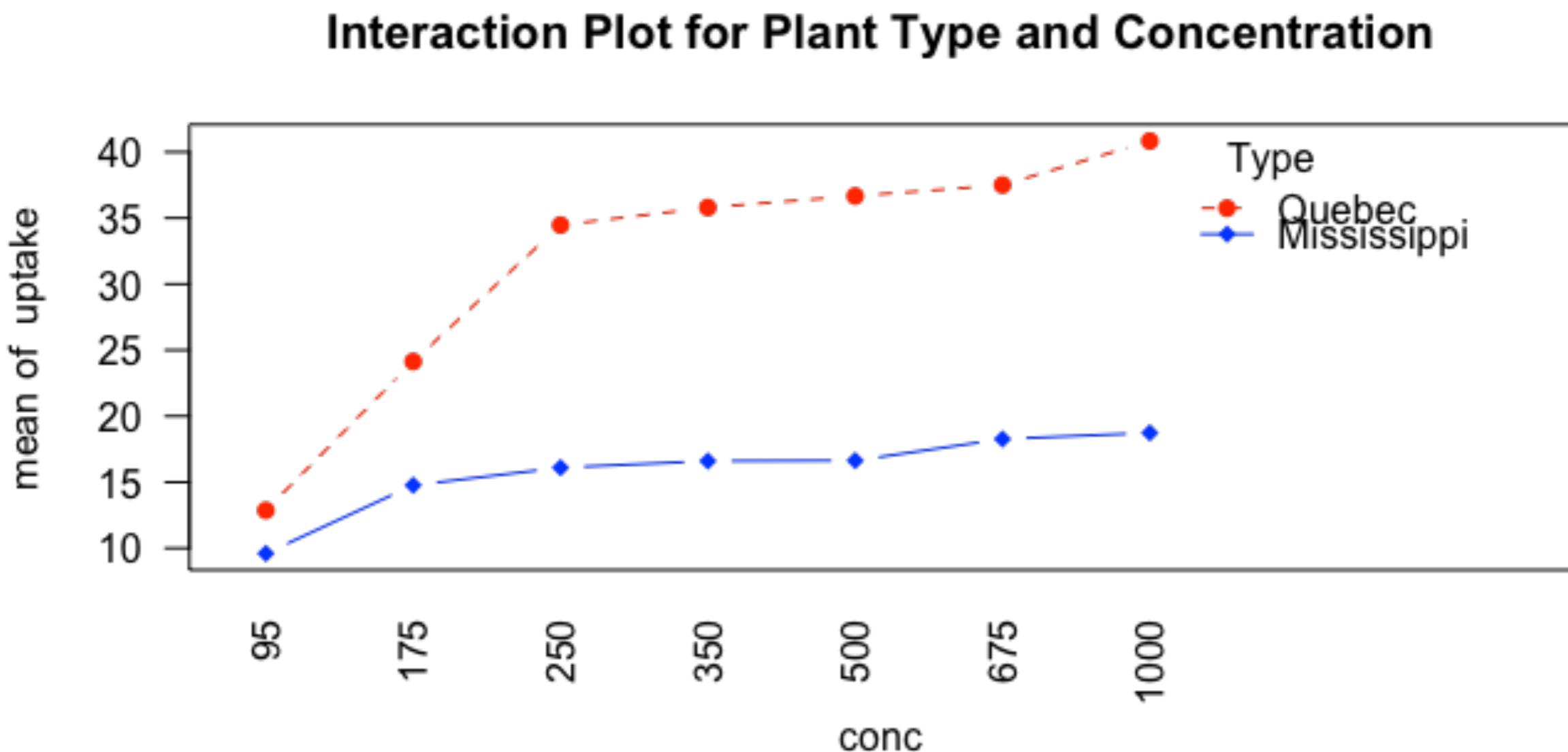
Error: Within

	Df	Sum Sq	Mean Sq	F value	Pr(>F)
Residuals	30	869	28.97		

.

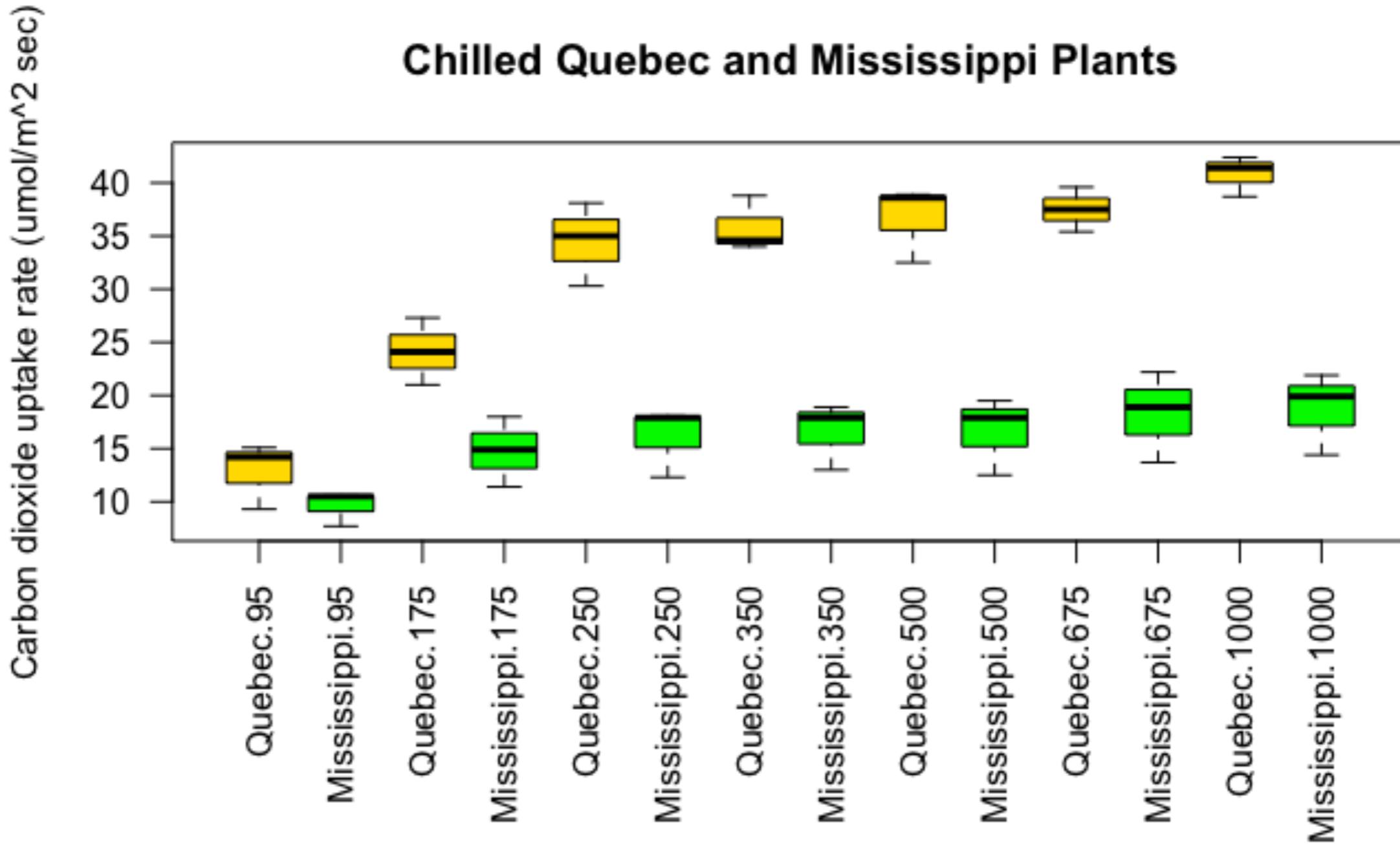
重复测量方差分析

```
par(las = 2)
par(mar = c(10, 4, 4, 2))
with(w1b1, interaction.plot(conc, Type, uptake, type = "b",
  col = c("red", "blue"), pch = c(16, 18),
  main = "Interaction Plot for Plant Type and Concentration"))
```



重复测量方差分析例子

```
boxplot(uptake ~ Type * conc, data = w1b1,  
        col = (c("gold", "green")),  
        main = "Chilled Quebec and Mississippi Plants",  
        ylab = "Carbon dioxide uptake rate (umol/m^2 sec)")
```



缺失值处理

缺失值处理方法

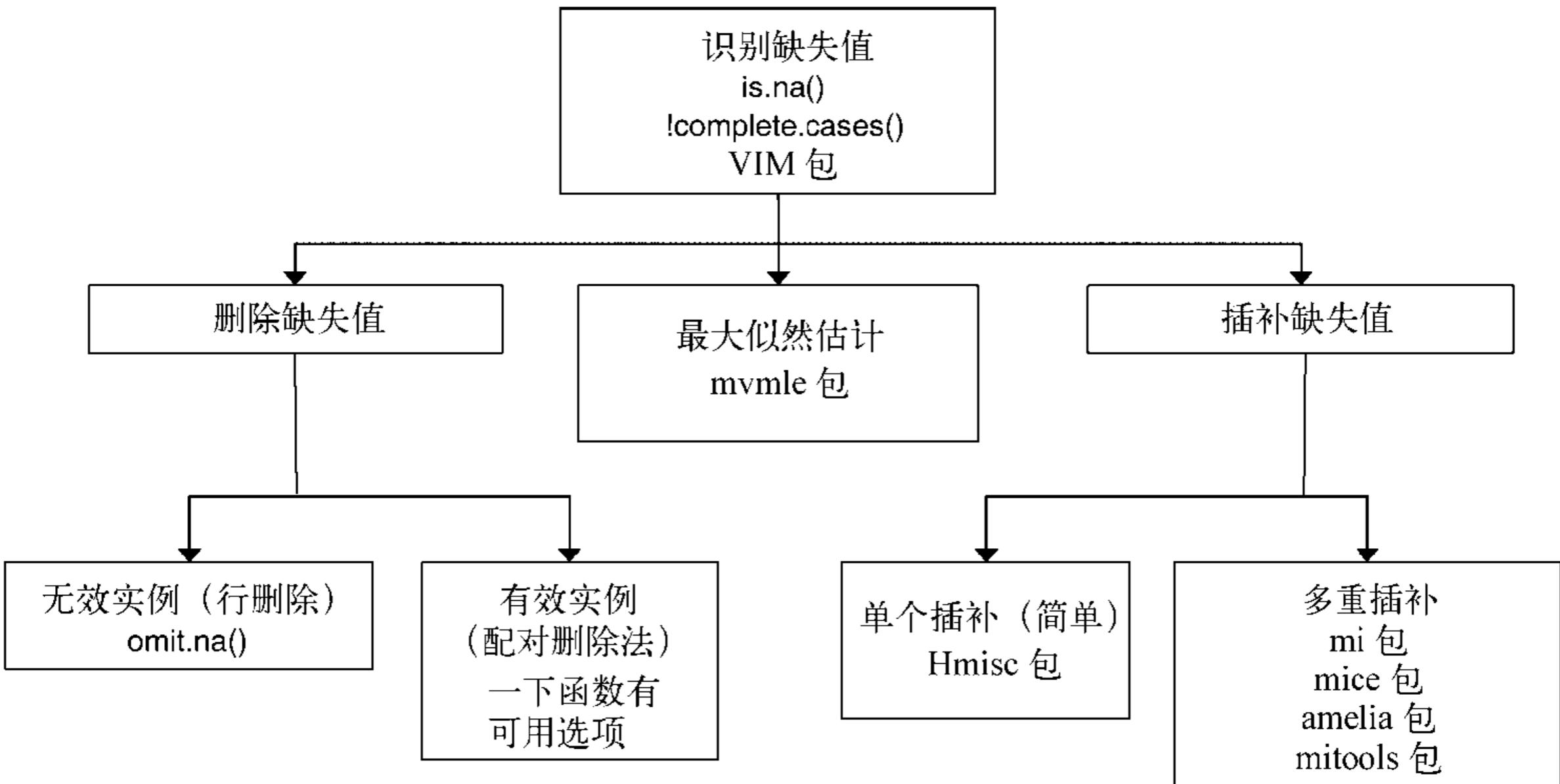


图18-1 处理不完整数据的方法，以及R中相关的包和函数

缺失值处理方法

表18-1 `is.na()`、`is.nan()`和`is.infinite()`函数的返回值示例

x	is.na(x)	is.nan(x)	is.infinite(x)
x <- NA	TRUE	FALSE	FALSE
x <- 0 / 0	TRUE	TRUE	FALSE
x <- 1 / 0	FALSE	FALSE	TRUE

`mice`包中的`md.pattern()`函数可生成一个以矩阵或数据框形式展示缺失值模式的表格。将函数应用到`sleep`数据集，可得到：

```
> library(mice)
> data(sleep, package="VIM")
> md.pattern(sleep)

  BodyWgt BrainWgt Pred Exp Danger Sleep Span Gest Dream NonD
42      1       1   1   1      1     1     1     1     1     1     1   0
  2      1       1   1   1      1     1     0     1     1     1     1   1
  3      1       1   1   1      1     1     1     0     1     1     1   1
  9      1       1   1   1      1     1     1     1     0     0     0   2
  2      1       1   1   1      1     0     1     1     1     0     0   2
  1      1       1   1   1      1     1     0     0     1     1     1   2
  2      1       1   1   1      1     0     1     1     0     0     0   3
  1      1       1   1   1      1     1     0     1     0     0     0   3
  0      0       0   0   0      0     4     4     4     12    14   38
```

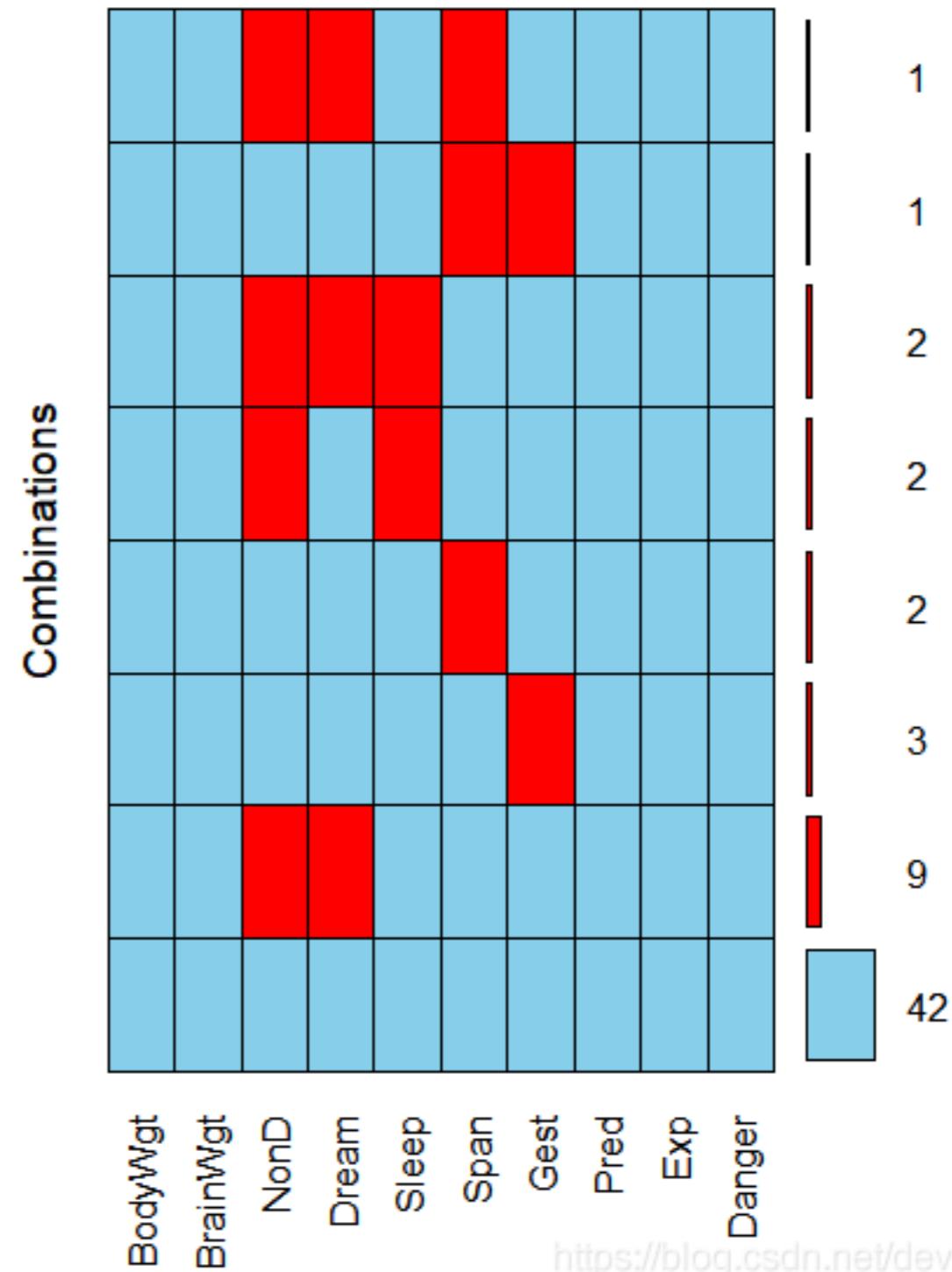
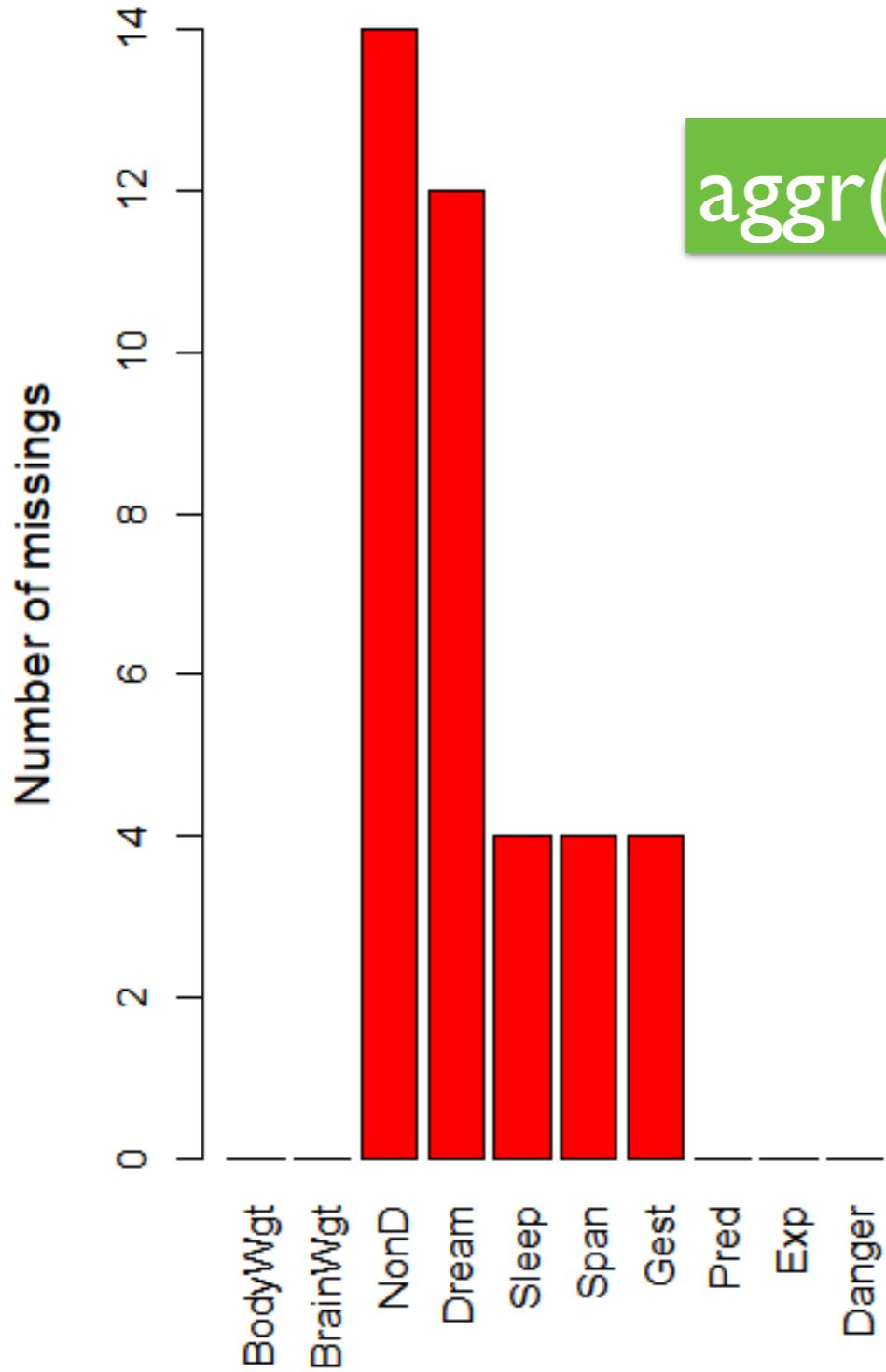
缺失值统计

mice包中的md.pattern()函数可生成一个以矩阵或数据框形式展示缺失值模式的表格。将函数应用到sleep数据集，可得到：

```
> library(mice)
> data(sleep, package="VIM")
> md.pattern(sleep)
```

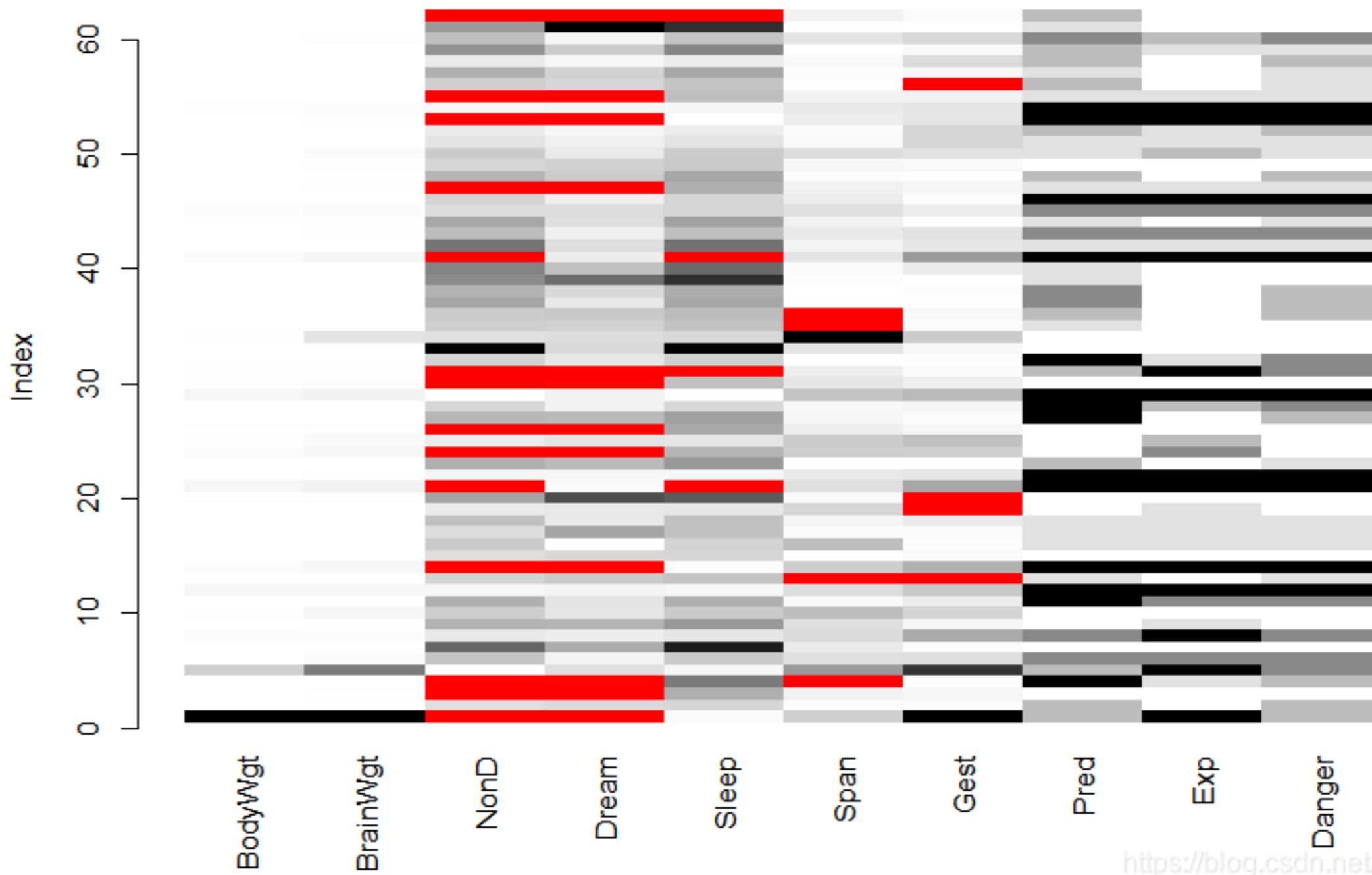
	BodyWgt	BrainWgt	Pred	Exp	Danger	Sleep	Span	Gest	Dream	NonD	
42	1	1	1	1	1	1	1	1	1	1	0
2	1	1	1	1	1	1	0	1	1	1	1
3	1	1	1	1	1	1	1	0	1	1	1
9	1	1	1	1	1	1	1	1	0	0	2
2	1	1	1	1	1	0	1	1	1	0	2
1	1	1	1	1	1	1	0	0	1	1	2
2	1	1	1	1	1	0	1	1	0	0	3
1	1	1	1	1	1	1	0	1	0	0	3
0	0	0	0	0	0	4	4	4	12	14	38

缺失值图形显示

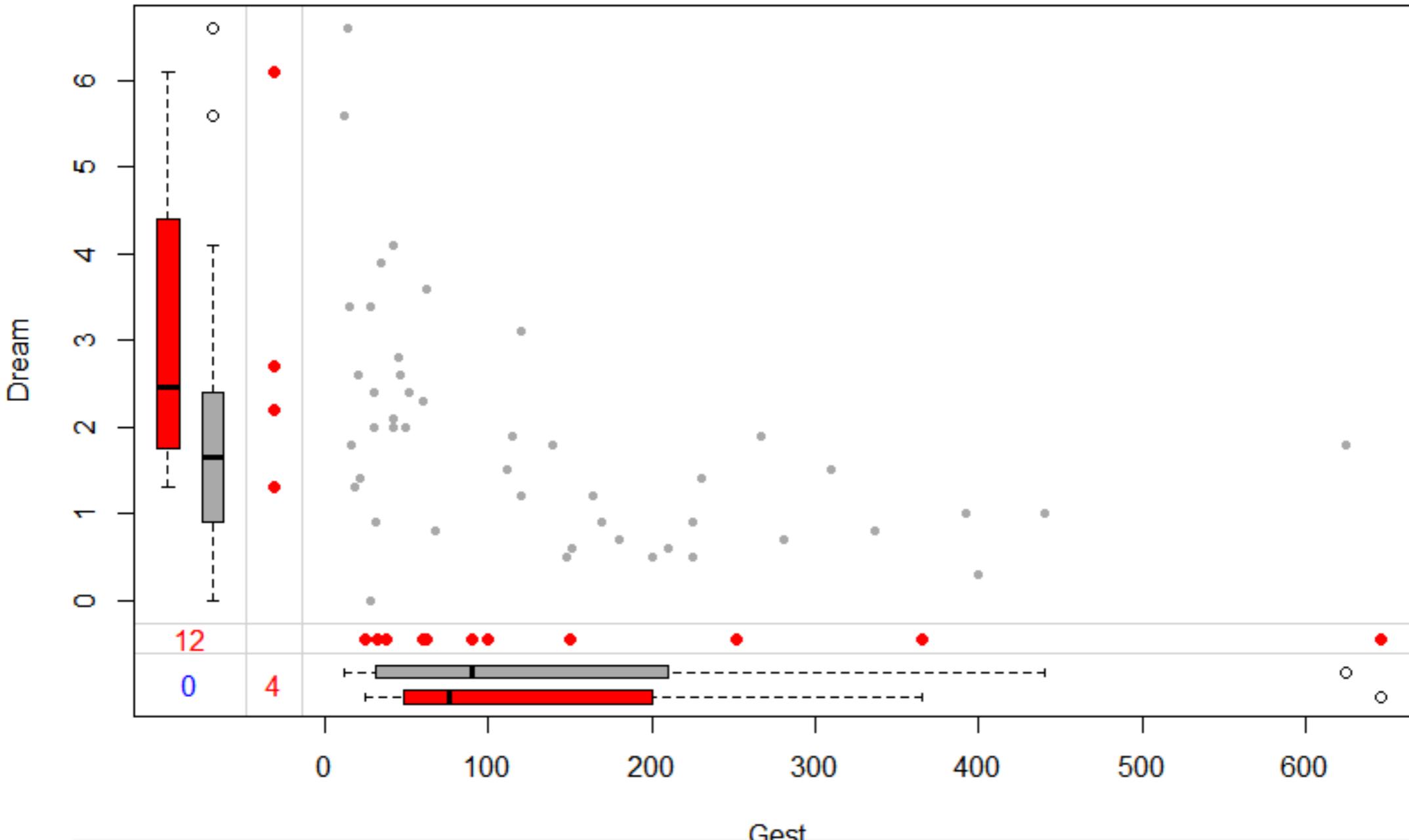


缺失值图形显示

matrixplot()



缺失值图形显示



```
marginplot(sleep[c("Gest", "Dream")], pch=c(20),  
          col=c("darkgray", "red", "blue"))
```

处理确实数据

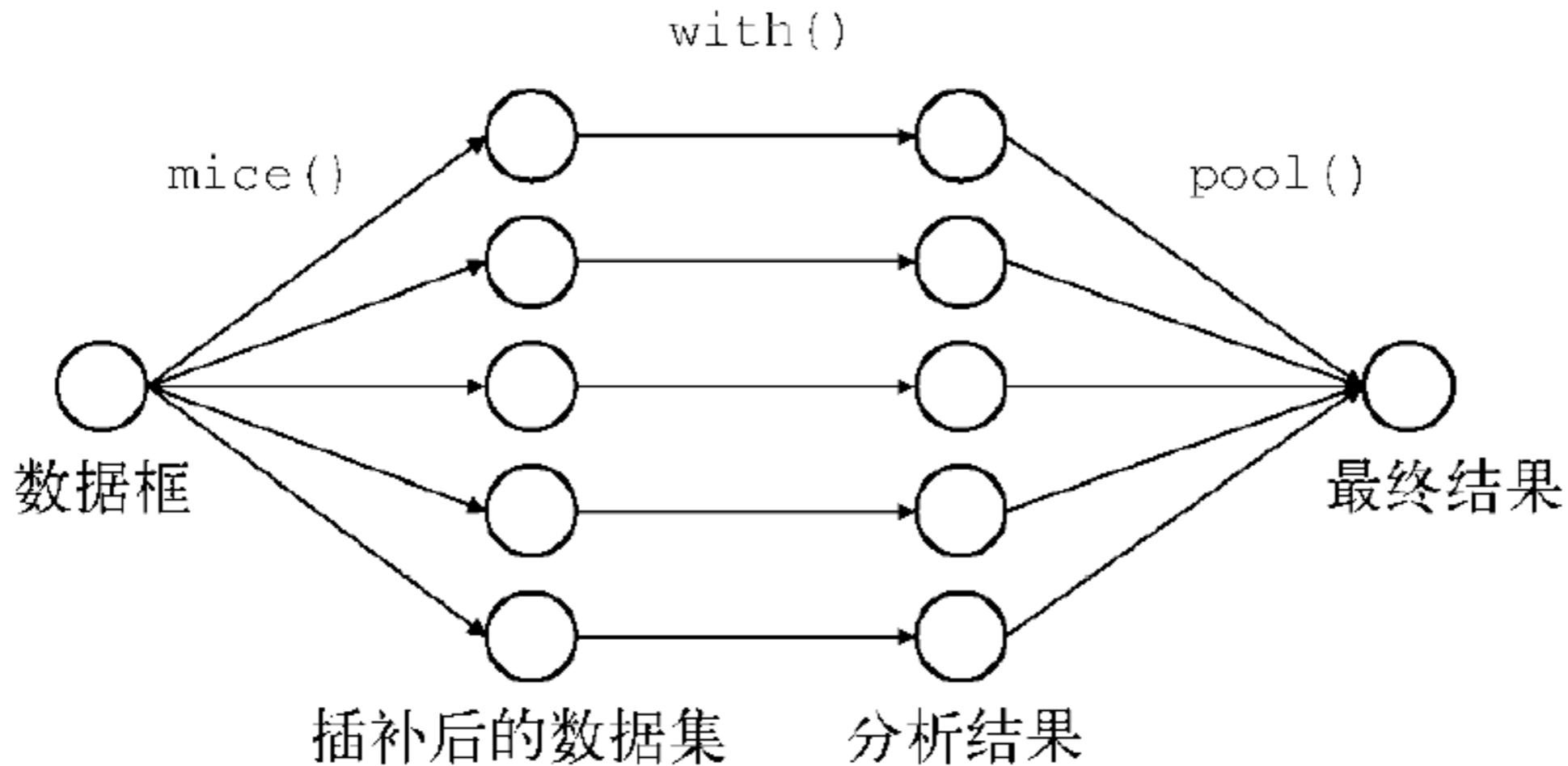


图18-5 通过mice包应用多重插补的步骤

完整行删除

多重插补

成对删除

简单插补

RCurl

Curl和RCurl

- ❑ curl：利用URL语法在命令行方式工作的开源文件传输工具
- ❑ libCurl：实现Curl的开源库，主要实现：获取页面、认证、上传、下载、信息搜索等

<https://curl.haxx.se/>



-
- ❑ RCurl：提供了R到libcurl的接口，从而实现了HTTP的一些功能。例如：服务器下载文件、保持连接、上传文件、采用二进制格式读取、句柄重定向、密码认证等

<http://anson.ucdavis.edu/~duncan/>



getURL()

```
> url.exists(url="www.baidu.com")
[1] TRUE
[1] "text"      "headerIn"   "headerOut"  "dataIn"      "dataOut"     "sslDataIn"
[7] "sslDataOut"

> cat(d$value()[1])
Rebuilt URL to: www.baidu.com/
  Trying 119.75.218.70...
Connected to www.baidu.com (119.75.218.70) port 80 (#0)
Connection #0 to host www.baidu.com left intact
```

getURL()

```
> cat(d$value()[2])
HTTP/1.1 200 OK
Date: Wed, 25 May 2016 02:58:38 GMT
Content-Type: text/html
Content-Length: 14613
Last-Modified: Wed, 03 Sep 2014 02:48:32 GMT
Connection: Keep-Alive
Vary: Accept-Encoding
Set-Cookie: BAIDUID=2CB1A15F480DC65312CC359A6DE5ADBC:FG=1; expires=Thu, 31-Dec-37 23:55:55 GMT
max-age=2147483647; path=/; domain=.baidu.com
Set-Cookie: BIDUPSID=2CB1A15F480DC65312CC359A6DE5ADBC; expires=Thu, 31-Dec-37 23:55:55 GMT; max-
-age=2147483647; path=/; domain=.baidu.com
Set-Cookie: PSTM=1464145118; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; d
omain=.baidu.com
P3P: CP=" OTI DSP COR IVA OUR IND COM "
Server: BWS/1.1
X-UA-Compatible: IE=Edge,chrome=1
Pragma: no-cache
Cache-control: no-cache
Accept-Ranges: bytes
```

getURL()

```
> cat(d$value()[3])
```

```
GET / HTTP/1.1
```

```
Host: www.baidu.com
```

```
Accept: */*
```

```
> d$reset()
```

```
> d$value()
```

text	headerIn	headerOut	dataIn	dataOut	sslDataIn	sslDataOut
""	""	""	""	""	""	""

getURL()

```
> h <- basicTextGatherer()
> txt <- getURL("http://www.baidu.com", headerfunction = h$update)
> names(h$value())
NULL
> h$value()
[1] "HTTP/1.1 200 OK\r\nDate: Wed, 25 May 2016 03:35:42 GMT\r\nContent-Type: text/html\r\nContent-Length: 14613\r\nLast-Modified: Wed, 03 Sep 2014 02:48:32 GMT\r\nConnection: Keep-Alive\r\nVary: Accept-Encoding\r\nSet-Cookie: BAIDUID=E077EAC981524636A14600D379007DBF:FG=1; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com\r\nSet-Cookie: BIDUPSID=E077EAC981524636A14600D379007DBF; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com\r\nSet-Cookie: PSTM=1464147342; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com\r\nP3P: CP=\" OTI DSP COR IVA OUR IND COM \"\r\nServer: BWS/1.1\r\nX-UA-Compatible: IE=Edge,chrome=1\r\nPragma: no-cache\r\nCache-control: no-cache\r\nAccept-Ranges: bytes\r\n\r\n"
> cat(h$value())
HTTP/1.1 200 OK
Date: Wed, 25 May 2016 03:35:42 GMT
Content-Type: text/html
Content-Length: 14613
Last-Modified: Wed, 03 Sep 2014 02:48:32 GMT
Connection: Keep-Alive
Vary: Accept-Encoding
Set-Cookie: BAIDUID=E077EAC981524636A14600D379007DBF:FG=1; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com
Set-Cookie: BIDUPSID=E077EAC981524636A14600D379007DBF; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com
Set-Cookie: PSTM=1464147342; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com
P3P: CP=" OTI DSP COR IVA OUR IND COM "
Server: BWS/1.1
X-UA-Compatible: IE=Edge,chrome=1
Pragma: no-cache
Cache-control: no-cache
Accept-Ranges: bytes
```

getURL()

```
> cat(h$value())
HTTP/1.1 200 OK
Date: Wed, 25 May 2016 03:35:42 GMT
Content-Type: text/html
Content-Length: 14613
Last-Modified: Wed, 03 Sep 2014 02:48:32 GMT
Connection: Keep-Alive
Vary: Accept-Encoding
Set-Cookie: BAIDUID=E077EAC981524636A14600D379007DBF:FG=1; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com
Set-Cookie: BIDUPSID=E077EAC981524636A14600D379007DBF; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com
Set-Cookie: PSTM=1464147342; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com
P3P: CP=" OTI DSP COR IVA OUR IND COM "
Server: BWS/1.1
X-UA-Compatible: IE=Edge,chrome=1
Pragma: no-cache
Cache-control: no-cache
Accept-Ranges: bytes
```

getURL()

```
> h <- basicHeaderGatherer()
> txt <- getURL(url="http://www.baidu.com", headerfunction = h$update)
> names(h$value())
[1] "Date"           "Content-Type"    "Content-Length"   "Last-Modified"   "Connection"
[6] "Vary"          "Set-Cookie"      "Set-Cookie"      "Set-Cookie"      "P3P"
[11] "Server"        "X-UA-Compatible" "Pragma"         "Cache-control"  "Accept-Ranges"
[16] "status"        "statusMessage" 

> h$value()
Date
"Wed, 25 May 2016 03:34:03 GMT"
Content-Type
"text/html"
Content-Length
"14613"
Last-Modified
"Wed, 03 Sep 2014 02:48:32 GMT"
Connection
"Keep-Alive"
Vary
"Accept-Encoding"
Set-Cookie
"BAIDUID=D7ACEDAEAA0295848111148DA18F535C;FG=1; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com"
Set-Cookie
"BIDUPSID=D7ACEDAEAA0295848111148DA18F535C; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com"
Set-Cookie
"PSTM=1464147243; expires=Thu, 31-Dec-37 23:55:55 GMT; max-age=2147483647; path=/; domain=.baidu.com"
P3P
"CP=\\" OTI DSP COR IVA OUR IND COM \\""
Server
"BWS/1.1"
X-UA-Compatible
"IE=Edge,chrome=1"
Pragma
"no-cache"
```

Curl参数

```
> listCurlOptions()
[1] "address.scope"
[4] "buffersize"
[7] "certinfo"
[10] "connecttimeout"
[13] "conv.from.utf8.function"
[16] "cookiefile"
[19] "cookiesession"
[22] "crlfile"
[25] "debugfunction"
[28] "dns.use.global.cache"
[31] "errorbuffer"
[34] "filetime"
[37] "fresh.connect"
[40] "ftp.create.missing.dirs"
[43] "ftp.skip.pasv.ip"
[46] "ftp.use.eprt"
[49] "ftplistonly"
[52] "header"
[55] "http.content.decoding"
[58] "http.timeout.ms"
[163] "unrestricted.auth"
[166] "use.ssl"
[169] "userpwd"
[172] "writefunction"

[1]      C U R L O P T I O N S
[4]      C U R L O P T I O N S
[7]      C U R L O P T I O N S
[10]     C U R L O P T I O N S
[13]     C U R L O P T I O N S
[16]     C U R L O P T I O N S
[19]     C U R L O P T I O N S
[22]     C U R L O P T I O N S
[25]     C U R L O P T I O N S
[28]     C U R L O P T I O N S
[31]     C U R L O P T I O N S
[34]     C U R L O P T I O N S
[37]     C U R L O P T I O N S
[40]     C U R L O P T I O N S
[43]     C U R L O P T I O N S
[46]     C U R L O P T I O N S
[49]     C U R L O P T I O N S
[52]     C U R L O P T I O N S
[55]     C U R L O P T I O N S
[58]     C U R L O P T I O N S
[163]    C U R L O P T I O N S
[166]    C U R L O P T I O N S
[169]    C U R L O P T I O N S
[172]    C U R L O P T I O N S

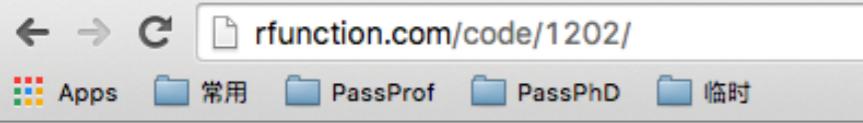
[1]      C U R L V A L U E
[4]      C U R L V A L U E
[7]      C U R L V A L U E
[10]     C U R L V A L U E
[13]     C U R L V A L U E
[16]     C U R L V A L U E
[19]     C U R L V A L U E
[22]     C U R L V A L U E
[25]     C U R L V A L U E
[28]     C U R L V A L U E
[31]     C U R L V A L U E
[34]     C U R L V A L U E
[37]     C U R L V A L U E
[40]     C U R L V A L U E
[43]     C U R L V A L U E
[46]     C U R L V A L U E
[49]     C U R L V A L U E
[52]     C U R L V A L U E
[55]     C U R L V A L U E
[58]     C U R L V A L U E
[163]    C U R L V A L U E
[166]    C U R L V A L U E
[169]    C U R L V A L U E
[172]    C U R L V A L U E

[1]      C U R L E X T E N D E R
[4]      C U R L E X T E N D E R
[7]      C U R L E X T E N D E R
[10]     C U R L E X T E N D E R
[13]     C U R L E X T E N D E R
[16]     C U R L E X T E N D E R
[19]     C U R L E X T E N D E R
[22]     C U R L E X T E N D E R
[25]     C U R L E X T E N D E R
[28]     C U R L E X T E N D E R
[31]     C U R L E X T E N D E R
[34]     C U R L E X T E N D E R
[37]     C U R L E X T E N D E R
[40]     C U R L E X T E N D E R
[43]     C U R L E X T E N D E R
[46]     C U R L E X T E N D E R
[49]     C U R L E X T E N D E R
[52]     C U R L E X T E N D E R
[55]     C U R L E X T E N D E R
[58]     C U R L E X T E N D E R
[163]    C U R L E X T E N D E R
[166]    C U R L E X T E N D E R
[169]    C U R L E X T E N D E R
[172]    C U R L E X T E N D E R
```

其余

- `getForm()`
- `postForm()`

批量下载



Index of /code/1202

- [Parent Directory](#)
- [120201.R](#)
- [120202.R](#)
- [120203.R](#)
- [120204.R](#)
- [120205.R](#)
- [120206.R](#)
- [120207.R](#)
- [120208.R](#)
- [120209.R](#)
- [120210.R](#)
- [120211.R](#)
- [120212.R](#)
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- [120225.R](#)
- [120226.R](#)
- [120227.R](#)
- [120228.R](#)
- [120229.R](#)
- [BarackObamaTweets.txt](#)
- [data1.txt](#)
- [par-120208.pdf](#)

```
html=getURL("http://rfunction.com/code/1202/")
temp =strsplit(html, "<li><a href=\"\">[[1]]
files =strsplit(temp, "\"")
files=lapply(files, function(x){ x[1] })
files= unlist(files)

files = files[-(1:2)]

base="http://rfunction.com/code/1202/"
for(i in 1:length(files)){
  URL = paste(base, files[i], sep="")
  bin <- getBinaryURL(URL)
  con <- file(paste("1202", files[i], sep="."))
  writeBin(bin, con)
  close(con)
  Sys.sleep(2)
}
```

使用XML

The screenshot shows the homepage of the AnimalTFDB. At the top, there's a logo with a DNA helix and a fish, followed by the text "AnimalTFDB" and "Animal Transcription Factor Database". Below the header, there are navigation links: HOME, BROWSE FAMILY, BROWSE SPECIES, SEARCH, and DOWNLOAD. A sub-header "Mus musculus" is displayed. The main content is a table with 11 rows, each representing a gene entry. The columns are labeled: No., Ensembl ID, Gene ID, Symbol, and Family. The data includes entries for genes like Aff1, Aff2, Aff3, Aff4, Ar, Tcfap2a, Tcfap2b, Tcfap2c, Tcfap2e, Tcfap2d, and Arid3b, categorized under AF-4, Androgen receptor, and AP-2 families.

No.	Ensembl ID	Gene ID	Symbol	Family
1	ENSMUSG00000029313	17355	Aff1	AF-4
2	ENSMUSG00000031189	14266	Aff2	AF-4
3	ENSMUSG00000037138	16764	Aff3	AF-4
4	ENSMUSG00000049470	93736	Aff4	AF-4
5	ENSMUSG00000046532	11835	Ar	Androgen receptor
6	ENSMUSG00000021359	21418	Tcfap2a	AP-2
7	ENSMUSG00000025927	21419	Tcfap2b	AP-2
8	ENSMUSG00000028640	21420	Tcfap2c	AP-2
9	ENSMUSG00000042477	332937	Tcfap2e	AP-2
10	ENSMUSG00000042596	226896	Tcfap2d	AP-2
11	ENSMUSG00000004661	56380	Arid3b	ARID

> head(tables)

No.		Ensembl ID	Gene ID	Symbol	Family
1	1	ENSMUSG00000029313	17355	Aff1	AF-4
2	2	ENSMUSG00000031189	14266	Aff2	AF-4
3	3	ENSMUSG00000037138	16764	Aff3	AF-4
4	4	ENSMUSG00000049470	93736	Aff4	AF-4
5	5	ENSMUSG00000046532	11835	Ar	Androgen receptor
6	6	ENSMUSG00000021359	21418	Tcfap2a	AP-2

```
url="http://www.bioguo.org/AnimalTFDB/BrowseAllTF.php?spe=Mus_musculus"
wp <- getURL(url)
doc <- htmlParse(wp, asText= TRUE)
tables <- readHTMLTable(doc,which=5)
```

其余

- 使用正则表达式来匹配文本

https://en.wikipedia.org/wiki/Regular_expression

- 其余包

→ Rvest

→ httr

HTTP: https://en.wikipedia.org/wiki/Hypertext_Transfer_Protocol

- 天眼查每个企业均有详细信息和企业评分
- 写一个爬虫，下载1000个企业信息
- 根据企业信息项，分析预测天眼查现有企业评分的计算方法，用下载的数据进行回归分析，再爬一些企业作为测试数据，分析方法的有效性
- 自己设计一个企业评分算法，说明设计原理，并用数据检验

企业最好针对一个行业

法定代表人	季昊 任职 8 家企业，分布如下 北京 (共8家) 北京国际大数据交易... 等	经营状态	存续	天眼评分	评分 68 
		成立日期	2020-04-29		
		注册资本 ②	625万人民币		
		实缴资本	-		
统一社会信用代码 ②	91110108MA01R2FT36	纳税人识别号 ②	91110108MA01R2FT36	工商注册号	110108028700442
营业期限	2020-04-29 至 2070-04-28	纳税人资质	-	组织机构代码 ②	MA01R2FT3
公司类型	其他有限责任公司	行业	软件和信息技术服务业	核准日期	2021-01-06
参保人数	-	登记机关	北京市海淀区市场监督管理局	人员规模	-
用户名	-	英文名称	-		
地址 ②	北京市海淀区丹棱街1号院1号楼26层2601室 附近公司				

下周上课前提交
方式和以前一样

谢谢！

孙惠平

sunhp@ss.pku.edu.cn