

The_apply_family.R

015004

Thu Feb 21 13:44:41 2019

```
#####
# Load packages #
#####

#install.packages("JM")
library(JM)

## Warning: package 'JM' was built under R version 3.5.2
## Loading required package: MASS
## Loading required package: nlme
## Loading required package: splines
## Loading required package: survival
## Warning: package 'survival' was built under R version 3.5.2
#####
# The apply family #
#####

## apply
apply(pbc2.id[, c(2,5)], 2, mean)

##      years      age
## 6.411239 50.020376
apply(pbc2.id[, c("years", "age")], 2, mean)

##      years      age
## 6.411239 50.020376
X <- sample(0:200, 100)
Mat <- matrix(X, 50, 50)
apply(Mat, 1, mean)

## [1] 34.5 76.5 160.0 134.0 101.5 107.0 154.0 149.0 68.5 95.5 114.5
## [12] 126.0 169.0 151.5 73.0 136.5 174.5 127.5 28.5 175.5 168.0 94.0
## [23] 73.5 153.5 109.5 128.0 36.0 83.0 151.0 195.5 98.5 85.5 86.5
## [34] 94.0 38.5 76.5 119.0 83.0 50.5 136.5 152.5 12.5 101.0 144.0
## [45] 94.0 8.5 90.5 93.0 25.5 45.5
apply(Mat, 2, mean)

## [1] 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90
## [11] 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90
## [21] 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90
## [31] 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90
## [41] 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90 110.48 96.90
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```
apply(Mat, 2, function(x) x^2)
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##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11]
## [1,] 3600   81 3600   81 3600   81 3600   81 3600   81 3600
## [2,] 20449  100 20449  100 20449  100 20449  100 20449  100 20449
## [3,] 23409 27889 23409 27889 23409 27889 23409 27889 23409 27889 23409
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## [11,] 13689 12544 13689 12544 13689 12544 13689 12544 13689 12544 13689
## [12,]  6084 30276  6084 30276  6084 30276  6084 30276  6084 30276  6084
## [13,] 39204 19600 39204 19600 39204 19600 39204 19600 39204 19600 39204
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## [15,]  8649  2809  8649  2809  8649  2809  8649  2809  8649  2809  8649
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## [19,]    4  3025    4  3025    4  3025    4  3025    4  3025    4
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## [27,]   576  2304   576  2304   576  2304   576  2304   576  2304   576
## [28,] 19044   784 19044   784 19044   784 19044   784 19044   784 19044
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## [32,]   676 21025   676 21025   676 21025   676 21025   676 21025   676
## [33,]  1936 16641  1936 16641  1936 16641  1936 16641  1936 16641  1936
## [34,]   289 29241   289 29241   289 29241   289 29241   289 29241   289
## [35,]    36  5041    36  5041    36  5041    36  5041    36  5041    36
## [36,]  5476  6241  5476  6241  5476  6241  5476  6241  5476  6241  5476
## [37,] 27556  5184 27556  5184 27556  5184 27556  5184 27556  5184 27556
## [38,] 23716   144 23716   144 23716   144 23716   144 23716   144 23716
## [39,]  3844  1521  3844  1521  3844  1521  3844  1521  3844  1521  3844
## [40,] 32761  8464 32761  8464 32761  8464 32761  8464 32761  8464 32761
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## [42,]    0  625    0  625    0  625    0  625    0  625    0
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## [45,] 24964   900 24964   900 24964   900 24964   900 24964   900 24964
## [46,]    9  196    9  196    9  196    9  196    9  196    9
## [47,] 17424  2401 17424  2401 17424  2401 17424  2401 17424  2401 17424
## [48,] 28224   324 28224   324 28224   324 28224   324 28224   324 28224
## [49,]   225  1296   225  1296   225  1296   225  1296   225  1296   225
## [50,]   441  4900   441  4900   441  4900   441  4900   441  4900   441
##      [,12] [,13] [,14] [,15] [,16] [,17] [,18] [,19] [,20] [,21] [,22]
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## [48,] 28224 324 28224 324 28224 324
## [49,] 225 1296 225 1296 225 1296
## [50,] 441 4900 441 4900 441 4900
```

```
## lapply
```

```
lapply(1:3, function(x) x^2)
```

```
## [[1]]
## [1] 1
##
## [[2]]
## [1] 4
##
## [[3]]
## [1] 9
```

```
X <- list(Mat, Mat^2)
lapply(X, mean)
```

```
## [[1]]
## [1] 103.69
##
## [[2]]
## [1] 14233.07
```

```
A <- matrix(1:9, 3,3)
B <- matrix(4:15, 4,3)
C <- matrix(8:10, 3,2)
MyList <- list(A,B,C)
lapply(MyList,"[, 1, )
```

```
## [[1]]
## [1] 1 4 7
##
## [[2]]
## [1] 4 8 12
##
## [[3]]
## [1] 8 8
```

```
lapply(MyList,"[, , 2)
```

```
## [[1]]
## [1] 4 5 6
##
## [[2]]
## [1] 8 9 10 11
##
## [[3]]
## [1] 8 9 10
```

```
## sapply
```

```
sapply(1:3, function(x) x^2)
```

```
## [1] 1 4 9
```

```
X <- list(Mat, Mat^2)
sapply(X, mean)
```

```
## [1] 103.69 14233.07
```

```
sapply(MyList,"[, 2, 1)
```

```
## [1] 2 5 9
```

```
## tapply
```

```
tapply(pbc2.id$age, pbc2.id$sex, mean)
```

```
##      male      female
## 56.20346 49.21389
```

```
tapply(pbc2.id$years, pbc2.id$sex, mean)
```

```

##      male      female
## 5.796721 6.491394

tapply(pbc2.id$age, pbc2.id$sex, function(x) mean(x/2))

##      male      female
## 28.10173 24.60694

## mapply

mapply(rep, 1:4, 4:1)

## [[1]]
## [1] 1 1 1 1
##
## [[2]]
## [1] 2 2 2
##
## [[3]]
## [1] 3 3
##
## [[4]]
## [1] 4

#### list(rep(1, 4), rep(2, 3), rep(3, 2), rep(4, 1))

mapply(rep, times = 1:4, x = 4)

## [[1]]
## [1] 4
##
## [[2]]
## [1] 4 4
##
## [[3]]
## [1] 4 4 4
##
## [[4]]
## [1] 4 4 4 4

#### list(rep(4, times = 1), rep(4, times = 2), rep(4, times = 3), rep(4, times = 4))

X = matrix(c(rep(1, 4), rep(2, 4), rep(3, 4), rep(4, 4)), 4, 4)
mapply(rep, 1:4, 4)

##      [,1] [,2] [,3] [,4]
## [1,]    1    2    3    4
## [2,]    1    2    3    4
## [3,]    1    2    3    4
## [4,]    1    2    3    4

mapply(function(x,y) seq_len(x) + y,
        c(a = 1, b = 2, c = 3),
        c(A = 10, B = 0, C = -10))

## $a
## [1] 11
##

```



```

## $b
## [1] 1 2
##
## $c
## [1] -9 -8 -7

#### list(c(1) + 10, c(1, 2) + 0, c(1, 2, 3) - 10)

X <- list(Mat, Mat^2)
mapply(mean, X)

## [1] 103.69 14233.07

mapply(mean, MyList)

## [1] 5.0 9.5 9.0

sapply(MyList, mean)

## [1] 5.0 9.5 9.0

mapply(function(x,y) {x^y}, x = c(2, 3), y = c(4))

## [1] 16 81

#### list(2^4, 3^4)

#####
# Long format data #
#####

## Let's assume that only the long format of the data set is available.
## We want to obtain the mean serum bilirubin of the last follow-up
## measurements per event group.
## Each patient is counted once!
head(pbc2)

##   id   years status   drug   age   sex   year ascites
## 1  1  1.09517   dead D-penicil 58.76684 female 0.0000000    Yes
## 2  1  1.09517   dead D-penicil 58.76684 female 0.5256817    Yes
## 3  2 14.15234   alive D-penicil 56.44782 female 0.0000000     No
## 4  2 14.15234   alive D-penicil 56.44782 female 0.4983025     No
## 5  2 14.15234   alive D-penicil 56.44782 female 0.9993429     No
## 6  2 14.15234   alive D-penicil 56.44782 female 2.1027270     No
##   hepatomegaly spiders           edema serBilir serChol albumin
## 1           Yes      Yes edema despite diuretics    14.5    261    2.60
## 2           Yes      Yes edema despite diuretics    21.3     NA    2.94
## 3           Yes      Yes           No edema         1.1    302    4.14
## 4           Yes      Yes           No edema         0.8     NA    3.60
## 5           Yes      Yes           No edema         1.0     NA    3.55
## 6           Yes      Yes           No edema         1.9     NA    3.92
##   alkaline  SGOT platelets prothrombin histologic status2
## 1    1718 138.0      190      12.2         4         1
## 2    1612   6.2      183      11.2         4         1
## 3    7395 113.5      221      10.6         3         0
## 4    2107 139.5      188      11.0         3         0

```

```

## 5      1711 144.2      161      11.6      3      0
## 6      1365 144.2      122      10.6      3      0

## sort data
pbc2 <- pbc2[order(pbc2$id, pbc2$year), ]

## select the last follow-up measurement of each patient
pbc2.idNEW2 <- pbc2[!duplicated(pbc2[c("id")], fromLast = T), ]

## obtain the mean serum bilirubin per event group
tapply(pbc2.idNEW2$serBilir, pbc2.idNEW2$status, mean)

##           alive transplanted           dead
##      2.190909      9.172414      11.800000

## Let's assume that we want to obtain the mean serum bilirubin
## of the last stage of edema per event group.
## Each patient and edema stage is counted once!

## sort data
pbc2 <- pbc2[order(pbc2$id, pbc2$edema), ]

## select the last stage of edema of each patient
pbc2.idNEW3 <- pbc2[!duplicated(pbc2[c("id")], fromLast = T), ]

## obtain the mean serum bilirubin per event group
tapply(pbc2.idNEW3$serBilir, pbc2.idNEW3$status, mean)

##           alive transplanted           dead
##      2.182517      8.651724      11.425000

```