```
data(BP1)
X < -BP1[,-1]
y < -BP1[,1]
library(mctest)
mctest(x, y, type=c("b"), na.rm = TRUE,
                Inter=TRUE, method=NULL, corr=TRUE, detr=0.01, red=0.5,
                theil=0.5, cn=30, vif=5, tol=0.1, conf=0.95, cvif=10,
                leamer=0.1, all=FALSE)
eigprop(x)
Overall Multicollinearity Diagnostics
                       MC Results detection
Determinant |X'X|:
                           0.0307
Farrar Chi-Square:
                          56.2967
                                           1
Red Indicator:
                           0.4424
                                           0
                          22.9943
Sum of Lambda Inverse:
                                           0
Theil's Method:
                          -1.4343
                                           0
                         201.4958
Condition Number:
1 --> COLLINEARITY is detected by the test
0 --> COLLINEARITY is not detected by the test
 All Individual Multicollinearity Diagnostics Result
            VIF
                   TOL
                            Wi
                                    Fi Leamer
                                                  CVIF Klein
         1.7628 0.5673 2.1359 2.8605 0.7532 -0.0039
Age
                                                           0
         8.4170 0.1188 20.7677 27.8139 0.3447 -0.0188
                                                           0
Weight
         5.3288 0.1877 12.1205 16.2328 0.4332 -0.0119
                                                           0
BSA
Duration 1.2373 0.8082 0.6645 0.8899 0.8990 -0.0028 Pulse 4.4136 0.2266 9.5580 12.8009 0.4760 -0.0099
                                                           0
                                                           0
Stress
         1.8348 0.5450 2.3376 3.1307 0.7382 -0.0041
1 --> COLLINEARITY is detected by the test
0 --> COLLINEARITY is not detected by the test
Duration , Pulse , Stress , coefficient(s) are non-significant may be due
to multicollinearity
R-square of y on all x: 0.9962
* use method argument to check which regressors may be the reason of colli
nearity
```

```
call:
eigprop(x = x)
  Eigenvalues
                                                 BSA Duration Pulse Stre
                    CI Intercept
                                   Age Weight
SS
                          0.0000 0.0000 0.0000 0.0000
                                                        0.0016 0.0000 0.00
       6.6558
                1.0000
1
30
               4.9842
                          0.0002 0.0001 0.0000 0.0001
                                                        0.0000 0.0000 0.55
2
       0.2679
12
                          0.0005 0.0004 0.0001 0.0003
                                                        0.9285 0.0002 0.05
       0.0714
              9.6536
3
24
                          0.1027 0.0890 0.0052 0.1698
                                                        0.0007 0.0040 0.02
4
       0.0027 50.0668
54
5
       0.0011 77.3965
                          0.3723 0.7775 0.0092 0.0113
                                                        0.0259 0.0050 0.03
91
                          0.2900 0.0381 0.0061 0.0736
       0.0009 83.7514
                                                        0.0432 0.4376 0.14
6
33
                          0.2343 0.0949 0.9794 0.7450
                                                        0.0000 0.5532 0.18
       0.0002 201.4958
7
56
```

\_\_\_\_\_

Row 5==> Age, proportion 0.777502 >= 0.50

Row 7==> Weight, proportion 0.979397 >= 0.50

Row 7==> BSA, proportion 0.744958 >= 0.50

Row 3==> Duration, proportion 0.928489 >= 0.50

Row 7==> Pulse, proportion 0.553205 >= 0.50

Row 2==> Stress, proportion 0.551209 >= 0.50