## betaMC: Staging

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Staging...

```
object <- lm(QUALITY ~ NARTIC + PCTGRT + PCTSUPP, data = nas1982)
```

```
dif(BetaMC(object))
#> Difference between standardized regression coefficients with type = " HC3 "
                 est se R 0.05%
                                             0.5% 2.5% 97.5% 99.5% 99.95%
#> NARTIC-PCTGRT 0.1037 0.1435 20000 -0.3736 -0.2704 -0.1805 0.3768 0.4654 0.5537
#> NARTIC-PCTSUPP 0.2319 0.1335 20000 -0.2192 -0.1274 -0.0409 0.4821 0.5645 0.6514
#> PCTGRT-PCTSUPP 0.1282 0.1368 20000 -0.3172 -0.2255 -0.1448 0.3895 0.4691 0.5658
dif(BetaMC(object, type = "mvn"))
#> Difference between standardized regression coefficients with type = " MVN "
                         se
                                  R 0.05%
                                               0.5%
                                                       2.5% 97.5% 99.5% 99.95%
                    est
#> NARTIC-PCTGRT 0.1037 0.1352 20000 -0.3629 -0.2574 -0.1653 0.3604 0.4399 0.5225
#> NARTIC-PCTSUPP 0.2319 0.1257 20000 -0.1858 -0.0997 -0.0250 0.4677 0.5416 0.6222
#> PCTGRT-PCTSUPP 0.1282 0.1215 20000 -0.2641 -0.1920 -0.1172 0.3622 0.4402 0.5201
dif(BetaMC(object, type = "adf"))
#> Difference between standardized regression coefficients with type = " ADF "
                                                      2.5% 97.5% 99.5% 99.95%
                    est se
                                 R 0.05%
                                             0.5%
#> NARTIC-PCTGRT 0.1037 0.1215 20000 -0.2828 -0.2070 -0.1337 0.3395 0.4146 0.4924
#> NARTIC-PCTSUPP 0.2319 0.1180 20000 -0.1810 -0.0766 -0.0056 0.4581 0.5342 0.6137
#> PCTGRT-PCTSUPP 0.1282 0.1213 20000 -0.2848 -0.1939 -0.1176 0.3623 0.4320 0.5128
out <- rsq(BetaMC(object))</pre>
print(out)
#> Multiple correlation
#> type = "HC3"
#>
                        R 0.05% 0.5% 2.5% 97.5% 99.5% 99.95%
         est
                 se
#> rsq 0.8045 0.0619 20000 0.4969 0.5772 0.6461 0.8881 0.9109 0.9327
#> adj 0.7906 0.0663 20000 0.4609 0.5470 0.6209 0.8801 0.9045 0.9279
```

```
summary(out)
#> Multiple correlation
#> type = "HC3"
#> est se
                     R 0.05% 0.5% 2.5% 97.5% 99.5% 99.95%
#> rsq 0.8045 0.0619 20000 0.4969 0.5772 0.6461 0.8881 0.9109 0.9327
#> adj 0.7906 0.0663 20000 0.4609 0.5470 0.6209 0.8801 0.9045 0.9279
coef(out)
      rsq
                 adj
#> 0.8045263 0.7905638
vcov(out)
             rsq
#> rsq 0.003826412 0.004099727
#> adj 0.004099727 0.004392564
confint(out)
           2.5%
                97.5%
#> rsq 0.6461316 0.8881363
#> adj 0.6208553 0.8801461
rsq(BetaMC(object, type = "mvn"))
#> Multiple correlation
#> type = "MVN"
#> est
                se R 0.05% 0.5% 2.5% 97.5% 99.5% 99.95%
#> rsq 0.8045 0.0567 20000 0.4984 0.5973 0.6601 0.8806 0.9029 0.9286
#> adj 0.7906 0.0607 20000 0.4626 0.5685 0.6358 0.8721 0.8960 0.9235
rsq(BetaMC(object, type = "adf"))
#> Multiple correlation
#> type = "ADF"
        est
                       R 0.05% 0.5% 2.5% 97.5% 99.5% 99.95%
               se
#> rsq 0.8045 0.0550 20000 0.5129 0.6102 0.6637 0.8799 0.9025 0.9297
#> adj 0.7906 0.0589 20000 0.4781 0.5824 0.6396 0.8713 0.8955 0.9247
```

## References

R Core Team. (2022). R: A language and environment for statistical computing. R Foundation for Statistical Computing. Vienna, Austria. https://www.R-project.org/