
Subject: hypertension other biomarkers doesn't exist error

Posted by [krittika03](#) on Sun, 03 Mar 2024 01:54:13 GMT

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hello, i am trying into the women's file (IR) and want to look into how education can impact health behaviors. since my R is not able to load the whole dataset, I had to select columns to import. the hypertension, cancer or other similar biomarkers are coming as doesn't exist in R. (cc below)

```
DHSIR <- import("D:/Thesis 101/data/Individual Recode/IAIR7EFL.DTA", col_select = c("v106",  
"s116", "v025", "v101", "v130", "v190", "s728c", "v463a"))
```

```
Error in `skip_cols()`:
```

```
! Can't subset columns that don't exist.
```

Backtrace:

1. rio::import(...)
3. rio:::import.rio_dta(file = file, ...)
5. rio:::docall(haven::read_dta, ..., args = list(file = file))
9. haven (local) ``<fn>`` (col_select = ``<chr>``, file = ``<rio_dta>``)
10. haven:::skip_cols(...)

>

what is happening? is there any other way? am i doing something wrong?
