

GENETICS

NHANES R DATA PARSER

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[NHANES](#) is a great resource for doing epidemiological research. As the NIH website provides only data import for commercial software here is my rewrite in [R](#). First load from their site

[adult.exe](#)

[youth.exe](#)

[lab.exe](#)

[lab2.exe](#)

[exam.exe](#)

put everything in one directory and expand the self-extracting archives. Then create from each SAS file a new variable content file that will only contain variable name and tab separated start position in the .dat file. Adult.var for example would read like this:

```
SEQN 1
DMPFSEQ 6
DMPSTAT 11
DMARETHN 12
DMARACER 13
...
HAZN0K5R 3345
```

Then start the following R job with the datasets and variables that you are interested in

```
|wj_nhanes.R|
```

