```
R version 3.6.3 (2020-02-29) -- "Holding the Windsock"
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Platform: x86_64-w64-mingw32/x64 (64-bit)

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  Natural language support but running in an English locale

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Type 'contributors()' for more information and
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Type 'demo()' for some demos, 'help()' for on-line help, or
'help.start()' for an HTML browser interface to help.
Type 'q()' to quit R.

> library(memisc)
Loading required package: lattice
Loading required package: MASS

Attaching package: 'memisc'

The following objects are masked from 'package:stats':

    contr.sum, contr.treatment, contrasts

The following object is masked from 'package:base':

    as.array

> # options(error=recover)
> project_folder<-"D:\\OPS\\gss30_analysis"
> new_dir<-setwd(project_folder)
> print(paste("Current directory is",getwd()))
[1] "Current directory is D:/OPS/gss30_analysis"
>
> gss30 <- spss.fixed.file(file="D:\\STATCAN\\gss_c30\\CYCLE30\\Data_Données\\GSS30PUMFM.txt",
+                columns.file="gss30_columns.txt"
+                # varlab.file="D:/STATCAN/NGS/2010/NGS 2010 PUMF SPSS CARDS/NGS 2010 PUMF_vare.
sps",
+                # codes.file="D:/STATCAN/NGS/2010/NGS 2010 PUMF SPSS CARDS/NGS 2010 PUMF_vale.s
ps",
+                # missval.file="D:/STATCAN/NGS/2010/NGS 2010 PUMF SPSS CARDS/NGS 2010 PUMF_miss
.sps",
+                # count.cases=TRUE,
+                # to.lower=TRUE
+ )
Error in rofseek(fptr, pos = 0) : not an rofile
>
```