

## Validate Statsomat/edapy

```
# Import
library(pastecs)
library(Hmisc)
```

```
## Loading required package: lattice
```

```
## Loading required package: survival
```

```
## Loading required package: Formula
```

```
## Loading required package: ggplot2
```

```
##
```

```
## Attaching package: 'Hmisc'
```

```
## The following objects are masked from 'package:base':
```

```
##
```

```
##      format.pval, units
```

```
library(knitr)
library(data.table)
```

```
##
```

```
## Attaching package: 'data.table'
```

```
## The following objects are masked from 'package:pastecs':
```

```
##
```

```
##      first, last
```

```
library(psych)
```

```
##
```

```
## Attaching package: 'psych'
```

```
## The following object is masked from 'package:Hmisc':
```

```
##
```

```
##      describe
```

```
## The following objects are masked from 'package:ggplot2':
```

```
##
```

```
##      %+%, alpha
```

```
# Upload and prepare dfs
filepath = "HolzingerSwineford1939.csv"
df <- fread(filepath, data.table=FALSE)

# Data frame of the continuous variables
cols_continuous = c(0,1,7,8,9,10,11,12,13,14,15)
cols_continuous <- cols_continuous+1
df_num <- df[,cols_continuous]

# Validate table for continuous variables
kable(stat.desc(df_num),digits=2)
```

	V1	id	x1	x2	x3	x4	x5	x6	x7	x8	x9
nbr.val	301.00	301.00	301.00	301.00	301.00	301.00	301.00	301.00	301.00	301.00	301.00
nbr.null	0.00	0.00	0.00	0.00	0.00	1.00	0.00	0.00	0.00	0.00	0.00
nbr.na	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00	0.00
min	1.00	1.00	0.67	2.25	0.25	0.00	1.00	0.14	1.30	3.05	2.78
max	301.00	351.00	8.50	9.25	4.50	6.33	7.00	6.14	7.43	10.00	9.25
range	300.00	350.00	7.83	7.00	4.25	6.33	6.00	6.00	6.13	6.95	6.47
sum	45451.00	53143.00	1485.67	1832.50	677.38	921.33	1306.50	657.86	1259.96	1663.65	1617.61
median	151.00	163.00	5.00	6.00	2.12	3.00	4.50	2.00	4.09	5.50	5.42
mean	151.00	176.55	4.94	6.09	2.25	3.06	4.34	2.19	4.19	5.53	5.37
SE.mean	5.02	6.11	0.07	0.07	0.07	0.07	0.07	0.06	0.06	0.06	0.06
CI.mean.0.95	9.87	12.02	0.13	0.13	0.13	0.13	0.15	0.12	0.12	0.11	0.11
var	7575.17	11222.96	1.36	1.39	1.28	1.36	1.67	1.20	1.19	1.03	1.02
std.dev	87.04	105.94	1.17	1.18	1.13	1.16	1.29	1.10	1.09	1.01	1.01
coef.var	0.58	0.60	0.24	0.19	0.50	0.38	0.30	0.50	0.26	0.18	0.19

```
psych::describe(df_num)
```

```
##      vars   n  mean      sd median trimmed   mad  min    max range skew
## V1      1 301 151.00  87.04 151.00  151.00 111.19 1.00 301.00 300.00  0.00
## id      2 301 176.55 105.94 163.00  176.78 140.85 1.00 351.00 350.00 -0.01
## x1      3 301   4.94   1.17   5.00   4.96   1.24 0.67   8.50   7.83 -0.25
## x2      4 301   6.09   1.18   6.00   6.02   1.11 2.25   9.25   7.00  0.47
## x3      5 301   2.25   1.13   2.12   2.20   1.30 0.25   4.50   4.25  0.38
## x4      6 301   3.06   1.16   3.00   3.02   0.99 0.00   6.33   6.33  0.27
## x5      7 301   4.34   1.29   4.50   4.40   1.48 1.00   7.00   6.00 -0.35
## x6      8 301   2.19   1.10   2.00   2.09   1.06 0.14   6.14   6.00  0.86
## x7      9 301   4.19   1.09   4.09   4.16   1.10 1.30   7.43   6.13  0.25
## x8     10 301   5.53   1.01   5.50   5.49   0.96 3.05  10.00   6.95  0.53
## x9     11 301   5.37   1.01   5.42   5.37   0.99 2.78   9.25   6.47  0.20
##      kurtosis   se
## V1      -1.21 5.02
## id      -1.36 6.11
## x1       0.31 0.07
## x2       0.33 0.07
## x3      -0.91 0.07
## x4       0.08 0.07
## x5      -0.55 0.07
## x6       0.82 0.06
```

```
## x7      -0.31 0.06
## x8       1.17 0.06
## x9       0.29 0.06
```

```
# Data frame of the discrete variables
cols_discrete <- c(2,3,4,5,6)
df_cat = df[,cols_discrete]

# Validate tables for discrete variables
Hmisc::describe(df_cat)
```

```
## df_cat
##
##  5 Variables      301 Observations
## -----
## id
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    301      0      301      1    176.6    122.2      17      33
##    .25      .50      .75      .90      .95
##    82      163      272      318      335
##
## lowest : 1 2 3 4 5, highest: 346 347 348 349 351
## -----
## sex
##      n missing distinct      Info      Mean      Gmd
##    301      0      2    0.749    1.515    0.5012
##
## Value      1      2
## Frequency   146   155
## Proportion 0.485 0.515
## -----
## ageyr
##      n missing distinct      Info      Mean      Gmd
##    301      0      6    0.907      13    1.123
##
## lowest : 11 12 13 14 15, highest: 12 13 14 15 16
##
## Value      11      12      13      14      15      16
## Frequency    8    101    110    55    20      7
## Proportion 0.027 0.336 0.365 0.183 0.066 0.023
## -----
## agemo
##      n missing distinct      Info      Mean      Gmd      .05      .10
##    301      0      12    0.993    5.375    3.976      0      1
##    .25      .50      .75      .90      .95
##    2      5      8      10      11
##
## lowest : 0 1 2 3 4, highest: 7 8 9 10 11
##
## Value      0      1      2      3      4      5      6      7      8      9      10
## Frequency   22    31    26    26    27    27    21    25    26    23    19
## Proportion 0.073 0.103 0.086 0.086 0.090 0.090 0.070 0.083 0.086 0.076 0.063
##
## Value      11
```

```

## Frequency      28
## Proportion 0.093
## -----
## school
##      n missing distinct
##    301      0        2
##
## Value      Grant-White      Pasteur
## Frequency      145      156
## Proportion      0.482      0.518
## -----

```