

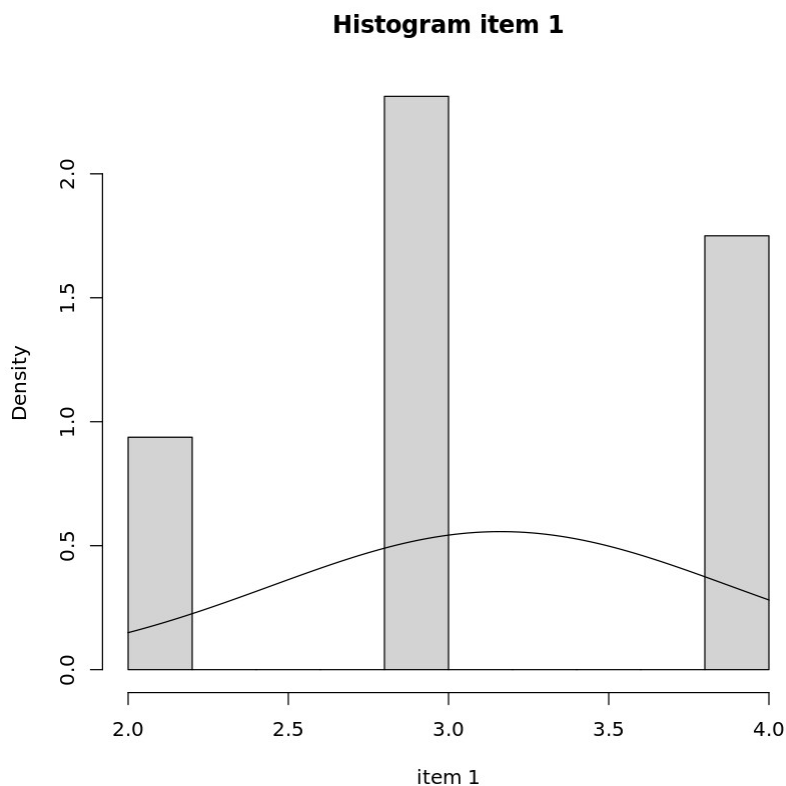
Histogramm:

Ein erster Hinweis auf die Güte der Items ist sicherlich eine grobe Annäherung der Rohdaten an eine Normalverteilung

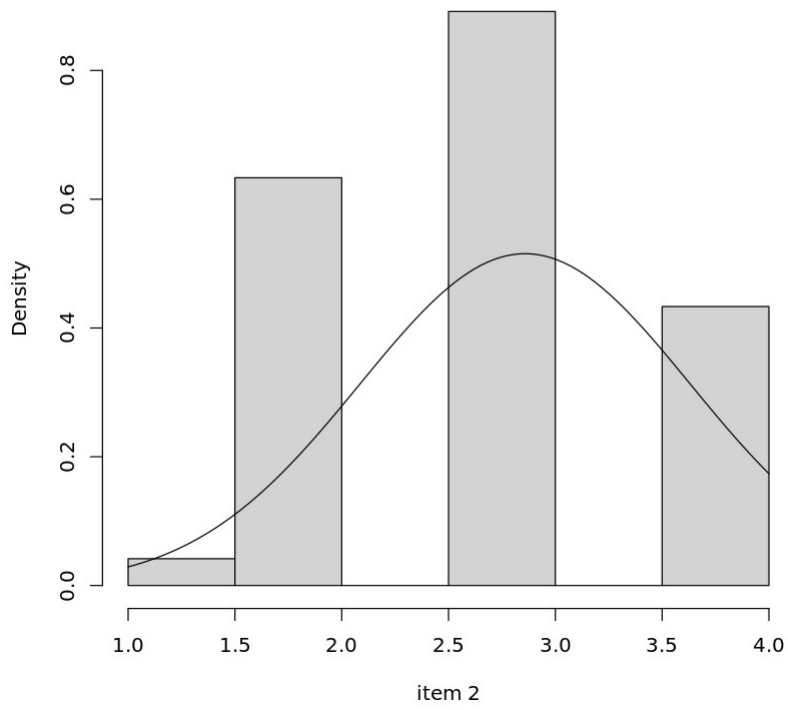
```
fallprozeile <- read.csv2
("https://paul-koop.org/fallprozeilenurdaten.csv", header=FALSE,
dec=",");

for (i in 1:36){
  hist(fallprozeile[,i],
      freq=FALSE,
      main=paste("Histogram item",i),
      xlab=paste("item",i)
      )
  x <- seq(1,4,0.01)

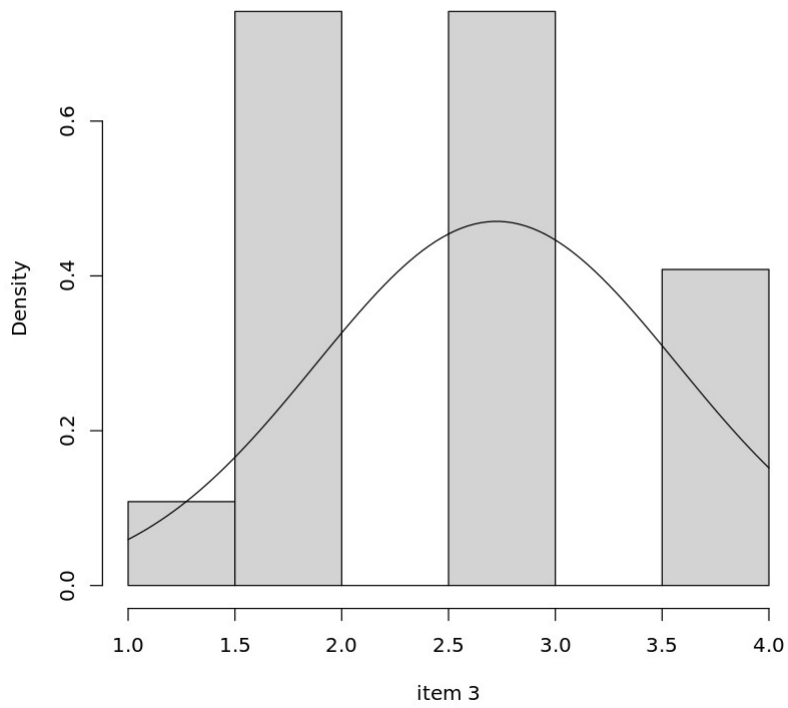
  curve(dnorm(x,mean=mean(fallprozeile[,i]),sd=sd(fallprozeile[,i])),add
=TRUE)
}
```



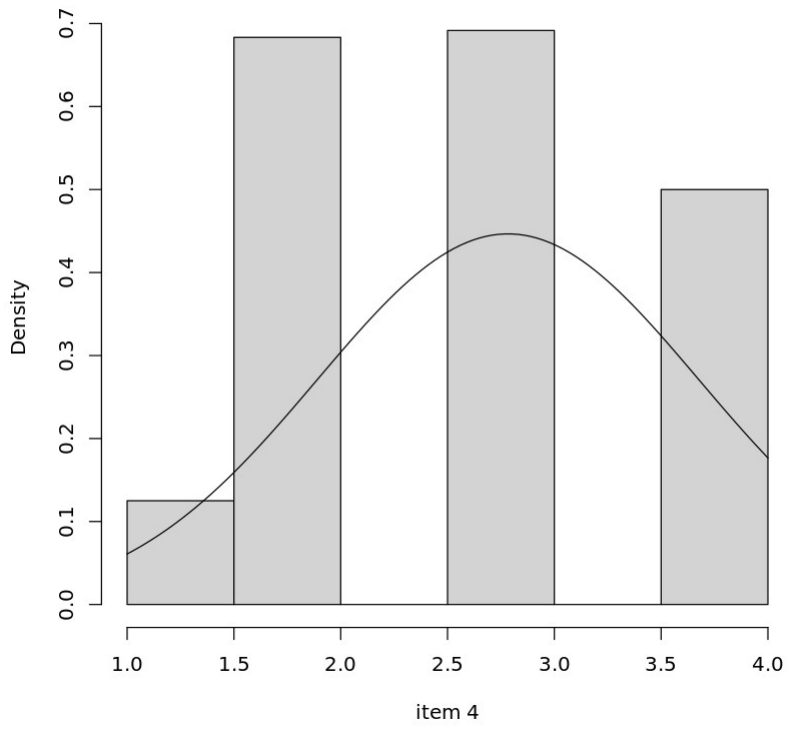
Histogram item 2



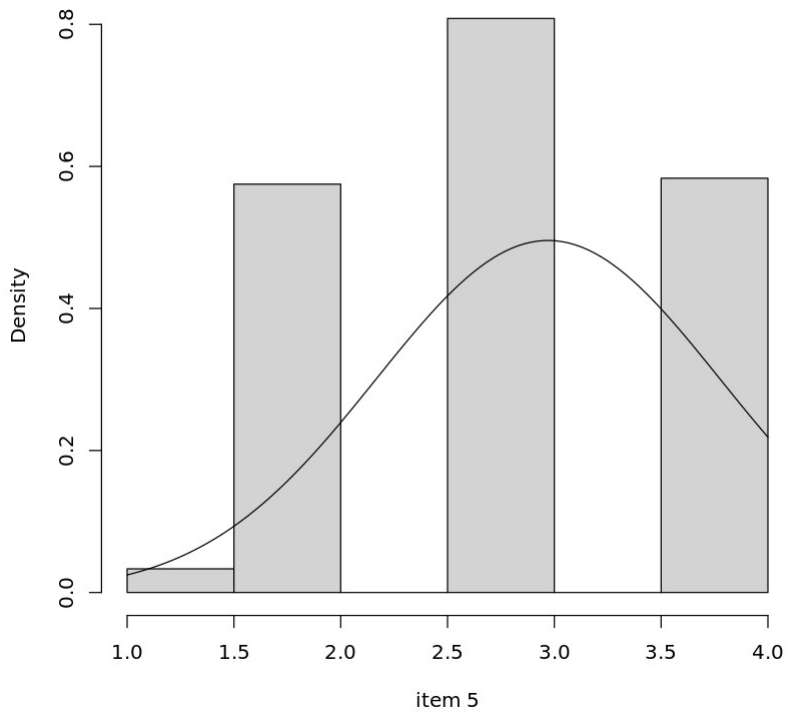
Histogram item 3



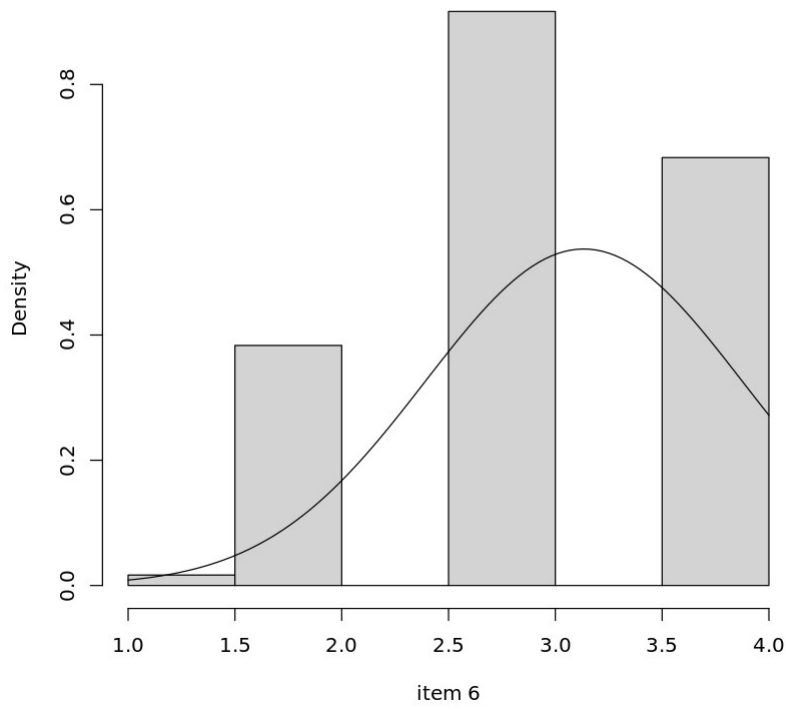
Histogram item 4



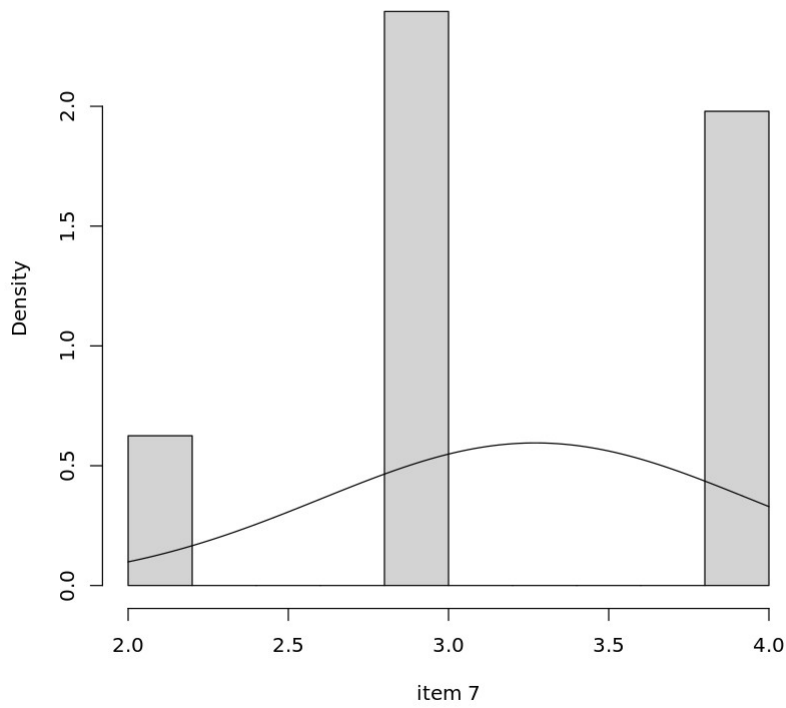
Histogram item 5



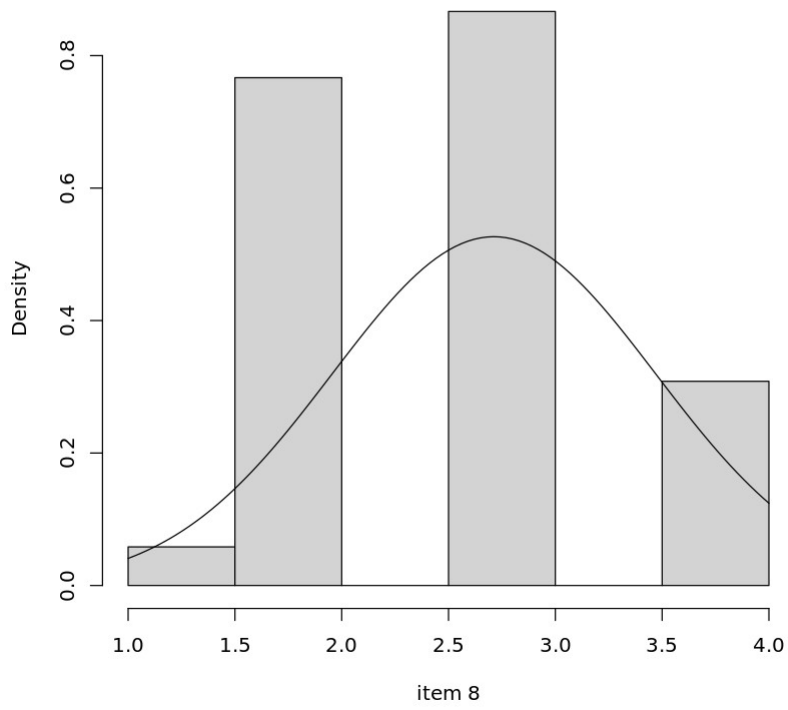
Histogram item 6



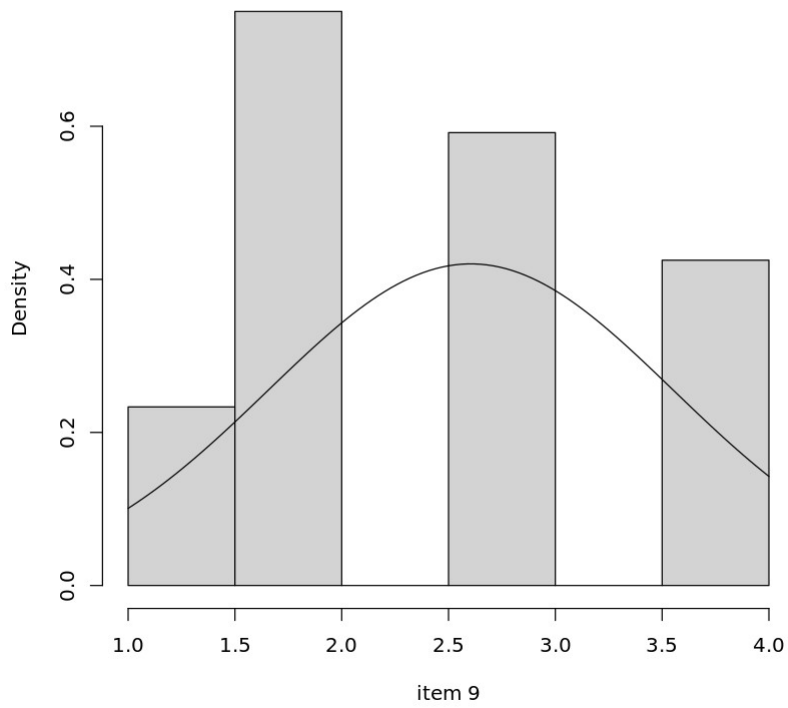
Histogram item 7



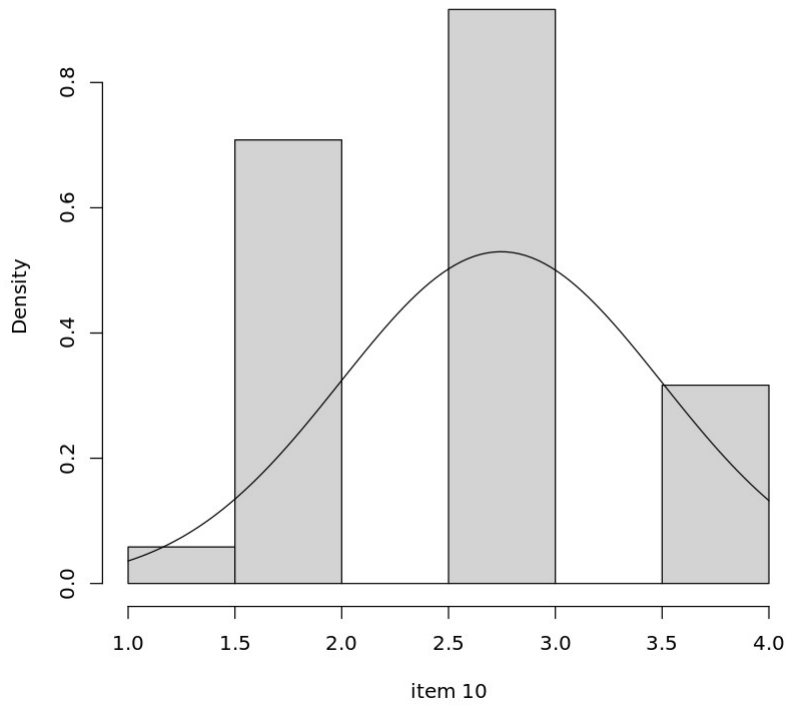
Histogram item 8



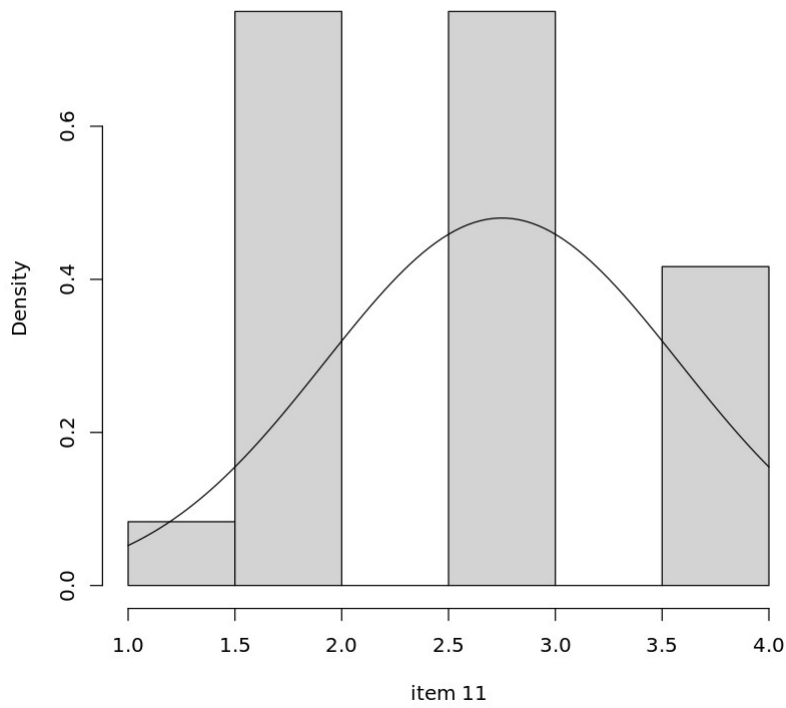
Histogram item 9



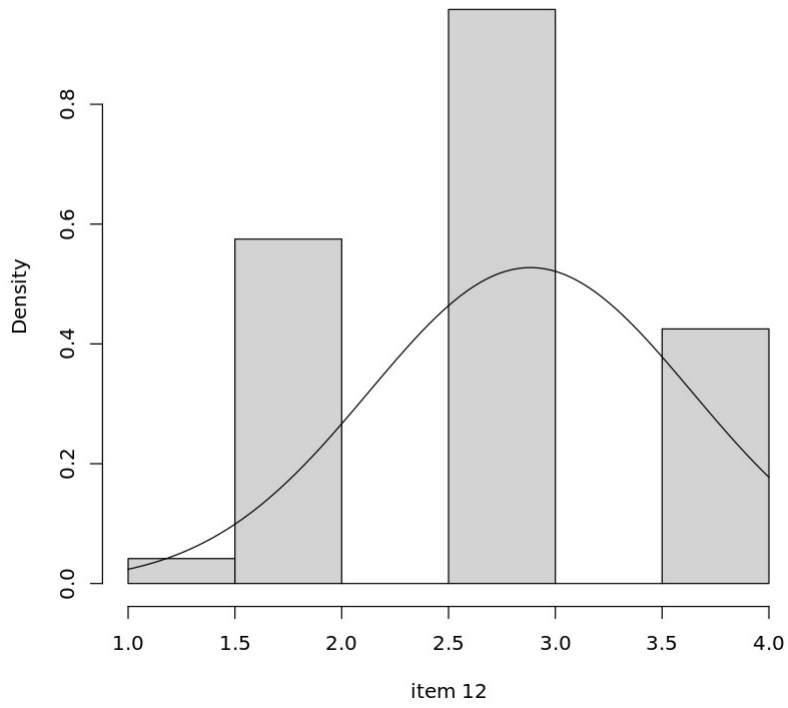
Histogram item 10



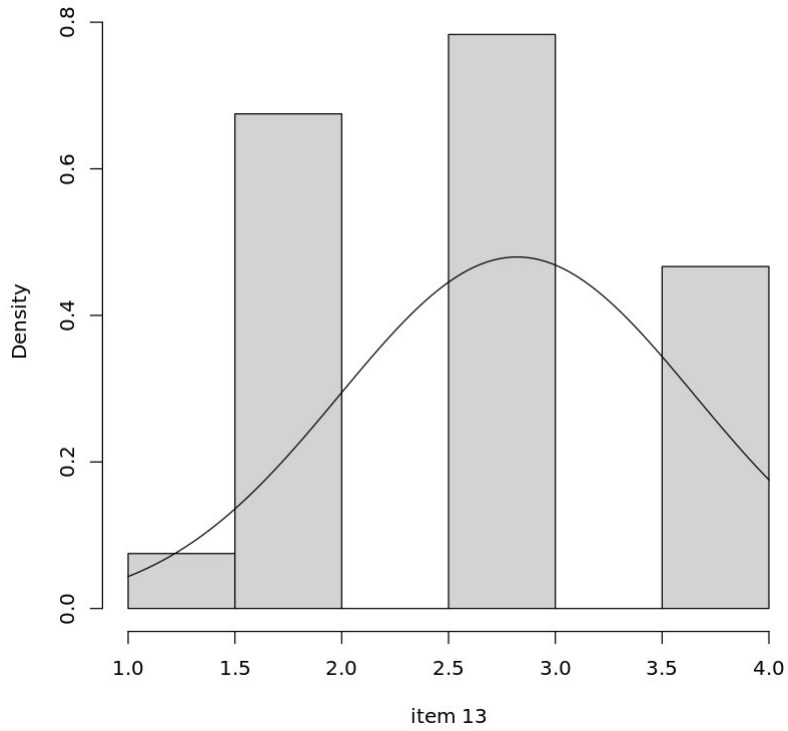
Histogram item 11



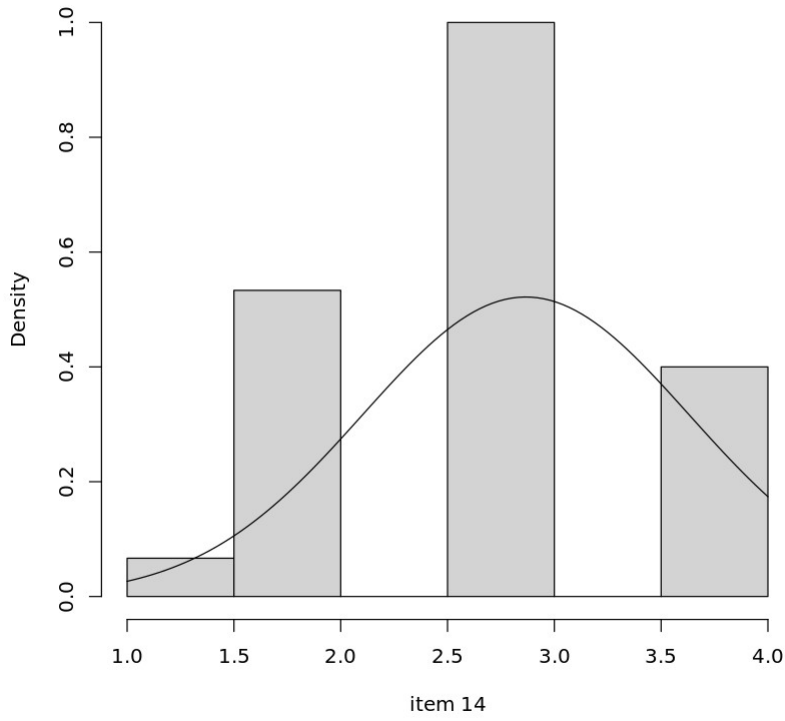
Histogram item 12



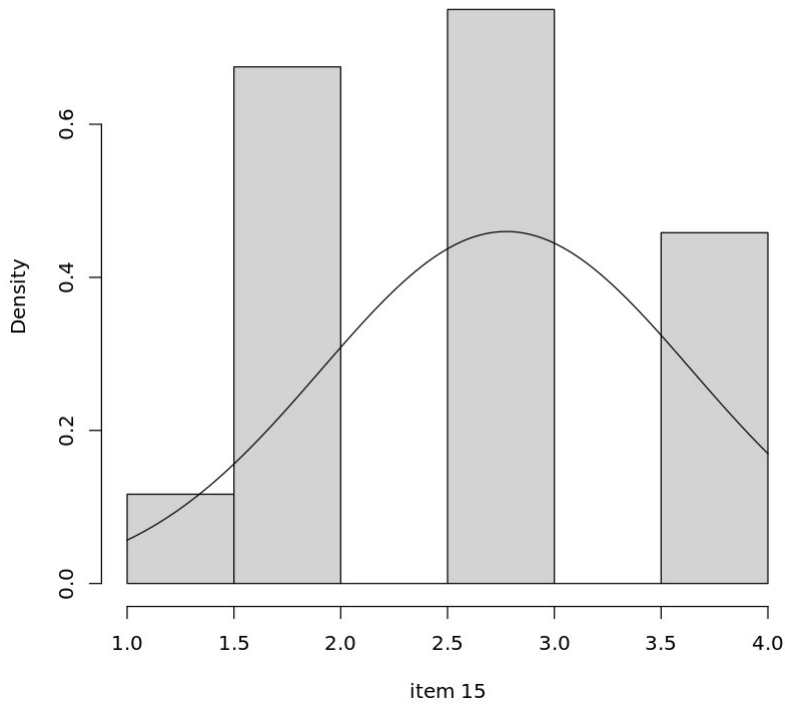
Histogram item 13



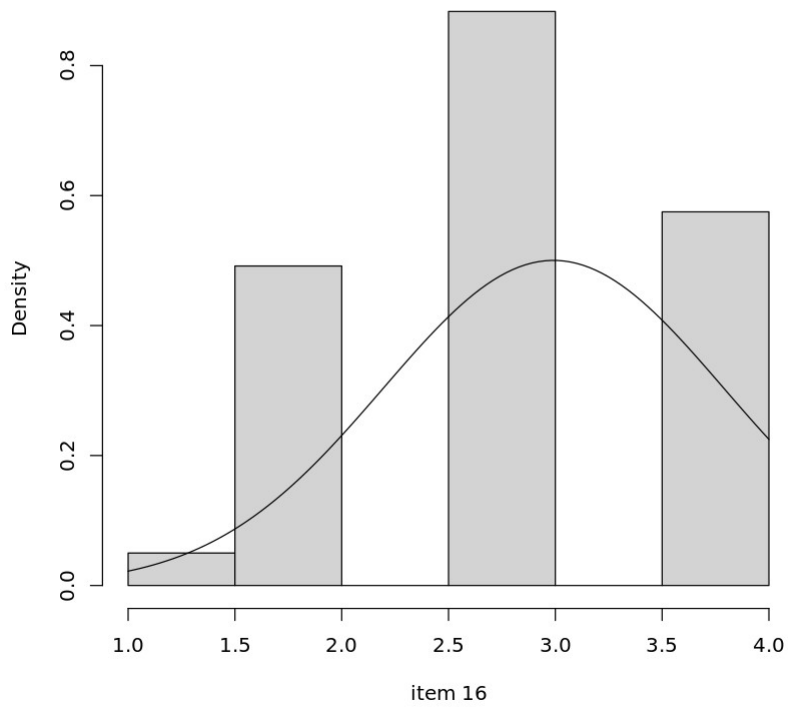
Histogram item 14



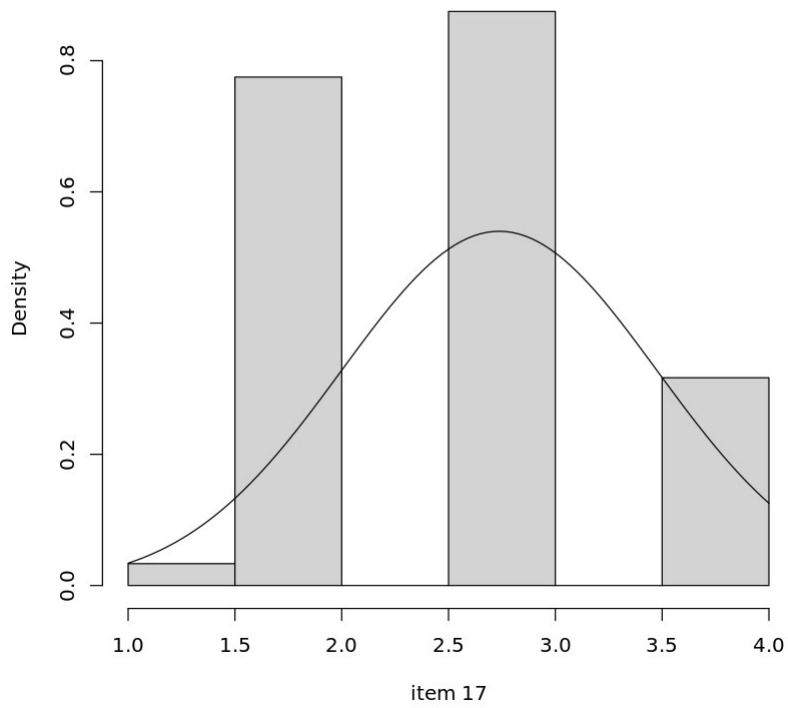
Histogram item 15



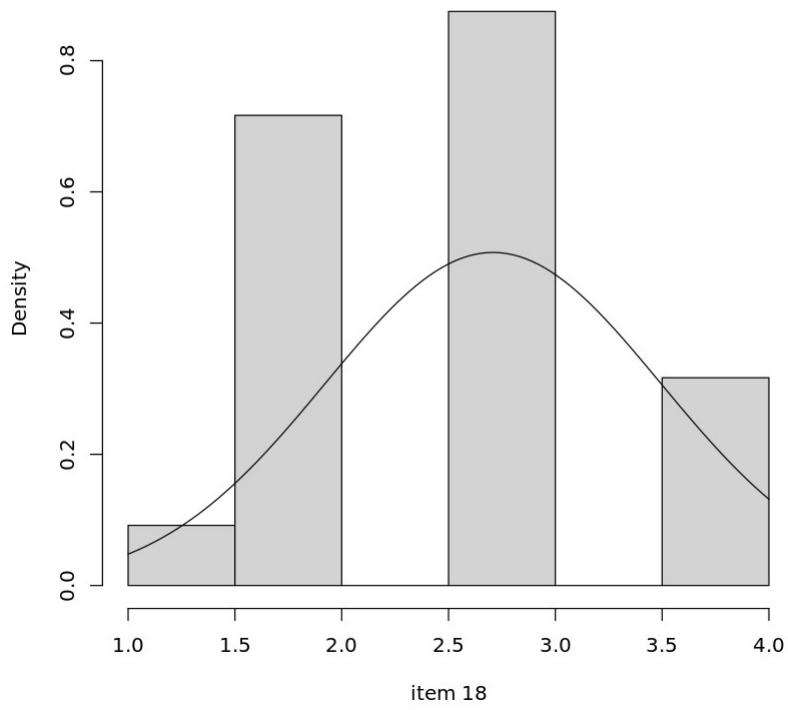
Histogram item 16



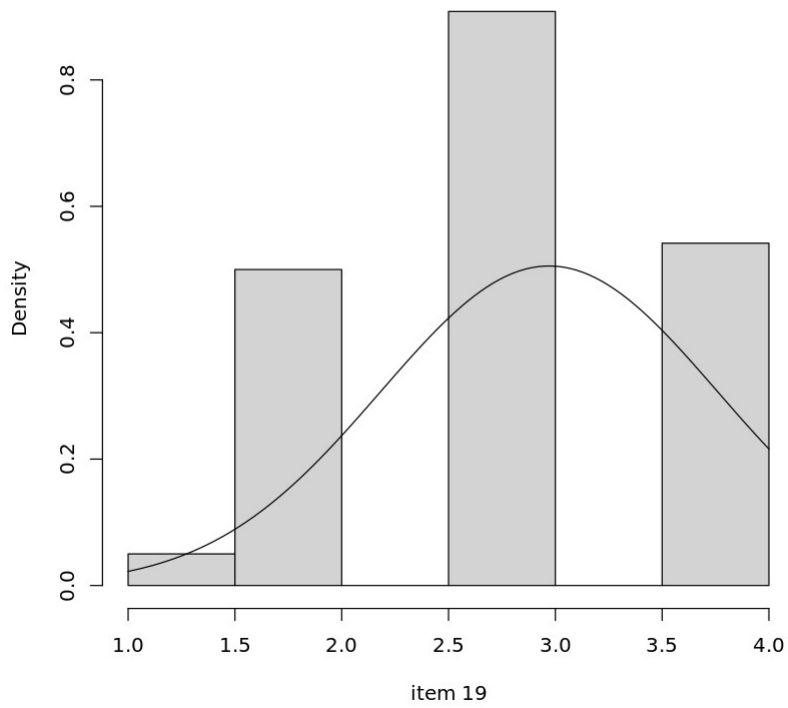
Histogram item 17



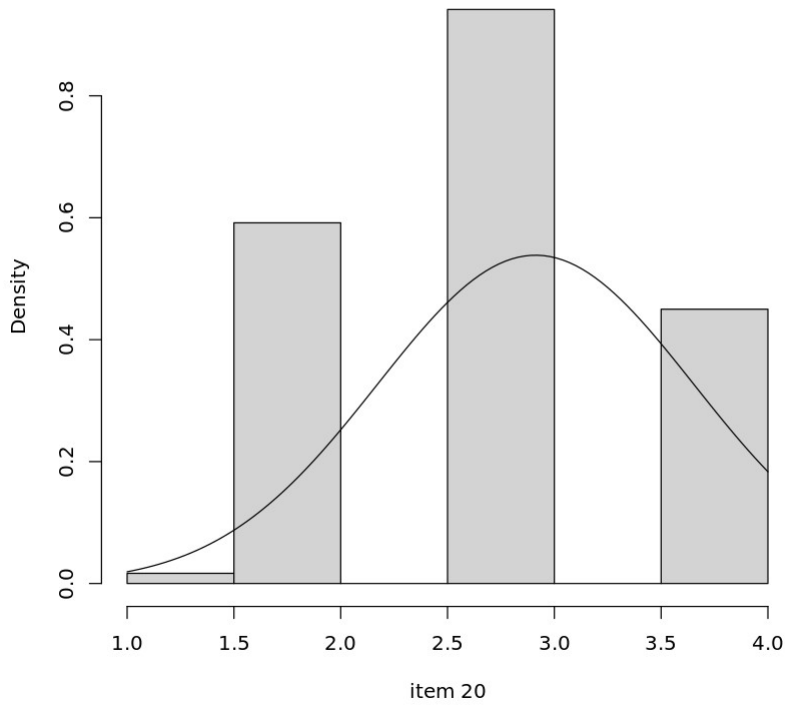
Histogram item 18



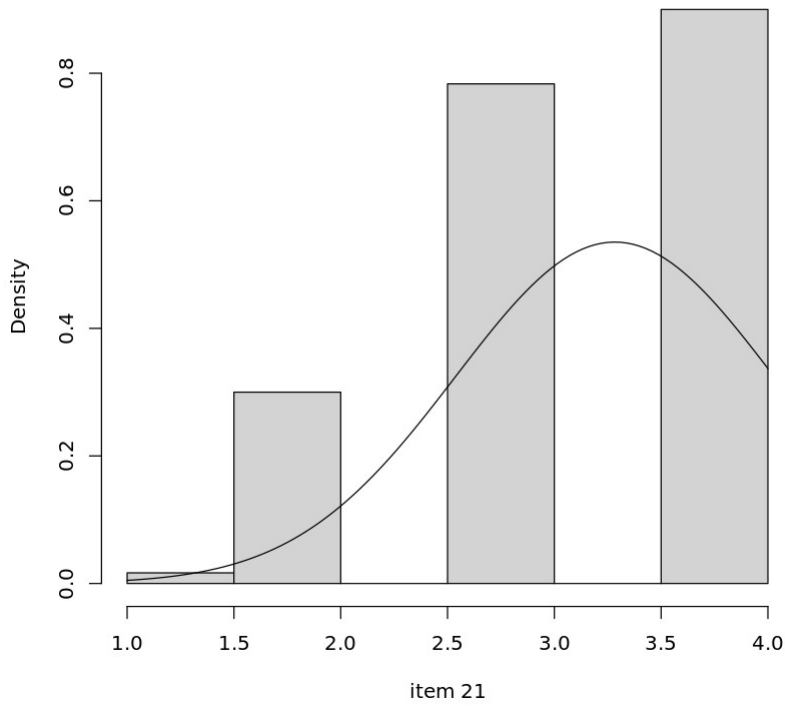
Histogram item 19



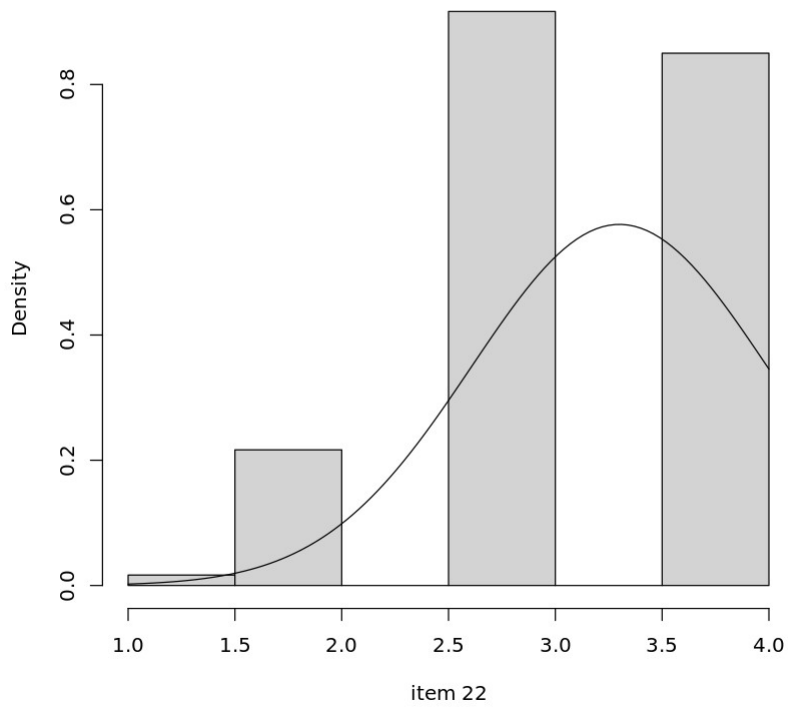
Histogram item 20



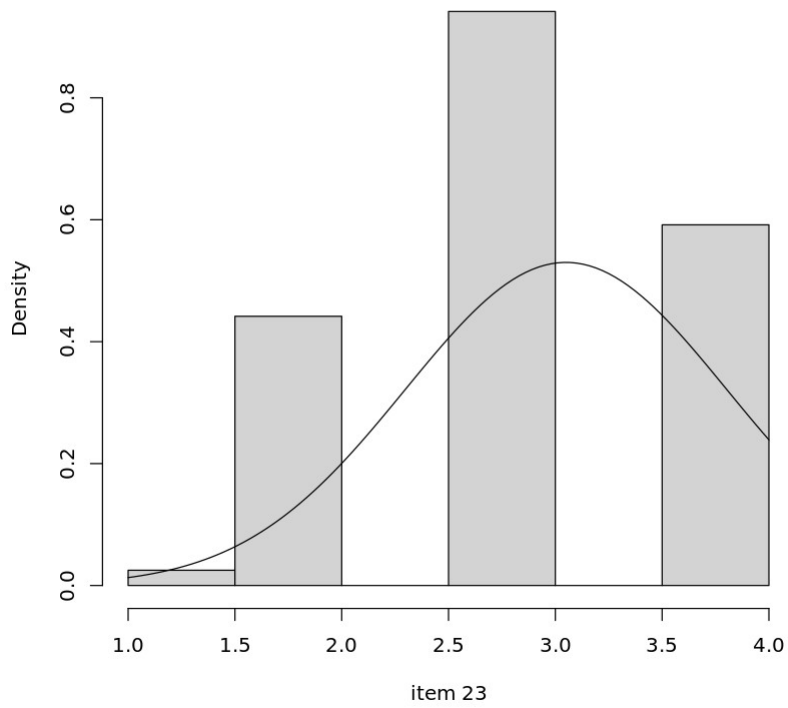
Histogram item 21



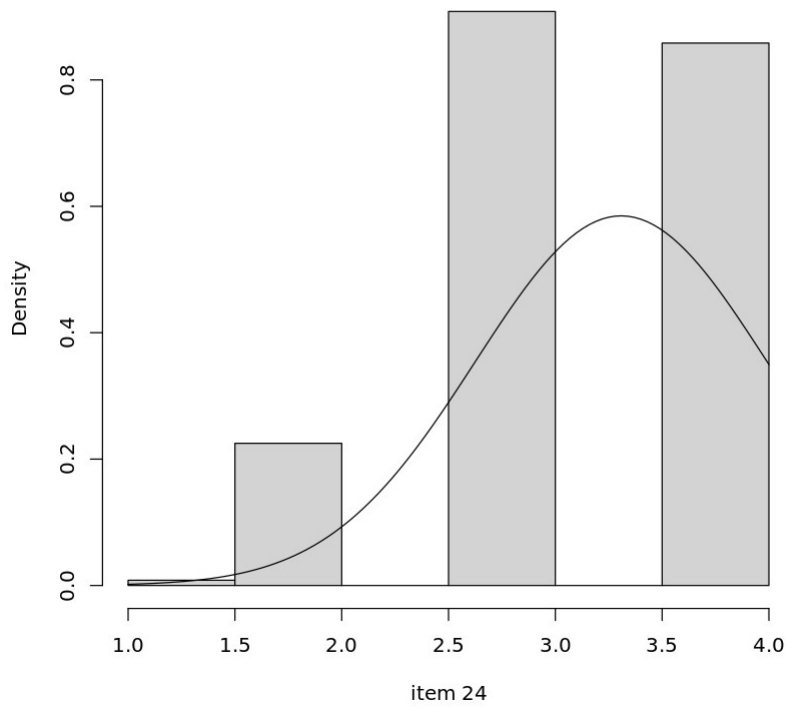
Histogram item 22



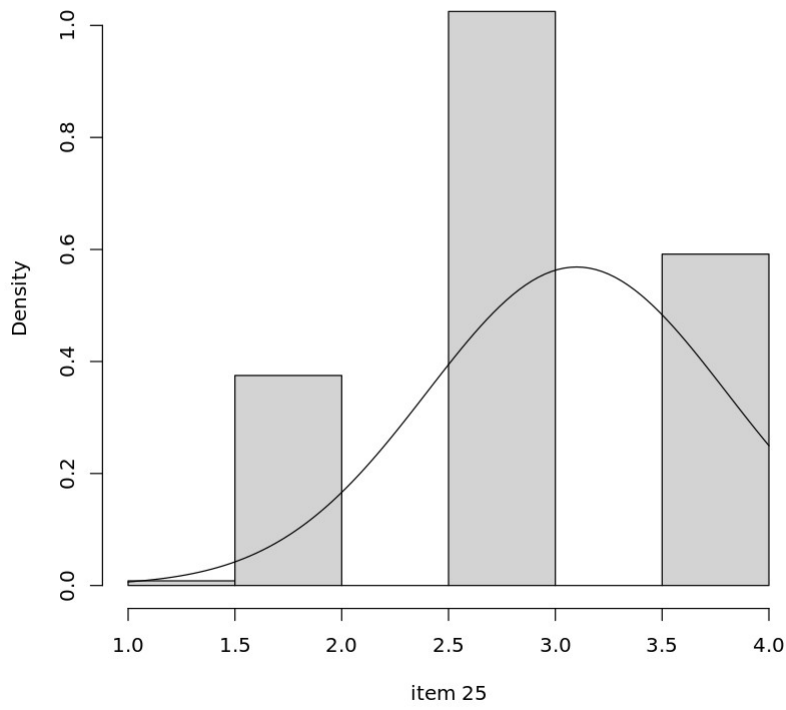
Histogram item 23



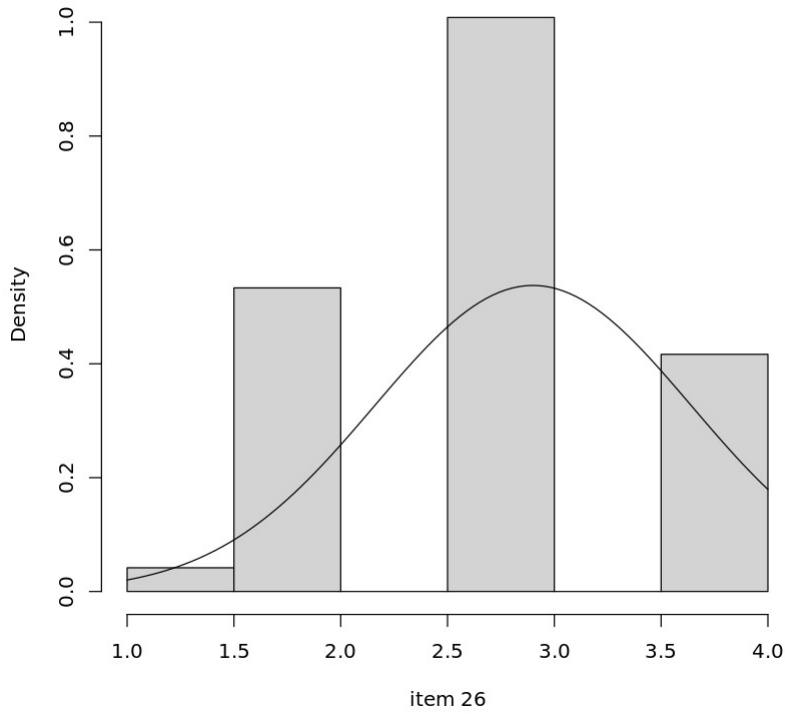
Histogram item 24



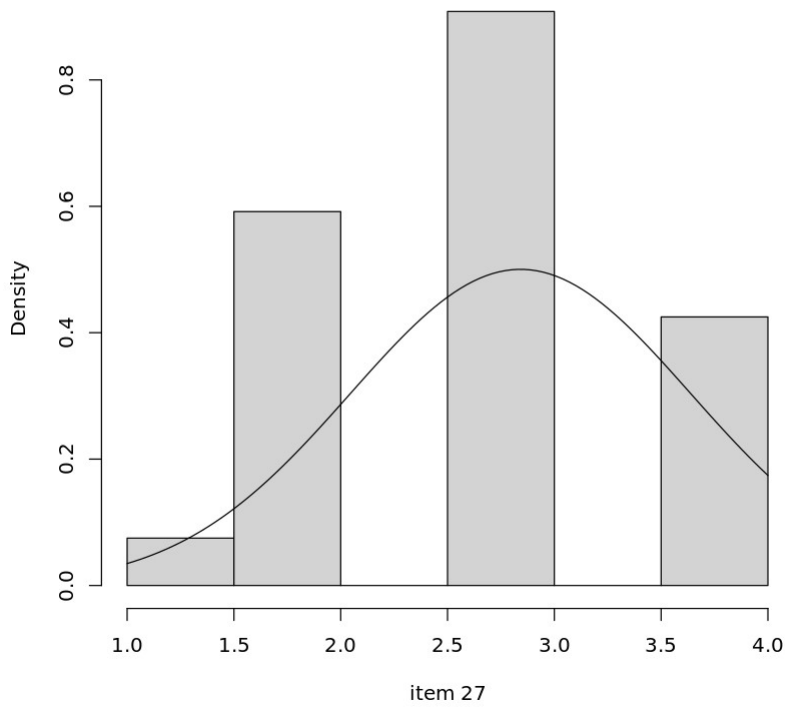
Histogram item 25



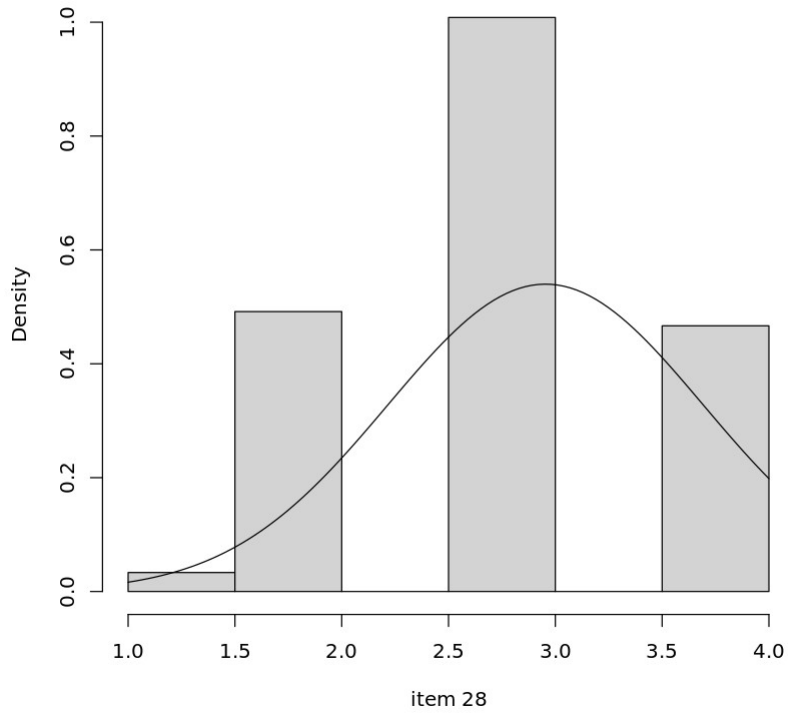
Histogram item 26



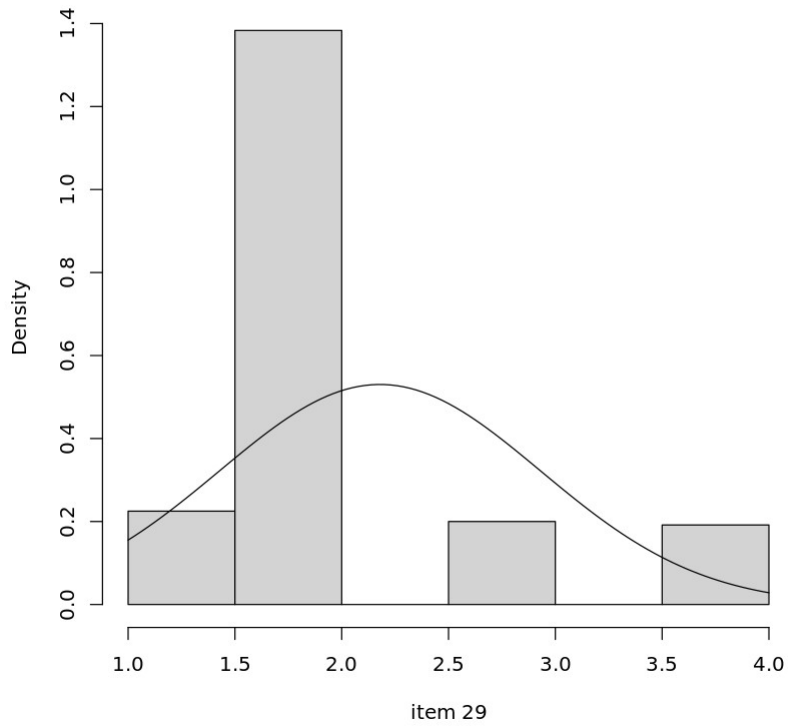
Histogram item 27



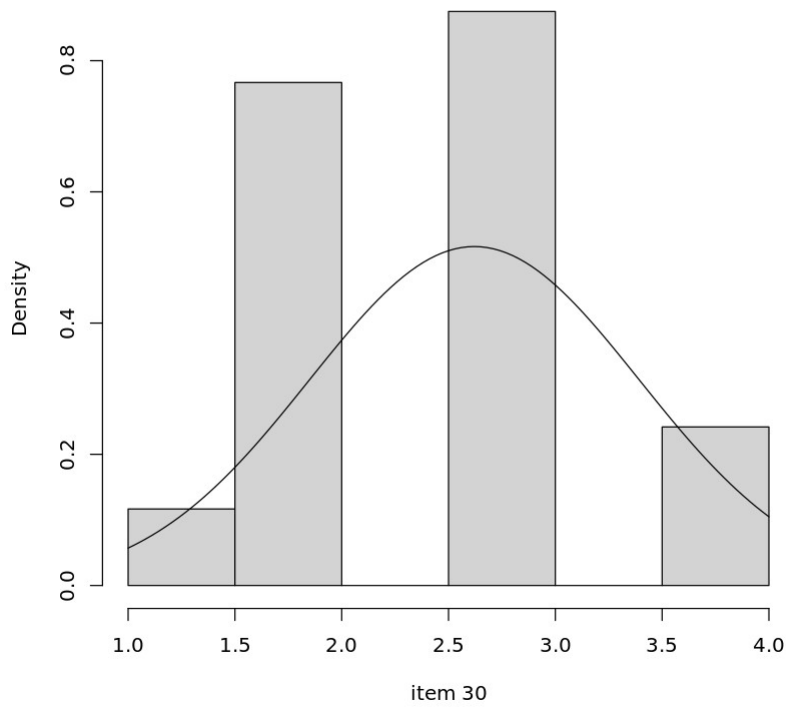
Histogram item 28



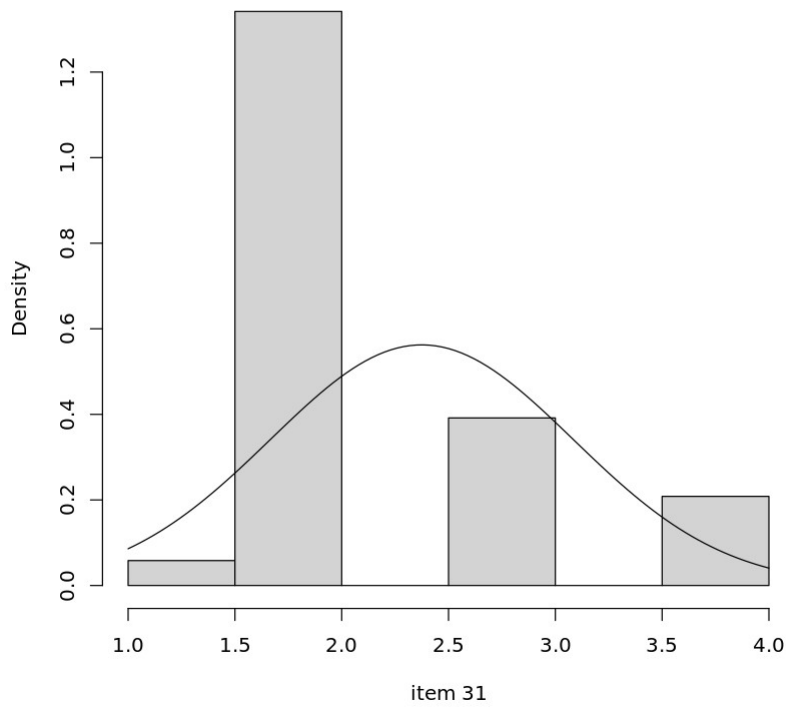
Histogram item 29



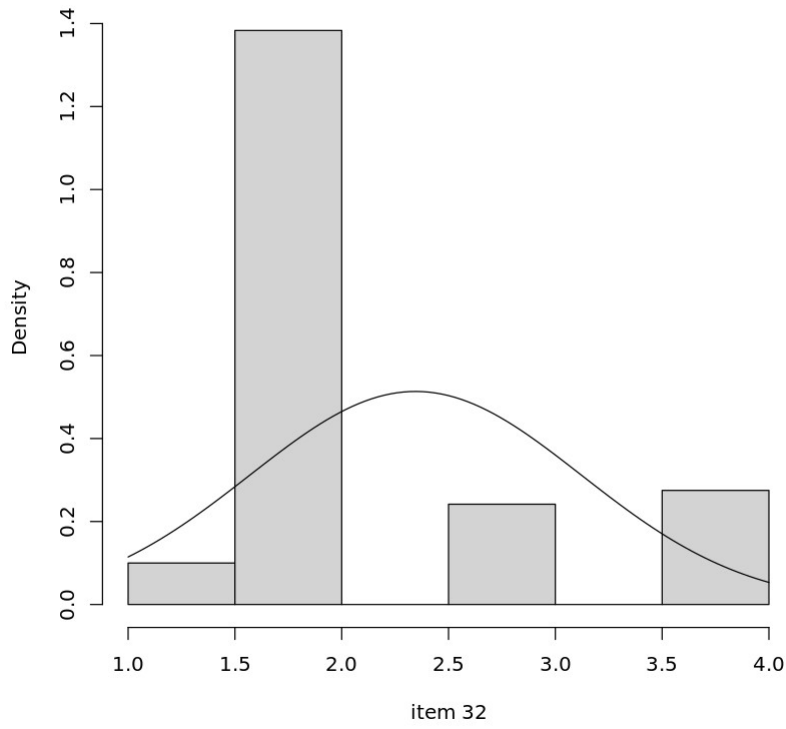
Histogram item 30



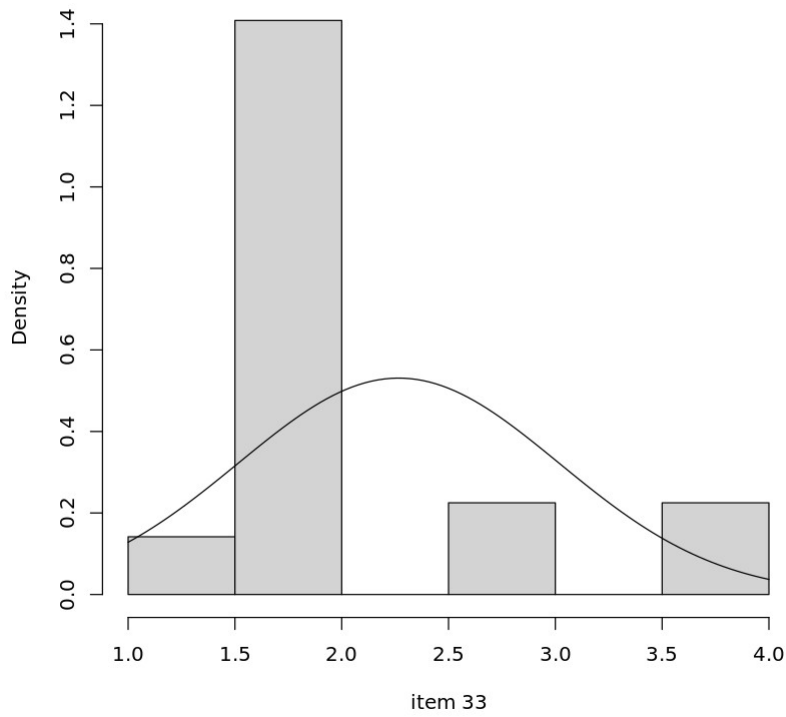
Histogram item 31



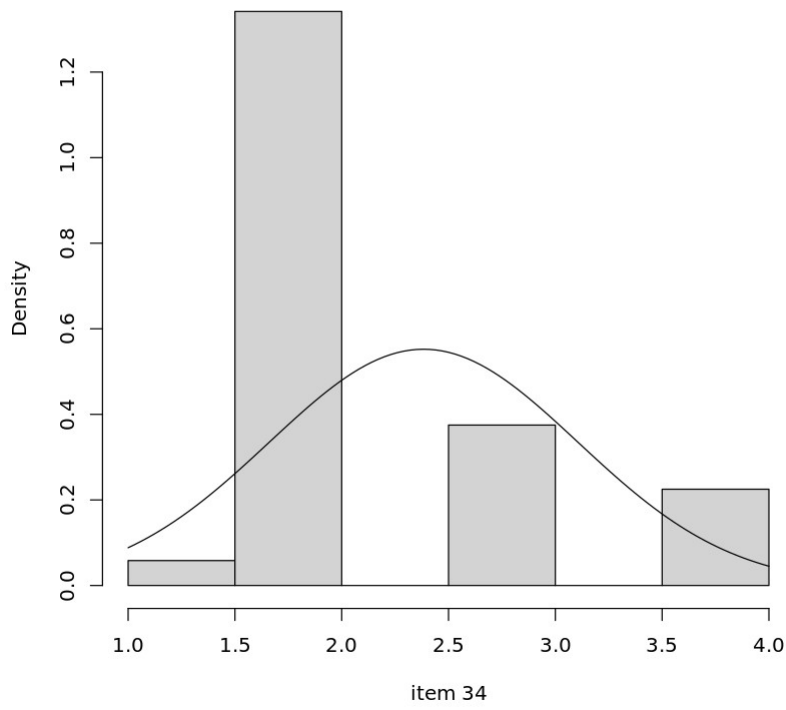
Histogram item 32



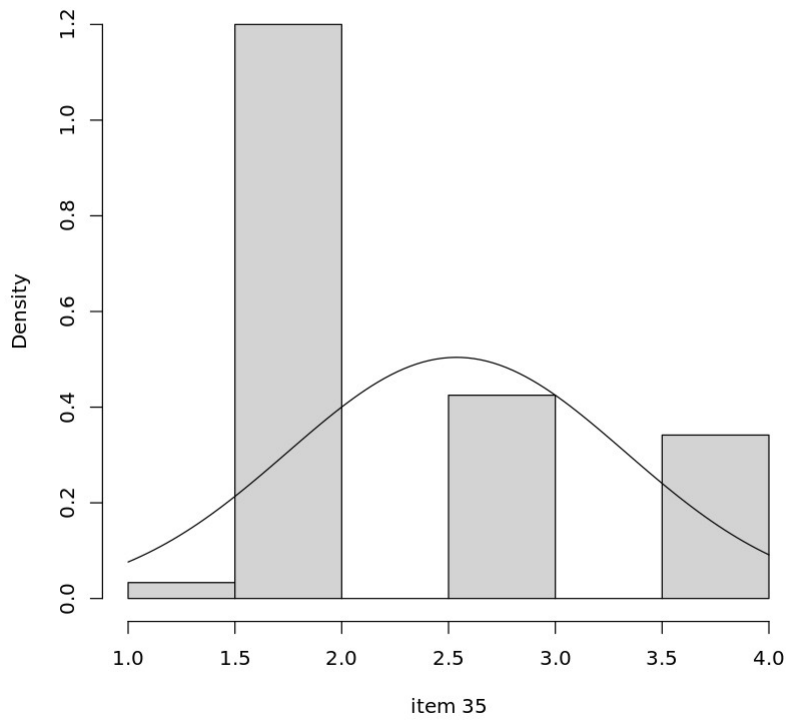
Histogram item 33

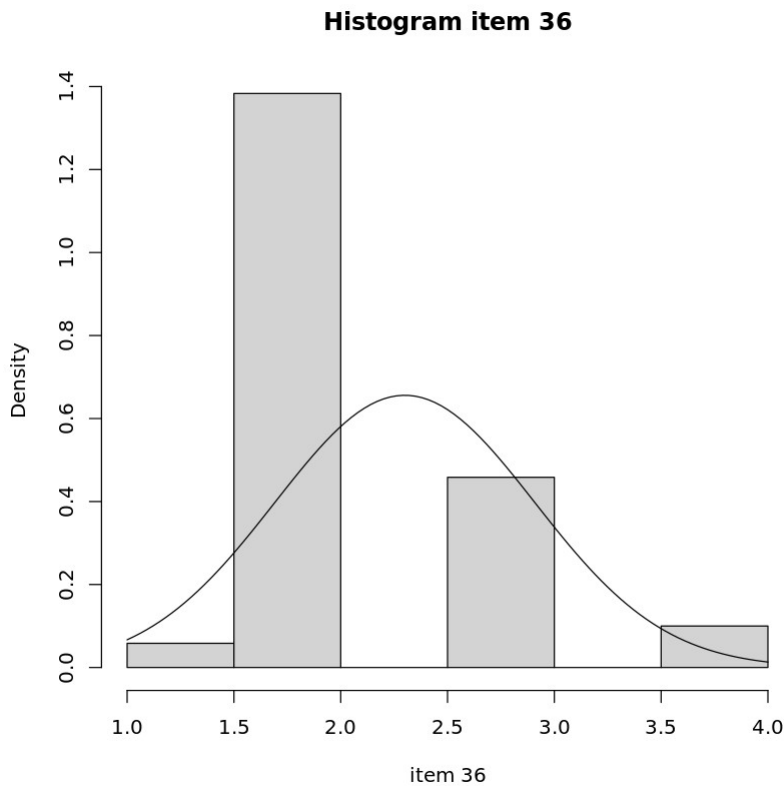


Histogram item 34



Histogram item 35





Ein erstes objektives Merkmal der Validität eines Tests ist die Trennschärfe der Items. Valide ist ein Test dann, wenn er auch tatsächlich die Variable misst, die er auch vorgibt zu messen. Unter der Trennschärfe eines Items versteht man die Korrelation des Items mit dem Gesamtergebnis der jeweils gemessenen Dimension eines Tests.

Fachkompetenz:

```
fachkompetenz <- read.csv2("https://paul-koop.org/SV.csv",
header=TRUE, dec=",");
fachkompetenz$ts <- rowSums(fachkompetenz[, -1])
round(cor(fachkompetenz[, -1]), 2)
```

	X21	X22	X23	X24	X25	X26	X27	X28	ts
X21	1.00	0.61	0.47	0.49	0.48	0.39	0.46	0.42	0.75
X22	0.61	1.00	0.45	0.50	0.47	0.50	0.47	0.36	0.75
X23	0.47	0.45	1.00	0.42	0.56	0.42	0.44	0.41	0.72
X24	0.49	0.50	0.42	1.00	0.54	0.28	0.46	0.27	0.68
X25	0.48	0.47	0.56	0.54	1.00	0.45	0.43	0.40	0.75
X26	0.39	0.50	0.42	0.28	0.45	1.00	0.52	0.46	0.70
X27	0.46	0.47	0.44	0.46	0.43	0.52	1.00	0.56	0.76

```
X28 0.42 0.36 0.41 0.27 0.40 0.46 0.56 1.00 0.68
ts 0.75 0.75 0.72 0.68 0.75 0.70 0.76 0.68 1.00
```

Arbeitsverhalten:

```
arbeitsverhalten <- read.csv2 ("https://paul-koop.org/AV.csv",
header=TRUE, dec=",");
arbeitsverhalten$ts <- rowSums(arbeitsverhalten[, -1])
round(cor(arbeitsverhalten[, -1]),2)
```

	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	ts
X1	1.00	0.41	0.45	0.40	0.41	0.43	0.55	0.40	0.39	0.43	0.69
X2	0.41	1.00	0.44	0.56	0.36	0.35	0.35	0.49	0.51	0.35	0.69
X3	0.45	0.44	1.00	0.60	0.42	0.35	0.34	0.54	0.56	0.41	0.75
X4	0.40	0.56	0.60	1.00	0.43	0.31	0.36	0.61	0.46	0.49	0.76
X5	0.41	0.36	0.42	0.43	1.00	0.29	0.45	0.53	0.43	0.43	0.68
X6	0.43	0.35	0.35	0.31	0.29	1.00	0.64	0.27	0.31	0.26	0.59
X7	0.55	0.35	0.34	0.36	0.45	0.64	1.00	0.33	0.28	0.43	0.66
X8	0.40	0.49	0.54	0.61	0.53	0.27	0.33	1.00	0.55	0.52	0.76
X9	0.39	0.51	0.56	0.46	0.43	0.31	0.28	0.55	1.00	0.37	0.72
X10	0.43	0.35	0.41	0.49	0.43	0.26	0.43	0.52	0.37	1.00	0.67
ts	0.69	0.69	0.75	0.76	0.68	0.59	0.66	0.76	0.72	0.67	1.00

Sozialverhalten:

```
sozialverhalten <- read.csv2 ("https://paul-koop.org/SV.csv",
header=TRUE, dec=",");
sozialverhalten$ts <- rowSums(sozialverhalten[, -1])
round(cor(sozialverhalten[, -1]),2)
```

	X21	X22	X23	X24	X25	X26	X27	X28	ts
X21	1.00	0.61	0.47	0.49	0.48	0.39	0.46	0.42	0.75
X22	0.61	1.00	0.45	0.50	0.47	0.50	0.47	0.36	0.75
X23	0.47	0.45	1.00	0.42	0.56	0.42	0.44	0.41	0.72
X24	0.49	0.50	0.42	1.00	0.54	0.28	0.46	0.27	0.68
X25	0.48	0.47	0.56	0.54	1.00	0.45	0.43	0.40	0.75
X26	0.39	0.50	0.42	0.28	0.45	1.00	0.52	0.46	0.70
X27	0.46	0.47	0.44	0.46	0.43	0.52	1.00	0.56	0.76
X28	0.42	0.36	0.41	0.27	0.40	0.46	0.56	1.00	0.68
ts	0.75	0.75	0.72	0.68	0.75	0.70	0.76	0.68	1.00

Lernverhalten:

```
lernverhalten <- read.csv2 ("https://paul-koop.org/LV.csv",
header=TRUE, dec=",");
```

```
lernverhalten$ts <- rowSums(lernverhalten[,-1])
round(cor(lernverhalten[,-1]),2)
```

	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20	ts
X11	1.00	0.54	0.52	0.47	0.55	0.47	0.53	0.50	0.46	0.46	0.77
X12	0.54	1.00	0.53	0.48	0.50	0.49	0.45	0.29	0.53	0.42	0.73
X13	0.52	0.53	1.00	0.35	0.51	0.45	0.48	0.44	0.48	0.46	0.74
X14	0.47	0.48	0.35	1.00	0.50	0.38	0.43	0.28	0.52	0.48	0.68
X15	0.55	0.50	0.51	0.50	1.00	0.39	0.34	0.34	0.44	0.44	0.71
X16	0.47	0.49	0.45	0.38	0.39	1.00	0.49	0.33	0.52	0.42	0.69
X17	0.53	0.45	0.48	0.43	0.34	0.49	1.00	0.45	0.48	0.42	0.71
X18	0.50	0.29	0.44	0.28	0.34	0.33	0.45	1.00	0.46	0.45	0.64
X19	0.46	0.53	0.48	0.52	0.44	0.52	0.48	0.46	1.00	0.55	0.76
X20	0.46	0.42	0.46	0.48	0.44	0.42	0.42	0.45	0.55	1.00	0.71
ts	0.77	0.73	0.74	0.68	0.71	0.69	0.71	0.64	0.76	0.71	1.00

Interkorrelation:

Einen weiteren ersten qualitativen Hinweis auf die Güte der Items bieten ihre Interkorrelationen innerhalb der Dimensionen, denen die Items zugeordnet sind. Denn wenn die Items eine gemeinsame Dimension messen, müssen sie positiv miteinander korreliert sein.

```
install.packages("psych", repos='http://cran.us.r-project.org')
library(psych)
options(max.print = 9999)
interkorrelation <- read.csv2 ("https://paul-koop.org/rohdaten.csv",
header=TRUE, dec=",");
round(cor(interkorrelation[,-1]),2)
```

Installing package into ‘/srv/rlibs’
(as ‘lib’ is unspecified)

	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	...	X27	X28
X29	X30												
X1	1.00	0.41	0.45	0.40	0.41	0.43	0.55	0.40	0.39	0.43	...	0.28	0.22
X2	0.41	1.00	0.44	0.56	0.36	0.35	0.35	0.49	0.51	0.35	...	0.18	0.33
X3	0.45	0.44	1.00	0.60	0.42	0.35	0.34	0.54	0.56	0.41	...	0.24	0.30
X4	0.40	0.56	0.60	1.00	0.43	0.31	0.36	0.61	0.46	0.49	...	0.30	0.31
X5	0.41	0.36	0.42	0.43	1.00	0.29	0.45	0.53	0.43	0.43	...	0.08	0.10
X6	0.43	0.35	0.35	0.31	0.29	1.00	0.64	0.27	0.31	0.26	...	0.32	0.21

0.31	0.29												
X7	0.55	0.35	0.34	0.36	0.45	0.64	1.00	0.33	0.28	0.43	...	0.21	0.18
0.30	0.31												
X8	0.40	0.49	0.54	0.61	0.53	0.27	0.33	1.00	0.55	0.52	...	0.21	0.18
0.06	0.44												
X9	0.39	0.51	0.56	0.46	0.43	0.31	0.28	0.55	1.00	0.37	...	0.20	0.30
0.15	0.38												
X10	0.43	0.35	0.41	0.49	0.43	0.26	0.43	0.52	0.37	1.00	...	0.30	0.27
0.08	0.36												
X11	0.44	0.56	0.48	0.49	0.41	0.33	0.34	0.52	0.48	0.49	...	0.27	0.23
0.04	0.42												
X12	0.47	0.49	0.43	0.38	0.45	0.34	0.43	0.45	0.54	0.36	...	0.26	0.36
0.08	0.25												
X13	0.55	0.42	0.44	0.38	0.49	0.42	0.43	0.42	0.48	0.49	...	0.29	0.37
0.09	0.34												
X14	0.33	0.39	0.45	0.51	0.37	0.33	0.36	0.46	0.37	0.45	...	0.38	0.32
0.13	0.37												
X15	0.48	0.44	0.54	0.49	0.44	0.38	0.36	0.52	0.40	0.46	...	0.27	0.28
0.07	0.27												
X16	0.46	0.44	0.44	0.53	0.38	0.38	0.44	0.38	0.42	0.38	...	0.24	0.31
0.06	0.31												
X17	0.41	0.54	0.42	0.46	0.43	0.39	0.41	0.47	0.53	0.47	...	0.23	0.30
0.15	0.44												
X18	0.30	0.39	0.32	0.39	0.33	0.32	0.36	0.41	0.46	0.50	...	0.29	0.25
0.13	0.44												
X19	0.44	0.49	0.45	0.54	0.46	0.41	0.50	0.47	0.48	0.45	...	0.33	0.35
0.15	0.39												
X20	0.37	0.40	0.41	0.43	0.43	0.42	0.48	0.46	0.37	0.48	...	0.33	0.21
0.19	0.42												
X21	0.43	0.37	0.37	0.38	0.29	0.34	0.43	0.29	0.24	0.24	...	0.46	0.42
0.27	0.34												
X22	0.30	0.29	0.30	0.34	0.18	0.37	0.37	0.27	0.19	0.24	...	0.47	0.36
0.19	0.33												
X23	0.21	0.26	0.30	0.38	0.26	0.25	0.26	0.36	0.27	0.29	...	0.44	0.41
0.14	0.34												
X24	0.41	0.21	0.22	0.25	0.22	0.40	0.40	0.30	0.21	0.19	...	0.46	0.27
0.20	0.25												
X25	0.40	0.27	0.32	0.32	0.23	0.32	0.33	0.31	0.18	0.31	...	0.43	0.40
0.11	0.38												
X26	0.31	0.25	0.30	0.39	0.07	0.15	0.15	0.23	0.19	0.29	...	0.52	0.46
0.09	0.31												
X27	0.28	0.18	0.24	0.30	0.08	0.32	0.21	0.21	0.20	0.30	...	1.00	0.56
0.19	0.24												
X28	0.22	0.33	0.30	0.31	0.10	0.21	0.18	0.18	0.30	0.27	...	0.56	1.00
0.07	0.28												
X29	0.12	0.12	0.08	0.06	0.12	0.31	0.30	0.06	0.15	0.08	...	0.19	0.07
1.00	0.28												
X30	0.32	0.36	0.43	0.43	0.27	0.29	0.31	0.44	0.38	0.36	...	0.24	0.28
0.28	1.00												
X31	0.02	0.15	-0.07	0.03	0.13	0.06	0.16	0.06	0.06	0.11	...	-0.13	-0.07

0.12	0.17												
X32	0.10	0.06	0.14	0.23	0.10	0.22	0.24	0.11	0.08	0.20	...	0.11	0.07
0.12	0.28												
X33	0.00	0.07	0.10	0.10	0.02	0.14	0.20	0.05	0.08	0.25	...	0.01	0.01
0.20	0.27												
X34	0.07	0.05	0.18	0.11	0.16	0.31	0.31	0.13	0.08	0.18	...	0.09	0.03
0.33	0.35												
X35	-0.01	0.11	0.12	0.11	0.25	0.08	0.27	0.15	0.00	0.15	...	-0.18	-0.09
0.08	0.27												
X36	0.19	0.09	0.11	0.21	0.16	0.22	0.30	0.21	0.00	0.22	...	0.05	-0.05
0.12	0.28												

	X31	X32	X33	X34	X35	X36
X1	0.02	0.10	0.00	0.07	-0.01	0.19
X2	0.15	0.06	0.07	0.05	0.11	0.09
X3	-0.07	0.14	0.10	0.18	0.12	0.11
X4	0.03	0.23	0.10	0.11	0.11	0.21
X5	0.13	0.10	0.02	0.16	0.25	0.16
X6	0.06	0.22	0.14	0.31	0.08	0.22
X7	0.16	0.24	0.20	0.31	0.27	0.30
X8	0.06	0.11	0.05	0.13	0.15	0.21
X9	0.06	0.08	0.08	0.08	0.00	0.00
X10	0.11	0.20	0.25	0.18	0.15	0.22
X11	0.08	0.06	0.05	0.16	0.00	0.17
X12	0.10	0.15	-0.13	0.14	0.09	0.08
X13	-0.02	0.09	0.01	0.11	0.01	0.09
X14	0.03	0.16	0.08	0.27	0.18	0.19
X15	0.03	0.05	-0.05	0.20	0.10	0.19
X16	-0.08	0.23	-0.07	-0.01	0.01	0.01
X17	0.10	0.15	0.12	0.12	0.07	0.18
X18	0.23	0.12	0.22	0.18	0.09	0.25
X19	-0.03	0.22	0.02	0.13	0.05	0.11
X20	0.07	0.21	0.19	0.30	0.23	0.23
X21	0.01	0.04	0.10	0.25	0.09	0.15
X22	0.04	0.20	0.03	0.30	0.16	0.14
X23	-0.12	0.06	0.08	0.10	-0.06	0.08
X24	-0.16	0.11	0.08	0.13	-0.06	0.08
X25	-0.03	0.14	0.14	0.16	0.08	0.15
X26	0.04	0.17	0.00	0.15	-0.04	0.11
X27	-0.13	0.11	0.01	0.09	-0.18	0.05
X28	-0.07	0.07	0.01	0.03	-0.09	-0.05
X29	0.12	0.12	0.20	0.33	0.08	0.12
X30	0.17	0.28	0.27	0.35	0.27	0.28
X31	1.00	0.26	0.24	0.26	0.47	0.36
X32	0.26	1.00	0.14	0.37	0.46	0.37
X33	0.24	0.14	1.00	0.44	0.31	0.30
X34	0.26	0.37	0.44	1.00	0.49	0.49
X35	0.47	0.46	0.31	0.49	1.00	0.42
X36	0.36	0.37	0.30	0.49	0.42	1.00

Cronbachs Alpha:

Ein Test muss eine Variable aber auch möglichst genau messen. Ein Maß für die Genauigkeit der Messung ist die Reliabilität. Wenn es nicht möglich ist, an derselben Testgruppe einen Wiederholungstest zu machen oder die Testergebnisse mit anderen bereits als valide und reliabel eingestuften Tests zu korrelieren, wird häufig der Split-Half Test und die Konsistenzanalyse nach Cronbach durchgeführt. Bei der Split-Half Analyse wird der Test über alle Dimensionen in zwei Hälften aufgeteilt und diese beiden Hälften werden miteinander korreliert.

```
install.packages("psych", repos='http://cran.us.r-project.org')
library(psych)
cronalpha <- read.csv2 ("https://paul-koop.org/rohdaten.csv",
header=TRUE, dec=",");
alpha(cronalpha[,-1])#paket psych
```

```
cronalpha <- cronalpha[,-1]
```

```
erste_haelfte <- cronalpha[,1:18]
zweite_haelfte <- cronalpha[,19:36]
```

```
cor(rowSums(erste_haelfte),rowSums(zweite_haelfte))
```

Installing package into '/srv/rlibs'
(as 'lib' is unspecified)

Reliability analysis

Call: alpha(x = cronalpha[, -1])

raw_alpha	std.alpha	G6(smc)	average_r	S/N	ase	mean	sd	median_r
0.94	0.94	0.96	0.29	15	0.0057	2.8	0.43	0.31

95% confidence boundaries

	lower	alpha	upper
Feldt	0.92	0.94	0.95
Duhachek	0.93	0.94	0.95

Reliability if an item is dropped:

	raw_alpha	std.alpha	G6(smc)	average_r	S/N	alpha	se	var.r	med.r
X1	0.93	0.93	0.96	0.29	14	0.0059	0.025	0.30	
X2	0.93	0.93	0.96	0.29	14	0.0060	0.025	0.30	
X3	0.93	0.93	0.96	0.29	14	0.0060	0.025	0.30	
X4	0.93	0.93	0.96	0.28	14	0.0060	0.025	0.30	
X5	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.31	
X6	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.30	
X7	0.93	0.93	0.96	0.29	14	0.0060	0.026	0.30	
X8	0.93	0.93	0.96	0.29	14	0.0060	0.025	0.31	
X9	0.93	0.93	0.96	0.29	14	0.0059	0.025	0.31	

X10	0.93	0.93	0.96	0.29	14	0.0060	0.026	0.31
X11	0.93	0.93	0.96	0.28	14	0.0060	0.025	0.30
X12	0.93	0.93	0.96	0.29	14	0.0060	0.025	0.31
X13	0.93	0.93	0.96	0.29	14	0.0060	0.025	0.30
X14	0.93	0.93	0.96	0.29	14	0.0060	0.026	0.30
X15	0.93	0.93	0.96	0.29	14	0.0060	0.025	0.30
X16	0.93	0.93	0.96	0.29	14	0.0059	0.025	0.30
X17	0.93	0.93	0.96	0.29	14	0.0060	0.026	0.30
X18	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.31
X19	0.93	0.93	0.96	0.28	14	0.0060	0.025	0.30
X20	0.93	0.93	0.96	0.28	14	0.0060	0.026	0.30
X21	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.30
X22	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.31
X23	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.31
X24	0.94	0.93	0.96	0.29	14	0.0059	0.026	0.31
X25	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.30
X26	0.94	0.93	0.96	0.29	14	0.0058	0.026	0.31
X27	0.94	0.93	0.96	0.29	14	0.0058	0.025	0.31
X28	0.94	0.94	0.96	0.29	14	0.0058	0.025	0.31
X29	0.94	0.94	0.96	0.30	15	0.0057	0.025	0.32
X30	0.93	0.93	0.96	0.29	14	0.0059	0.026	0.30
X31	0.94	0.94	0.96	0.30	15	0.0056	0.023	0.32
X32	0.94	0.94	0.96	0.30	15	0.0057	0.025	0.32
X33	0.94	0.94	0.96	0.30	15	0.0056	0.024	0.32
X34	0.94	0.94	0.96	0.29	15	0.0057	0.025	0.31
X35	0.94	0.94	0.96	0.30	15	0.0056	0.024	0.32
X36	0.94	0.94	0.96	0.30	15	0.0057	0.025	0.32

Item statistics

	n	raw.r	std.r	r.cor	r.drop	mean	sd
X1	240	0.62	0.63	0.62	0.60	3.2	0.72
X2	240	0.64	0.63	0.62	0.60	2.9	0.77
X3	240	0.66	0.65	0.64	0.63	2.7	0.85
X4	240	0.70	0.69	0.69	0.67	2.8	0.89
X5	240	0.59	0.58	0.57	0.55	3.0	0.80
X6	240	0.59	0.60	0.59	0.56	3.1	0.74
X7	240	0.65	0.66	0.66	0.63	3.3	0.67
X8	240	0.66	0.66	0.65	0.64	2.7	0.76
X9	240	0.61	0.60	0.59	0.57	2.6	0.95
X10	240	0.65	0.64	0.64	0.62	2.7	0.75
X11	240	0.69	0.68	0.68	0.66	2.8	0.83
X12	240	0.64	0.64	0.63	0.61	2.9	0.76
X13	240	0.66	0.65	0.64	0.62	2.8	0.83
X14	240	0.67	0.67	0.66	0.64	2.9	0.76
X15	240	0.65	0.64	0.63	0.61	2.8	0.87
X16	240	0.62	0.61	0.60	0.58	3.0	0.80
X17	240	0.66	0.66	0.65	0.63	2.7	0.74
X18	240	0.61	0.61	0.60	0.58	2.7	0.79
X19	240	0.71	0.71	0.70	0.68	3.0	0.79
X20	240	0.70	0.70	0.69	0.67	2.9	0.74

X21	240	0.61	0.62	0.61	0.58	3.3	0.75
X22	240	0.60	0.61	0.60	0.57	3.3	0.69
X23	240	0.54	0.55	0.53	0.51	3.0	0.75
X24	240	0.52	0.53	0.51	0.48	3.3	0.68
X25	240	0.59	0.59	0.58	0.55	3.1	0.70
X26	240	0.51	0.51	0.50	0.47	2.9	0.74
X27	240	0.49	0.49	0.48	0.45	2.8	0.80
X28	240	0.48	0.48	0.46	0.44	3.0	0.74
X29	240	0.29	0.30	0.27	0.25	2.2	0.75
X30	240	0.63	0.63	0.62	0.60	2.6	0.77
X31	240	0.18	0.18	0.16	0.13	2.4	0.71
X32	240	0.33	0.34	0.32	0.29	2.3	0.78
X33	240	0.23	0.24	0.22	0.19	2.3	0.75
X34	240	0.40	0.41	0.39	0.35	2.4	0.72
X35	240	0.27	0.27	0.26	0.22	2.5	0.79
X36	240	0.35	0.36	0.34	0.31	2.3	0.61

Non missing response frequency for each item

	1	2	3	4	miss
X1	0.00	0.19	0.46	0.35	0
X2	0.02	0.32	0.45	0.22	0
X3	0.05	0.37	0.37	0.20	0
X4	0.06	0.34	0.35	0.25	0
X5	0.02	0.29	0.40	0.29	0
X6	0.01	0.19	0.46	0.34	0
X7	0.00	0.12	0.48	0.40	0
X8	0.03	0.38	0.43	0.15	0
X9	0.12	0.38	0.30	0.21	0
X10	0.03	0.35	0.46	0.16	0
X11	0.04	0.38	0.38	0.21	0
X12	0.02	0.29	0.48	0.21	0
X13	0.04	0.34	0.39	0.23	0
X14	0.03	0.27	0.50	0.20	0
X15	0.06	0.34	0.38	0.23	0
X16	0.03	0.25	0.44	0.29	0
X17	0.02	0.39	0.44	0.16	0
X18	0.05	0.36	0.44	0.16	0
X19	0.03	0.25	0.45	0.27	0
X20	0.01	0.30	0.47	0.22	0
X21	0.01	0.15	0.39	0.45	0
X22	0.01	0.11	0.46	0.42	0
X23	0.01	0.22	0.47	0.30	0
X24	0.00	0.11	0.45	0.43	0
X25	0.00	0.19	0.51	0.30	0
X26	0.02	0.27	0.50	0.21	0
X27	0.04	0.30	0.45	0.21	0
X28	0.02	0.25	0.50	0.23	0
X29	0.11	0.69	0.10	0.10	0
X30	0.06	0.38	0.44	0.12	0
X31	0.03	0.67	0.20	0.10	0

```
X32 0.05 0.69 0.12 0.14 0
X33 0.07 0.70 0.11 0.11 0
X34 0.03 0.67 0.19 0.11 0
X35 0.02 0.60 0.21 0.17 0
X36 0.03 0.69 0.23 0.05 0
```

```
[1] 0.674236
```

Hauptkomponentenanalyse PCA

Die Hauptkomponentenanalyse (abgekürzt: PCA) ist eine Methode der multivariaten Statistik und strukturiert Datensätze durch Approximation einer großen Anzahl statistischer Variablen mit einer kleineren Anzahl korrelierter linearer Hauptkomponenten.

```
install.packages("stats", repos='http://cran.us.r-project.org')
library(psych)
tabellePCAMVS <- read.csv2 ("https://paul-koop.org/PCAMVS.csv",
header=TRUE, dec=",");
PCAbeobachtung <- tabellePCAMVS;

PCAMVSnr <- PCAbeobachtung[,1]; PCAbeobachtung <- PCAbeobachtung[, -
1];
PCAMVSart <- PCAbeobachtung[,1];PCAbeobachtung <- PCAbeobachtung[, -1];

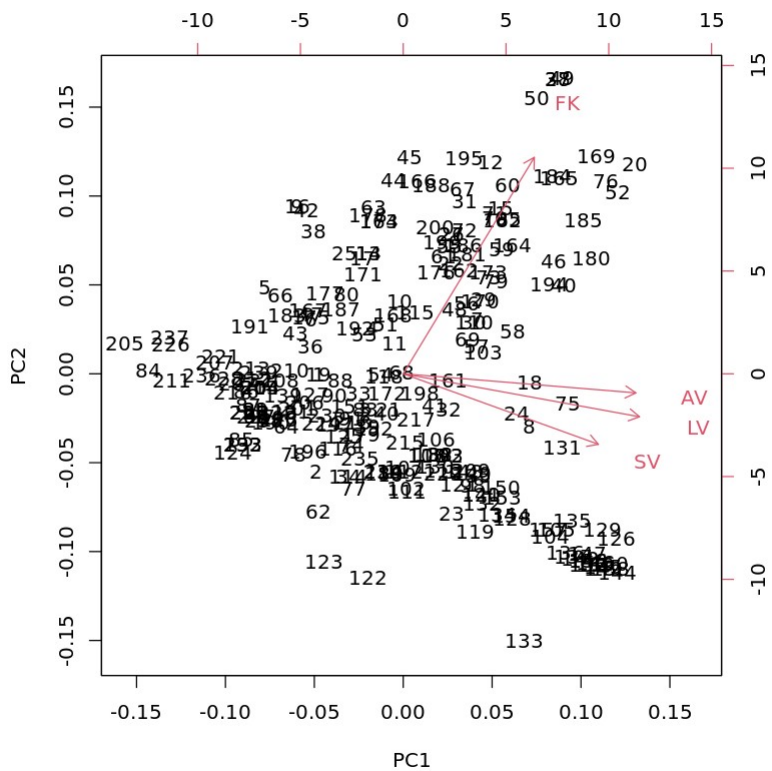
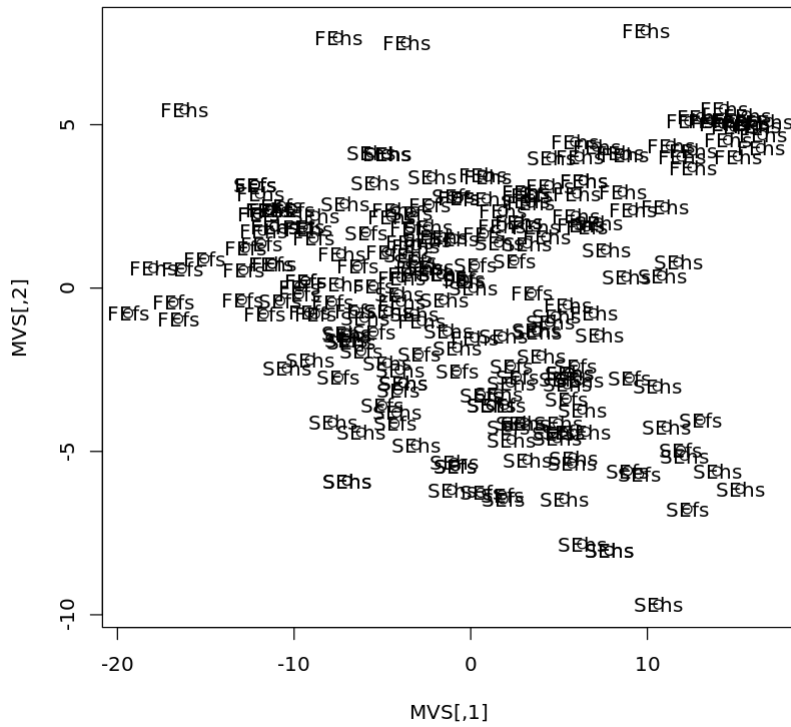
MVS <- cmdscale(dist(PCAbeobachtung));
plot (MVS, type = "p", col = 1);
text(MVS[,1],
      MVS[,2],
      PCAMVSart,
      col=1)

PCA<-prcomp(scale(PCAbeobachtung))
biplot(PCA,choices=c(1,2))
```

```
Installing package into '/srv/rlibs'
(as 'lib' is unspecified)
```

Warning message:

```
"package 'stats' is a base package, and should not be updated"
```



Faktorenanalyse:

Mithilfe der Faktorenanalyse wird überprüft, ob sich die Items in mehrere Subskalen unterteilen lassen.

```
install.packages("psych", repos='http://cran.us.r-project.org')
library(psych)
rohdaten <- read.csv2 ("http://paul-koop.org/rohdaten.csv",
header=TRUE, dec=",");
rohdaten <- rohdaten[,-1]
KMO(rohdaten)
scree(rohdaten)
print(factanal(rohdaten, factors=4, rotation="varimax",
scores="Bartlett"), digits=2, cutoff=.3)
print(factanal(rohdaten, factors=6, rotation="varimax",
scores="Bartlett"), digits=2, cutoff=.3)
```

Installing package into '/srv/rlibs'
(as 'lib' is unspecified)

Kaiser-Meyer-Olkin factor adequacy

Call: KMO(r = rohdaten)

Overall MSA = 0.9

MSA for each item =

X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14
X15	X16												
0.91	0.92	0.94	0.93	0.93	0.88	0.89	0.93	0.91	0.91	0.92	0.91	0.95	0.95
0.94	0.94												
X17	X18	X19	X20	X21	X22	X23	X24	X25	X26	X27	X28	X29	X30
X31	X32												
0.96	0.90	0.94	0.95	0.90	0.90	0.92	0.87	0.90	0.85	0.89	0.85	0.76	0.93
0.68	0.70												
X33	X34	X35	X36										
0.68	0.82	0.68	0.85										

Call:

```
factanal(x = rohdaten, factors = 4, scores = "Bartlett", rotation =
"varimax")
```

Uniquenesses:

X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14
X15	X16												
0.50	0.53	0.51	0.45	0.53	0.48	0.24	0.44	0.48	0.56	0.45	0.52	0.50	0.53
0.55	0.53												
X17	X18	X19	X20	X21	X22	X23	X24	X25	X26	X27	X28	X29	X30

X31	X32												
0.52	0.63	0.46	0.52	0.49	0.45	0.56	0.51	0.52	0.48	0.45	0.59	0.83	0.57
0.69	0.73												
X33	X34	X35	X36										
0.75	0.46	0.42	0.60										

Loadings:

	Factor1	Factor2	Factor3	Factor4
X1	0.53			0.43
X2	0.66			
X3	0.66			
X4	0.67			
X5	0.62			
X6	0.33			0.58
X7	0.40			0.70
X8	0.72			
X9	0.71			
X10	0.60			
X11	0.69			
X12	0.63			
X13	0.62			
X14	0.51	0.40		
X15	0.62			
X16	0.57			
X17	0.65			
X18	0.53			
X19	0.61	0.30		
X20	0.51			
X21		0.58		0.31
X22		0.65		
X23		0.59		
X24		0.52		0.43
X25		0.60		
X26		0.67		
X27		0.71		
X28		0.57		
X29				
X30	0.44	0.30	0.38	
X31			0.53	
X32			0.49	
X33			0.50	
X34			0.70	
X35			0.75	
X36			0.62	

	Factor1	Factor2	Factor3	Factor4
SS loadings	7.88	4.19	2.89	2.01
Proportion Var	0.22	0.12	0.08	0.06
Cumulative Var	0.22	0.34	0.42	0.47

Test of the hypothesis that 4 factors are sufficient.
 The chi square statistic is 1022.46 on 492 degrees of freedom.
 The p-value is 2.15e-39

Call:
 factanal(x = rohdaten, factors = 6, scores = "Bartlett", rotation =
 "varimax")

Uniquenesses:

	X1	X2	X3	X4	X5	X6	X7	X8	X9	X10	X11	X12	X13	X14	X15	X16	X17	X18	X19	X20	X21	X22	X23	X24	X25	X26	X27	X28	X29	X30	X31	X32	X33	X34	X35	X36
	0.50	0.52	0.47	0.39	0.51	0.47	0.26	0.37	0.46	0.52	0.44	0.31	0.47	0.48																						
	0.51	0.53																																		
	0.49	0.41	0.46	0.51	0.47	0.37	0.53	0.48	0.51	0.46	0.43	0.55	0.81	0.53																						
	0.53	0.72																																		
	0.59	0.49	0.30	0.61																																

Loadings:

	Factor1	Factor2	Factor3	Factor4	Factor5	Factor6
X1	0.53			0.42		
X2	0.65					
X3	0.66					
X4	0.68					
X5	0.63					
X6	0.33			0.60		
X7	0.40			0.69		
X8	0.74					
X9	0.71					
X10	0.59					
X11	0.68					
X12	0.65				-0.34	
X13	0.61					
X14	0.52	0.42				
X15	0.62					
X16	0.58					
X17	0.63					
X18	0.51				0.45	
X19	0.61	0.30				
X20	0.51					
X21		0.57		0.32		
X22		0.67				
X23		0.57				
X24		0.49		0.46		
X25		0.58				
X26		0.69				
X27		0.71				
X28		0.58				

X29			0.32			
X30	0.42	0.35				
X31		0.57			0.34	
X32		0.50				
X33		0.45		0.44		
X34		0.67				
X35		0.81				
X36		0.58				

	Factor1	Factor2	Factor3	Factor4	Factor5	Factor6
SS loadings	7.83	4.17	2.87	2.14	0.91	0.60
Proportion Var	0.22	0.12	0.08	0.06	0.03	0.02
Cumulative Var	0.22	0.33	0.41	0.47	0.50	0.51

Test of the hypothesis that 6 factors are sufficient.
The chi square statistic is 810.71 on 429 degrees of freedom.
The p-value is 7.64e-26

Scree plot

