

# EXPLORATORY FACTOR ANALYSIS

```

> cm <- matrix(c(1, 0.62, 0.54, 0.32, 0.284, 0.37,
+               0.62, 1, 0.51, 0.38, 0.351, 0.43,
+               0.54, 0.51, 1, 0.36, 0.336, 0.405,
+               0.32, 0.38, 0.36, 1, 0.686, 0.730,
+               0.284, 0.351, 0.336, 0.686, 1, 0.735,
+               0.37, 0.43, 0.405, 0.730, 0.735, 1),nrow = 6, ncol = 6)
> rownames(cm) <- c("m","p","c","e","h","f")
> colnames(cm) <- c("m","p","c","e","h","f")
> cm

      m      p      c      e      h      f
m 1.000 0.620 0.540 0.320 0.284 0.370
p 0.620 1.000 0.510 0.380 0.351 0.430
c 0.540 0.510 1.000 0.360 0.336 0.405
e 0.320 0.380 0.360 1.000 0.686 0.730
h 0.284 0.351 0.336 0.686 1.000 0.735
f 0.370 0.430 0.405 0.730 0.735 1.000

> eigen(cm)

$values
[1] 3.36709 1.19418 0.50701 0.37185 0.31316
[6] 0.24673

$vectors

      [,1]      [,2]      [,3]      [,4]
[1,] -0.36777 -0.50984  0.2669856  0.727680
[2,] -0.39135 -0.40921  0.4859166 -0.664645
[3,] -0.37195 -0.38258 -0.8316262 -0.152044
[4,] -0.43219  0.37482  0.0215319  0.065318
[5,] -0.42196  0.42146  0.0027301  0.011745
[6,] -0.45652  0.32882  0.0230327  0.034735

      [,5]      [,6]
[1,]  0.0475840 -0.041785
[2,] -0.0053920 -0.038728
[3,] -0.0033314 -0.023524
[4,] -0.7429703 -0.340567
[5,]  0.6651097 -0.449230
[6,]  0.0576177  0.823655

> library(psych)
> library(GPArotation)
> library(MASS)
> #fm="pa" gives principal factoring and by default SMC=FALSE such that initial communalities are 1
> #SMC=FALSE, max.iter=1 yields our princomp solution
> #SMC=TRUE, max.iter=100 yields our prinit solution
> faout <- fa(cm,nfactors=2, fm="pa", SMC=FALSE, max.iter=1,residuals = TRUE, rotate = "none")
> faout #h2 are communalities, u2 are specificities

Factor Analysis using method = pa
Call: fa(r = cm, nfactors = 2, rotate = "none", residuals = TRUE, SMC = FALSE,
      max.iter = 1, fm = "pa")

```

Standardized loadings (pattern matrix) based upon correlation matrix

	PA1	PA2	h2	u2
m	0.67	0.56	0.77	0.23
p	0.72	0.45	0.72	0.28
c	0.68	0.42	0.64	0.36
e	0.79	-0.41	0.80	0.20
h	0.77	-0.46	0.81	0.19
f	0.84	-0.36	0.83	0.17

	PA1	PA2
SS loadings	3.37	1.19
Proportion Var	0.56	0.20
Cumulative Var	0.56	0.76
Proportion Explained	0.74	0.26
Cumulative Proportion	0.74	1.00

Test of the hypothesis that 2 factors are sufficient.

The degrees of freedom for the null model are 15 and the objective function was 2.84  
The degrees of freedom for the model are 4 and the objective function was 0.39

The root mean square of the residuals (RMSR) is 0.08  
The df corrected root mean square of the residuals is 0.21

Fit based upon off diagonal values = 0.97

Measures of factor score adequacy

	PA1
Correlation of scores with factors	1
Multiple R square of scores with factors	1
Minimum correlation of possible factor scores	1

  

	PA2
Correlation of scores with factors	1
Multiple R square of scores with factors	1
Minimum correlation of possible factor scores	1

> faout\$residual #asks for the residual correlations with uniqueness on the diagonal

	m	p	c
m	0.2341798	-0.11374940	-1.5352e-01
p	-0.1137494	0.28435560	-1.6707e-01
c	-0.1535197	-0.16707420	3.5938e-01
e	0.0130286	-0.00632967	-1.0020e-02
h	0.0180897	0.00093934	9.8314e-05
f	0.0048839	-0.01087784	-1.6516e-02

  

	e	h	f
m	0.0130286	1.8090e-02	0.0048839
p	-0.0063297	9.3934e-04	-0.0108778
c	-0.0100198	9.8314e-05	-0.0165159
e	0.2033017	-1.1669e-01	-0.0815195
h	-0.1166858	1.8838e-01	-0.0791067
f	-0.0815195	-7.9107e-02	0.1691378

> faoutrot <- fa(cm,nfactors=2, fm="pa", SMC=FALSE, max.iter=1,residuals = TRUE, rotate = "varimax")  
> faoutrot

```
Factor Analysis using method = pa
Call: fa(r = cm, nfactors = 2, rotate = "varimax", residuals = TRUE,
      SMC = FALSE, max.iter = 1, fm = "pa")
```

Standardized loadings (pattern matrix) based upon correlation matrix

	PA1	PA2	h2	u2
m	0.15	0.86	0.77	0.23
p	0.25	0.81	0.72	0.28
c	0.24	0.76	0.64	0.36
e	0.87	0.21	0.80	0.20
h	0.89	0.16	0.81	0.19
f	0.87	0.28	0.83	0.17

	PA1	PA2
SS loadings	2.44	2.12
Proportion Var	0.41	0.35
Cumulative Var	0.41	0.76
Proportion Explained	0.53	0.47
Cumulative Proportion	0.53	1.00

Test of the hypothesis that 2 factors are sufficient.

The degrees of freedom for the null model are 15 and the objective function was 2.84  
 The degrees of freedom for the model are 4 and the objective function was 0.39

The root mean square of the residuals (RMSR) is 0.08  
 The df corrected root mean square of the residuals is 0.21

Fit based upon off diagonal values = 0.97

Measures of factor score adequacy

	PA1
Correlation of scores with factors	1
Multiple R square of scores with factors	1
Minimum correlation of possible factor scores	1

  

	PA2
Correlation of scores with factors	1
Multiple R square of scores with factors	1
Minimum correlation of possible factor scores	1

```
> princfa <-fa(cm,nfactors=2, fm="pa", SMC=TRUE, max.iter=100,residuals = TRUE, rotate = "none") #pr
> princfa
```

```
Factor Analysis using method = pa
Call: fa(r = cm, nfactors = 2, rotate = "none", residuals = TRUE, SMC = TRUE,
      max.iter = 100, fm = "pa")
```

Standardized loadings (pattern matrix) based upon correlation matrix

	PA1	PA2	h2	u2
m	0.63	0.52	0.67	0.33
p	0.66	0.39	0.58	0.42
c	0.60	0.31	0.45	0.55
e	0.76	-0.31	0.68	0.32
h	0.75	-0.37	0.70	0.30
f	0.83	-0.30	0.79	0.21

	PA1	PA2
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SS loadings	3.03	0.84
Proportion Var	0.50	0.14
Cumulative Var	0.50	0.64
Proportion Explained	0.78	0.22
Cumulative Proportion	0.78	1.00

Test of the hypothesis that 2 factors are sufficient.

The degrees of freedom for the null model are 15 and the objective function was 2.84  
The degrees of freedom for the model are 4 and the objective function was 0

The root mean square of the residuals (RMSR) is 0  
The df corrected root mean square of the residuals is 0

Fit based upon off diagonal values = 1  
Measures of factor score adequacy

	PA1
Correlation of scores with factors	0.95
Multiple R square of scores with factors	0.91
Minimum correlation of possible factor scores	0.81
	PA2
Correlation of scores with factors	0.84
Multiple R square of scores with factors	0.71
Minimum correlation of possible factor scores	0.42

```
> mlfa <-fa(cm,nfactors=2, residuals = TRUE,fm="ml") #maximum likelihood solution
> mlfa
```

Factor Analysis using method = ml

Call: fa(r = cm, nfactors = 2, residuals = TRUE, fm = "ml")

Standardized loadings (pattern matrix) based upon correlation matrix

	ML1	ML2	h2	u2
m	-0.07	0.86	0.68	0.32
p	0.08	0.72	0.58	0.42
c	0.11	0.60	0.45	0.55
e	0.82	0.01	0.68	0.32
h	0.86	-0.05	0.70	0.30
f	0.86	0.05	0.79	0.21

	ML1	ML2
SS loadings	2.21	1.66
Proportion Var	0.37	0.28
Cumulative Var	0.37	0.65
Proportion Explained	0.57	0.43
Cumulative Proportion	0.57	1.00

With factor correlations of

	ML1	ML2
ML1	1.00	0.52
ML2	0.52	1.00

Test of the hypothesis that 2 factors are sufficient.

The degrees of freedom for the null model are 15 and the objective function was 2.84

The degrees of freedom for the model are 4 and the objective function was 0

The root mean square of the residuals (RMSR) is 0

The df corrected root mean square of the residuals is 0

Fit based upon off diagonal values = 1

Measures of factor score adequacy

	ML1
Correlation of scores with factors	0.94
Multiple R square of scores with factors	0.89
Minimum correlation of possible factor scores	0.79
	ML2
Correlation of scores with factors	0.91
Multiple R square of scores with factors	0.82
Minimum correlation of possible factor scores	0.64