

More on items

Examining the effect of skew

Standard scales and tests

- I. SEM/CFA tests are based upon normal theory
- II. Scales are thought to
 - A. Be normally distributed
 - B. Have continuous responses
 - C. Joint distributions of scales are multivariate normal distributed
- III. In addition, scales are hoped to be reliable measures of one construct

But, we work with items

- I. Items are
 - A.discrete
 - B.skewed
 - C.unreliable
 - D.possibly multi-vocal

Consider emotion words

Consider a simulation

I. Base line case

A. 24 items with continuous scores

B. simple structured on two factors

C. average loadings of .6

D. Sample size of 500

E. (Default value of sim.item, except for number of variables)

II. Items with discrete scores

III. Skewed discrete items

Base line simulation

- I. Using the sim.item function with default values except for number of items

Simple descriptives

```
> describe(simple)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	skew	kurtosis	se
V1	1	500	0.01	1.01	0.03	0.01	0.97	-3.26	3.44	6.69	0.05	0.13	0.04
V2	2	500	0.00	1.00	0.00	0.00	1.05	-3.16	3.12	6.28	0.02	-0.14	0.04
V3	3	500	-0.03	1.05	-0.04	-0.04	1.06	-2.95	2.80	5.76	0.10	-0.28	0.05
V4	4	500	-0.04	1.03	-0.05	-0.04	1.07	-3.95	2.90	6.85	0.03	0.08	0.05
V5	5	500	-0.03	0.99	-0.01	-0.02	0.95	-4.04	4.07	8.12	-0.08	0.62	0.04
V6	6	500	-0.04	1.09	-0.07	-0.04	1.06	-3.34	2.96	6.30	0.02	-0.24	0.05
V7	7	500	-0.03	0.95	-0.04	-0.01	0.93	-2.79	2.16	4.95	-0.19	-0.17	0.04
V8	8	500	-0.02	1.07	-0.02	-0.02	1.12	-3.36	3.45	6.81	0.00	-0.17	0.05
V9	9	500	0.00	1.02	-0.01	0.00	1.05	-3.06	3.37	6.43	0.05	-0.27	0.05
V10	10	500	0.00	0.96	0.00	0.00	0.98	-3.37	2.98	6.35	-0.10	0.09	0.04
V11	11	500	-0.06	0.97	-0.09	-0.07	0.98	-3.16	2.92	6.08	0.12	0.07	0.04
V12	12	500	-0.06	0.97	-0.05	-0.07	0.97	-2.61	2.68	5.29	0.08	-0.19	0.04
V13	13	500	-0.05	1.07	-0.08	-0.07	1.11	-2.93	3.74	6.66	0.13	-0.01	0.05
V14	14	500	0.07	0.98	0.08	0.07	0.97	-2.97	2.81	5.78	-0.04	-0.19	0.04
V15	15	500	0.02	1.03	-0.05	0.01	1.03	-2.89	3.22	6.11	0.08	-0.10	0.05
V16	16	500	0.00	1.05	0.00	-0.01	1.02	-3.43	3.15	6.58	0.06	-0.06	0.05
V17	17	500	-0.03	1.05	0.01	-0.01	1.04	-3.27	3.09	6.36	-0.14	0.05	0.05
V18	18	500	0.03	1.04	-0.01	0.03	1.03	-3.04	3.40	6.44	-0.02	0.02	0.05
V19	19	500	-0.01	0.97	-0.02	-0.01	1.02	-2.99	2.62	5.62	-0.01	-0.23	0.04
V20	20	500	0.03	0.99	0.01	0.03	0.96	-3.17	3.15	6.33	0.04	0.26	0.04
V21	21	500	-0.02	1.05	-0.05	-0.02	1.05	-2.92	3.08	6.00	0.06	-0.08	0.05
V22	22	500	0.05	1.03	0.00	0.04	1.01	-3.15	3.06	6.21	0.04	-0.05	0.05
V23	23	500	0.03	0.98	0.02	0.02	0.95	-2.62	3.03	5.65	0.18	0.15	0.04
V24	24	500	0.02	0.98	-0.03	0.02	0.97	-3.58	3.23	6.81	-0.02	0.28	0.04

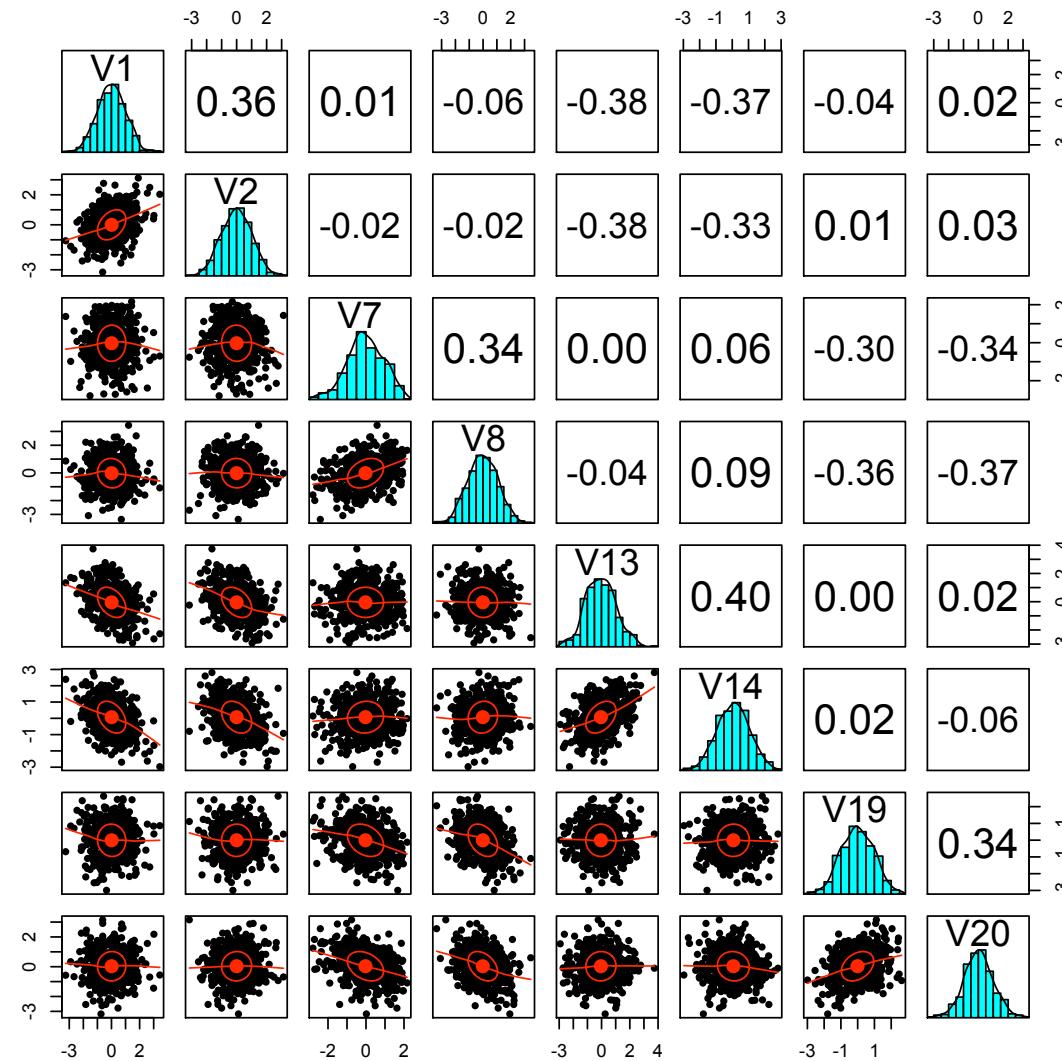
Continuous simple structured

```
> set.seed(42)
> simple <- sim.item(24)
> colnames(simple) <- paste("V", 1:24, sep = "")
> round(cor(simple)[, 1:12], 2)
```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	1.00	0.36	0.38	0.37	0.45	0.39	0.01	-0.06	-0.07	0.02	-0.01	-0.02
V2	0.36	1.00	0.37	0.36	0.40	0.40	-0.02	-0.02	-0.04	-0.01	-0.05	-0.07
V3	0.38	0.37	1.00	0.33	0.35	0.33	0.01	-0.02	0.06	0.05	0.09	-0.07
V4	0.37	0.36	0.33	1.00	0.39	0.43	-0.02	-0.03	0.01	0.05	0.00	0.01
V5	0.45	0.40	0.35	0.39	1.00	0.41	0.02	0.00	-0.01	0.04	0.00	0.00
V6	0.39	0.40	0.33	0.43	0.41	1.00	0.06	0.10	0.02	0.00	0.06	0.01
V7	0.01	-0.02	0.01	-0.02	0.02	0.06	1.00	0.34	0.29	0.29	0.34	0.37
V8	-0.06	-0.02	-0.02	-0.03	0.00	0.10	0.34	1.00	0.39	0.34	0.33	0.39
V9	-0.07	-0.04	0.06	0.01	-0.01	0.02	0.29	0.39	1.00	0.30	0.36	0.32
V10	0.02	-0.01	0.05	0.05	0.04	0.00	0.29	0.34	0.30	1.00	0.39	0.35
V11	-0.01	-0.05	0.09	0.00	0.00	0.06	0.34	0.33	0.36	0.39	1.00	0.38
V12	-0.02	-0.07	-0.07	0.01	0.00	0.01	0.37	0.39	0.32	0.35	0.38	1.00
V13	-0.38	-0.38	-0.39	-0.40	-0.41	-0.36	0.00	-0.04	0.02	-0.05	0.00	0.01
V14	-0.37	-0.33	-0.36	-0.42	-0.39	-0.30	0.06	0.09	0.07	0.01	0.02	0.03
V15	-0.38	-0.35	-0.29	-0.37	-0.32	-0.35	0.00	0.00	0.04	-0.06	0.03	0.00
V16	-0.38	-0.35	-0.36	-0.41	-0.42	-0.37	-0.03	-0.01	-0.01	-0.01	-0.02	0.04
V17	-0.37	-0.40	-0.32	-0.42	-0.39	-0.39	0.01	0.02	0.03	-0.02	0.04	0.06
V18	-0.37	-0.38	-0.37	-0.35	-0.39	-0.38	0.00	-0.03	0.08	-0.02	-0.01	-0.01
V19	-0.04	0.01	-0.01	-0.03	-0.02	-0.03	-0.30	-0.36	-0.29	-0.36	-0.33	-0.36
V20	0.02	0.03	-0.04	0.00	0.00	-0.01	-0.34	-0.37	-0.39	-0.35	-0.35	-0.33
V21	0.00	0.00	0.00	-0.05	0.02	-0.01	-0.34	-0.39	-0.38	-0.37	-0.31	-0.39
V22	0.05	0.03	-0.05	-0.03	0.03	-0.03	-0.35	-0.37	-0.35	-0.31	-0.34	-0.34
V23	-0.02	0.10	-0.04	0.03	0.03	0.01	-0.28	-0.37	-0.31	-0.36	-0.36	-0.37
V24	0.04	-0.01	-0.06	0.00	-0.03	-0.05	-0.33	-0.42	-0.34	-0.33	-0.37	-0.37

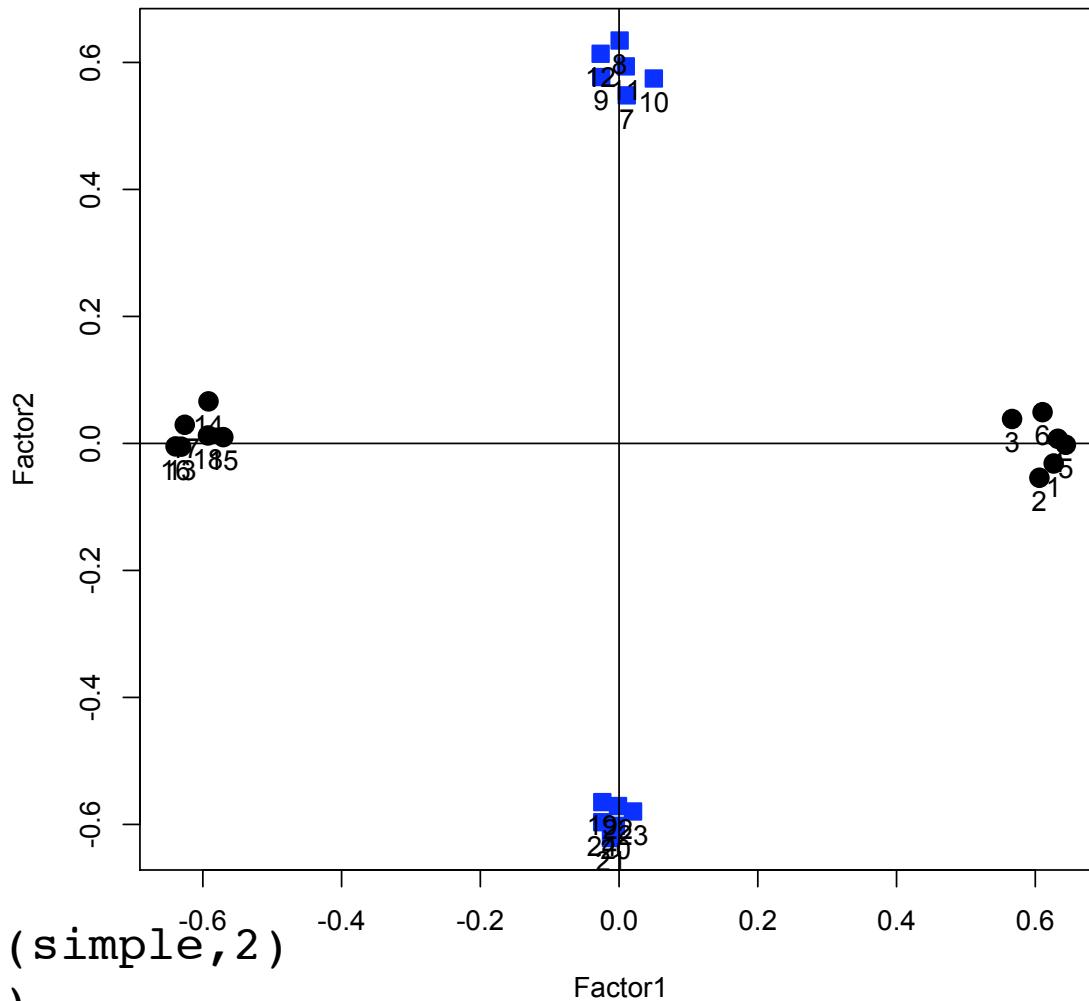
SPLOM

```
pairs.panels(simple[,c(1,2,7,8,13,14,19,20)])
```



Clear two factor structure

Cluster plot



```
> f2 <- factanal(simple, 2)
> factor.plot(f2)
```

Clear 2 factor solution

Call:

```
factanal(x = simple, factors = 2)
```

Uniquenesses:

V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12	V13	V14
V15	V16	V17	V18	V19	V20	V21	V22	V23	V24				
0.607	0.630	0.678	0.600	0.585	0.625	0.699	0.597	0.666	0.667	0.647	0.623	0.602	0.645
0.674	0.591	0.607	0.649	0.680	0.637	0.614	0.674	0.664	0.644				

Loadings:

	Factor1	Factor2	
V1	0.626		V13 -0.631
V2	0.606		V14 -0.592
V3	0.567		V15 -0.571
V4	0.632		V16 -0.639
V5	0.644		V17 -0.626
V6	0.610		V18 -0.593
V7	0.548		V19 -0.565
V8	0.635		V20 -0.602
V9	0.577		V21 -0.622
V10	0.575		V22 -0.571
V11	0.594		V23 -0.580
V12	0.614		V24 -0.597

Test of the hypothesis that 2 factors are sufficient.
The chi square statistic is 217.99 on 229 degrees of freedom.
The p-value is 0.689

	Factor1	Factor2
SS loadings	4.500	4.195
Proportion Var	0.188	0.175
Cumulative Var	0.188	0.362

Categorical items

- I. using the sim.item function with categorical=TRUE
 - A. This just rounds items to integer values
 - B. Setting other parameters to default values

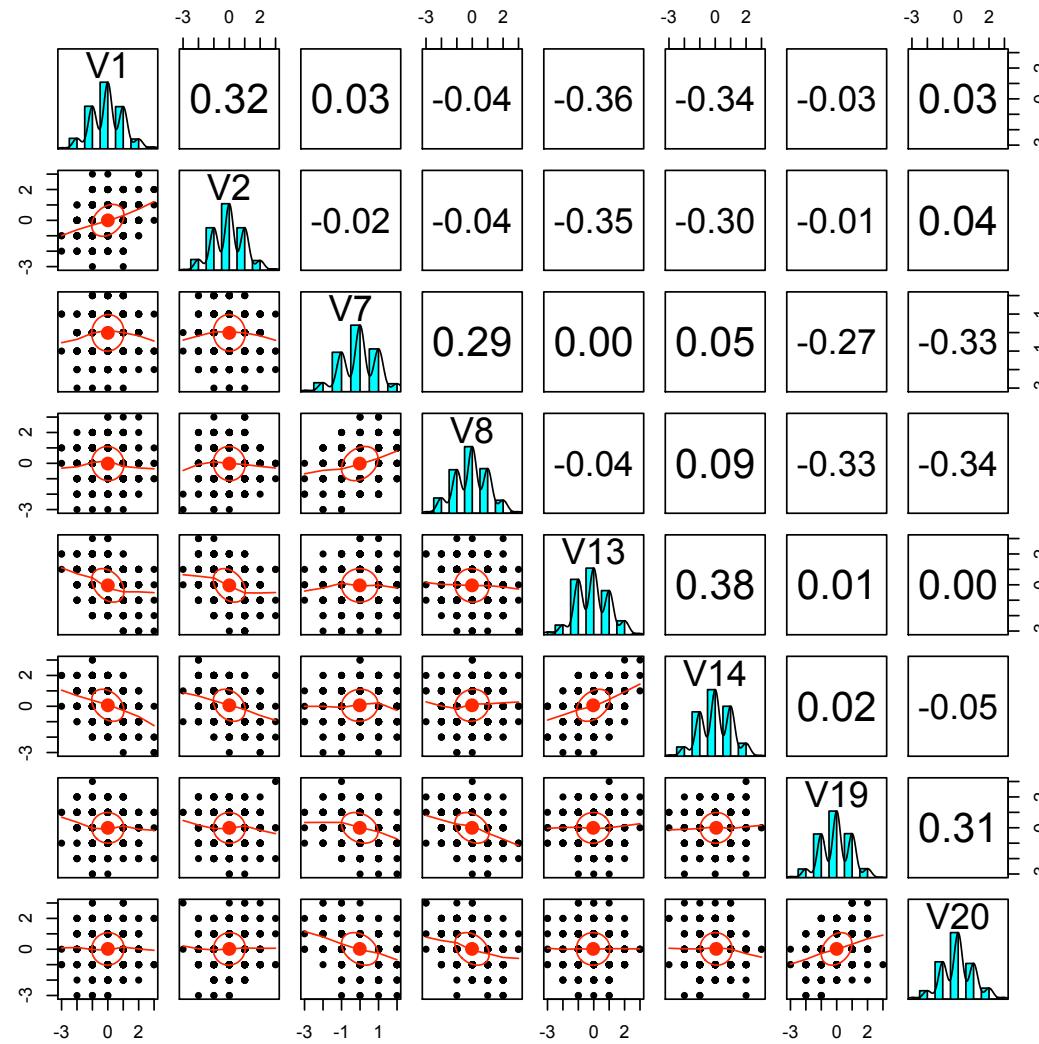
Categorical items

```
> set.seed(42)
> cate <- sim.item(24,categorical=TRUE)
> colnames(cate) <- paste("V",1:24,sep="")
> describe(cate)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	skew	kurtosis	se
V1	1	500	0.00	1.05	0	0.00	1.48	-3	3	6	0.06	0.03	0.05
V2	2	500	0.00	1.03	0	0.00	1.48	-3	3	6	0.05	-0.08	0.05
V3	3	500	-0.04	1.07	0	-0.06	1.48	-3	3	6	0.15	-0.23	0.05
V4	4	500	-0.04	1.07	0	-0.04	1.48	-3	3	6	0.11	-0.05	0.05
V5	5	500	-0.01	1.00	0	-0.01	1.48	-3	3	6	-0.03	0.04	0.04
V6	6	500	-0.04	1.13	0	-0.06	1.48	-3	3	6	0.02	-0.34	0.05
V7	7	500	-0.02	0.99	0	0.01	1.48	-3	2	5	-0.21	-0.09	0.04
V8	8	500	-0.02	1.11	0	-0.01	1.48	-3	3	6	-0.02	-0.22	0.05
V9	9	500	0.01	1.06	0	0.00	1.48	-3	3	6	0.04	-0.37	0.05
V10	10	500	0.00	1.01	0	0.02	1.48	-3	3	6	-0.14	-0.10	0.05
V11	11	500	-0.04	1.00	0	-0.06	1.48	-3	3	6	0.10	-0.01	0.04
V12	12	500	-0.08	1.02	0	-0.08	1.48	-3	3	6	0.08	-0.22	0.05
V13	13	500	-0.04	1.09	0	-0.06	1.48	-3	3	6	0.03	-0.08	0.05
V14	14	500	0.06	1.04	0	0.06	1.48	-3	3	6	-0.06	-0.22	0.05
V15	15	500	0.02	1.05	0	0.01	1.48	-3	3	6	0.05	-0.30	0.05
V16	16	500	0.00	1.08	0	-0.02	1.48	-3	3	6	0.15	-0.05	0.05
V17	17	500	-0.03	1.08	0	-0.03	1.48	-3	3	6	-0.08	-0.01	0.05
V18	18	500	0.04	1.08	0	0.05	1.48	-3	3	6	-0.14	-0.11	0.05
V19	19	500	-0.01	0.99	0	0.00	1.48	-3	3	6	-0.07	-0.15	0.04
V20	20	500	0.02	1.05	0	0.01	1.48	-3	3	6	0.04	0.22	0.05
V21	21	500	-0.02	1.11	0	-0.03	1.48	-3	3	6	0.09	-0.20	0.05
V22	22	500	0.04	1.05	0	0.02	1.48	-3	3	6	0.00	-0.13	0.05
V23	23	500	0.02	1.02	0	-0.01	1.48	-3	3	6	0.17	0.03	0.05
V24	24	500	0.05	1.02	0	0.04	1.48	-3	3	6	-0.01	-0.10	0.05

SPLOM (selected variables)

```
pairs.panels(cate[,c(1,2,7,8,13,14,19,20)])
```



Correlations similar to original

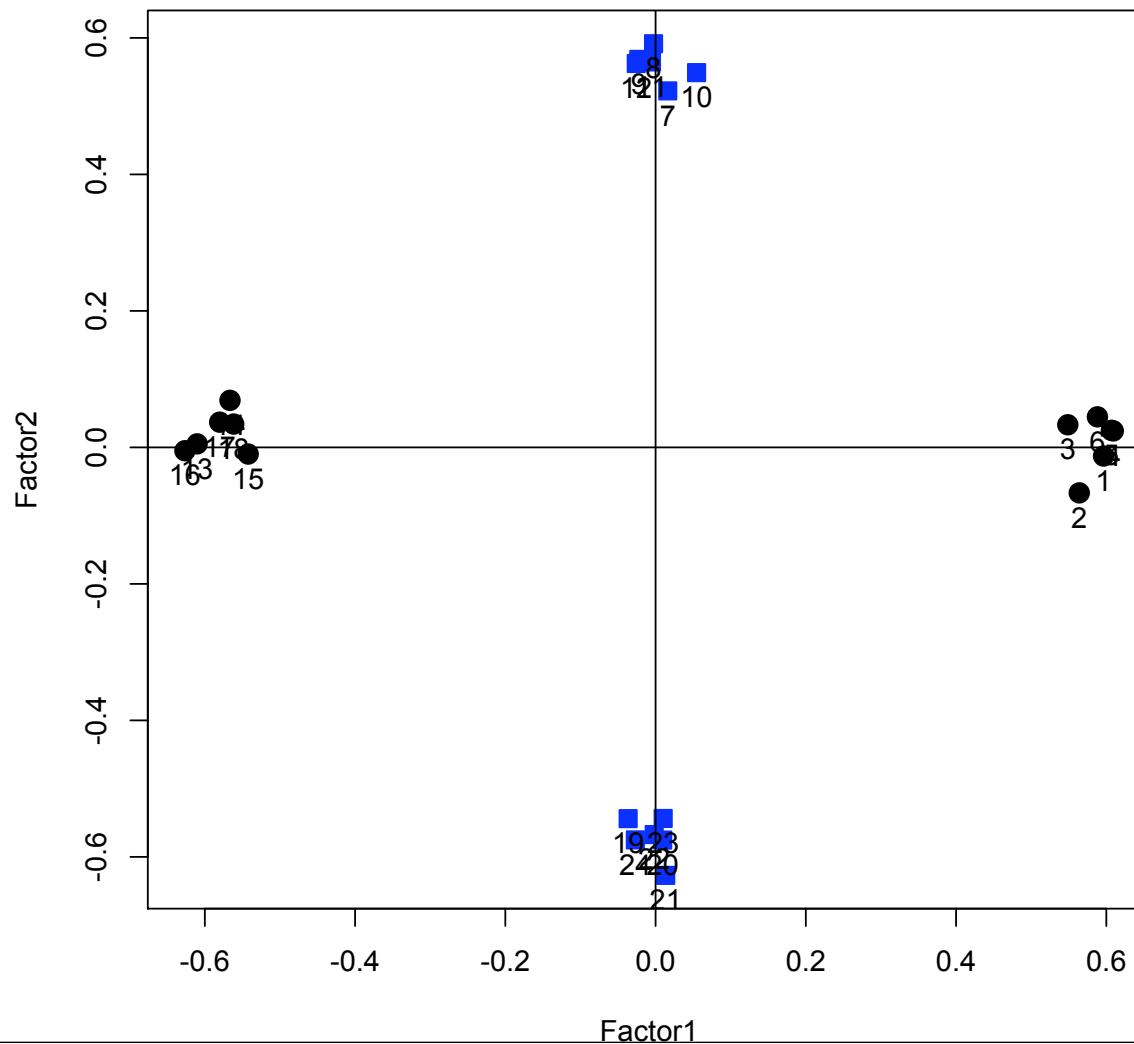
```
> colnames(cate) <- paste("V",1:24,sep=" ")
> round(cor(cate)[,1:12],2)
```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	1.00	0.32	0.35	0.33	0.42	0.35	0.03	-0.04	-0.05	0.03	0.01	-0.03
V2	0.32	1.00	0.33	0.33	0.34	0.35	-0.02	-0.04	-0.05	0.00	-0.07	-0.05
V3	0.35	0.33	1.00	0.31	0.32	0.30	0.01	-0.01	0.07	0.06	0.07	-0.10
V4	0.33	0.33	0.31	1.00	0.35	0.40	0.01	-0.02	0.03	0.05	0.01	0.02
V5	0.42	0.34	0.32	0.35	1.00	0.39	0.05	0.01	0.01	0.07	0.00	0.01
V6	0.35	0.35	0.30	0.40	0.39	1.00	0.06	0.09	0.03	0.00	0.04	0.02
V7	0.03	-0.02	0.01	0.01	0.05	0.06	1.00	0.29	0.27	0.25	0.33	0.32
V8	-0.04	-0.04	-0.01	-0.02	0.01	0.09	0.29	1.00	0.37	0.29	0.28	0.33
V9	-0.05	-0.05	0.07	0.03	0.01	0.03	0.27	0.37	1.00	0.32	0.33	0.26
V10	0.03	0.00	0.06	0.05	0.07	0.00	0.25	0.29	0.32	1.00	0.36	0.31
V11	0.01	-0.07	0.07	0.01	0.00	0.04	0.33	0.28	0.33	0.36	1.00	0.34
V12	-0.03	-0.05	-0.10	0.02	0.01	0.02	0.32	0.33	0.26	0.31	0.34	1.00
V13	-0.36	-0.35	-0.37	-0.38	-0.36	-0.31	0.00	-0.04	0.04	-0.06	0.03	0.02
V14	-0.34	-0.30	-0.32	-0.37	-0.36	-0.29	0.05	0.09	0.09	0.02	0.01	0.03
V15	-0.36	-0.34	-0.29	-0.34	-0.26	-0.30	-0.02	-0.02	0.06	-0.07	0.03	-0.02
V16	-0.34	-0.32	-0.34	-0.37	-0.38	-0.36	-0.01	0.00	-0.02	-0.02	0.02	0.03
V17	-0.33	-0.33	-0.28	-0.40	-0.32	-0.34	0.03	0.03	0.03	0.01	0.04	0.05
V18	-0.33	-0.34	-0.33	-0.31	-0.36	-0.36	0.02	0.01	0.05	-0.02	0.02	0.00
V19	-0.03	-0.01	0.00	-0.02	-0.05	-0.03	-0.27	-0.33	-0.27	-0.33	-0.32	-0.32
V20	0.03	0.04	-0.02	-0.01	0.02	-0.01	-0.33	-0.34	-0.36	-0.32	-0.31	-0.30
V21	0.00	0.04	0.02	-0.05	0.01	0.00	-0.31	-0.36	-0.38	-0.35	-0.31	-0.37
V22	0.02	0.06	-0.04	-0.04	0.01	-0.02	-0.34	-0.35	-0.34	-0.29	-0.32	-0.30
V23	-0.04	0.09	-0.06	0.03	0.03	0.01	-0.24	-0.33	-0.28	-0.33	-0.31	-0.33
V24	0.01	-0.02	-0.06	-0.01	-0.06	-0.04	-0.31	-0.40	-0.33	-0.29	-0.32	-0.34

Factor structure similar

```
f2 <- factanal(cate, 2)  
factor.plot(f2)
```

Cluster plot



```

Call:
factanal(x = cate, factors = 2)
Uniquenesses:
      V1     V2     V3     V4     V5     V6     V7     V8     V9     V10    V11    V12    V13    V14    V15    V16    V17
V18   V19   V20   V21   V22   V23   V24
0.644 0.677 0.698 0.628 0.631 0.652 0.727 0.650 0.676 0.695 0.681 0.683 0.628 0.674 0.706 0.607 0.662
0.683 0.703 0.669 0.607 0.678 0.704 0.669
Loadings:
  Factor1 Factor2
V1     0.596
V2     0.564
V3     0.549
V4     0.609
V5     0.607
V6     0.588
V7     0.522
V8     0.591
V9     0.569
V10    0.549
V11    0.565
V12    0.563
V13   -0.610
V14   -0.567
V15   -0.542
V16   -0.626
V17   -0.580
V18   -0.562
V19     -0.544
V20   -0.575
V21   -0.627
V22   -0.567
V23   -0.544
V24   -0.575
  Factor1 Factor2
SS loadings   4.099   3.867
Proportion Var  0.171   0.161
Cumulative Var  0.171   0.332
Test of the hypothesis that 2 factors are sufficient.
The chi square statistic is 235.68 on 229 degrees of freedom.
The p-value is 0.367

```

Chi Squares are similar

Skewed items

- I. Simulate 24 items representing two factors
- II. 6 positive and negative on each factor
- III. Items differ in skew
 - A. skew created by truncating items to positive values
 - B. items differing in base value (add .5 to y coordinate before cutting)
 - C. > `set.seed(42)`

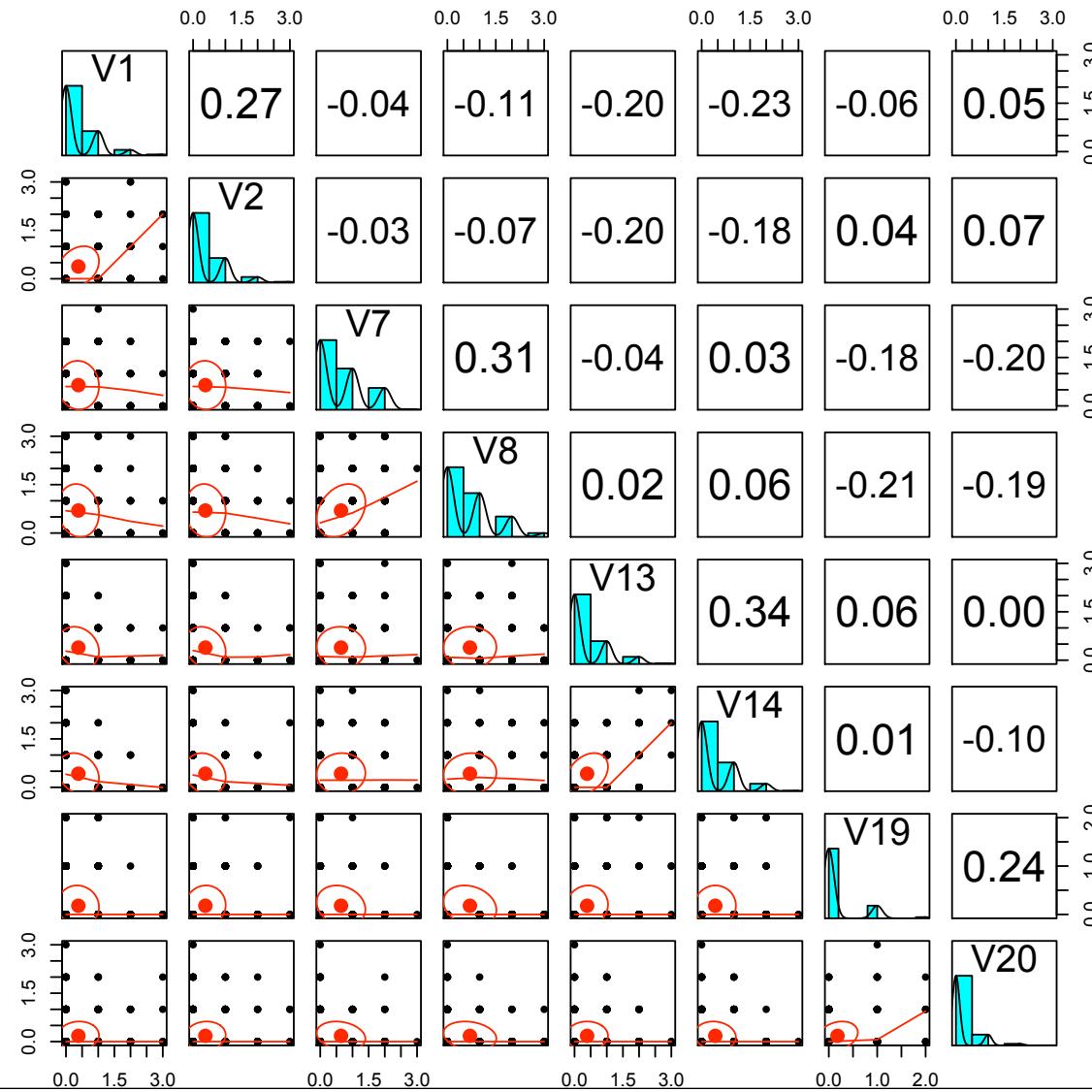
```
> simp <- sim.item(24,ybias=.5,categorical=TRUE,low=0,truncate=TRUE)
> colnames(simp) <- paste("V",1:24,sep="")
```

Descriptive statistics

```
> describe(simp)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	skew	kurtosis	se
V1	1	500	0.38	0.64	0.0	0.26	0.00	0	3	3	1.65	2.37	0.03
V2	2	500	0.38	0.63	0.0	0.26	0.00	0	3	3	1.61	2.18	0.03
V3	3	500	0.38	0.64	0.0	0.25	0.00	0	3	3	1.63	2.09	0.03
V4	4	500	0.38	0.64	0.0	0.24	0.00	0	3	3	1.70	2.44	0.03
V5	5	500	0.36	0.60	0.0	0.25	0.00	0	3	3	1.62	2.29	0.03
V6	6	500	0.41	0.67	0.0	0.26	0.00	0	3	3	1.44	1.05	0.03
V7	7	500	0.64	0.76	0.0	0.55	0.00	0	3	3	0.72	-0.81	0.03
V8	8	500	0.70	0.82	0.0	0.59	0.00	0	3	3	0.91	-0.08	0.04
V9	9	500	0.69	0.81	0.0	0.58	0.00	0	3	3	0.92	0.01	0.04
V10	10	500	0.67	0.77	0.5	0.56	0.74	0	3	3	0.89	0.00	0.03
V11	11	500	0.62	0.80	0.0	0.49	0.00	0	3	3	1.18	0.77	0.04
V12	12	500	0.65	0.77	0.0	0.54	0.00	0	3	3	0.95	0.12	0.03
V13	13	500	0.39	0.65	0.0	0.26	0.00	0	3	3	1.54	1.62	0.03
V14	14	500	0.42	0.64	0.0	0.30	0.00	0	3	3	1.33	0.96	0.03
V15	15	500	0.41	0.64	0.0	0.28	0.00	0	3	3	1.45	1.45	0.03
V16	16	500	0.40	0.67	0.0	0.26	0.00	0	3	3	1.65	2.14	0.03
V17	17	500	0.38	0.64	0.0	0.26	0.00	0	3	3	1.51	1.42	0.03
V18	18	500	0.41	0.65	0.0	0.29	0.00	0	3	3	1.38	1.03	0.03
V19	19	500	0.18	0.42	0.0	0.08	0.00	0	2	2	2.21	4.21	0.02
V20	20	500	0.18	0.45	0.0	0.07	0.00	0	3	3	2.65	7.34	0.02
V21	21	500	0.18	0.47	0.0	0.06	0.00	0	3	3	2.67	7.16	0.02
V22	22	500	0.21	0.50	0.0	0.10	0.00	0	3	3	2.39	5.43	0.02
V23	23	500	0.19	0.47	0.0	0.07	0.00	0	3	3	2.63	6.87	0.02
V24	24	500	0.17	0.44	0.0	0.06	0.00	0	3	3	2.70	7.67	0.02

Skew differentially affects r



correlations with skew

```
> round(cor(simp)[,1:12],2)
```

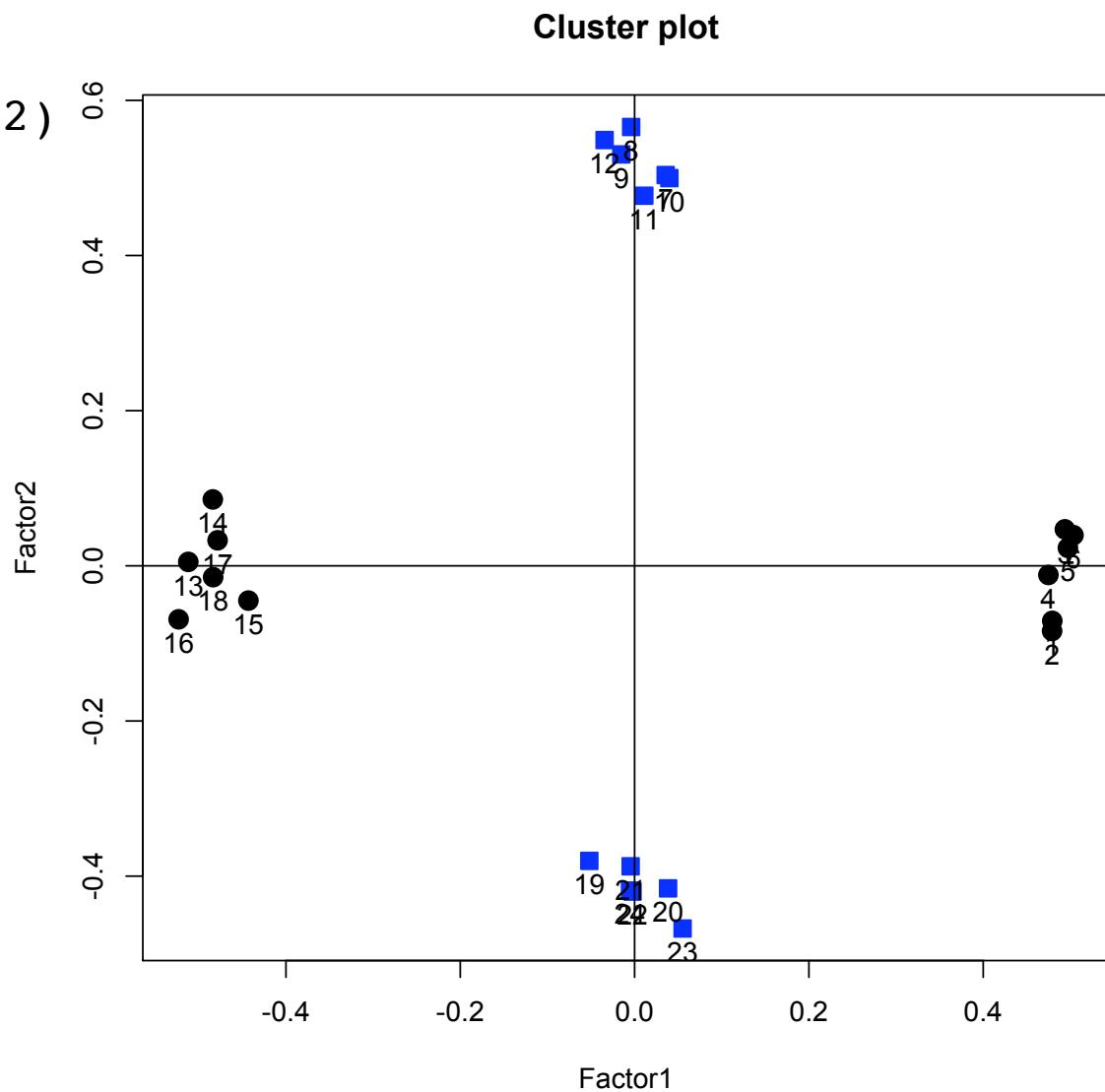
	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	1.00	0.27	0.26	0.30	0.33	0.29	-0.04	-0.11	-0.08	-0.01	-0.02	-0.05
V2	0.27	1.00	0.28	0.25	0.28	0.33	-0.03	-0.07	-0.07	0.01	-0.05	-0.06
V3	0.26	0.28	1.00	0.27	0.31	0.26	0.01	0.01	0.10	0.07	0.09	-0.07
V4	0.30	0.25	0.27	1.00	0.21	0.29	-0.05	-0.05	0.06	0.05	-0.02	-0.05
V5	0.33	0.28	0.31	0.21	1.00	0.30	0.05	0.01	0.00	0.03	0.03	0.01
V6	0.29	0.33	0.26	0.29	0.30	1.00	0.05	0.05	0.01	-0.01	0.07	-0.01
V7	-0.04	-0.03	0.01	-0.05	0.05	0.05	1.00	0.31	0.26	0.24	0.26	0.32
V8	-0.11	-0.07	0.01	-0.05	0.01	0.05	0.31	1.00	0.34	0.31	0.26	0.33
V9	-0.08	-0.07	0.10	0.06	0.00	0.01	0.26	0.34	1.00	0.25	0.32	0.36
V10	-0.01	0.01	0.07	0.05	0.03	-0.01	0.24	0.31	0.25	1.00	0.28	0.27
V11	-0.02	-0.05	0.09	-0.02	0.03	0.07	0.26	0.26	0.32	0.28	1.00	0.30
V12	-0.05	-0.06	-0.07	-0.05	0.01	-0.01	0.32	0.33	0.36	0.27	0.30	1.00
V13	-0.20	-0.20	-0.23	-0.24	-0.24	-0.19	-0.04	0.02	0.01	-0.02	0.01	0.00
V14	-0.23	-0.18	-0.21	-0.20	-0.22	-0.17	0.03	0.06	0.06	0.07	0.01	0.01
V15	-0.19	-0.19	-0.20	-0.18	-0.12	-0.23	-0.04	-0.07	0.02	-0.07	0.02	-0.03
V16	-0.18	-0.18	-0.24	-0.18	-0.23	-0.23	-0.08	-0.08	-0.02	-0.01	-0.05	0.03
V17	-0.17	-0.22	-0.16	-0.25	-0.22	-0.19	-0.04	0.00	0.01	-0.01	0.02	0.03
V18	-0.17	-0.18	-0.22	-0.20	-0.21	-0.23	-0.03	-0.03	0.08	-0.04	0.03	-0.03
V19	-0.06	0.04	-0.01	-0.04	-0.06	0.00	-0.18	-0.21	-0.15	-0.20	-0.11	-0.20
V20	0.05	0.07	-0.01	0.03	0.05	0.00	-0.20	-0.19	-0.16	-0.18	-0.16	-0.17
V21	0.04	-0.02	-0.02	-0.01	-0.03	-0.01	-0.19	-0.20	-0.22	-0.16	-0.12	-0.20
V22	0.04	0.00	-0.02	-0.01	0.03	0.02	-0.16	-0.23	-0.21	-0.23	-0.14	-0.23
V23	-0.03	0.08	-0.02	0.02	0.02	-0.02	-0.22	-0.22	-0.18	-0.22	-0.22	-0.19
V24	0.00	0.04	0.00	-0.04	-0.02	-0.05	-0.19	-0.19	-0.17	-0.22	-0.18	-0.18

Alternative models

- I. Factor model (2-4 factors)
- II. Cluster model (most appropriate for items)
- III. sem model based upon factor model
- IV.sem model with correlated factors
- V.sem model with bi-factor solution

Clear 2 dimensional structure

```
> f2 <- factanal(simp,2)
> factor.plot(f2)
```



Call:

```
factanal(x = simp, factors = 2)
```

Uniquenesses:

V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11
V12	V13	V14	V15	V16	V17	V18	V19	V20	V21	V22
0.765	0.763	0.754	0.774	0.752	0.745	0.745	0.680	0.719	0.749	0.772
0.698	0.738	0.758	0.802	0.722	0.770	0.766	0.853	0.826	0.850	0.824
V23	V24		V13	-0.512						
0.778	0.825		V14	-0.484						
			V15	-0.443						
Loadings:			V16	-0.523						

Factor1 Factor2 V17 -0.478

V1 0.479 V18 -0.484

V2 0.479 V19 -0.380

V3 0.494 V20 -0.416

V4 0.475 V21 -0.387

V5 0.497 V22 -0.420

V6 0.504 V23 -0.468

V7 0.504 V24 -0.419

V8 0.566

V9 0.530

Factor1 Factor2

V10 0.500 SS loadings 2.87 2.703

V11 0.477 Proportion Var 0.12 0.113

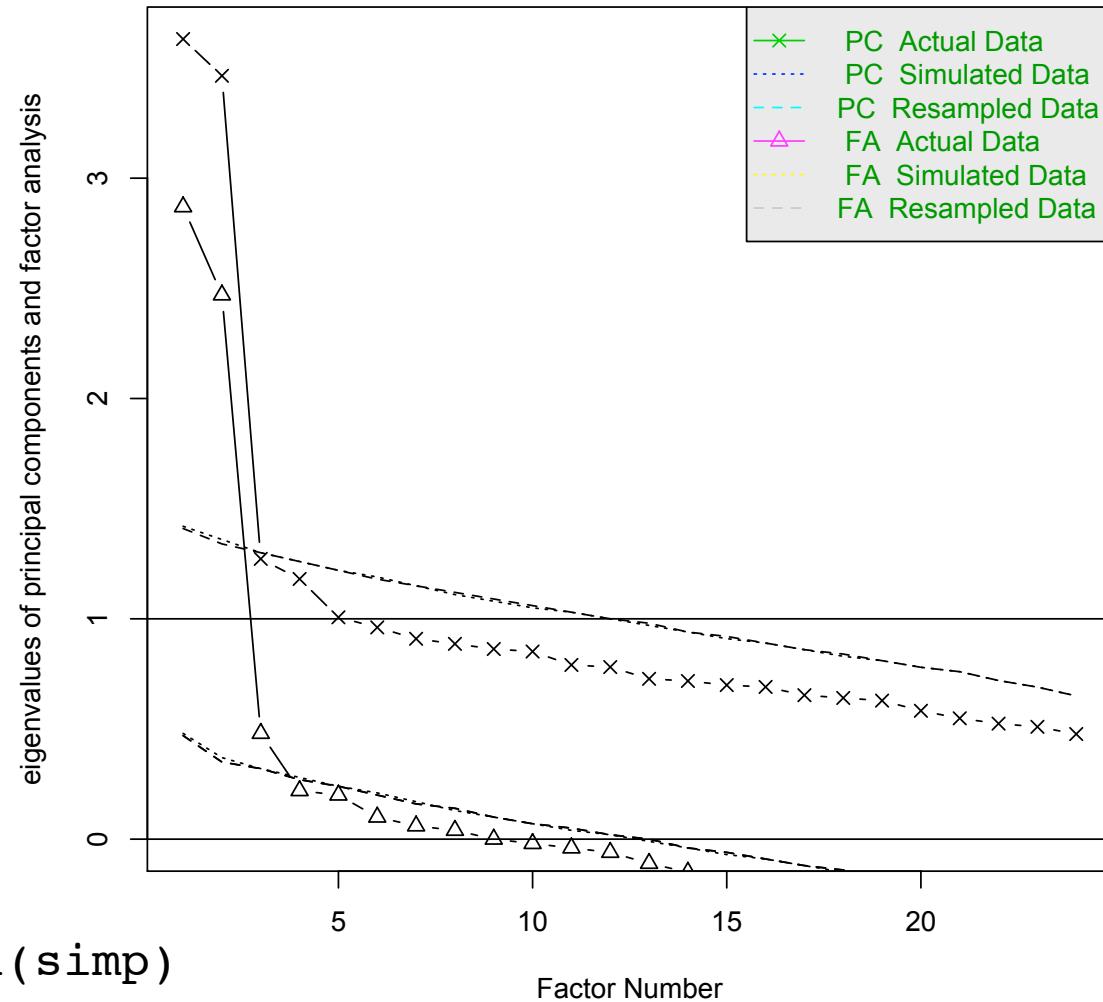
V12 0.549 Cumulative Var 0.12 0.232

Test of the hypothesis that
2 factors are sufficient.
The chi square statistic is
350.52 on 229 degrees of
freedom.

The p-value is 4.06e-07

Maybe 3 factors?

Parallel Analysis Scree Plots



Call:
 factanal(x = simp, factors = 3)

Uniquenesses:

V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12	V13	V14	V15	V16	V17
V18	V19	V20	V21	V22	V23	V24										
0.666	0.733	0.725	0.747	0.719	0.708	0.736	0.656	0.721	0.749	0.771	0.690	0.712	0.740	0.763	0.649	0.744
0.703	0.849	0.823	0.848	0.823	0.735	0.805										

Loadings:

	Factor1	Factor2	Factor3
V1	-0.143	0.557	
V2	-0.219	0.462	
V3	-0.238	0.463	
V4	-0.223	0.452	
V5	-0.234	0.475	
V6	-0.234	0.484	
V7	0.500	-0.102	
V8	0.563	-0.109	-0.123
V9	0.528		
V10	0.500		
V11	0.478		
V12	0.543	-0.110	
V13	0.478	-0.245	
V14	0.439	-0.244	
V15	0.452	-0.177	
V16	0.556	-0.195	
V17	0.454	-0.220	
V18	0.515	-0.178	
V19	-0.383		
V20	-0.418		
V21	-0.389		
V22	-0.418		
V23	-0.482	-0.161	
V24	-0.427		

Test of the hypothesis that 3 factors are sufficient.
 The chi square statistic is 262.01 on 207 degrees of freedom.
 The p-value is 0.00574

	Factor1	Factor2	Factor3
SS loadings	2.711	1.751	1.722
Proportion Var	0.113	0.073	0.072
Cumulative Var	0.113	0.186	0.258

3 factors

```

> f4 <- factanal(simp,4)
> f4
Call:
factanal(x = simp, factors = 4)
Uniquenesses:
   V1    V2    V3    V4    V5    V6    V7    V8    V9    V10   V11   V12   V13   V14   V15   V16   V17
V18   V19   V20   V21   V22   V23   V24
0.667 0.731 0.712 0.747 0.709 0.706 0.740 0.647 0.598 0.755 0.731 0.660 0.716 0.747 0.745 0.650 0.750
0.674 0.806 0.731 0.843 0.809 0.649 0.758
Loadings:
  Factor1 Factor2 Factor3 Factor4
V1    0.541 -0.129 -0.143
V2    0.472           -0.199
V3    0.492           -0.201
V4    0.461           -0.203
V5    0.499           -0.200
V6    0.497           -0.211
V7          0.421      -0.272
V8          0.523      -0.254
V9          0.615      -0.127
V10         0.388      -0.304
V11         0.477      -0.184
V12         0.537      -0.214
V13     -0.269      0.459
V14     -0.272      0.413
V15     -0.174      0.471
V16     -0.219      0.546
V17     -0.252      0.427
V18     -0.172      0.544
V19          -0.158     0.407
V20          -0.139     0.495
V21          -0.242     0.314
V22          -0.251     0.350
V23     -0.176 -0.129     0.549
V24     -0.174           0.456
  Factor1 Factor2 Factor3 Factor4
SS loadings  1.818   1.757   1.655   1.492
Proportion Var 0.076   0.073   0.069   0.062
Cumulative Var 0.076   0.149   0.218   0.280

```

Four factors are messy

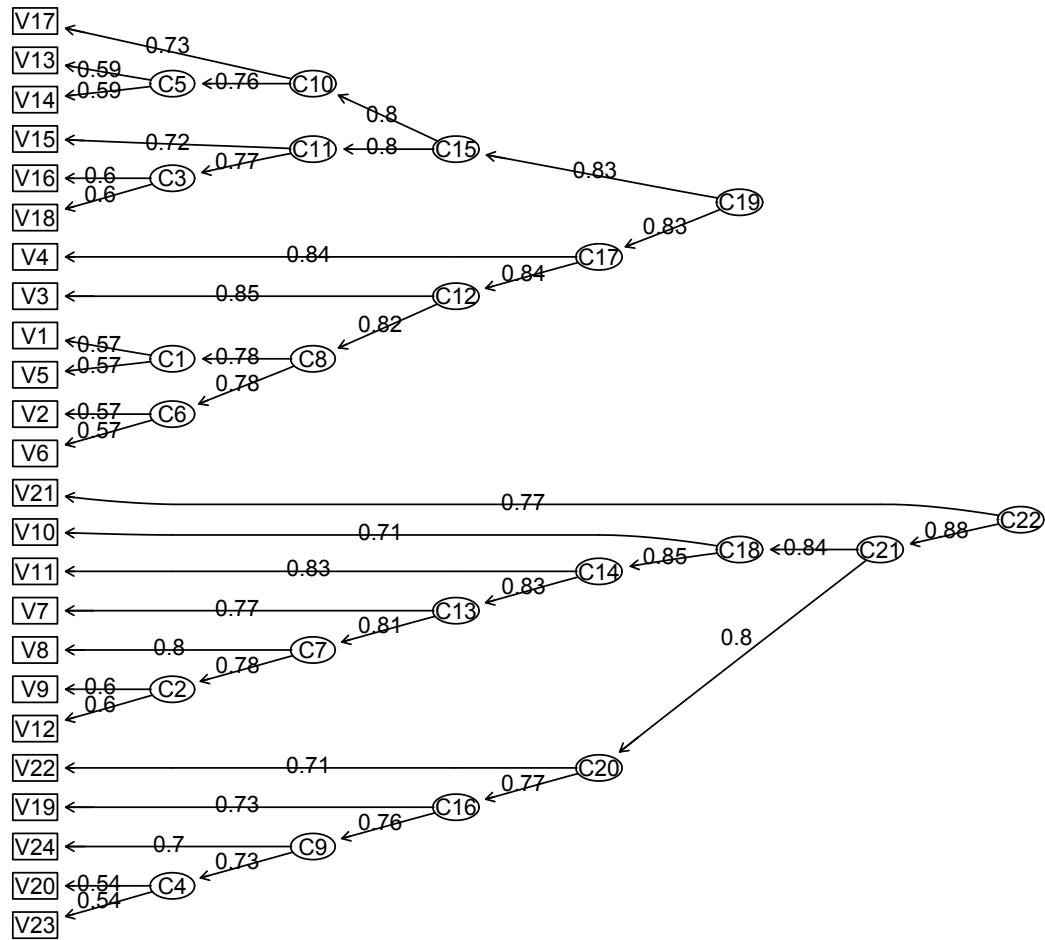
Test of the hypothesis that 4 factors are sufficient.
The chi square statistic is 196.54 on 186 degrees of freedom.
The p-value is 0.284

Item Cluster analysis (ICLUST)

- I. Hierarchical cluster analysis meant for items
 - A. Find most similar item pair
 - B. Combine them
 - C. Repeat steps A,B until coefficients alpha and beta do not increase

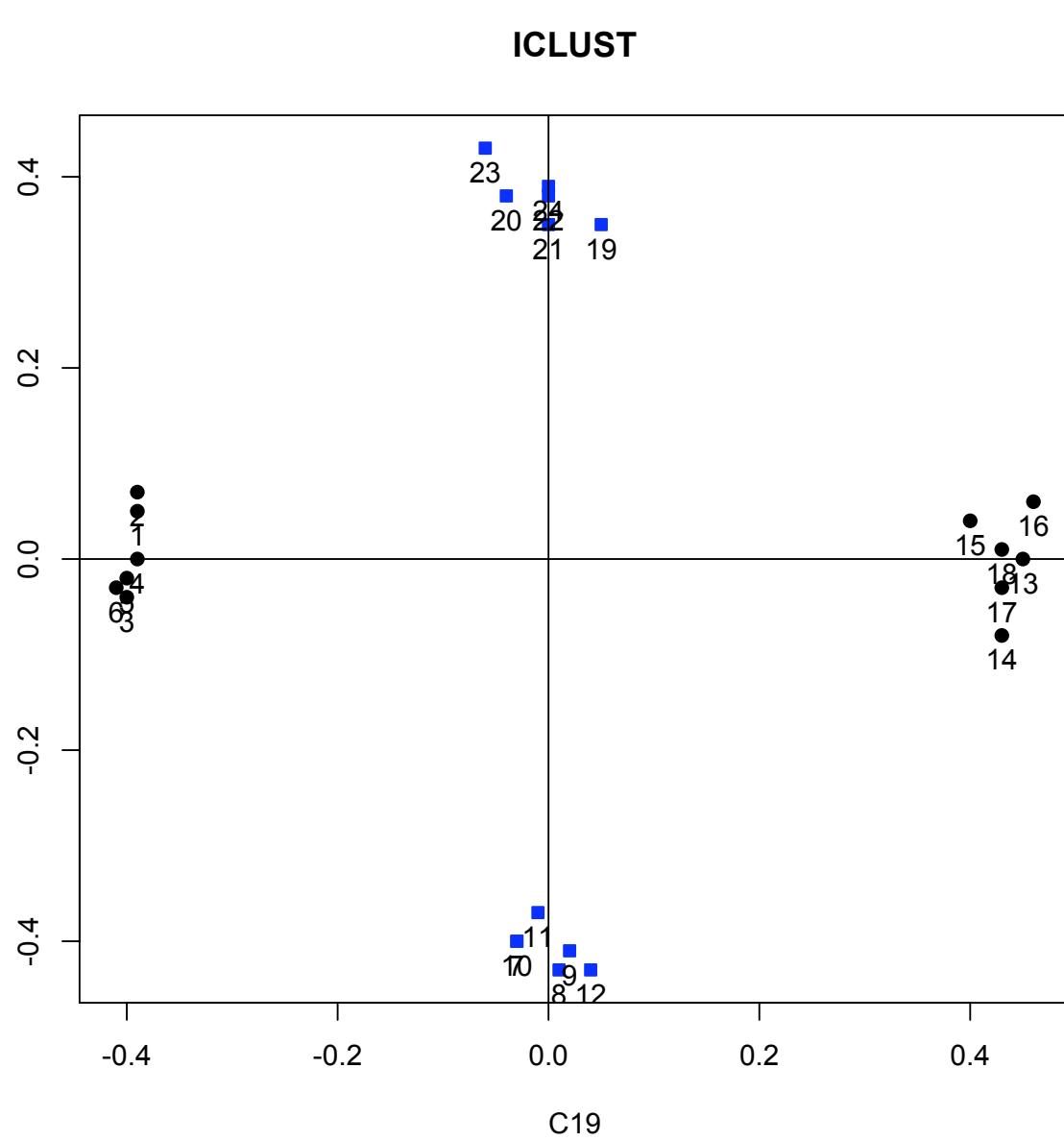
ICLUST shows 2 with sub structure

ICLUST



```
> ic <- ICLUST(simp)
```

Items match factoring



2 clusters

```
> summary(ic)
```

ICLUST

Purified Alpha:

C19	C22
0.79	0.77

Original Beta:

C19	C22
0.67	0.51

Cluster size:

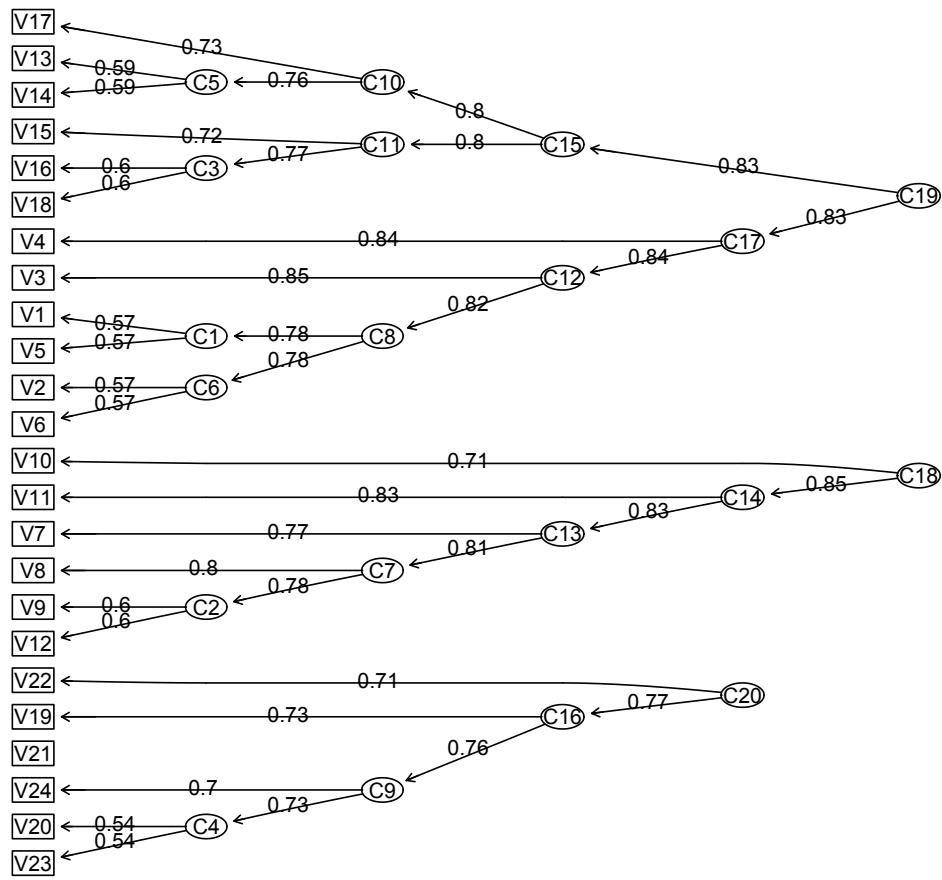
C19	C22
12	12

Purified scale intercorrelations:

	C19	C22
C19	1.00	-0.01
C22	-0.01	1.00

4 clusters

ICLUST

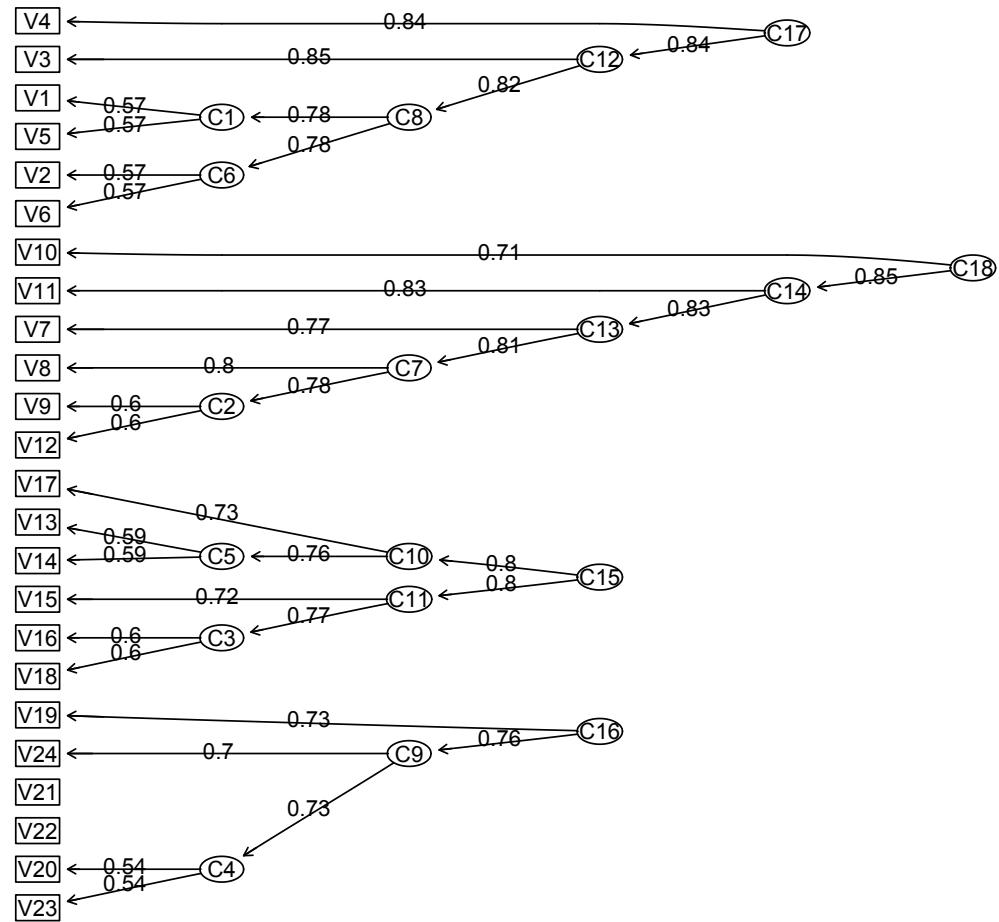


```
ic4 <- ICLUST(simp, 4)
```

Note how item 21
did not cluster

6 clusters

ICLUST



```
> ic6 <- ICLUST(simp,6)
> summary(ic6)
ICLUST
Purified Alpha:
  C15   C17   C18   V22   V21   C16
0.70  0.70  0.71  1.00  1.00  0.58
Original Beta:
  C15   C17   C18   V22   V21   C16
0.63  0.57  0.57    NA    NA  0.48
```

4 clusters
+ 2 items

Custer size:

C15	C17	C18	V22	V21	C16
6	6	6	1	1	4

Purified scale intercorrelations:

	C15	C17	C18	V22	V21	C16
C15	1.00	-0.50	-0.01	0.02	-0.01	-0.03
C17	-0.50	1.00	-0.01	0.02	-0.01	0.00
C18	-0.01	-0.01	1.00	-0.31	-0.29	-0.44
V22	0.02	0.02	-0.31	1.00	0.21	0.30
V21	-0.01	-0.01	-0.29	0.21	1.00	0.28
C16	-0.03	0.00	-0.44	0.30	0.28	1.00

```
> ic4 <- ICLUST(simp,4)
> summary(ic4)
ICLUST
```

4 clusters

Purified Alpha:

C19	C20	V21	C13
0.79	0.71	1.00	0.47

Original Beta:

C19	C20	V21	C13
0.64	0.53	NA	0.43

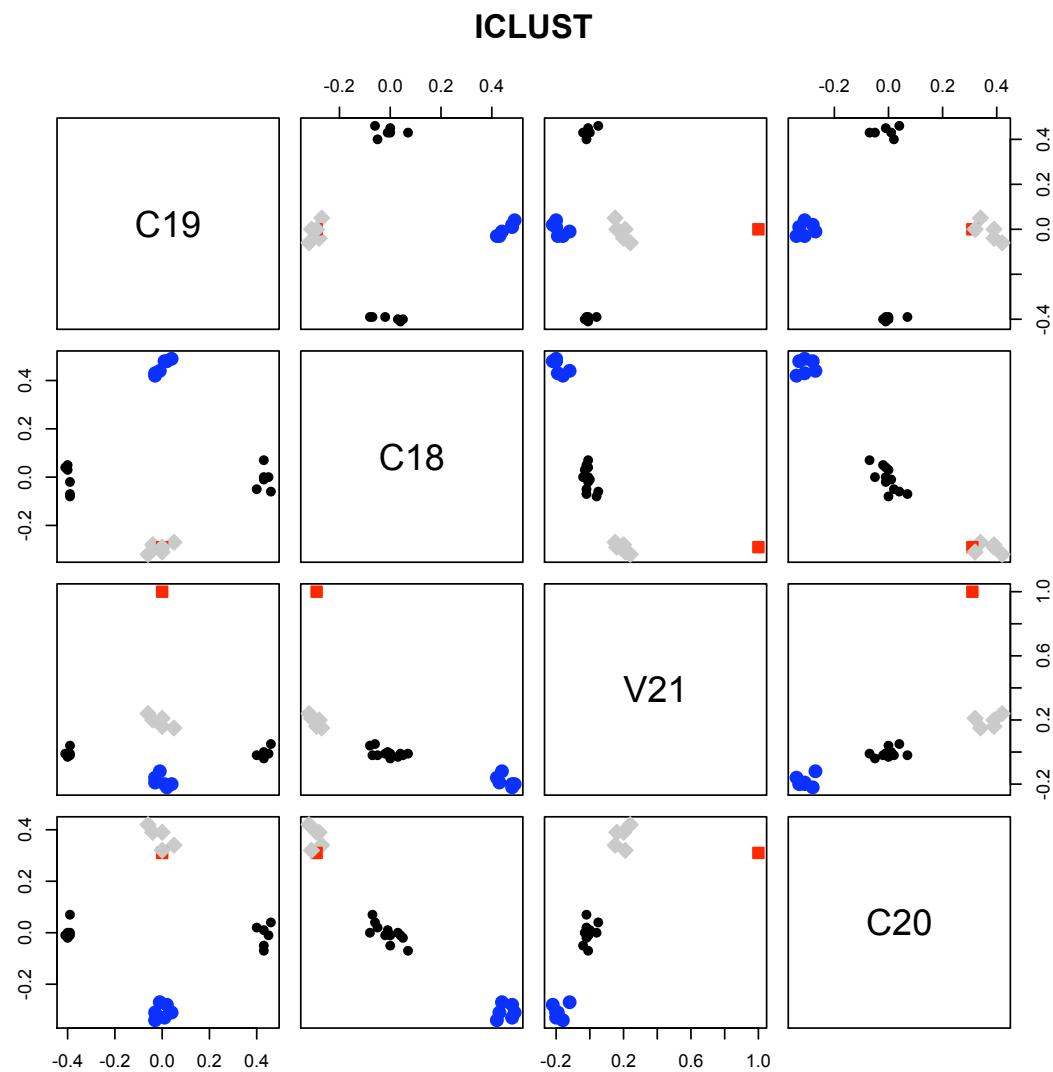
Cluster size:

C19	C20	V21	C13
12	8	1	3

Purified scale intercorrelations:

	C19	C20	V21	C13
C19	1.00	0.07	-0.03	0.10
C20	0.07	1.00	0.28	0.42
V21	-0.03	0.28	1.00	0.18
C13	0.10	0.42	0.18	1.00

Clusters are overlapping



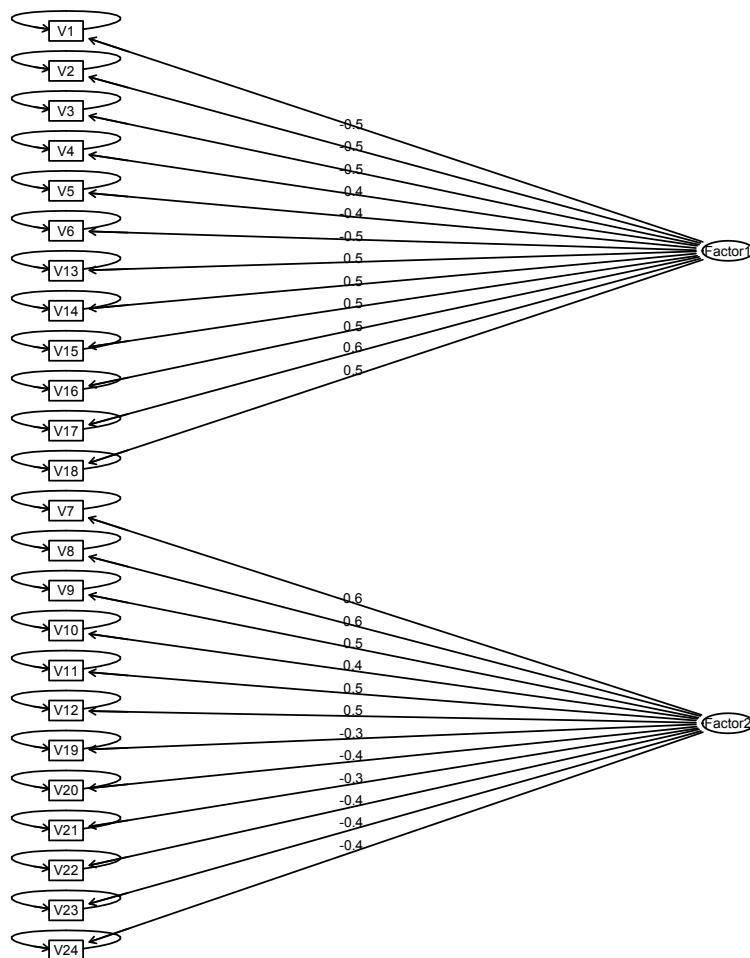
Sem for alternative sets

- I. First do it with the continuous data set
- II. Then the categorical
- III. Then the skewed categorical

Try sem for 2 factors

```
mod2 <- structure.graph(f2)
```

Structural model



Model

```
> mod2
      Path          Parameter Value [ 1, ] "Factor1->V1" "F1V1"     NA [ 2, ] "Factor1->V2" "F1V2"     NA [ 3, ] "Factor1->V3" "F1V3"     NA [ 4, ] "Factor1->V4" "F1V4"     NA [ 5, ] "Factor1->V5" "F1V5"     NA [ 6, ] "Factor1->V6" "F1V6"     NA [ 7, ] "Factor2->V7" "F2V7"     NA [ 8, ] "Factor2->V8" "F2V8"     NA [ 9, ] "Factor2->V9" "F2V9"     NA [10, ] "Factor2->V10" "F2V10"    NA [11, ] "Factor2->V11" "F2V11"    NA [12, ] "Factor2->V12" "F2V12"    NA [13, ] "Factor1->V13" "F1V13"    NA [14, ] "Factor1->V14" "F1V14"    NA [15, ] "Factor1->V15" "F1V15"    NA [16, ] "Factor1->V16" "F1V16"    NA [17, ] "Factor1->V17" "F1V17"    NA [18, ] "Factor1->V18" "F1V18"    NA [19, ] "Factor2->V19" "F2V19"    NA [20, ] "Factor2->V20" "F2V20"    NA [21, ] "Factor2->V21" "F2V21"    NA [22, ] "Factor2->V22" "F2V22"    NA [23, ] "Factor2->V23" "F2V23"    NA [24, ] "Factor2->V24" "F2V24"    NA [ 5, ] "V1<->V1"   NA [ 6, ] "V2<->V2"   NA [ 7, ] "V3<->V3"   NA [ 8, ] "V4<->V4"   NA [ 9, ] "V5<->V5"   NA [10, ] "V6<->V6"  NA [11, ] "V7<->V7"  NA [12, ] "V8<->V8"  NA [13, ] "V9<->V9"  NA [14, ] "V10<->V10" NA [15, ] "V11<->V11" NA [16, ] "V12<->V12" NA [17, ] "V13<->V13" NA [18, ] "V14<->V14" NA [19, ] "V15<->V15" NA [20, ] "V16<->V16" NA [21, ] "V17<->V17" NA [22, ] "V18<->V18" NA [23, ] "V19<->V19" NA [24, ] "V20<->V20" NA [25, ] "V21<->V21" NA [26, ] "V22<->V22" NA [27, ] "V23<->V23" NA [28, ] "V24<->V24" NA [29, ] "Factor1<->Factor1" NA [30, ] "Factor2<->Factor2" NA [31, ] "x1e"      NA [32, ] "x2e"      NA [33, ] "x3e"      NA [34, ] "x4e"      NA [35, ] "x5e"      NA [36, ] "x6e"      NA [37, ] "x7e"      NA [38, ] "x8e"      NA [39, ] "x9e"      NA [40, ] "x10e"     NA [41, ] "x11e"     NA [42, ] "x12e"     NA [43, ] "x13e"     NA [44, ] "x14e"     NA [45, ] "x15e"     NA [46, ] "x16e"     NA [47, ] "x17e"     NA [48, ] "x18e"     NA [49, ] "x19e"     NA [50, ] "x20e"     NA [51, ] "x21e"     NA [52, ] "x22e"     NA [53, ] "x23e"     NA [54, ] "x24e"     NA [55, ] "1"        NA
>
```

Continuous two factor

```
> summary(sem2,digits=2)
```

```
Model Chisquare = 235    Df = 252 Pr(>Chisq) = 0.78
Chisquare (null model) = 3568    Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.96
RMSEA index = 0    90% CI: (NA, 0.013)
Bentler-Bonnett NFI = 0.93
Tucker-Lewis NNFI = 1
Bentler CFI = 1
SRMR = 0.029
BIC = -1331
```

Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-1.6e+00	-4.1e-01	-3.1e-05	8.8e-03	3.9e-01	2.2e+00

Relevant paths

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)	
F1V1	0.63	0.043	15	0	V1 <--- Factor1
F1V2	0.61	0.043	14	0	V2 <--- Factor1
F1V3	0.57	0.044	13	0	V3 <--- Factor1
F1V4	0.63	0.042	15	0	V4 <--- Factor1
F1V5	0.64	0.042	15	0	V5 <--- Factor1
F1V6	0.61	0.043	14	0	V6 <--- Factor1
F2V7	0.55	0.044	12	0	V7 <--- Factor2
F2V8	0.63	0.043	15	0	V8 <--- Factor2
F2V9	0.58	0.044	13	0	V9 <--- Factor2
F2V10	0.58	0.044	13	0	V10 <--- Factor2
F2V11	0.59	0.043	14	0	V11 <--- Factor2
F2V12	0.61	0.043	14	0	V12 <--- Factor2
F1V13	-0.63	0.042	-15	0	V13 <--- Factor1
F1V14	-0.59	0.043	-14	0	V14 <--- Factor1
F1V15	-0.57	0.043	-13	0	V15 <--- Factor1
F1V16	-0.64	0.042	-15	0	V16 <--- Factor1
F1V17	-0.63	0.043	-15	0	V17 <--- Factor1
F1V18	-0.59	0.043	-14	0	V18 <--- Factor1
F2V19	-0.57	0.044	-13	0	V19 <--- Factor2
F2V20	-0.60	0.043	-14	0	V20 <--- Factor2
F2V21	-0.62	0.043	-15	0	V21 <--- Factor2
F2V22	-0.57	0.044	-13	0	V22 <--- Factor2
F2V23	-0.58	0.044	-13	0	V23 <--- Factor2
F2V24	-0.60	0.043	-14	0	V24 <--- Factor2

Categorical 2 factor

- I. Original model does not converge
- II. Reset some paths to give initial estimates closer to the factor solution
- III. Use the edit command

```

> mod2.c <- edit(mod2)
> mod2.c
      Path          Parameter Value
[1,] "Factor1->V1"    "F1V1"    ".6"
[2,] "Factor1->V2"    "F1V2"    ".6"
[3,] "Factor1->V3"    "F1V3"    ".6"
[4,] "Factor1->V4"    "F1V4"    ".6"
[5,] "Factor1->V5"    "F1V5"    ".6"
[6,] "Factor1->V6"    "F1V6"    ".6"
[7,] "Factor2->V7"    "F2V7"    ".6"
[8,] "Factor2->V8"    "F2V8"    ".6"
[9,] "Factor2->V9"    "F2V9"    ".6"
[10,] "Factor2->V10"   "F2V10"   ".6"
[11,] "Factor2->V11"   "F2V11"   ".6"
[12,] "Factor2->V12"   "F2V12"   ".6"
[13,] "Factor1->V13"   "F1V13"   "-.6"
[14,] "Factor1->V14"   "F1V14"   "-.6"
[15,] "Factor1->V15"   "F1V15"   "-.6"
[16,] "Factor1->V16"   "F1V16"   NA
[17,] "Factor1->V17"   "F1V17"   NA
[18,] "Factor1->V18"   "F1V18"   NA
[19,] "Factor2->V19"   "F2V19"   NA
[20,] "Factor2->V20"   "F2V20"   NA
[21,] "Factor2->V21"   "F2V21"   NA
[22,] "Factor2->V22"   "F2V22"   NA
[23,] "Factor2->V23"   "F2V23"   NA
[24,] "Factor2->V24"   "F2V24"   "-.6"
[25,] "V1<->V1"      "x1e"     NA
[26,] "V2<->V2"      "x2e"     NA
...
[49,] "Factor1<->Factor1" NA        "1"
[50,] "Factor2<->Factor2" NA        "1"

```

Modified paths in model

Fit is very good

```
> summary(sem.c,digits=2)

Model Chisquare = 255    Df = 252 Pr(>Chisq) = 0.44
Chisquare (null model) = 3143    Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.95
RMSEA index = 0.0049 90% CI: (NA, 0.019)
Bentler-Bonnett NFI = 0.92
Tucker-Lewis NNFI = 1
Bentler CFI = 1
SRMR = 0.032
BIC = -1311

Normalized Residuals
  Min. 1st Qu. Median      Mean 3rd Qu.      Max.
-2.2e+00 -4.3e-01  2.2e-05  4.1e-02  5.4e-01  2.1e+00
```

Path coefficients are fine

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)	
F1V1	0.60	0.043	14	0	V1 <--- Factor1
F1V2	0.56	0.044	13	0	V2 <--- Factor1
F1V3	0.55	0.044	12	0	V3 <--- Factor1
F1V4	0.61	0.043	14	0	V4 <--- Factor1
F1V5	0.61	0.043	14	0	V5 <--- Factor1
F1V6	0.59	0.044	13	0	V6 <--- Factor1
F2V7	0.52	0.045	12	0	V7 <--- Factor2
F2V8	0.59	0.044	13	0	V8 <--- Factor2
F2V9	0.57	0.044	13	0	V9 <--- Factor2
F2V10	0.55	0.045	12	0	V10 <--- Factor2
F2V11	0.56	0.044	13	0	V11 <--- Factor2
F2V12	0.56	0.044	13	0	V12 <--- Factor2
F1V13	-0.61	0.043	-14	0	V13 <--- Factor1
F1V14	-0.57	0.044	-13	0	V14 <--- Factor1
F1V15	-0.54	0.044	-12	0	V15 <--- Factor1
F1V16	-0.63	0.043	-15	0	V16 <--- Factor1
F1V17	-0.58	0.044	-13	0	V17 <--- Factor1
F1V18	-0.56	0.044	-13	0	V18 <--- Factor1
F2V19	-0.55	0.045	-12	0	V19 <--- Factor2
F2V20	-0.58	0.044	-13	0	V20 <--- Factor2

Now try skewed

- I. Once again, the initial model does not work (with a weird diagnostic), but the model with roughly the right answers (taken from the exploratory factor analysis) does.
- II. However, the model does not fit by some criteria (i.e., chi square), but does by other (RMSEA)

Fit statistics

```
> sem.sk <- sem(mod2.c,cor(simp),500)
> summary(sem.sk,digits=2)

Model Chisquare = 380    Df = 252 Pr(>Chisq) = 3.2e-07
Chisquare (null model) = 2002    Df = 276
Goodness-of-fit index = 0.94
Adjusted goodness-of-fit index = 0.93
RMSEA index = 0.032   90% CI: (0.025, 0.038)
Bentler-Bonnett NFI = 0.81
Tucker-Lewis NNFI = 0.92
Bentler CFI = 0.93
SRMR = 0.043
BIC = -1186

Normalized Residuals
  Min. 1st Qu. Median      Mean 3rd Qu.      Max.
-2.47 -0.26  0.30      0.31  1.02      2.51
```

Parameter Estimates						Path coefficients
	Estimate	Std Error	z value	P> z		
F1V1	0.48	0.048	10.1	0.0e+00	V1 <---	Factor1
F1V2	0.48	0.048	10.0	0.0e+00	V2 <---	Factor1
F1V3	0.49	0.047	10.4	0.0e+00	V3 <---	Factor1
F1V4	0.48	0.048	10.0	0.0e+00	V4 <---	Factor1
F1V5	0.50	0.047	10.5	0.0e+00	V5 <---	Factor1
F1V6	0.51	0.047	10.7	0.0e+00	V6 <---	Factor1
F2V7	0.50	0.047	10.6	0.0e+00	V7 <---	Factor2
F2V8	0.56	0.047	12.0	0.0e+00	V8 <---	Factor2
F2V9	0.53	0.047	11.2	0.0e+00	V9 <---	Factor2
F2V10	0.50	0.047	10.5	0.0e+00	V10 <---	Factor2
F2V11	0.48	0.048	9.9	0.0e+00	V11 <---	Factor2
F2V12	0.55	0.047	11.8	0.0e+00	V12 <---	Factor2
F1V13	-0.51	0.047	-10.8	0.0e+00	V13 <---	Factor1
F1V14	-0.48	0.048	-10.2	0.0e+00	V14 <---	Factor1
F1V15	-0.44	0.048	-9.2	0.0e+00	V15 <---	Factor1
F1V16	-0.52	0.047	-11.1	0.0e+00	V16 <---	Factor1
F1V17	-0.48	0.048	-10.0	0.0e+00	V17 <---	Factor1
F1V18	-0.48	0.048	-10.1	0.0e+00	V18 <---	Factor1
F2V19	-0.38	0.049	-7.8	5.1e-15	V19 <---	Factor2
F2V20	-0.41	0.049	-8.5	0.0e+00	V20 <---	Factor2
F2V21	-0.39	0.049	-8.0	1.6e-15	V21 <---	Factor2
F2V22	-0.42	0.049	-8.7	0.0e+00	V22 <---	Factor2
F2V23	-0.47	0.048	-9.7	0.0e+00	V23 <---	Factor2

Alternative models

- I. What about 4 unipolar factors
- II. Correlated in pairs to get the polarity
- III. Create the path model using symbolic notation with the structure.list and phi.list functions
 - A.

```
fx <-
structure.list(24,list(F1p=c(1:6),F2p=c(7:12),F1n
=c(13:18),F2n=c(19:24)))
```
 - B.

```
phi<- phi.list(4,list(F1=c(3),F2=c(4)))
```
 - C.

```
mod.4 <- structure.graph(fx,phi)
```

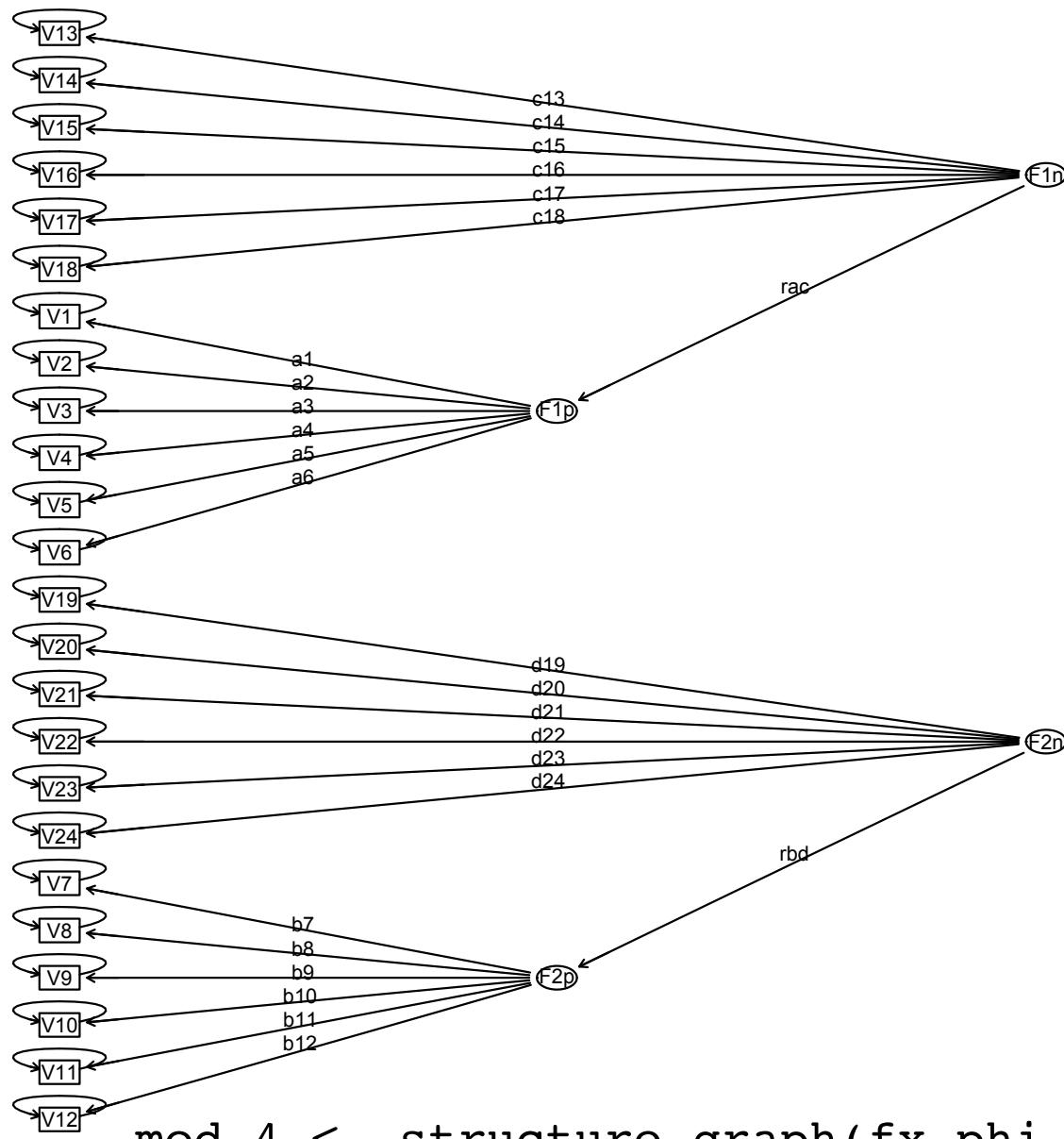
```
> fx <- structure.list(24,list(F1p=c(1:6),F2p=c(7:12),
F1n=c(13:18),F2n=c(19:24)))
> fx
```

	F1p	F2p	F1n	F2n
[1,]	"a1"	"0"	"0"	"0"
[2,]	"a2"	"0"	"0"	"0"
[3,]	"a3"	"0"	"0"	"0"
[4,]	"a4"	"0"	"0"	"0"
[5,]	"a5"	"0"	"0"	"0"
[6,]	"a6"	"0"	"0"	"0"
[7,]	"0"	"b7"	"0"	"0"
[8,]	"0"	"b8"	"0"	"0"
[9,]	"0"	"b9"	"0"	"0"
[10,]	"0"	"b10"	"0"	"0"
[11,]	"0"	"b11"	"0"	"0"
[12,]	"0"	"b12"	"0"	"0"
[13,]	"0"	"0"	"c13"	"0"
[14,]	"0"	"0"	"c14"	"0"
[15,]	"0"	"0"	"c15"	"0"
[16,]	"0"	"0"	"c16"	"0"
[17,]	"0"	"0"	"c17"	"0"
[18,]	"0"	"0"	"c18"	"0"
[19,]	"0"	"0"	"0"	"d19"
[20,]	"0"	"0"	"0"	"d20"
[21,]	"0"	"0"	"0"	"d21"
[22,]	"0"	"0"	"0"	"d22"
[23,]	"0"	"0"	"0"	"d23"

Symbolic structure

```
> phi<- phi.list(4,list(F1=c(3),
F2=c(4)))
> phi
      F1      F2      F3      F4
F1  "1"    "0"    "0"    "0"
F2  "0"    "1"    "0"    "0"
F3  "rac"  "0"    "1"    "0"
F4  "0"    "rbd"  "0"    "1"
```

Structural model



Create
the
model

Fit is fine

```
> mod.4c <- edit(mod.4) #make the paths bidirectional  
> sem.4 <- sem(mod.4c,cor(simp),500)  
> summary(sem.4,digits=2)
```

Model Chisquare = 265 Df = 250 Pr(>Chisq) = 0.25
Chisquare (null model) = 2002 Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.95
RMSEA index = 0.011 90% CI: (NA, 0.021)
Bentler-Bonnett NFI = 0.87
Tucker-Lewis NNFI = 1
Bentler CFI = 1
SRMR = 0.036
BIC = -1289

Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-2.470	-0.608	-0.069	-0.052	0.449	2.330

Paths

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)	
a1	0.53	0.048	11.0	0.0e+00	V1 <--- F1p
a2	0.53	0.049	10.8	0.0e+00	V2 <--- F1p
...					
a6	0.56	0.048	11.5	0.0e+00	V6 <--- F1p
b7	0.52	0.048	10.7	0.0e+00	V7 <--- F2p
b8	0.58	0.047	12.3	0.0e+00	V8 <--- F2p
...					
c13	0.55	0.048	11.4	0.0e+00	V13 <--- F1n
..					
c18	0.52	0.049	10.7	0.0e+00	V18 <--- F1n
d19	0.43	0.051	8.3	2.2e-16	V19 <--- F2n
..					
d24	0.48	0.051	9.5	0.0e+00	V24 <--- F2n
x1e	0.72	0.053	13.6	0.0e+00	V1 <--> V1
x2e	0.72	0.053	13.7	0.0e+00	V2 <--> V2
...					
x7e	0.73	0.053	13.9	0.0e+00	V7 <--> V7
x8e	0.66	0.050	13.1	0.0e+00	V8 <--> V8
...					
rF3F1	-0.72	0.044	-16.4	0.0e+00	F1p <--> F1n
rF4F2	-0.73	0.046	-15.7	0.0e+00	F2p <--> F2n

Consider a different model

- I. Let the factors be bipolar, but add a residual factor at each end to account for skew differences
- II. Basically a bifactor solution, but applied to each of the two main dimensions

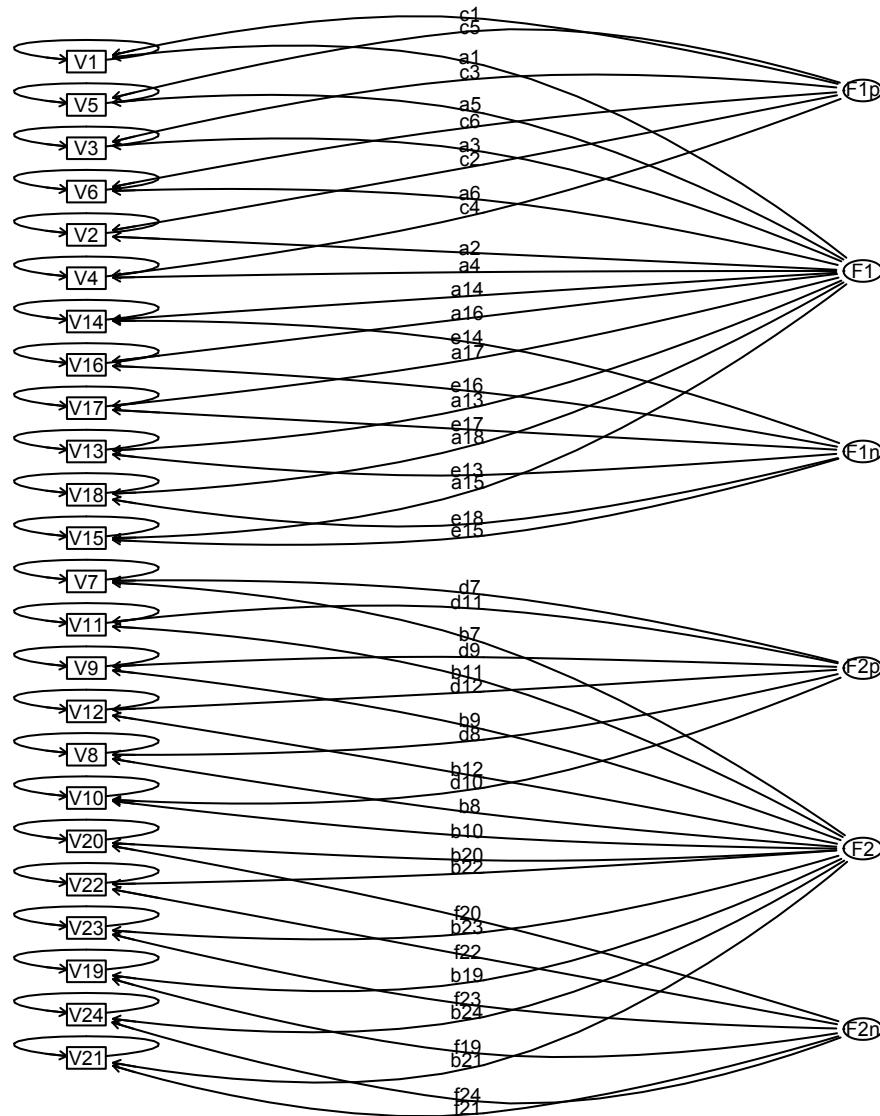
```

> fxb <-
structure.list(24,list(F1=c(1:6,13:18),F2=c(7:12,19:24),F1p=c(1:6),F2p=c(7:12),F1n
(13:18),F2n=c(19:24)))
> fxb
      F1     F2     F1p    F2p    F1n    F2n
[1,] "a1"   "0"    "c1"   "0"    "0"    "0"
[2,] "a2"   "0"    "c2"   "0"    "0"    "0"
[3,] "a3"   "0"    "c3"   "0"    "0"    "0"
[4,] "a4"   "0"    "c4"   "0"    "0"    "0"
[5,] "a5"   "0"    "c5"   "0"    "0"    "0"
[6,] "a6"   "0"    "c6"   "0"    "0"    "0"
[7,] "0"    "b7"   "0"    "d7"   "0"    "0"
[8,] "0"    "b8"   "0"    "d8"   "0"    "0"
[9,] "0"    "b9"   "0"    "d9"   "0"    "0"
[10,] "0"   "b10"  "0"    "d10"  "0"    "0"
[11,] "0"   "b11"  "0"    "d11"  "0"    "0"
[12,] "0"   "b12"  "0"    "d12"  "0"    "0"
[13,] "a13"  "0"    "0"    "0"    "e13"  "0"
[14,] "a14"  "0"    "0"    "0"    "e14"  "0"
[15,] "a15"  "0"    "0"    "0"    "e15"  "0"
[16,] "a16"  "0"    "0"    "0"    "e16"  "0"
[17,] "a17"  "0"    "0"    "0"    "e17"  "0"
[18,] "a18"  "0"    "0"    "0"    "e18"  "0"
[19,] "0"    "b19"  "0"    "0"    "0"    "f19"
[20,] "0"    "b20"  "0"    "0"    "0"    "f20"
[21,] "0"    "b21"  "0"    "0"    "0"    "f21"
[22,] "0"    "b22"  "0"    "0"    "0"    "f22"
[23,] "0"    "b23"  "0"    "0"    "0"    "f23"
[24,] "0"    "b24"  "0"    "0"    "0"    "f24"
> mod.2b <- structure.graph(fxb,labels=colnames(simp))

```

Create the
structure
symbolically

Structural model



Path
model

```
> sem.2b <- sem(mod.2b,cor(simp),500)
> summary(sem.2b,digits=2)

Model Chisquare = 228    Df = 228 Pr(>Chisq) = 0.49
Chisquare (null model) = 2002    Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.95
RMSEA index = 0 90% CI: (NA, 0.019)
Bentler-Bonnett NFI = 0.89
Tucker-Lewis NNFI = 1
Bentler CFI = 1
SRMR = 0.034
BIC = -1189

Normalized Residuals
  Min. 1st Qu. Median      Mean 3rd Qu.      Max.
-2.470 -0.481 -0.002 -0.040   0.380   2.330
```

Paths

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)		
a1	0.3530	0.051	6.884	5.8e-12	V1	<--- F1
c1	0.4170	0.064	6.475	9.5e-11	V1	<--- F1p
a2	0.3583	0.051	7.019	2.2e-12	V2	<--- F1
c2	0.3924	0.063	6.195	5.8e-10	V2	<--- F1p
...						
c6	0.4179	0.063	6.598	4.2e-11	V6	<--- F1p
b7	-0.5069	0.077	-6.580	4.7e-11	V7	<--- F2
d7	0.0800	0.227	0.353	7.2e-01	V7	<--- F2p
b8	-0.5632	0.063	-8.974	0.0e+00	V8	<--- F2
...						
d12	0.2523	0.238	1.059	2.9e-01	V12	<--- F2p
a13	-0.5628	0.050	-11.250	0.0e+00	V13	<--- F1
e13	-0.0985	0.087	-1.129	2.6e-01	V13	<--- F1n
a14	-0.5524	0.051	-10.847	0.0e+00	V14	<--- F1
e14	-0.1679	0.121	-1.383	1.7e-01	V14	<--- F1n
...						
a18	-0.5413	0.066	-8.140	4.4e-16	V18	<--- F1
e18	0.5160	0.248	2.079	3.8e-02	V18	<--- F1n
b19	0.3435	0.059	5.816	6.0e-09	V19	<--- F2
f19	0.2525	0.081	3.101	1.9e-03	V19	<--- F2n
b20	0.3495	0.061	5.730	1.0e-08	V20	<--- F2
f20	0.3791	0.085	4.460	8.2e-06	V20	<--- F2n
b21	0.3566	0.060	5.965	2.4e-09	V21	<--- F2
...						
b24	0.3664	0.062	5.937	2.9e-09	V24	<--- F2
f24	0.3304	0.085	3.909	9.3e-05	V24	<--- F2n
x1e	0.7015	0.057	12.322	0.0e+00	V1	<--> V1
x2e	0.7177	0.056	12.852	0.0e+00	V2	<--> V2
...						
x24e	0.7566	0.060	12.582	0.0e+00	V24	<--> V24

Do the models differ?

```
> anova(sem.2b,sem.4)
LR Test for Difference Between Models

          Model Df Model Chisq   Df LR Chisq Pr(>Chisq)
Model 1      228      227.960
Model 2      250      264.517    22      36.557      0.02644 *
---
Signif. codes:  0 '****' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' '
' 1
>
```

But they are not nested models

The bipolar model has a smaller chi square, but is penalized for 22 degrees of freedom (compare the BICs)

BIC = -1189 vs. BIC = -1289 vs. BIC = -1186 (for orig)

Comparisons of models and data

data	model	chi sq	df	BIC	RMSEA
Continuous	2 f	235	252	-1331	0.0000
Categorical	2 f	255	252	-1311	0.0049
Skewed	2 f	380	252	-1186	0.0320
	4 f	265	250	-1289	0.0110
	2 bi	228	228	-1189	0.0000

Consider Circumplex Structure

- I. Two dimensional model of data, with and without skew

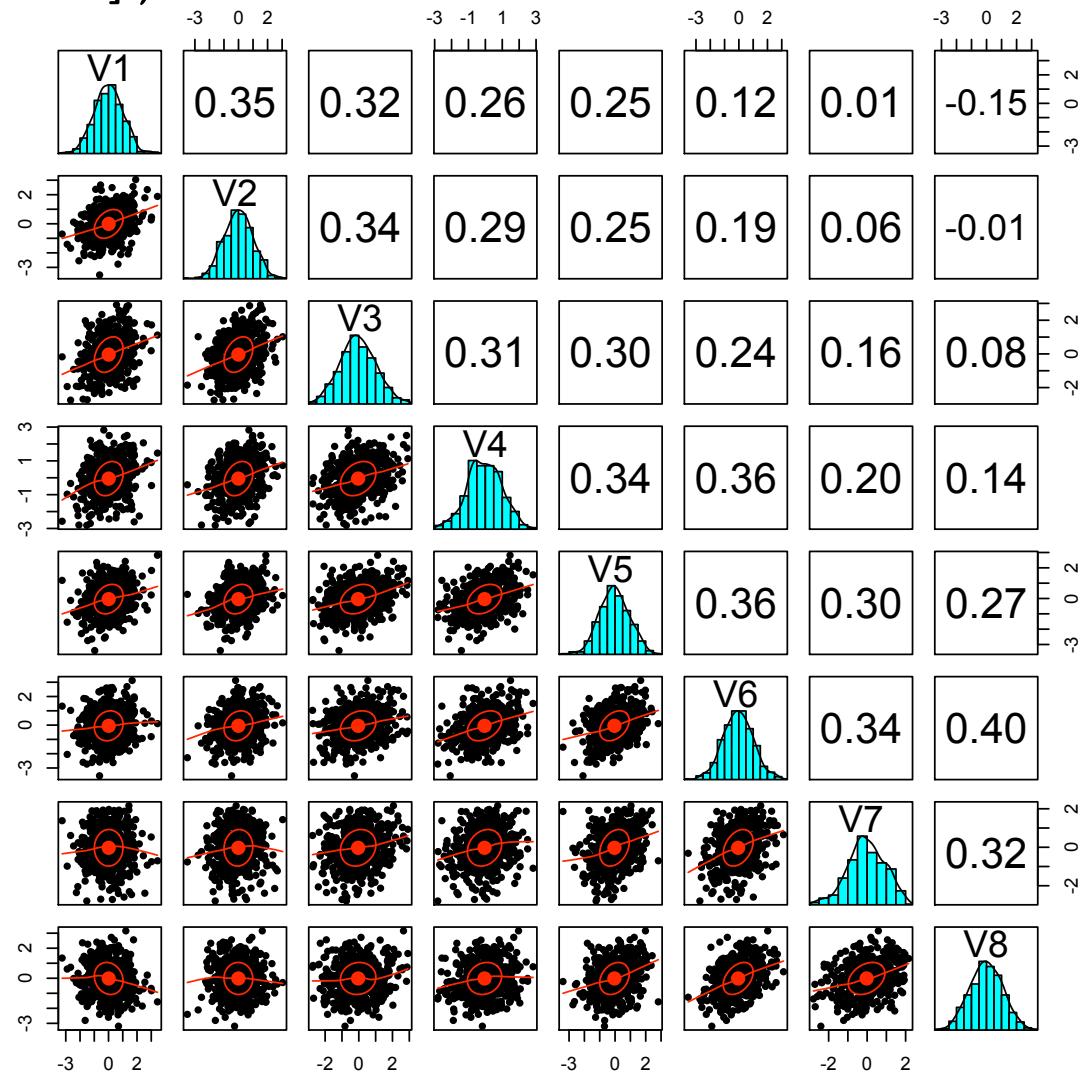
The data

```
> set.seed(42)
> circ <- sim.item(24,circum=TRUE)
> describe(circ)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	skew	kurtosis	se
V1	1	500	0.01	1.01	0.03	0.01	0.97	-3.26	3.44	6.69	0.05	0.13	0.04
V2	2	500	-0.01	0.99	-0.01	-0.01	1.02	-3.52	3.04	6.56	-0.03	-0.03	0.04
V3	3	500	-0.04	1.05	-0.07	-0.05	1.04	-2.75	2.92	5.66	0.12	-0.18	0.05
V4	4	500	-0.05	1.01	-0.06	-0.04	1.06	-2.82	2.83	5.64	-0.05	-0.17	0.05
V5	5	500	-0.04	0.96	-0.07	-0.05	0.95	-3.40	2.82	6.22	-0.02	-0.04	0.04
V6	6	500	-0.05	1.05	-0.05	-0.06	1.04	-3.55	3.12	6.67	0.03	0.09	0.05
V7	7	500	-0.03	0.95	-0.04	-0.01	0.93	-2.79	2.16	4.95	-0.19	-0.17	0.04
V8	8	500	-0.01	1.07	-0.01	-0.01	1.10	-3.19	3.15	6.35	0.01	-0.28	0.05
V9	9	500	0.01	1.02	0.04	0.02	1.07	-2.58	3.54	6.12	0.01	-0.28	0.05
V10	10	500	0.01	0.96	-0.04	0.01	0.94	-2.45	3.10	5.55	0.02	-0.02	0.04
V11	11	500	-0.03	0.98	-0.09	-0.05	0.99	-3.45	3.06	6.51	0.15	0.27	0.04
V12	12	500	-0.03	1.01	-0.04	-0.04	1.08	-3.36	3.26	6.62	0.03	-0.10	0.05
V13	13	500	-0.05	1.07	-0.08	-0.07	1.11	-2.93	3.74	6.66	0.13	-0.01	0.05
V14	14	500	0.08	0.97	0.09	0.08	0.99	-2.82	2.64	5.45	-0.07	-0.28	0.04
V15	15	500	0.03	1.01	-0.03	0.02	1.02	-2.64	3.26	5.90	0.13	-0.06	0.05
V16	16	500	0.01	1.02	-0.06	0.00	1.10	-2.90	2.99	5.89	0.08	-0.18	0.05
V17	17	500	-0.01	1.00	0.04	-0.01	1.09	-2.63	2.94	5.58	-0.01	-0.36	0.04
V18	18	500	0.04	1.00	0.04	0.04	1.03	-3.71	2.83	6.54	-0.05	0.09	0.04
V19	19	500	-0.01	0.97	-0.02	-0.01	1.02	-2.99	2.62	5.62	-0.01	-0.23	0.04
V20	20	500	0.03	0.98	-0.01	0.02	0.94	-2.96	2.94	5.90	0.05	0.25	0.04
V21	21	500	-0.03	1.04	-0.09	-0.04	1.05	-3.27	3.34	6.61	0.11	-0.03	0.05
V22	22	500	0.04	1.02	-0.03	0.03	1.02	-3.31	3.68	6.99	0.09	0.18	0.05
V23	23	500	0.01	1.00	0.00	0.00	1.01	-2.95	3.16	6.10	0.06	-0.09	0.04
V24	24	500	-0.01	1.00	0.07	-0.01	0.92	-2.71	3.11	5.82	0.01	0.06	0.04

SPLOM

```
pairs.panels(circ[,1:8])
```



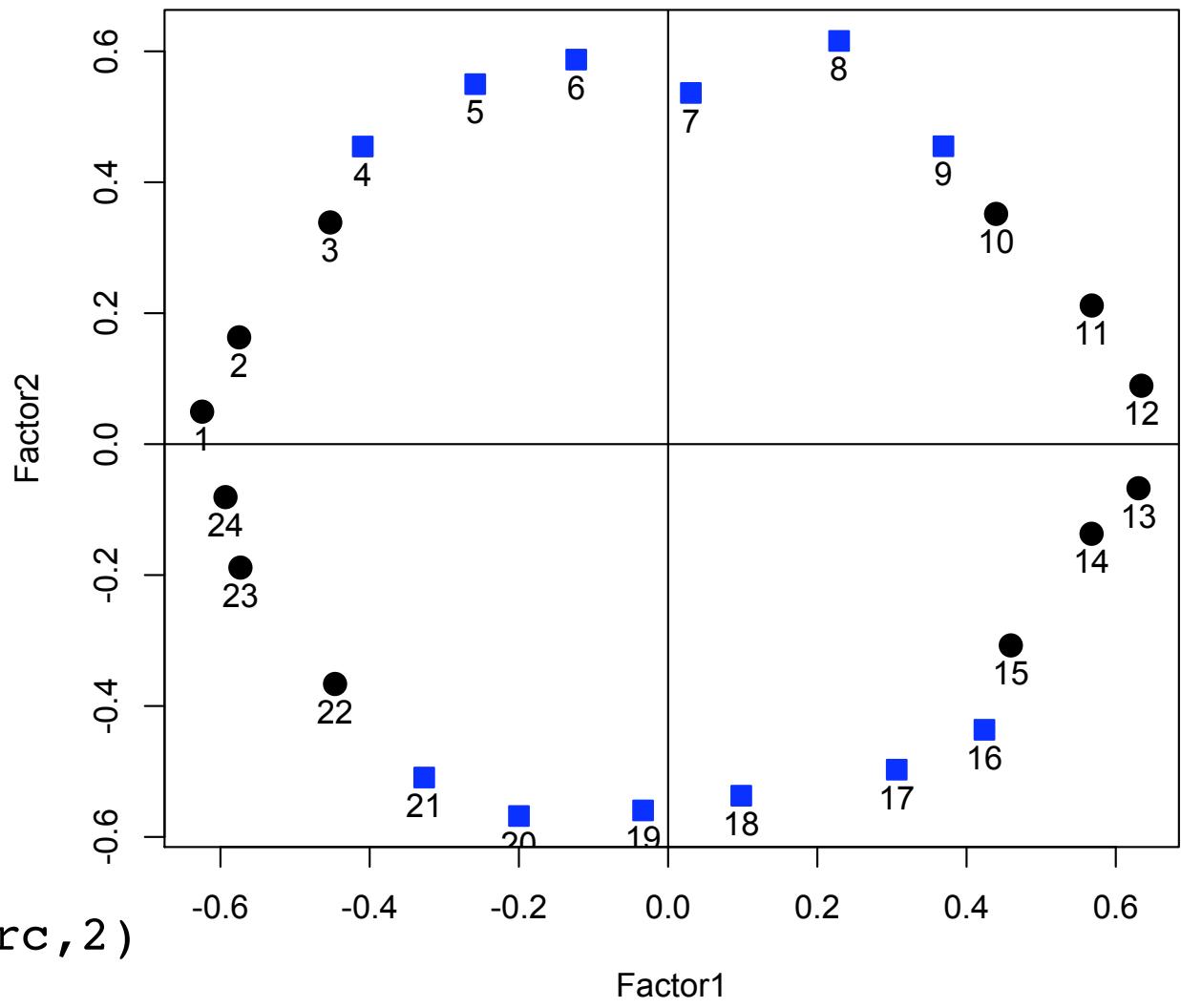
Circumplex correlations

```
> round(cor(circ)[,1:12],2)
```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	1.00	0.35	0.32	0.26	0.25	0.12	0.01	-0.15	-0.25	-0.25	-0.34	-0.37
V2	0.35	1.00	0.34	0.29	0.25	0.19	0.06	-0.01	-0.14	-0.20	-0.32	-0.36
V3	0.32	0.34	1.00	0.31	0.30	0.24	0.16	0.08	0.04	-0.07	-0.12	-0.33
V4	0.26	0.29	0.31	1.00	0.34	0.36	0.20	0.14	0.07	0.00	-0.15	-0.20
V5	0.25	0.25	0.30	0.34	1.00	0.36	0.30	0.27	0.16	0.11	-0.02	-0.08
V6	0.12	0.19	0.24	0.36	0.36	1.00	0.34	0.40	0.22	0.10	0.08	-0.03
V7	0.01	0.06	0.16	0.20	0.30	0.34	1.00	0.32	0.23	0.17	0.14	0.08
V8	-0.15	-0.01	0.08	0.14	0.27	0.40	0.32	1.00	0.38	0.29	0.21	0.21
V9	-0.25	-0.14	0.04	0.07	0.16	0.22	0.23	0.38	1.00	0.29	0.32	0.24
V10	-0.25	-0.20	-0.07	0.00	0.11	0.10	0.17	0.29	0.29	1.00	0.38	0.32
V11	-0.34	-0.32	-0.12	-0.15	-0.02	0.08	0.14	0.21	0.32	0.38	1.00	0.40
V12	-0.37	-0.36	-0.33	-0.20	-0.08	-0.03	0.08	0.21	0.24	0.32	0.40	1.00
V13	-0.38	-0.37	-0.33	-0.29	-0.21	-0.09	0.00	0.06	0.21	0.24	0.35	0.38
V14	-0.36	-0.31	-0.34	-0.35	-0.25	-0.10	-0.02	0.08	0.16	0.20	0.27	0.31
V15	-0.33	-0.32	-0.28	-0.34	-0.24	-0.22	-0.16	-0.10	0.06	0.06	0.22	0.25
V16	-0.28	-0.28	-0.35	-0.38	-0.35	-0.28	-0.25	-0.18	-0.06	0.06	0.13	0.25
V17	-0.19	-0.26	-0.27	-0.38	-0.34	-0.33	-0.26	-0.24	-0.11	-0.06	0.09	0.17
V18	-0.09	-0.16	-0.27	-0.26	-0.34	-0.33	-0.29	-0.35	-0.15	-0.16	-0.07	-0.01
V19	-0.04	-0.08	-0.16	-0.25	-0.30	-0.32	-0.30	-0.34	-0.24	-0.24	-0.15	-0.09
V20	0.11	0.03	-0.12	-0.16	-0.25	-0.29	-0.32	-0.37	-0.37	-0.29	-0.23	-0.14
V21	0.17	0.09	0.00	-0.13	-0.18	-0.23	-0.29	-0.38	-0.37	-0.34	-0.24	-0.26
V22	0.31	0.21	0.05	0.00	-0.05	-0.15	-0.23	-0.32	-0.35	-0.30	-0.32	-0.30
V23	0.30	0.35	0.16	0.17	0.05	-0.01	-0.08	-0.26	-0.28	-0.36	-0.38	-0.40
V24	0.39	0.29	0.22	0.22	0.08	0.03	-0.05	-0.23	-0.25	-0.29	-0.37	-0.40

2 factor solution

Cluster plot



```
> f2c <- factanal(circ, 2)
> factor.plot(f2c)
```

2 factors are great

```
factanal(x = circ, factors = 2)
```

Uniquenesses:

V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11
V12	V13	V14	V15	V16	V17	V18	V19	V20	V21	V22
0.607	0.643	0.680	0.626	0.631	0.640	0.711	0.568	0.657	0.683	0.633
0.590	0.598	0.659	0.694	0.630	0.659	0.702	0.686	0.637	0.634	0.667
V23	V24			[V13	0.630					
0.636	0.641			V14	0.568	-0.137				

Loadings:

	Factor1	Factor2								
V1	-0.625		V16	0.424	-0.436					
V2	-0.575	0.163	V17	0.306	-0.497					
V3	-0.453	0.339	V18		-0.537					
V4	-0.409	0.455	V19		-0.559					
V5	-0.259	0.550	V20	-0.200	-0.568					
V6	-0.123	0.587	V21	-0.327	-0.509					
V7		0.536	V22	-0.446	-0.366					
V8	0.229	0.616	V23	-0.573	-0.189					
V9	0.369	0.455	V24	-0.593						
V10	0.440	0.352				Factor1	Factor2			
V11	0.568	0.212	SS loadings			4.524	3.963			
V12	0.634		Proportion Var			0.189	0.165			
			Cumulative Var			0.189	0.354			

But, the simple structure model does not fit

```
> sem.c <- sem(mod2,cor(circ),500)
> summary(sem.c,digits=2)

Model Chisquare = 1204    Df = 252 Pr(>Chisq) = 0
Chisquare (null model) = 3449    Df = 276
Goodness-of-fit index = 0.8
Adjusted goodness-of-fit index = 0.75
RMSEA index = 0.087 90% CI: (NA, NA)
Bentler-Bonnett NFI = 0.65
Tucker-Lewis NNFI = 0.67
Bentler CFI = 0.7
SRMR = 0.15
BIC = -362

Normalized Residuals
  Min. 1st Qu. Median      Mean 3rd Qu.      Max.
-8.440 -1.850 -0.036 -0.054  1.690  8.960
```

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)	
F1V1	0.51	0.046	11.0	0.0e+00	V1 <--- Factor1
F1V2	0.54	0.045	11.8	0.0e+00	V2 <--- Factor1
F1V3	0.57	0.045	12.8	0.0e+00	V3 <--- Factor1
F1V4	0.61	0.044	13.8	0.0e+00	V4 <--- Factor1
F1V5	0.54	0.045	12.0	0.0e+00	V5 <--- Factor1
F1V6	0.44	0.047	9.3	0.0e+00	V6 <--- Factor1
F2V7	0.38	0.048	7.9	2.2e-15	V7 <--- Factor2
F2V8	0.57	0.045	12.7	0.0e+00	V8 <--- Factor2
F2V9	0.59	0.045	13.1	0.0e+00	V9 <--- Factor2
F2V10	0.57	0.045	12.7	0.0e+00	V10 <--- Factor2
F2V11	0.54	0.046	11.9	0.0e+00	V11 <--- Factor2
F2V12	0.50	0.047	10.6	0.0e+00	V12 <--- Factor2
F1V13	-0.52	0.046	-11.3	0.0e+00	V13 <--- Factor1
F1V14	-0.54	0.046	-11.8	0.0e+00	V14 <--- Factor1
F1V15	-0.55	0.045	-12.1	0.0e+00	V15 <--- Factor1
F1V16	-0.61	0.044	-13.8	0.0e+00	V16 <--- Factor1
F1V17	-0.54	0.045	-11.9	0.0e+00	V17 <--- Factor1
F1V18	-0.40	0.047	-8.5	0.0e+00	V18 <--- Factor1
F2V19	-0.41	0.047	-8.7	0.0e+00	V19 <--- Factor2
F2V20	-0.54	0.046	-11.7	0.0e+00	V20 <--- Factor2
F2V21	-0.61	0.045	-13.6	0.0e+00	V21 <--- Factor2
F2V22	-0.57	0.045	-12.6	0.0e+00	V22 <--- Factor2
F2V23	-0.54	0.046	-11.9	0.0e+00	V23 <--- Factor2
F2V24	-0.46	0.047	-9.9	0.0e+00	V24 <--- Factor2

Parameters
make sense

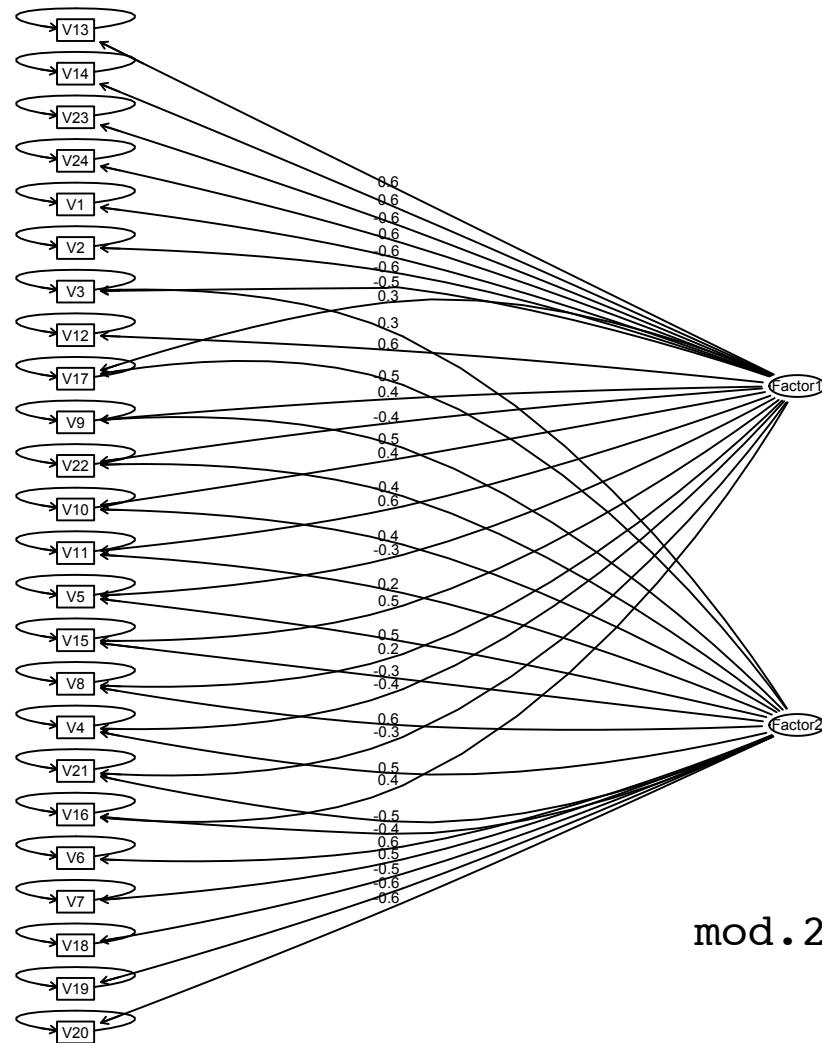
Examine residuals

```
> round(residuals(sem.c)[,1:12],2)
```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	0.00	0.08	0.03	-0.06	-0.03	-0.10	0.01	-0.15	-0.25	-0.25	-0.34	-0.37
V2	0.08	0.00	0.03	-0.04	-0.04	-0.05	0.06	-0.01	-0.14	-0.20	-0.32	-0.36
V3	0.03	0.03	0.00	-0.04	-0.01	-0.01	0.16	0.08	0.04	-0.07	-0.12	-0.33
V4	-0.06	-0.04	-0.04	0.00	0.01	0.09	0.20	0.14	0.07	0.00	-0.15	-0.20
V5	-0.03	-0.04	-0.01	0.01	0.00	0.12	0.30	0.27	0.16	0.11	-0.02	-0.08
V6	-0.10	-0.05	-0.01	0.09	0.12	0.00	0.34	0.40	0.22	0.10	0.08	-0.03
V7	0.01	0.06	0.16	0.20	0.30	0.34	0.00	0.11	0.01	-0.05	-0.06	-0.11
V8	-0.15	-0.01	0.08	0.14	0.27	0.40	0.11	0.00	0.04	-0.04	-0.10	-0.08
V9	-0.25	-0.14	0.04	0.07	0.16	0.22	0.01	0.04	0.00	-0.04	0.00	-0.05
V10	-0.25	-0.20	-0.07	0.00	0.11	0.10	-0.05	-0.04	-0.04	0.00	0.07	0.04
V11	-0.34	-0.32	-0.12	-0.15	-0.02	0.08	-0.06	-0.10	0.00	0.07	0.00	0.13
V12	-0.37	-0.36	-0.33	-0.20	-0.08	-0.03	-0.11	-0.08	-0.05	0.04	0.13	0.00
V13	-0.12	-0.09	-0.03	0.03	0.07	0.14	0.00	0.06	0.21	0.24	0.35	0.38
V14	-0.09	-0.03	-0.03	-0.02	0.04	0.13	-0.02	0.08	0.16	0.20	0.27	0.31
V15	-0.05	-0.02	0.03	0.00	0.06	0.02	-0.16	-0.10	0.06	0.06	0.22	0.25
V16	0.03	0.05	0.00	0.00	-0.02	-0.01	-0.25	-0.18	-0.06	0.06	0.13	0.25
V17	0.08	0.03	0.04	-0.05	-0.05	-0.09	-0.26	-0.24	-0.11	-0.06	0.09	0.17
V18	0.11	0.05	-0.04	-0.01	-0.12	-0.16	-0.29	-0.35	-0.15	-0.16	-0.07	-0.01
V19	-0.04	-0.08	-0.16	-0.25	-0.30	-0.32	-0.15	-0.11	0.00	-0.01	0.08	0.12
V20	0.11	0.03	-0.12	-0.16	-0.25	-0.29	-0.12	-0.07	-0.06	0.02	0.07	0.13
V21	0.17	0.09	0.00	-0.13	-0.18	-0.23	-0.06	-0.03	-0.02	0.01	0.09	0.04
V22	0.31	0.21	0.05	0.00	-0.05	-0.15	-0.02	0.00	-0.02	0.02	-0.01	-0.02
V23	0.30	0.35	0.16	0.17	0.05	-0.01	0.12	0.05	0.04	-0.05	-0.09	-0.13
V24	0.39	0.29	0.22	0.22	0.08	0.03	0.13	0.03	0.02	-0.02	-0.11	-0.17

Allow for cross loadings

Structural model



```
mod.2c <- structure.graph(f2,cut=.  
2,simple=FALSE)
```

Much better

```
Model Chisquare = 339.99 Df = 240 Pr(>Chisq) = 2.2571e-05
Chisquare (null model) = 3449.1 Df = 276
Goodness-of-fit index = 0.94666
Adjusted goodness-of-fit index = 0.93333
RMSEA index = 0.028894 90% CI: (0.021463, 0.035723)
Bentler-Bonnett NFI = 0.90143
Tucker-Lewis NNFI = 0.96376
Bentler CFI = 0.96849
SRMR = 0.054238
BIC = -1151.5
```

Normalized Residuals

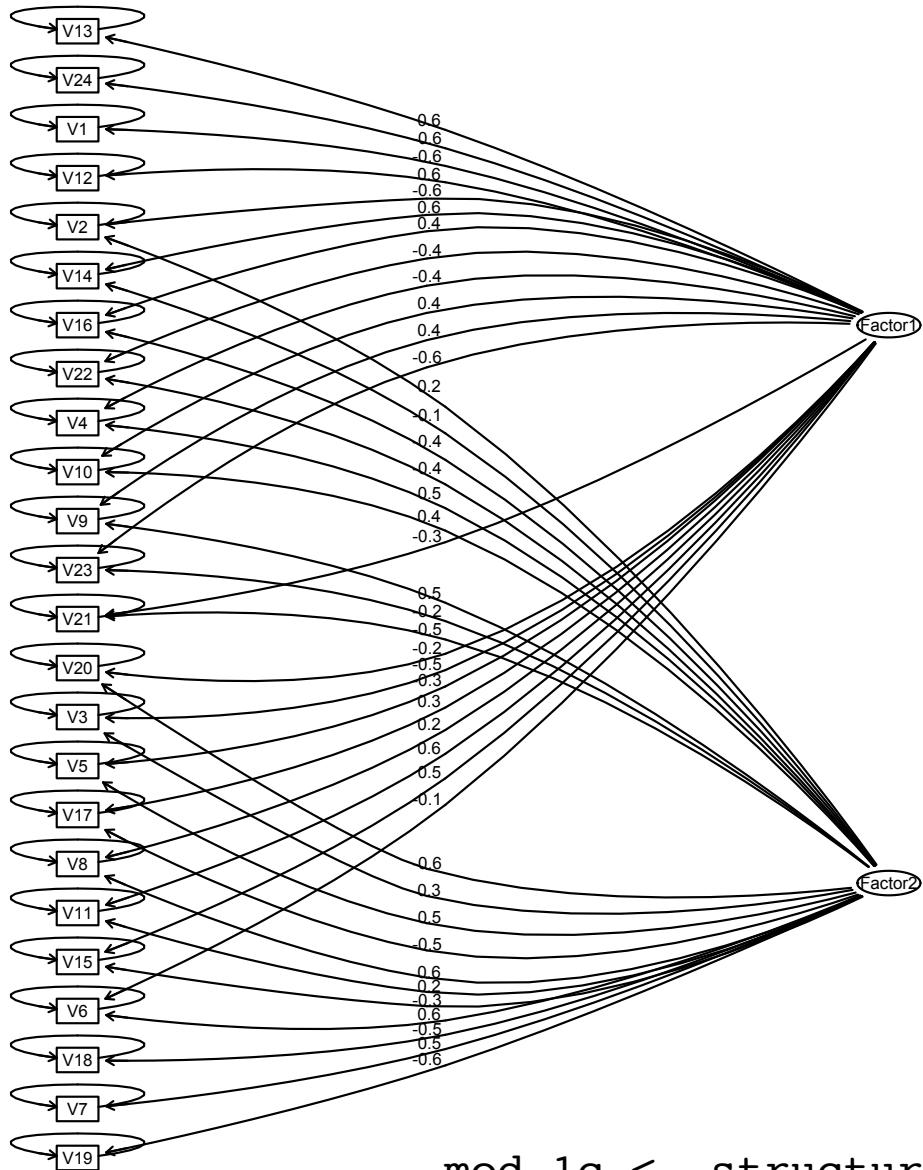
Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-3.63000	-0.72200	-0.01100	-0.00459	0.66900	5.55000

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)		
F1V1	-0.62510	0.042679	-14.6464	0.0000e+00	V1	<--- Factor1
F1V2	-0.57215	0.043609	-13.1200	0.0000e+00	V2	<--- Factor1
F1V3	-0.45852	0.043319	-10.5848	0.0000e+00	V3	<--- Factor1
F2V3	-0.32955	0.042673	-7.7225	1.1324e-14	V3	<--- Factor2
F1V4	-0.41496	0.042243	-9.8232	0.0000e+00	V4	<--- Factor1
F2V4	-0.44892	0.042254	-10.6244	0.0000e+00	V4	<--- Factor2
F1V5	-0.26585	0.042014	-6.3277	2.4878e-10	V5	<--- Factor1
F2V5	-0.54198	0.043029	-12.5957	0.0000e+00	V5	<--- Factor2
F2V6	-0.58679	0.043969	-13.3455	0.0000e+00	V6	<--- Factor2
F2V7	-0.54103	0.044515	-12.1540	0.0000e+00	V7	<--- Factor2
F1V8	0.22373	0.040683	5.4994	3.8119e-08	V8	<--- Factor1
F2V8	-0.61794	0.042222	-14.6355	0.0000e+00	V8	<--- Factor2
F1V9	0.35981	0.042647	8.4368	0.0000e+00	V9	<--- Factor1
F2V9	-0.46352	0.042975	-10.7856	0.0000e+00	V9	<--- Factor2
F1V10	0.43260	0.043362	9.9764	0.0000e+00	V10	<--- Factor1
F2V10	-0.35543	0.043014	-8.2631	2.2204e-16	V10	<--- Factor2
F1V11	0.56574	0.042811	13.2148	0.0000e+00	V11	<--- Factor1
F2V11	-0.21849	0.041302	-5.2900	1.2231e-07	V11	<--- Factor2
F1V12	0.63382	0.042565	14.8906	0.0000e+00	V12	<--- Factor1
F1V13	0.63250	0.042544	14.8671	0.0000e+00	V13	<--- Factor1
F1V14	0.56924	0.043669	13.0353	0.0000e+00	V14	<--- Factor1
F1V15	0.46708	0.043475	10.7437	0.0000e+00	V15	<--- Factor1
F2V15	0.30006	0.042985	6.9806	2.9388e-12	V15	<--- Factor2

Paths

Structural model



Allow for
even finer
resolution

More cross loadings

```
> mod.1cr <- edit(mod.1c) #specify a few to be correct  
> sem.1cr <- sem(mod.1cr,cor(circ),500)  
> summary(sem.1cr,digits=2)
```

Model Chisquare = 251 Df = 235 Pr(>Chisq) = 0.22
Chisquare (null model) = 3449 Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.95
RMSEA index = 0.012 90% CI: (NA, 0.022)
Bentler-Bonnett NFI = 0.93
Tucker-Lewis NNFI = 1
Bentler CFI = 1
SRMR = 0.033
BIC = -1209

Normalized Residuals

Min.	1st Qu.	Median	Mean	3rd Qu.	Max.
-2.1100	-0.4460	0.0253	0.0095	0.4770	2.1000

Paths

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)		
F1V1	-0.62	0.043	-14.6	0.0e+00	V1 <---	Factor1
F1V2	-0.57	0.043	-13.3	0.0e+00	V2 <---	Factor1
F2V2	-0.17	0.042	-4.1	4.0e-05	V2 <---	Factor2
F1V3	-0.45	0.044	-10.2	0.0e+00	V3 <---	Factor1
F2V3	-0.35	0.043	-8.0	8.9e-16	V3 <---	Factor2
F1V4	-0.40	0.043	-9.3	0.0e+00	V4 <---	Factor1
F2V4	-0.46	0.043	-10.8	0.0e+00	V4 <---	Factor2
F1V5	-0.25	0.043	-5.7	9.3e-09	V5 <---	Factor1
F2V5	-0.55	0.043	-12.8	0.0e+00	V5 <---	Factor2
F1V6	-0.11	0.043	-2.6	9.5e-03	V6 <---	Factor1
F2V6	-0.59	0.043	-13.6	0.0e+00	V6 <---	Factor2
F2V7	-0.54	0.045	-12.0	0.0e+00	V7 <---	Factor2
F1V8	0.24	0.042	5.7	1.2e-08	V8 <---	Factor1
F2V8	-0.61	0.042	-14.4	0.0e+00	V8 <---	Factor2
F1V9	0.38	0.044	8.7	0.0e+00	V9 <---	Factor1
F2V9	-0.45	0.043	-10.4	0.0e+00	V9 <---	Factor2
F1V10	0.45	0.044	10.2	0.0e+00	V10 <---	Factor1
F2V10	-0.34	0.043	-7.9	2.9e-15	V10 <---	Factor2
F1V11	0.57	0.043	13.3	0.0e+00	V11 <---	Factor1
F2V11	-0.20	0.042	-4.8	2.0e-06	V11 <---	Factor2

What about categorical?

```
> set.seed(42)
> circ.cat <- sim.item(24,circum=TRUE,categorical=TRUE)
> colnames(circ.cat) <- paste("V",1:24,sep="")
> describe(circ.cat)
```

	var	n	mean	sd	median	trimmed	mad	min	max	range	skew	kurtosis	se
V1	1	500	0.00	1.05	0	0.00	1.48	-3	3	6	0.06	0.03	0.05
V2	2	500	0.02	1.03	0	0.01	1.48	-3	3	6	0.03	-0.02	0.05
V3	3	500	-0.03	1.10	0	-0.04	1.48	-3	3	6	0.11	-0.12	0.05
V4	4	500	-0.06	1.07	0	-0.06	1.48	-3	3	6	-0.02	-0.22	0.05
V5	5	500	-0.04	1.00	0	-0.03	1.48	-3	3	6	-0.09	0.05	0.04
V6	6	500	-0.05	1.08	0	-0.04	1.48	-3	3	6	0.02	0.08	0.05
V7	7	500	-0.02	0.99	0	0.01	1.48	-3	2	5	-0.21	-0.09	0.04
V8	8	500	0.00	1.10	0	0.01	1.48	-3	3	6	-0.03	-0.25	0.05
V9	9	500	0.01	1.06	0	-0.01	1.48	-3	3	6	0.06	-0.34	0.05
V10	10	500	0.03	0.98	0	0.04	1.48	-2	3	5	-0.05	-0.28	0.04
V11	11	500	-0.02	1.03	0	-0.05	1.48	-3	3	6	0.18	0.04	0.05
V12	12	500	-0.03	1.05	0	-0.05	1.48	-3	3	6	0.07	-0.22	0.05
V13	13	500	-0.04	1.09	0	-0.06	1.48	-3	3	6	0.03	-0.08	0.05
V14	14	500	0.06	1.03	0	0.06	1.48	-3	3	6	-0.05	-0.25	0.05
V15	15	500	0.02	1.07	0	0.03	1.48	-3	3	6	0.02	-0.25	0.05
V16	16	500	0.03	1.05	0	0.01	1.48	-3	3	6	0.10	-0.09	0.05
V17	17	500	0.02	1.05	0	0.02	1.48	-3	3	6	0.02	-0.31	0.05
V18	18	500	0.04	1.03	0	0.03	1.48	-3	3	6	0.03	-0.02	0.05
V19	19	500	-0.01	0.99	0	0.00	1.48	-3	3	6	-0.07	-0.15	0.04
V20	20	500	0.01	1.03	0	-0.01	1.48	-3	3	6	0.06	0.32	0.05
V21	21	500	-0.04	1.07	0	-0.05	1.48	-3	3	6	0.12	0.04	0.05
V22	22	500	0.03	1.05	0	0.02	1.48	-3	3	6	0.04	0.08	0.05
V23	23	500	0.03	1.05	0	0.02	1.48	-3	3	6	0.05	-0.13	0.05
V24	24	500	-0.01	1.04	0	0.01	1.48	-3	3	6	-0.02	0.10	0.05

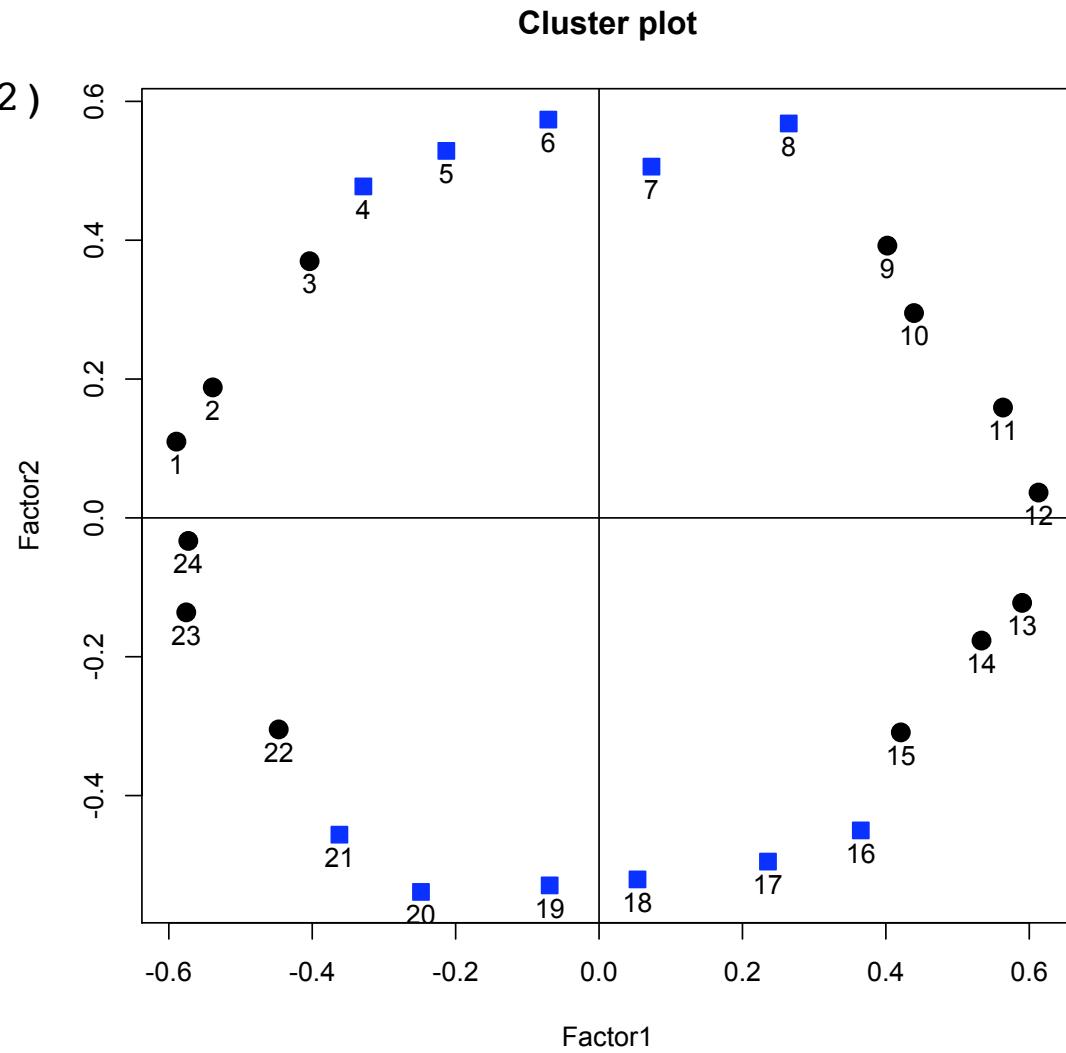
Correlations are similar

```
> round(cor(circ.cat)[,1:12],2)
```

	V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12
V1	1.00	0.32	0.30	0.21	0.24	0.13	0.03	-0.13	-0.21	-0.22	-0.31	-0.35
V2	0.32	1.00	0.31	0.26	0.23	0.18	0.06	-0.03	-0.13	-0.18	-0.30	-0.32
V3	0.30	0.31	1.00	0.29	0.28	0.22	0.14	0.08	0.03	-0.07	-0.09	-0.31
V4	0.21	0.26	0.29	1.00	0.29	0.33	0.20	0.16	0.07	0.01	-0.11	-0.19
V5	0.24	0.23	0.28	0.29	1.00	0.32	0.26	0.23	0.13	0.11	-0.04	-0.08
V6	0.13	0.18	0.22	0.33	0.32	1.00	0.31	0.39	0.19	0.08	0.06	-0.02
V7	0.03	0.06	0.14	0.20	0.26	0.31	1.00	0.28	0.22	0.15	0.15	0.07
V8	-0.13	-0.03	0.08	0.16	0.23	0.39	0.28	1.00	0.35	0.25	0.20	0.19
V9	-0.21	-0.13	0.03	0.07	0.13	0.19	0.22	0.35	1.00	0.30	0.29	0.23
V10	-0.22	-0.18	-0.07	0.01	0.11	0.08	0.15	0.25	0.30	1.00	0.35	0.28
V11	-0.31	-0.30	-0.09	-0.11	-0.04	0.06	0.15	0.20	0.29	0.35	1.00	0.35
V12	-0.35	-0.32	-0.31	-0.19	-0.08	-0.02	0.07	0.19	0.23	0.28	0.35	1.00
V13	-0.36	-0.34	-0.30	-0.25	-0.18	-0.07	0.00	0.03	0.20	0.20	0.32	0.33
V14	-0.33	-0.29	-0.29	-0.31	-0.26	-0.11	-0.01	0.07	0.17	0.19	0.23	0.30
V15	-0.30	-0.30	-0.23	-0.28	-0.20	-0.20	-0.13	-0.09	0.07	0.04	0.22	0.26
V16	-0.26	-0.24	-0.33	-0.35	-0.31	-0.24	-0.21	-0.17	-0.04	0.05	0.14	0.21
V17	-0.18	-0.22	-0.25	-0.35	-0.30	-0.29	-0.23	-0.20	-0.08	-0.05	0.07	0.12
V18	-0.08	-0.15	-0.27	-0.24	-0.31	-0.32	-0.26	-0.32	-0.12	-0.15	-0.08	-0.01
V19	-0.03	-0.07	-0.15	-0.22	-0.30	-0.29	-0.27	-0.31	-0.21	-0.20	-0.12	-0.07
V20	0.11	0.05	-0.12	-0.15	-0.23	-0.27	-0.31	-0.34	-0.35	-0.27	-0.22	-0.13
V21	0.15	0.11	0.02	-0.13	-0.14	-0.21	-0.27	-0.35	-0.34	-0.31	-0.22	-0.25
V22	0.25	0.21	0.03	-0.03	-0.04	-0.13	-0.22	-0.28	-0.30	-0.24	-0.31	-0.30
V23	0.27	0.35	0.16	0.13	0.07	-0.03	-0.06	-0.22	-0.27	-0.34	-0.36	-0.37
V24	0.36	0.25	0.22	0.20	0.08	0.03	-0.04	-0.24	-0.23	-0.26	-0.35	-0.37

Loadings are slightly smaller

```
> f2c <- factanal(circ.cat, 2)  
> factor.plot(f2c)
```



sem needs help

```
> mod.1crc <-edit(mod.1crc) #starting with the fa values
> summary(sem.2c,digits=2)

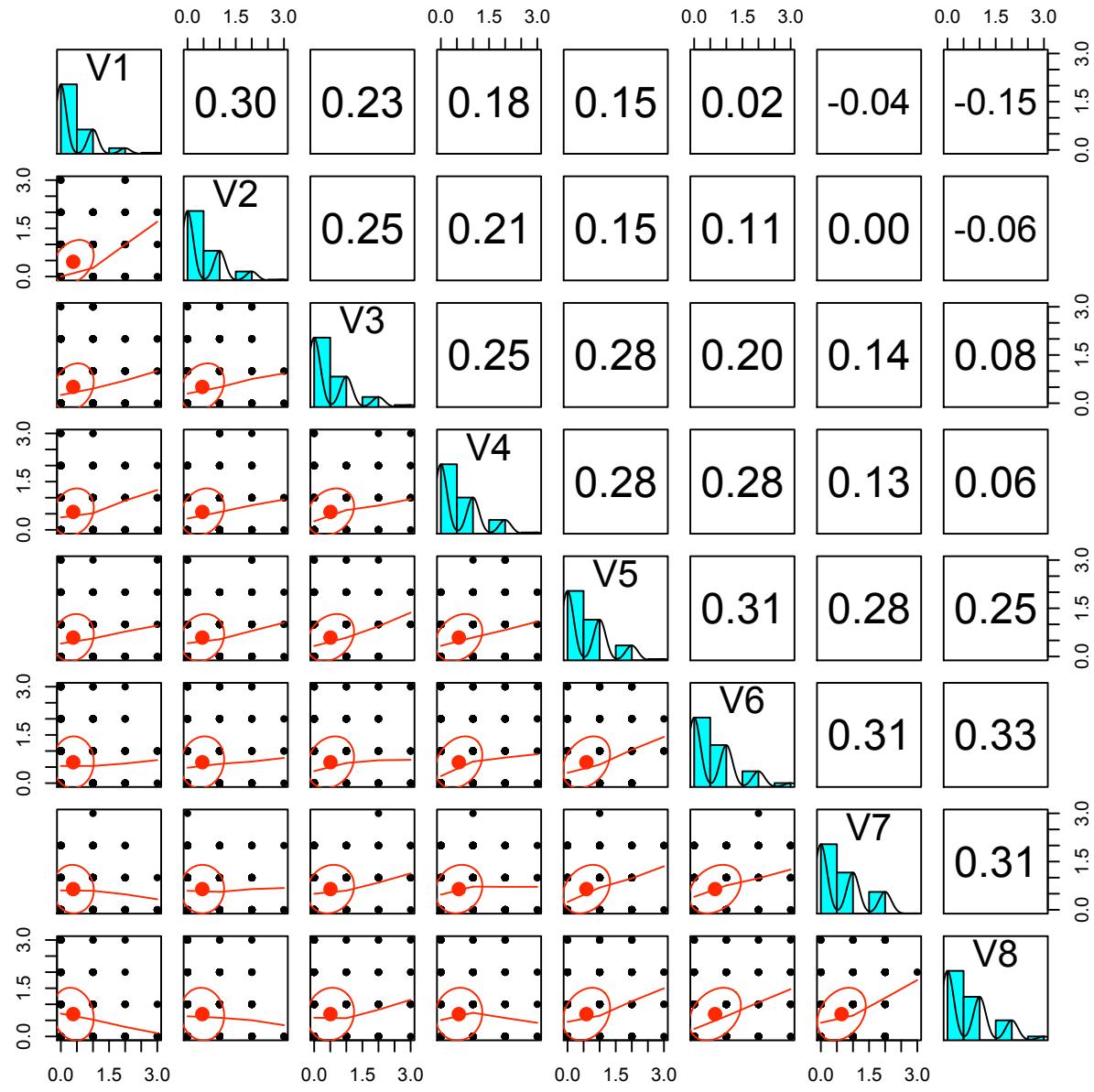
Model Chisquare = 258    Df = 235 Pr(>Chisq) = 0.15
Chisquare (null model) = 3021    Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.95
RMSEA index = 0.014  90% CI: (NA, 0.024)
Bentler-Bonnett NFI = 0.91
Tucker-Lewis NNFI = 1
Bentler CFI = 1
SRMR = 0.034
BIC = -1203

Normalized Residuals
  Min. 1st Qu. Median      Mean 3rd Qu.      Max.
-2.470 -0.427  0.045   0.025  0.463   2.190
```

skew categorical

```
> set.seed(42)
> circ.skewed <- sim.item(24,circum=TRUE,ybias=.5,categorical=TRUE,truncat=TRUE)
> colnames(circ.skewed) <- paste("V",1:24,sep="")
> describe(circ.skewed)

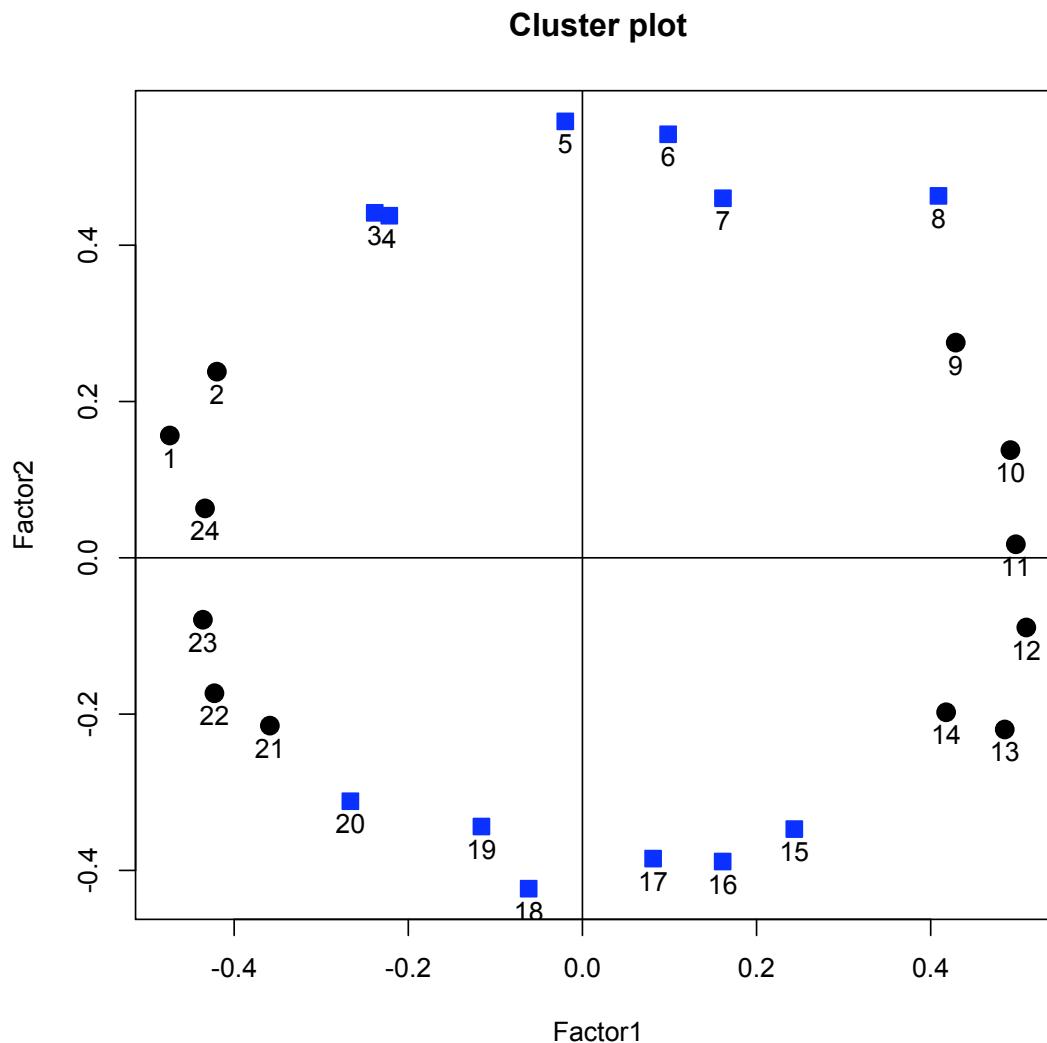
   var    n  mean    sd median trimmed  mad  min  max range skew kurtosis     se
V1    1 500 0.38 0.64      0    0.26  0   0   3    3 1.65    2.37  0.03
V2    2 500 0.46 0.68      0    0.33  0   0   3    3 1.31    0.97  0.03
V3    3 500 0.51 0.73      0    0.36  0   0   3    3 1.34    1.14  0.03
V4    4 500 0.56 0.73      0    0.44  0   0   3    3 1.05    0.19  0.03
V5    5 500 0.59 0.73      0    0.47  0   0   3    3 0.97    0.03  0.03
V6    6 500 0.65 0.81      0    0.52  0   0   3    3 1.06    0.36  0.04
V7    7 500 0.64 0.76      0    0.55  0   0   3    3 0.72   -0.81  0.03
V8    8 500 0.69 0.82      0    0.58  0   0   3    3 0.92   -0.05  0.04
V9    9 500 0.67 0.77      0    0.57  0   0   3    3 0.82   -0.26  0.03
V10  10 500 0.54 0.72      0    0.40  0   0   3    3 1.20    0.86  0.03
V11  11 500 0.47 0.71      0    0.32  0   0   3    3 1.38    1.15  0.03
V12  12 500 0.44 0.67      0    0.31  0   0   3    3 1.33    0.96  0.03
V13  13 500 0.39 0.65      0    0.26  0   0   3    3 1.54    1.62  0.03
V14  14 500 0.36 0.59      0    0.25  0   0   3    3 1.50    1.45  0.03
V15  15 500 0.30 0.57      0    0.18  0   0   3    3 1.97    3.67  0.03
V16  16 500 0.25 0.53      0    0.14  0   0   3    3 2.18    4.70  0.02
V17  17 500 0.20 0.46      0    0.09  0   0   3    3 2.42    6.06  0.02
V18  18 500 0.20 0.46      0    0.08  0   0   2    2 2.33    4.78  0.02
V19  19 500 0.18 0.42      0    0.08  0   0   2    2 2.21    4.21  0.02
V20  20 500 0.19 0.46      0    0.08  0   0   2    2 2.42    5.24  0.02
V21  21 500 0.21 0.49      0    0.10  0   0   3    3 2.38    5.46  0.02
V22  22 500 0.24 0.51      0    0.14  0   0   3    3 2.20    4.94  0.02
V23  23 500 0.28 0.56      0    0.16  0   0   3    3 2.03    3.78  0.02
V24  24 500 0.31 0.58      0    0.19  0   0   3    3 2.05    4.41  0.03
>
```



```
> pairs.panels(circ.skewed[,1:8])
```

Loadings are reduced

Angular
locations
are
preserved



```
> f2cs
```

Call:

```
factanal(x = circ.skewed, factors = 2)
```

Uniquenesses:

V1	V2	V3	V4	V5	V6	V7	V8	V9	V10	V11	V12	V13	V14
V15	V16	V17	V18	V19	V20	V21	V22						
0.751	0.767	0.748	0.759	0.688	0.697	0.762	0.618	0.740	0.739	0.752	0.732	0.716	0.786
0.820	0.823	0.845	0.817	0.868	0.832	0.825	0.791						
	V23	V24											
0.804	0.808												

Loadings:

	Factor1	Factor2											
V1	-0.474	0.157	V13	0.485	-0.220								
V2	-0.420	0.238	V14	0.418	-0.198								
V3	-0.239	0.441	V15	0.243	-0.347								
V4	-0.222	0.438	V16	0.161	-0.389								
V5		0.558	V17		-0.385								
V6		0.542	V18		-0.423								
V7	0.161	0.460	V19	-0.116	-0.344								
V8	0.409	0.463	V20	-0.267	-0.311								
V9	0.429	0.275	V21	-0.359	-0.215								
V10	0.492	0.138	V22	-0.423	-0.173								
V11	0.498		V23	-0.436									
V12	0.510		V24	-0.433									
				Factor1	Factor2								
			SS loadings	2.919	2.590								
			Proportion Var	0.122	0.108								
			Cumulative Var	0.122	0.230								

Test of the hypothesis that 2 factors are sufficient.

The chi square statistic is 238.55 on 229 degrees of freedom.

The p-value is 0.319

sem works as well

```
> sem.cs <- sem(mod.1crc,cor(circ.skewed),500)
> summary(sem.cs,digits=2)

Model Chisquare = 252    Df = 235 Pr(>Chisq) = 0.21
Chisquare (null model) = 1864    Df = 276
Goodness-of-fit index = 0.96
Adjusted goodness-of-fit index = 0.95
RMSEA index = 0.012  90% CI: (NA, 0.023)
Bentler-Bonnett NFI = 0.86
Tucker-Lewis NNFI = 0.99
Bentler CFI = 0.99
SRMR = 0.034
BIC = -1208

Normalized Residuals
  Min. 1st Qu. Median      Mean 3rd Qu.      Max.
-1.57   -0.23    0.25     0.27    0.80     2.31
```

Cross loadings helped

Parameter Estimates

	Estimate	Std Error	z value	Pr(> z)	
F1V1	-0.498	0.047	-10.5	0.0e+00	V1 <--- Factor1
F1V2	-0.468	0.048	-9.9	0.0e+00	V2 <--- Factor1
F2V2	0.114	0.048	2.4	1.8e-02	V2 <--- Factor2
F1V3	-0.349	0.048	-7.2	4.4e-13	V3 <--- Factor1
F2V3	0.359	0.048	7.5	5.7e-14	V3 <--- Factor2
F1V4	-0.333	0.048	-6.9	5.0e-12	V4 <--- Factor1
F2V4	0.360	0.048	7.5	6.8e-14	V4 <--- Factor2
F1V5	-0.172	0.048	-3.5	3.9e-04	V5 <--- Factor1
F2V5	0.530	0.047	11.3	0.0e+00	V5 <--- Factor2
F1V6	-0.054	0.049	-1.1	2.6e-01	V6 <--- Factor1
F2V6	0.550	0.047	11.7	0.0e+00	V6 <--- Factor2
F2V7	0.488	0.048	10.2	0.0e+00	V7 <--- Factor2
F1V8	0.267	0.048	5.6	2.0e-08	V8 <--- Factor1
F2V8	0.557	0.046	12.1	0.0e+00	V8 <--- Factor2
F1V9	0.337	0.048	7.0	3.1e-12	V9 <--- Factor1
F2V9	0.382	0.048	8.0	1.1e-15	V9 <--- Factor2
F1V10	0.436	0.047	9.2	0.0e+00	V10 <--- Factor1
F2V10	0.266	0.048	5.6	2.6e-08	V10 <--- Factor2
F1V11	0.475	0.047	10.0	0.0e+00	V11 <--- Factor1
F2V11	0.155	0.048	3.2	1.3e-03	V11 <--- Factor2
F1V12	0.515	0.047	11.0	0.0e+00	V12 <--- Factor1
F1V13	0.526	0.047	11.2	0.0e+00	V13 <--- Factor1
F1V14	0.456	0.048	9.6	0.0e+00	V14 <--- Factor1
F2V14	-0.074	0.049	-1.5	1.3e-01	V14 <--- Factor2
F1V15	0.327	0.049	6.7	1.6e-11	V15 <--- Factor1
F2V15	-0.267	0.049	-5.5	4.6e-08	V15 <--- Factor2
F1V16	0.260	0.049	5.3	8.9e-08	V16 <--- Factor1

Conclusions

- I. Characteristics of items will affect solutions
- II. In general, the multivariate methods will recover structure, but sometimes need significant help in starting values