

# UG structure learning

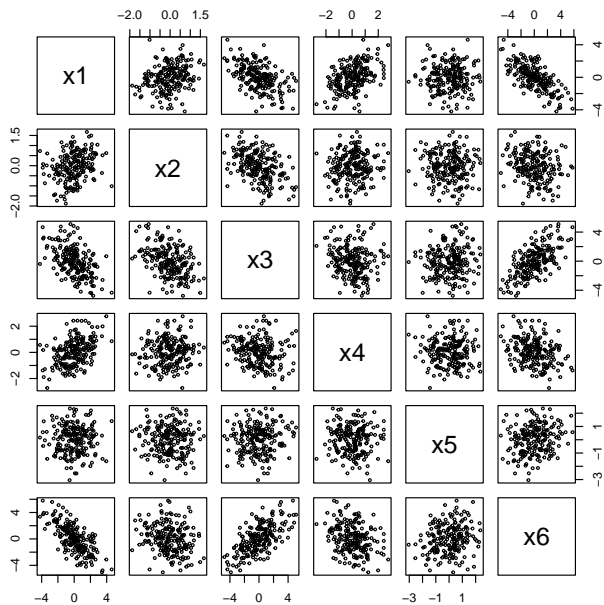
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## Data on six simulated variables

### The data set

```
source("simdat.R")  
  
pairs(simdat, pch = 1, cex = .5)
```



```
su <- apply(simdat, 2, function(x) rbind(mean(x), sd(x)))  
rownames(su) <- c("mean", "sd")  
print(round(su, 3))
```

	x1	x2	x3	x4	x5	x6
mean	-0.006	-0.036	-0.049	0.041	0.029	0.030
sd	1.576	0.699	2.010	1.004	1.046	2.137

```
round(cor(simdat), 3)
```

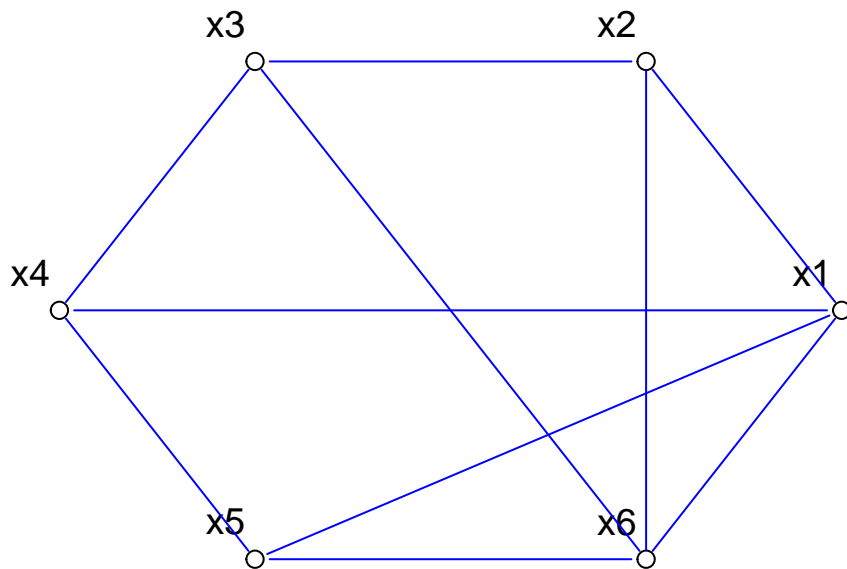
	x1	x2	x3	x4	x5	x6
x1	1.000	0.265	-0.514	0.440	0.002	-0.717
x2	0.265	1.000	-0.423	0.075	-0.032	-0.226
x3	-0.514	-0.423	1.000	-0.158	0.082	0.655
x4	0.440	0.075	-0.158	1.000	-0.045	-0.301
x5	0.002	-0.032	0.082	-0.045	1.000	0.187
x6	-0.717	-0.226	0.655	-0.301	0.187	1.000

## Partial correlations

```
S<- cov(simdat)
P <- parcor(S)
round(P, 3)
```

	x1	x2	x3	x4	x5	x6
x1	1.000	0.132	-0.046	0.347	0.205	-0.573
x2	0.132	1.000	-0.365	-0.018	-0.037	0.141
x3	-0.046	-0.365	1.000	0.074	-0.044	0.485
x4	0.347	-0.018	0.074	1.000	-0.057	-0.005
x5	0.205	-0.037	-0.044	-0.057	1.000	0.257
x6	-0.573	0.141	0.485	-0.005	0.257	1.000

```
G <- 0+(abs(P)>0.05)
diag(G) <- 0
drawGraph(G, layout = layout_in_circle)
```



## MLE fit graph of $G$

```
out <- fitConGraph(G,S, n = ncol(simdat))
out
```

\$Shat

	x1	x2	x3	x4	x5	x6
x1	2.485218932	0.291400308	-1.5457704	0.69693839	0.002859092	-2.4147010
x2	0.291400308	0.487919173	-0.5933878	0.06524697	-0.008774948	-0.3367596
x3	-1.545770407	-0.593387850	4.0395543	-0.31966182	0.240089720	2.8129414
x4	0.696938390	0.065246971	-0.3196618	1.00875647	-0.047098406	-0.6454886
x5	0.002859092	-0.008774948	0.2400897	-0.04709841	1.093722417	0.4179746
x6	-2.414701046	-0.336759557	2.8129414	-0.64548858	0.417974557	4.5688786

\$dev

[1] 0.03685723

\$df

[1] 5

\$it

[1] 4

Concentration matrix

```
Khat <- solve(out$Shat)
round(Khat,3)
```

	x1	x2	x3	x4	x5	x6
x1	0.997	-0.220	0.000	-0.380	-0.202	0.476
x2	-0.220	2.566	0.424	0.000	0.000	-0.188
x3	0.000	0.424	0.506	-0.051	0.000	-0.287
x4	-0.380	0.000	-0.051	1.241	0.066	0.000
x5	-0.202	0.000	0.000	0.066	0.990	-0.188
x6	0.476	-0.188	-0.287	0.000	-0.188	0.651

**Check decomposability**

```
is.triangulated(G)
```

```
[1] FALSE
```