The trouble with high-dimensional multi-level data

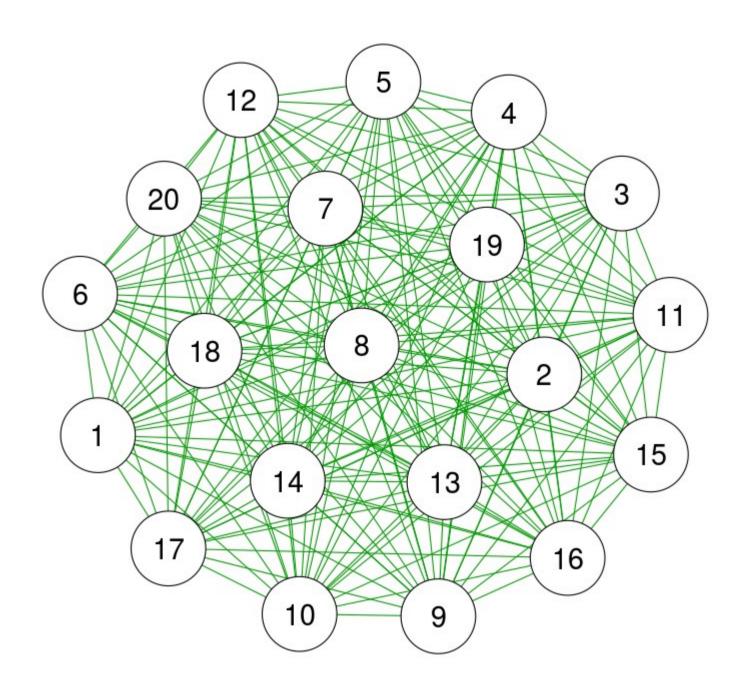
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#### The data

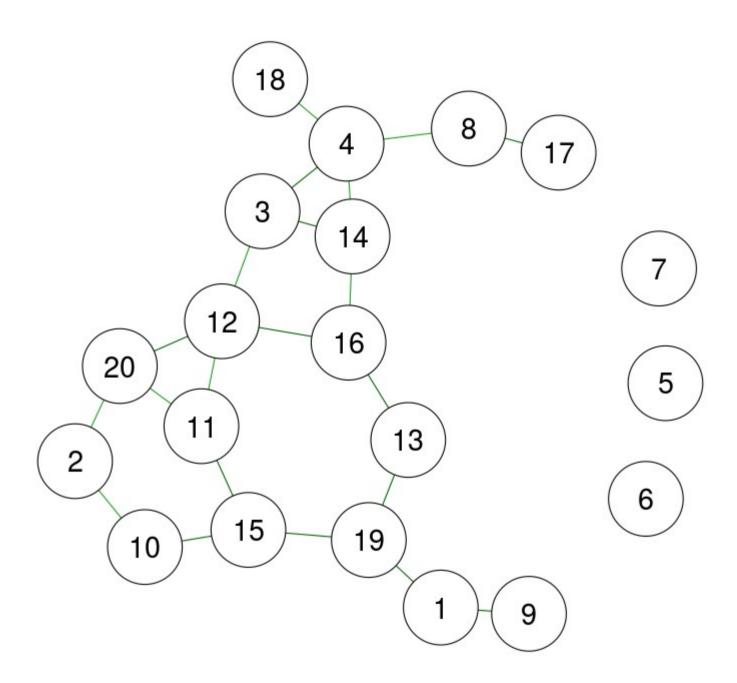
- 'High-dimensional' nested data
  - Say, 20 variables
  - People measured 5 times a day for 10 days
  - Random effects needed
  - Everything correlates with everything
  - Yet all associations **potentially** interesting

### The problem

- Univariate associations no go
  - Everything inter-correlating -> uninformative
  - Pervasive spuriousness -> distorted
- We want
  - A sparse pattern
  - Of unique associations

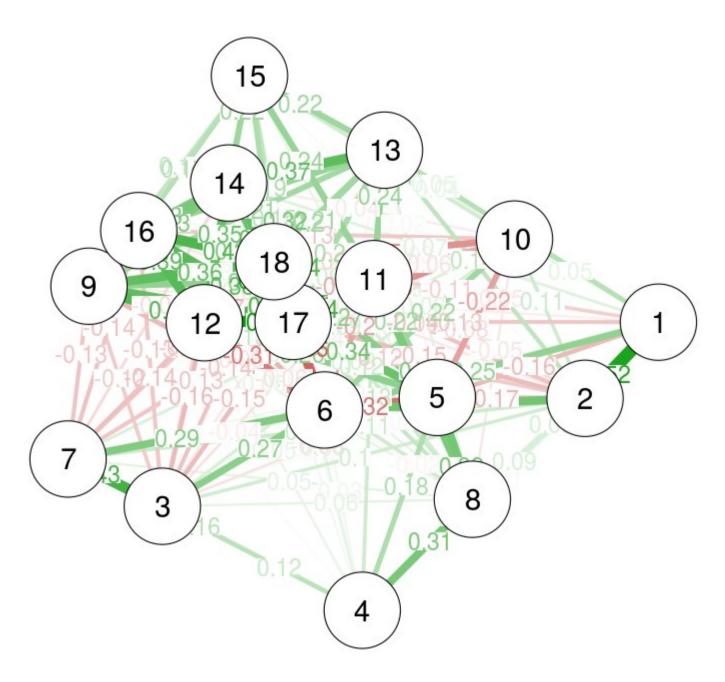


- > require(qgraph)
- > qgraph(yourMatrix)

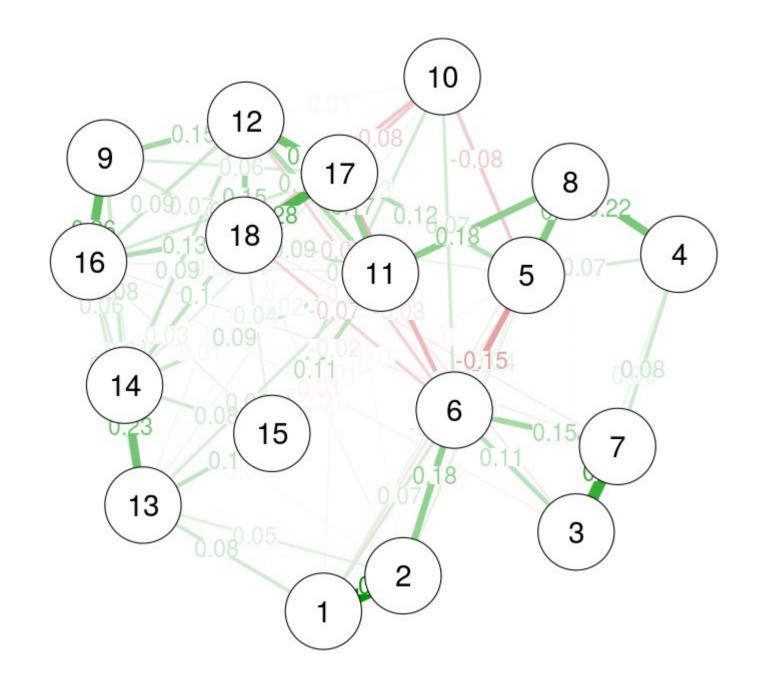


# Matrix of fixed effects, univariate lmer()

```
## d: 18 personality & situation items plus ID (8,188 observations)
## Standardized within individuals
> k = ncol(d) - 1
> r = matrix(ncol=k, nrow=k, 1)
> for(a in 1:k) {
+ for(b in a:k) {
+ if(b != a) {
        r[b,a] = fixef(lmer(d[,a] \sim d[,b] + (-1 + d[,b]|ID),d))[-1]
+
+ }
+ }
```



- > r[upper.tri(r)] = t(r)[upper.tri(r)]
- > qgraph(r, edge.labels=T)

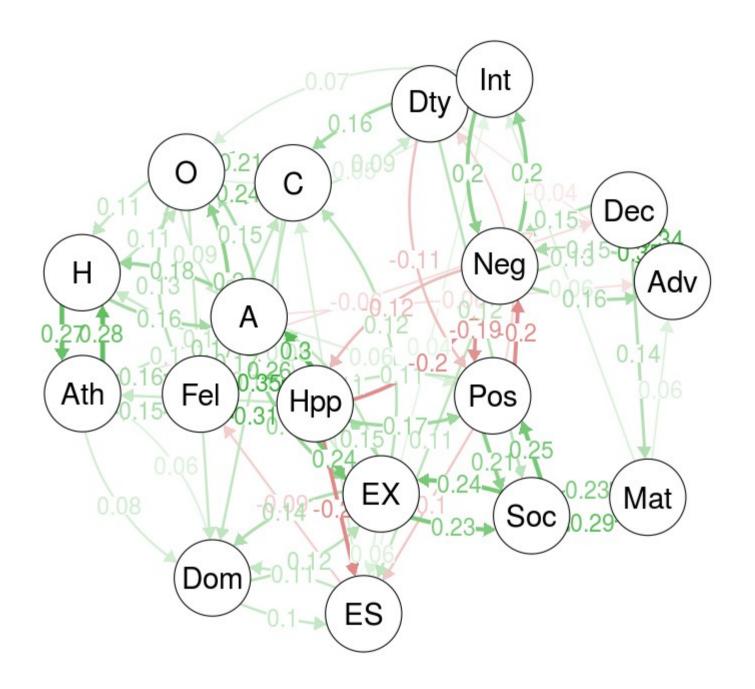


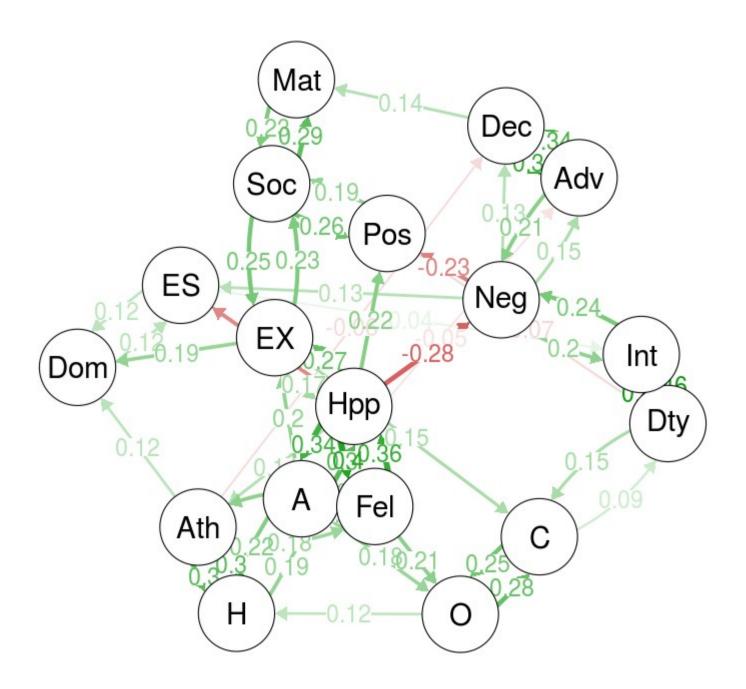
## Let every variable be predicted by all variables at once

- Hoping that most associations would become very small
  - Convergence issues unless random effects uncorrelated (not realistic)

# Step-wise lmer (really no package for this?!)

- Iterative model selection
  - AIC, BIC, R-squared
- Everything still predicts everything
  - Fit always improves
  - No sparsity, convergence problems
  - Manual limit for predictors (top 3, 5, 7)





### Ideas welcome!

• Pulling out most unique associations from high-dimensional multi-level data

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