

Breaking the Assumption of Group Homogeneity in Networks:

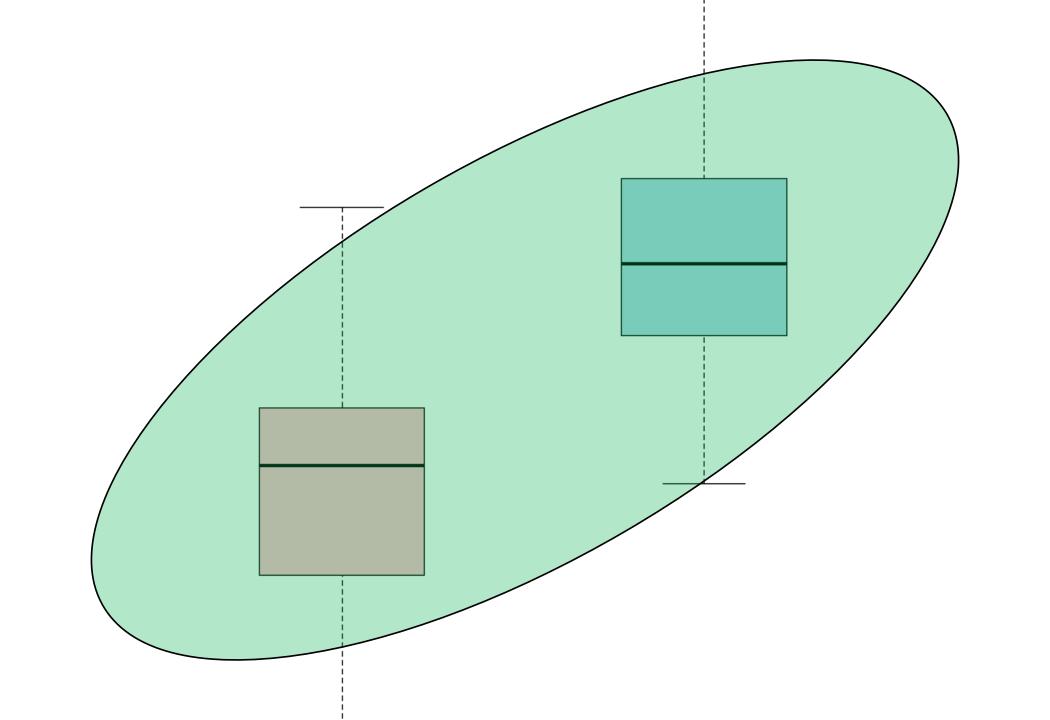
Partitioning Networks with Machine Learning

PAYTON JONES¹, PATRICK MAIR¹, THORSTEN SIMON², & ACHIM ZEILEIS²

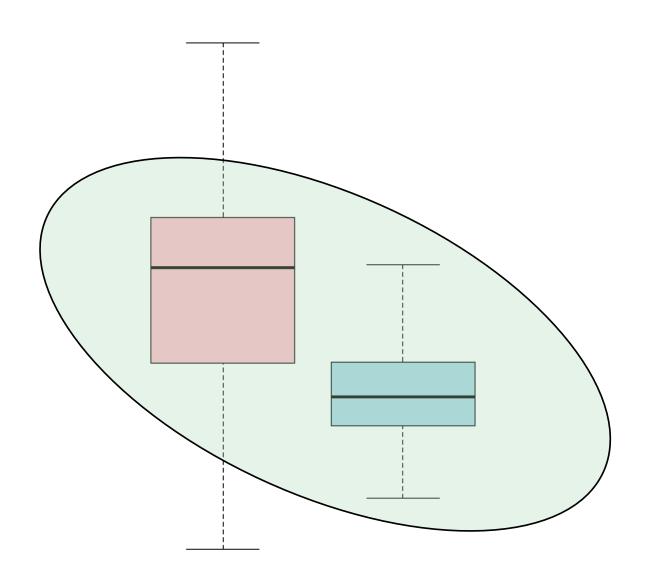
¹HARVARD UNIVERSITY

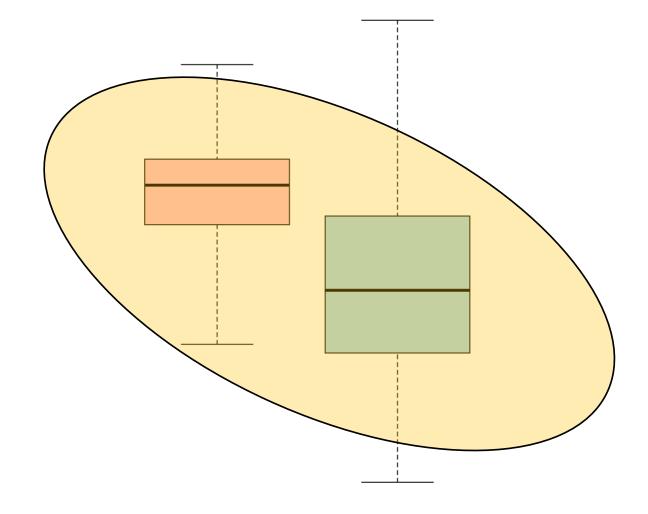
²UNIVERSITÄT INNSBRUCK



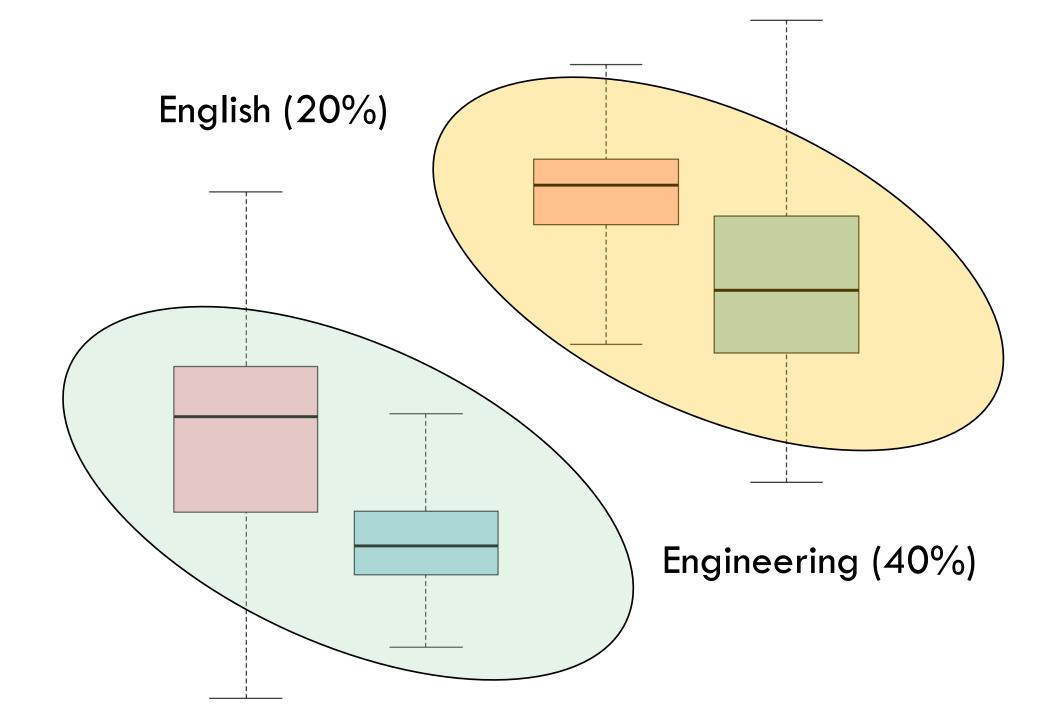


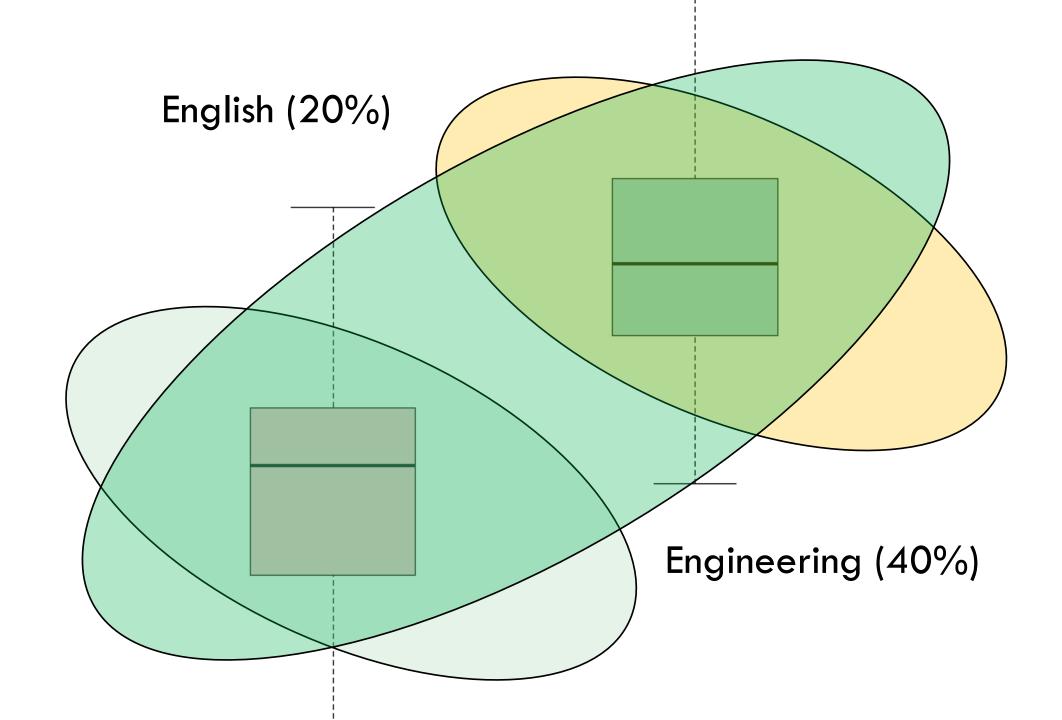
English

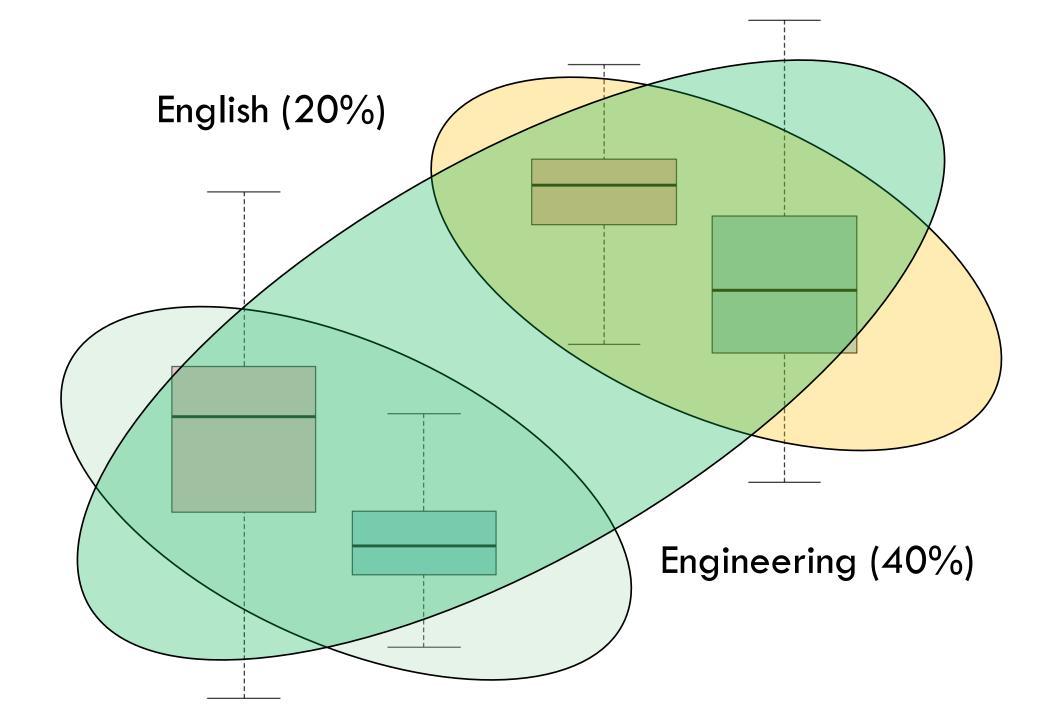




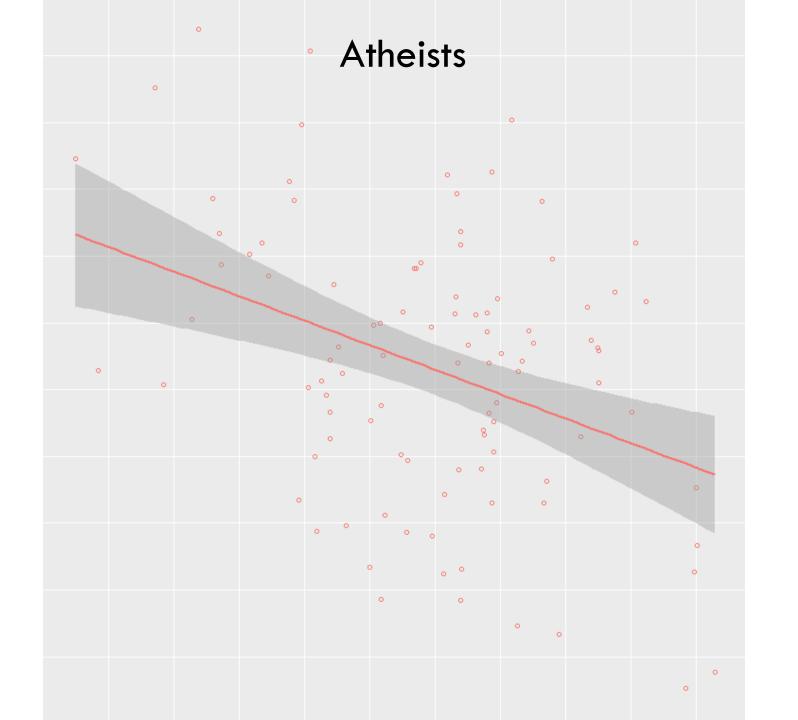
Engineering

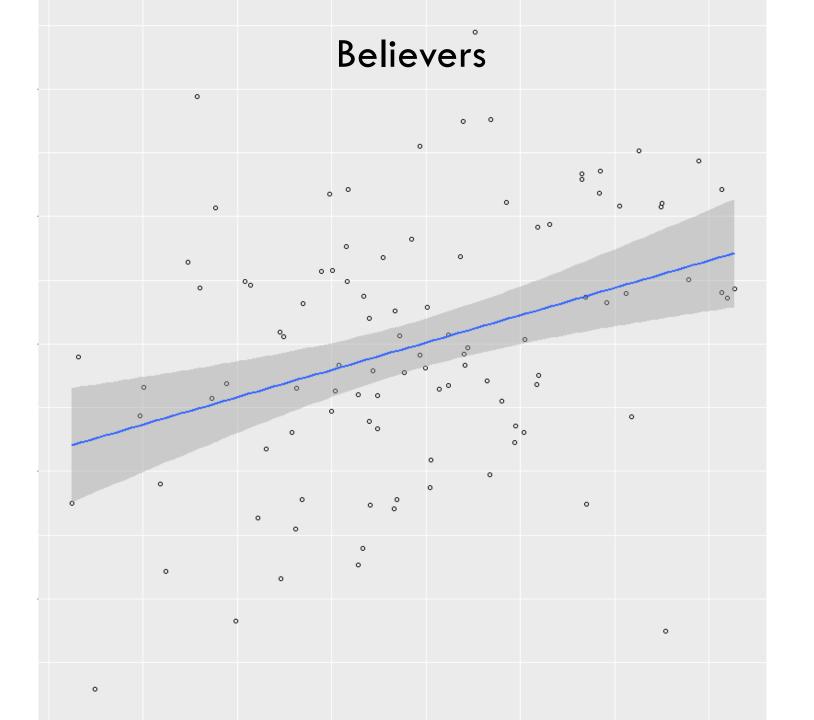


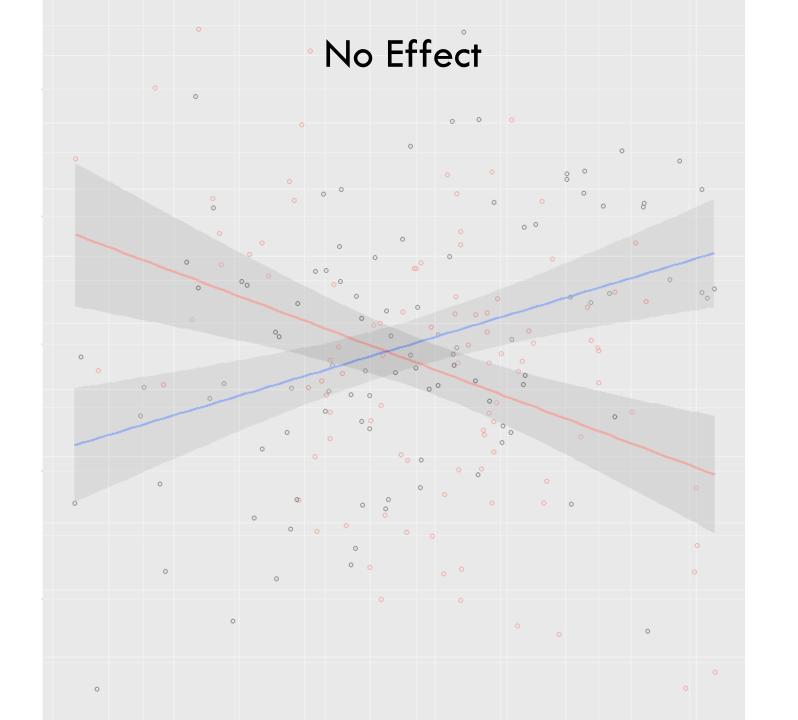


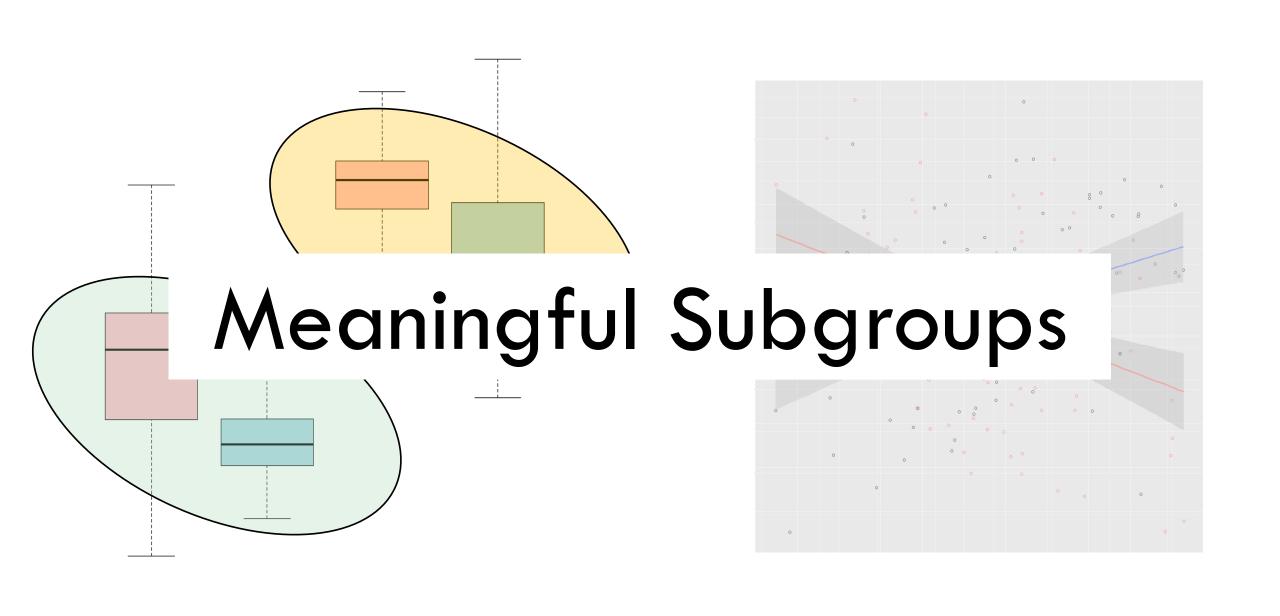


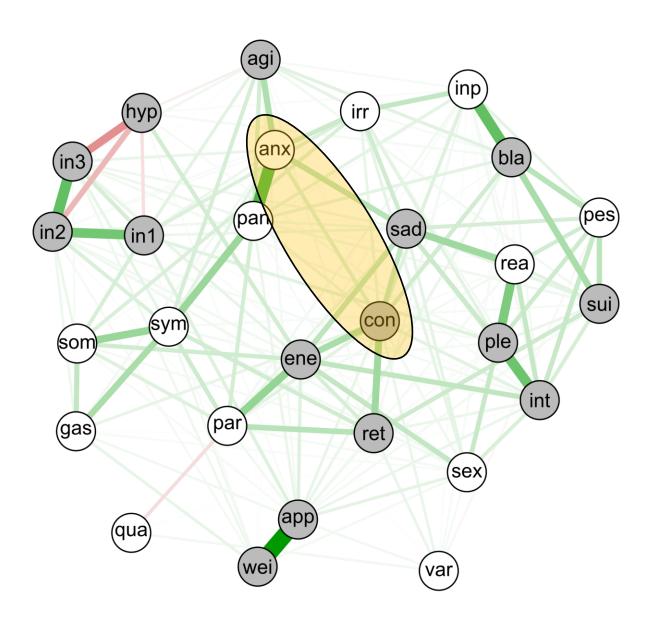






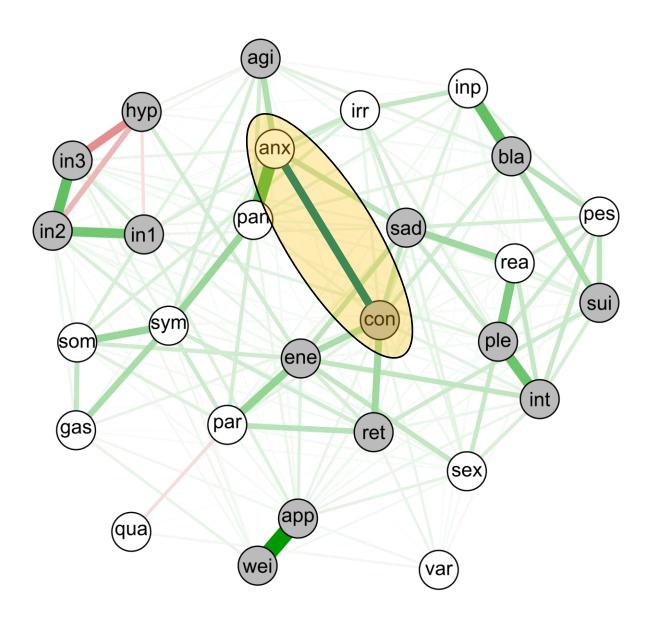






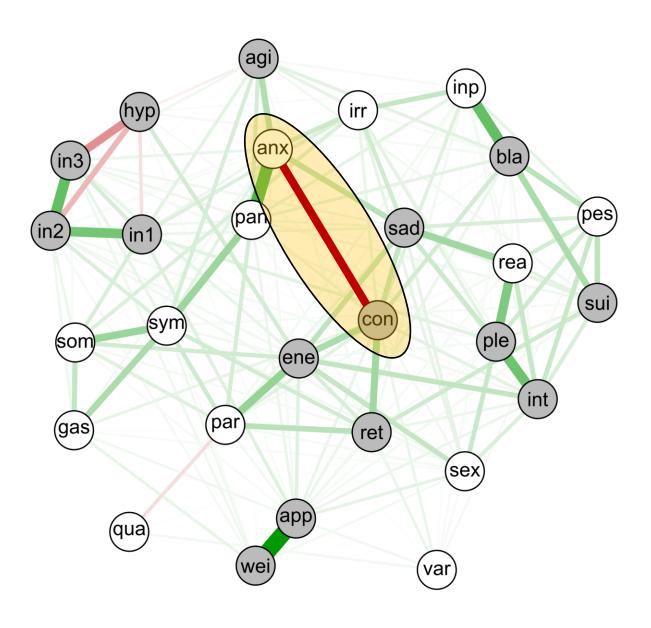
- DSM criterion symptom
- O Non-DSM symptom

Energy loss Sadness Sympathetic arousal Interest loss Pleasure loss Concentr. problems Panic / phobia Appetite probl. Mid insomnia Anxious / tense Self-blame Late insomnia Paralysis Early insomnia Retardation Interpers. sensitivity Mood reactivity Suicidal ideation Pessimism Somatic complaints Irritability Weight probl. Agitation Sexual disinterest Hypersomnia Gastroint. probl. Diurnal variation Mood quality



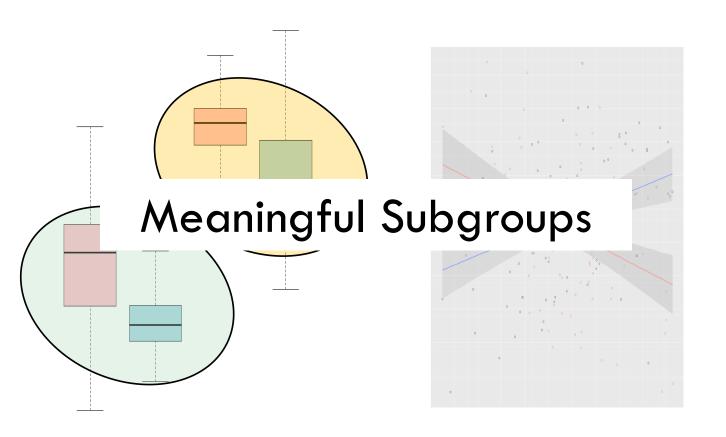
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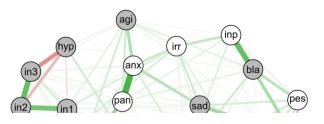
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Energy loss
Sadness
Sympathetic arousal
Interest loss
Pleasure loss
Concentr. problems
Panic / phobia
Appetite probl.
Mid insomnia

Meaningful Subgroups in Networks

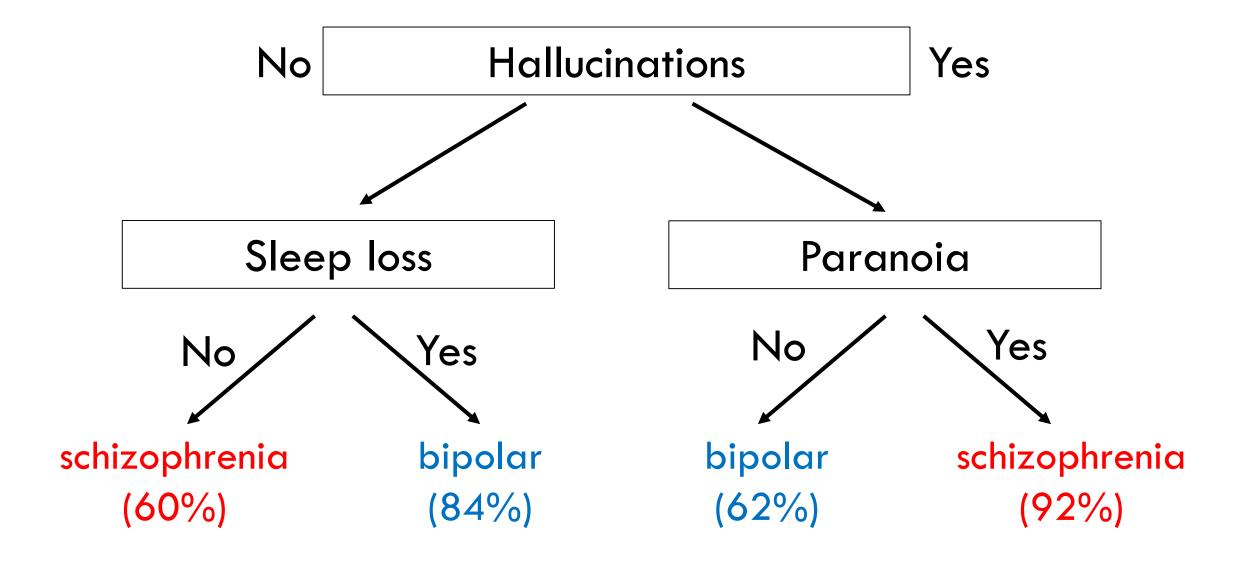


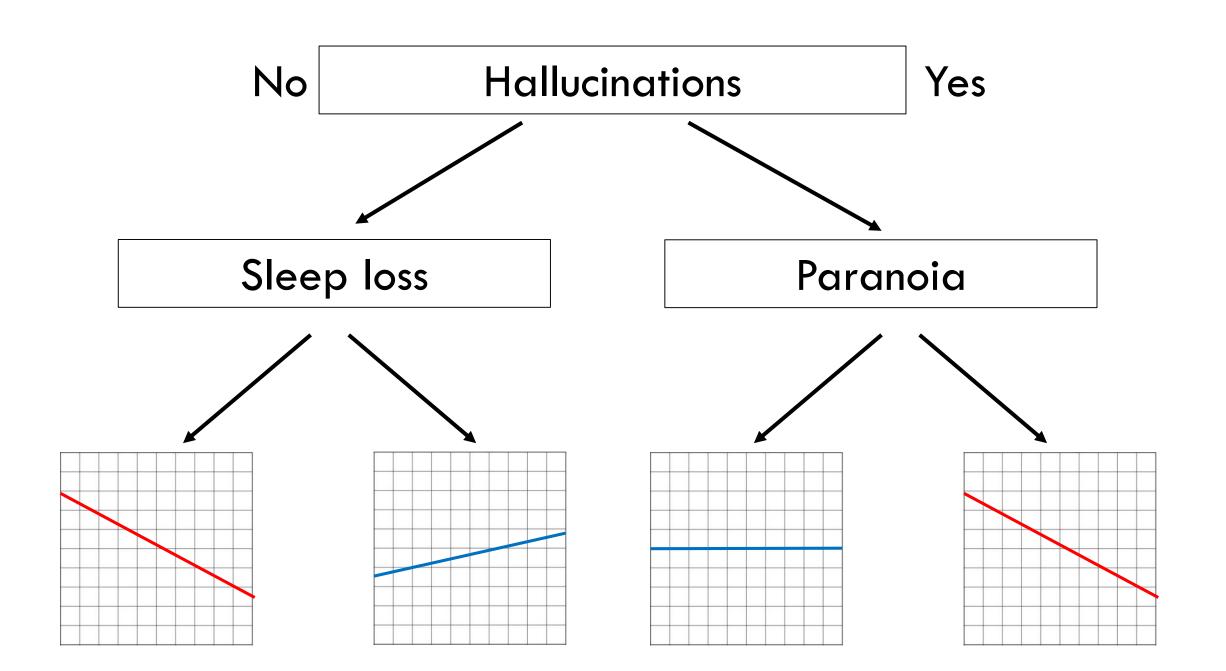
DSM criterion symptom

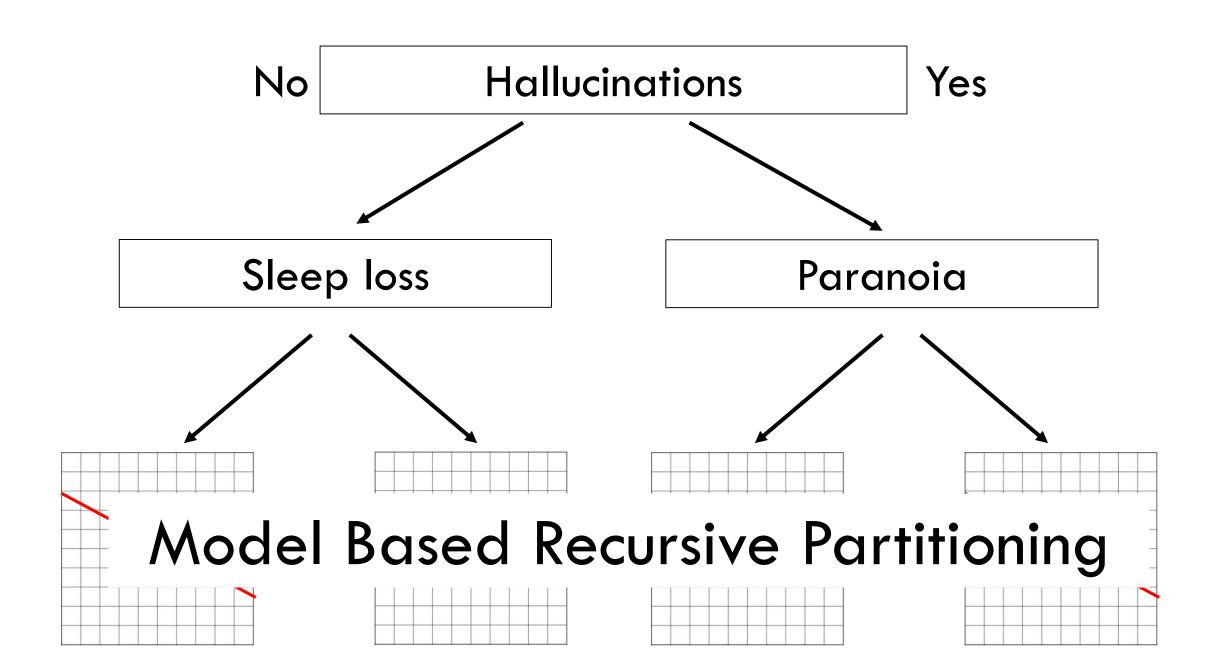
O Non-DSM symptom

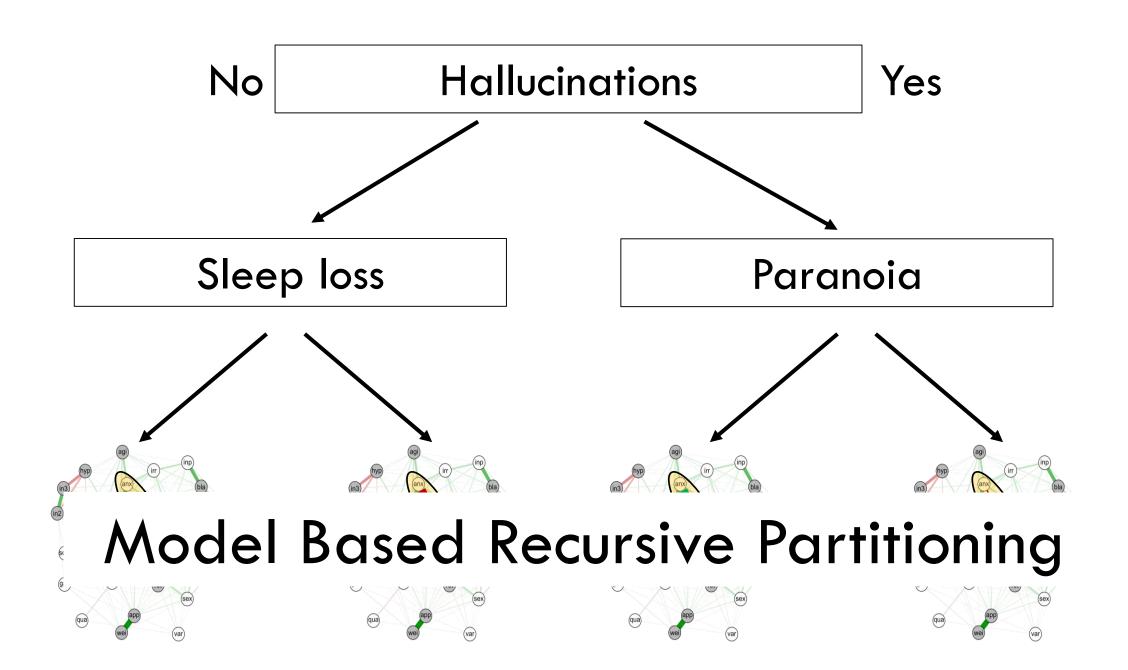
Somatic complaints
Irritability
Weight probl.
Agitation
Sexual disinterest
Hypersomnia
Gastroint. probl.
Diurnal variation
Mood quality

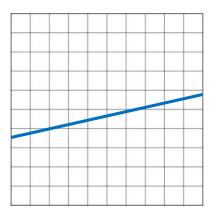
Decision Trees

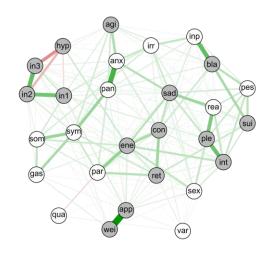








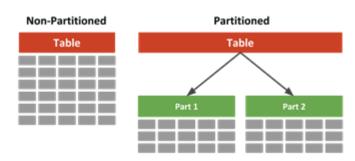


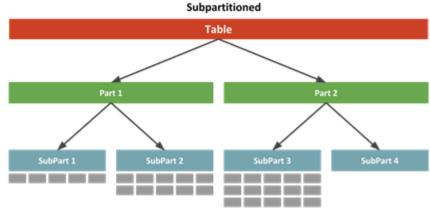


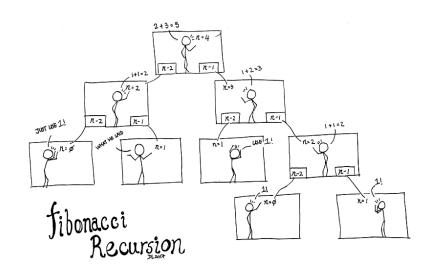
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Mood quality



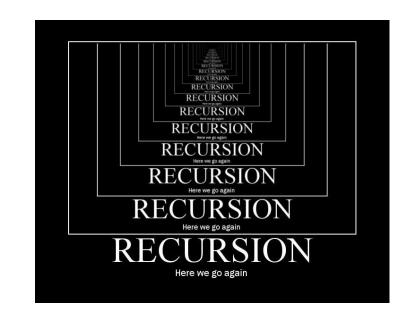


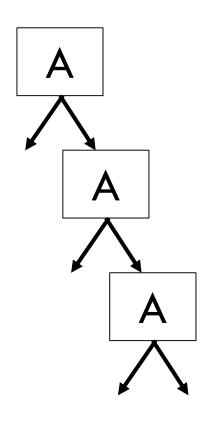


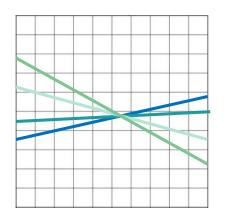
Model Based

Recursive

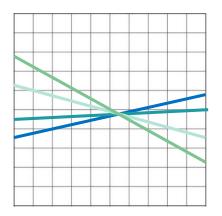
Partitioning



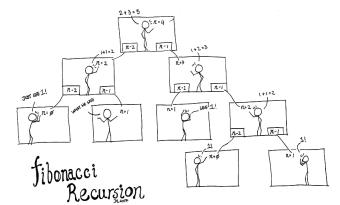




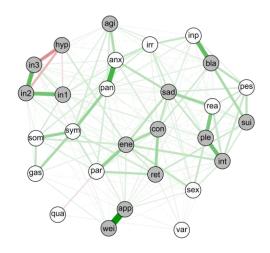
 $\frac{dy}{dx}$ f'(x)



RECURSION Here we go again RECURSION Here we go again



Model Based Recursive Partitioning

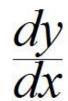


DSM criterion symptom

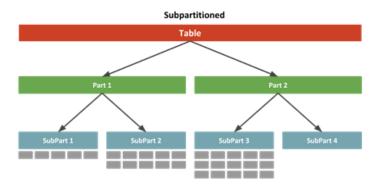
O Non-DSM symptom

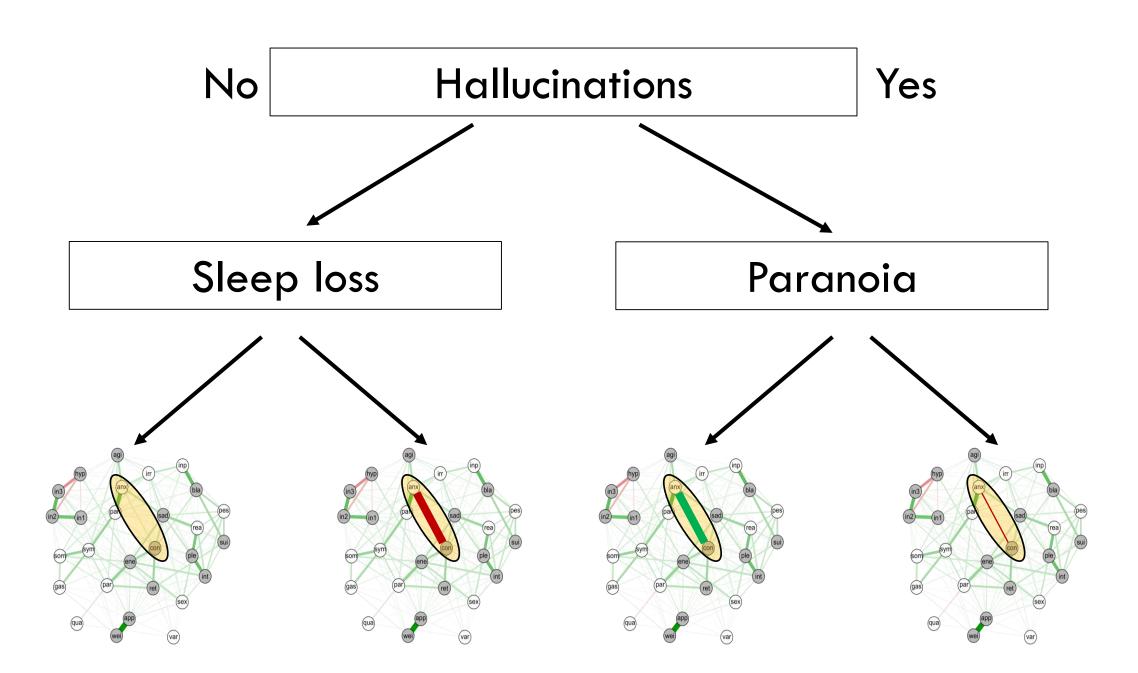
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Energy loss Sadness



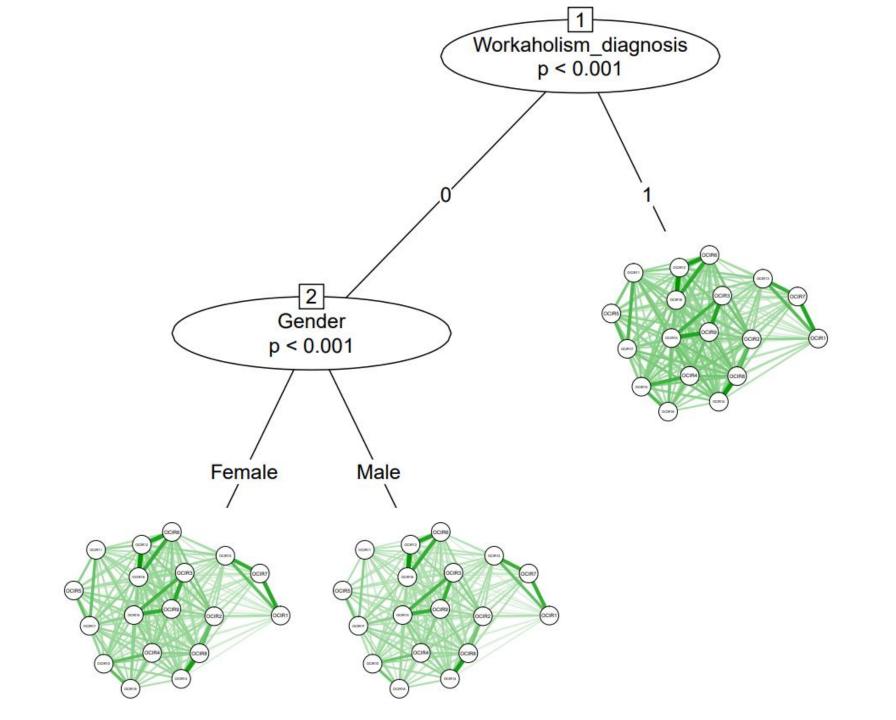
f'(x)

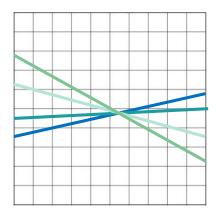


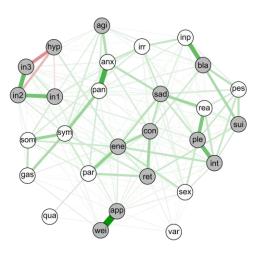


- > install.packages("networktree")
- > library(networktree)

- > data("workaholic")
- > nodevars <- paste("OCIR",1:18,sep="")
- > splitvars <- c("Workaholism_diagnosis","Gender")
- > myTree<- networktree(nodevars=workaholic[,nodevars]) splitvars=workaholic[,splitvars])
- > plot(myTree)

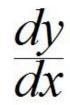


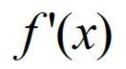


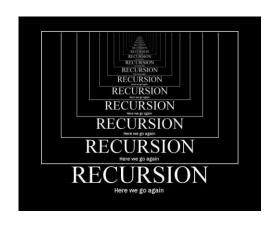


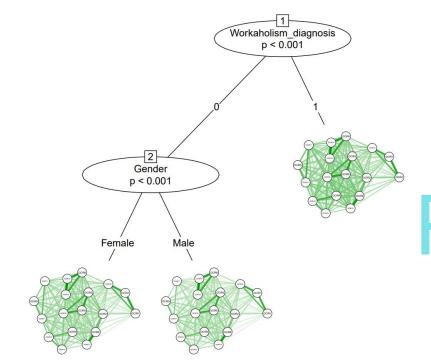
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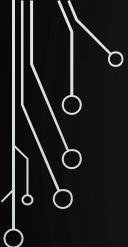
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THANK YOU!

Co-Authors:

Patrick Mair
Achim Zeileis
Thorsten Simon

Learn more about the method:
networktree R package
paytonjjones.wordpress.com/networktree



