

Tasks

Introduction to structural equation modeling and mixed models in

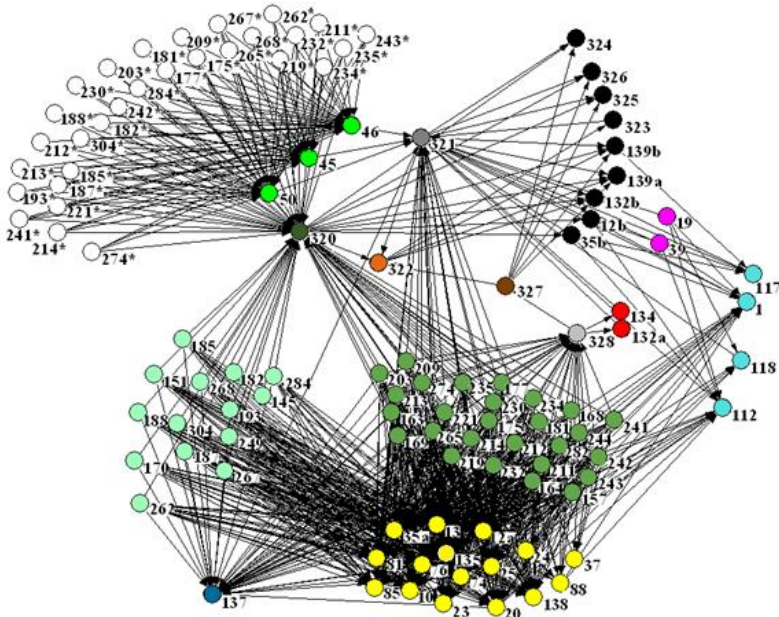
Day 9: SEM

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Day 9 Task 1

Effects of land use on arthropod food webs in grasslands



Food web length (1,2,3)



Net sampling of arthropods in grasslands

235 grasslands

Grazing type

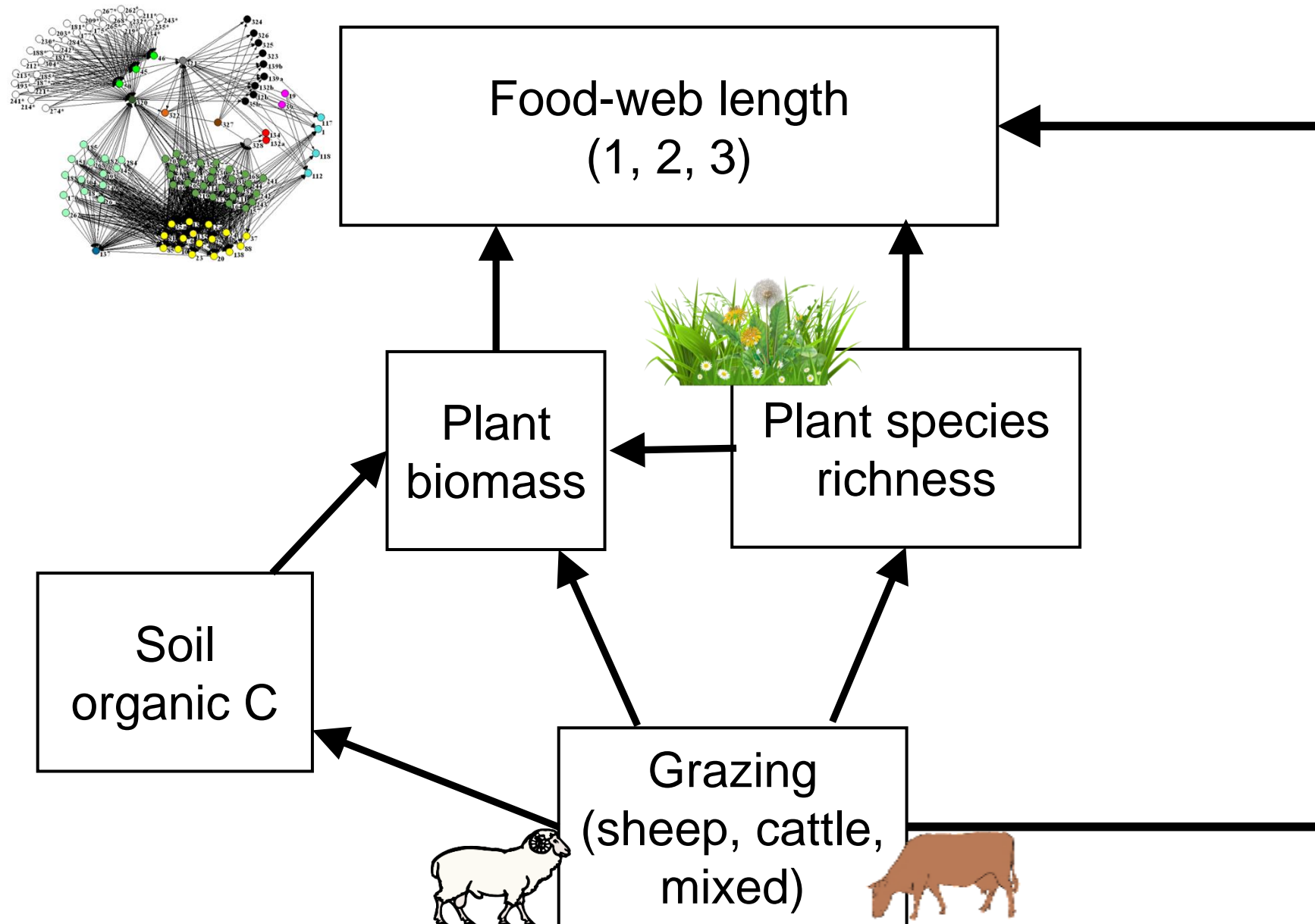
(“sheep”, “cattle”, or “mixed grazing”)

```
read.csv("Data/Food_web_data_2.csv")
```

```
'data.frame': 235 obs. of 5 variables:
 $ Gr_type   : chr  "sheep" "sheep" "sheep" ...
 $ soil_C    : num  1.336 1.631 1.577 ...
 $ plant_sr  : num  5.07 28.39 24.52 ...
 $ plant_biom: num  185 207 224 238 203 ...
 $ FW.length : int   1 1 1 1 1 1 1 1 1 ...
```

Day 9 Task 1

Effects of land use on food webs in grasslands



Day 9 Task 1

Effects of land use on food webs in grasslands

Gr_type (grazing type) is your exogenous nominal categorical variables. Test **Gr_type** (as a part of the SEM model on fig. 1) using marginal means in *piecewiseSEM*.

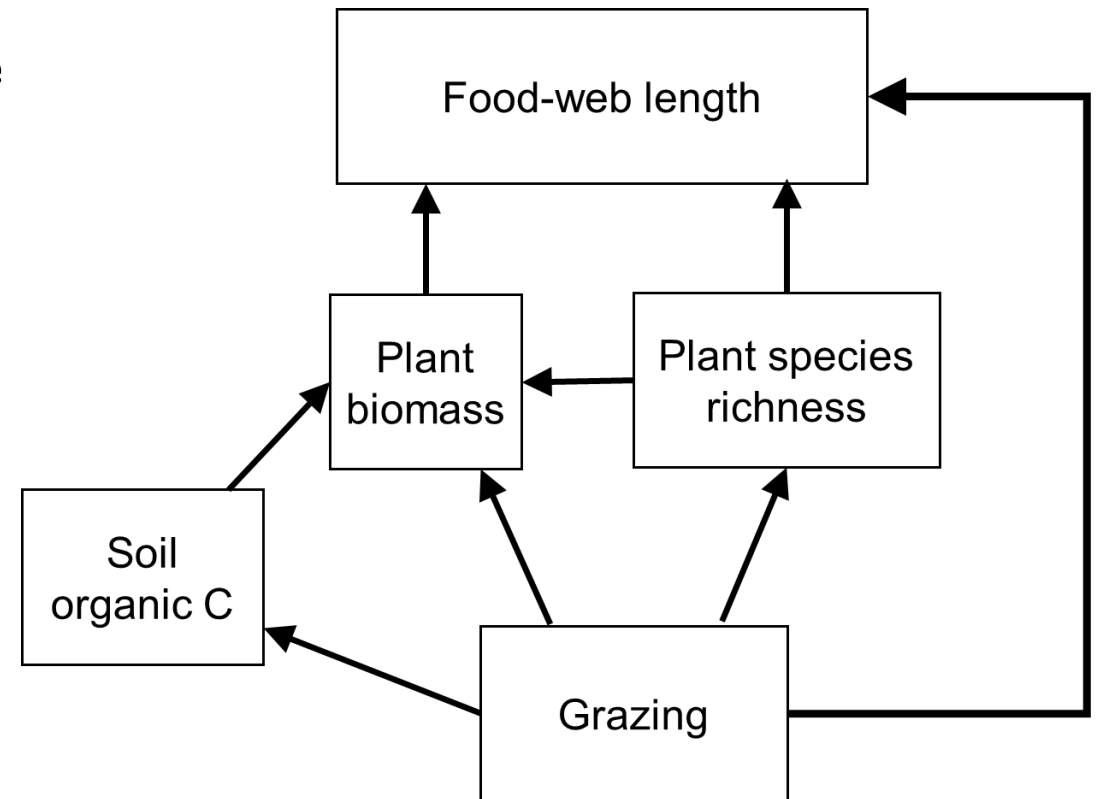


fig. 1

Day 9 Task 2

The Effects of Grazing on Finnish Coastal Meadows



Photo: Jorma Pessa

```
# meadow data
library(pieewiseSEM)
data(meadows)

> str(meadows)
'data.frame': 354 obs. of  4 variables:
 $ grazed: int  1 1 1 1 1 1 1 1 1 1 ...
 $ mass  : num  461.6 704.8 1278.8 76.4 ...
 $ elev  : num  19.2 18.8 12.5 38.2 37.4 ...
 $ rich  : int   6 2 1 6 14 12 6 10 10 17 ...
```

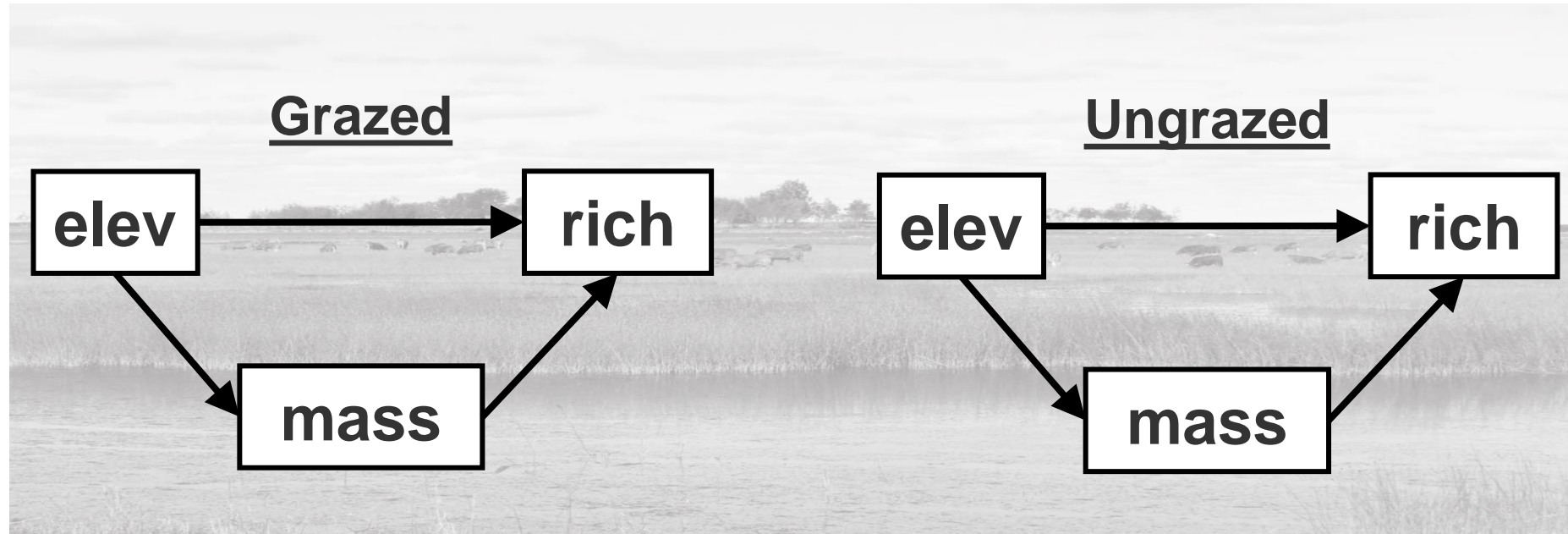
grazed is our grouping variable

Data:

Jutila, H. (1997) Vascular plant species richness in grazed and ungrazed coastal meadows, SW Finland. - Ann. Bot. Fenn. 34:245-263.

Grace, J.B. and Jutila, H. (1999) The relationship between species density and community biomass in grazed and ungrazed coastal meadows. Oikos, 85:398-408.

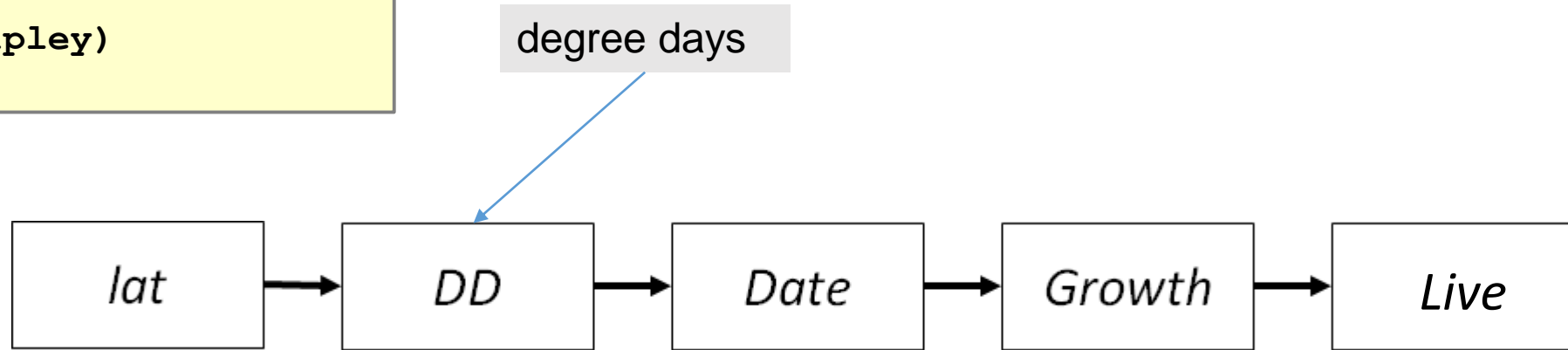
Hypothetical model



Task:

Perform the multigroup analysis for this hypothetical SEM model using *piecewiseSEM*

```
# Shipley data  
library(piecewiseSEM)  
data(shipley)
```



- Dataset: predicting latitude effect on survival of a tree species
- Repeated measures on 5 trees at 20 sites from 1970-2006
- Live (0/1) influenced by phenology (degree days until bud break, Julian days until bud break), size (stem diameter growth)

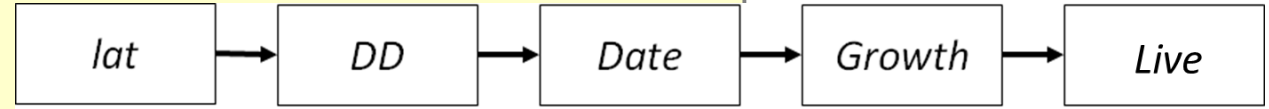
Day 9 Task 3

```
# Shipley data
library(piecewiseSEM)
data(shipley)
> str(shipley)
'data.frame': 1900 obs. of 9 variables:
 $ site      : int  1 1 1 1 1 1 1 1 1 1 ...
 $ tree      : int  1 2 3 4 5 1 2 3 4 5 ...
 $ lat       : num  40.4 40.4 40.4 40.4 40.4 ...
 $ year      : int  1970 1970 1970 1970 1970 1972 1972 1972 1972
1972 ...
 $ Date      : num  115 118 116 111 121 ...
 $ DD        : num  161 159 160 161 157 ...
 $ Growth    : num  61.4 43.8 44.7 48.2 50 ...
 $ Survival  : num  1 0.843 0.944 0.957 0.976 ...
 $ Live      : int  1 1 1 1 1 1 1 1 1 1 ...
```


Day 9 Task 3

LMM, and GLMM in SEM

```
library(nlme)
library(lme4)
```



```
lme(DD ~ lat, random = ~ 1 | site / tree, na.action = na.omit,  
    data = shipley),
```

```
lme(Date ~ DD, random = ~ 1 | site / tree, na.action = na.omit,  
    data = shipley),
```

```
lme(Growth ~ Date, random = ~ 1 | site / tree, na.action = na.omit,  
    data = shipley),
```

```
glmer(Live ~ Growth + (1 | site) + (1 | tree),  
      family = binomial(link = "logit"), data = shipley)
```

Task: Use these sub-models in piecewiseSEM as a part of the SEM model shown above. Think about the study design and explain the results.