

Introduction to structural equation modelling

Basic modelling

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Quick introduction of participants

- Who are you?

Quick introduction of participants

- Who are you?
- Why do you want to learn about SEM?

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- Who are you?
- Why do you want to learn about SEM?
- What is your research question for day 3?

General information

Introduction of the Swiss SEM team

- Dr. Frank Pennekamp (main instructor)



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- Dr. Frank Pennekamp (main instructor)
- Dr. James Grace (advanced topics and model clinic)



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- Dr. Rachel Korn (course development)



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- Dr. James Grace (advanced topics and model clinic)
- Dr. Rachel Korn (course development)
- Dr. Noémie Pichon, Dr. Fletcher Halliday, Dr. Eliane Meier, Dr. Hugo Saiz, Dr. Debra Zuppinger-Dingley, Rebecca Oester, Annabelle Constance, Fabienne Wiederkehr (course development)



- *Day 1:*

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 - General introduction to SEM to model ecological systems

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- *Day 3:*
 - Self-study with possibility to meet with instructor(s)

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 - Local estimation of SEMs (with piecewiseSEM)
 - Advanced topics like incorporating random effects, feedbacks, temporal autocorrelation

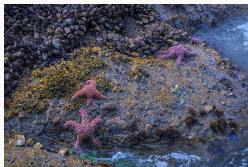
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- Participants are able to apply SEM to their own dataset

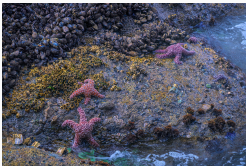
Getting started with Structural Equation Modeling

Research questions



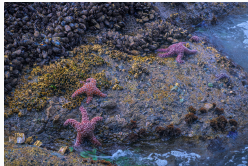
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Research questions



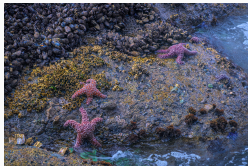
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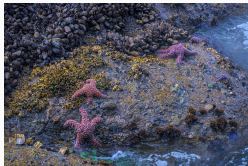
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- We often have ideas how things could be connected in ecological systems.
- To test hypotheses, we need a way to dissect when they occur for a reason versus randomness.
- We use statistics to understand what is signal and what is noise.
- Research questions are often about cause and effect.

Why do we need to use SEM in Ecology

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- Promotes system thinking:
 - “Understanding the whole rather than the parts in isolation”
- SEM unites multiple variables in a single causal network:
simultaneous tests of multiple hypotheses.
- Causality is central:
 - SEM assumes that the specified relationships among variables represent causal links.

Correlation Vs. Causation



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- Everything else being equal, seeing variation in X leading to variation in Y.
- Experiments to isolate effect of X on Y.
- Experiments not always feasible, hence development of SEM.

Differences and similarities between SEM and regression models

- We often have multiple observed variables.

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- Include terms to test for interactions.
- We want to test and evaluate multivariate causal relationship.
- Test direct and indirect effects on assumed causal relationships.
- Incorporate observed and latent variables.

Two goals of SEM:

- 1) Understand the underlying causal network driving the correlation/covariance among a set of variables.

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- 1) Understand the underlying causal network driving the correlation/covariance among a set of variables.
- 2) Explain as much of their variance as possible with the model specified.

A SEM is usually specified based on theory to determine and validate a proposed causal process and/or model.

Which variables to include?

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Which variables to include?

- Supported by theory.
- Ecologically meaningful.
- Garbage in - garbage out (both data quality and ecologically meaningful).

- Continuum between theory (hypothesis-driven) to exploratory (data-driven) modelling.

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- Important to be explicit about the approach taken.

Introduction to the dataset

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- Disturbance

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- Nitrogen deposition
- Changes in precipitation

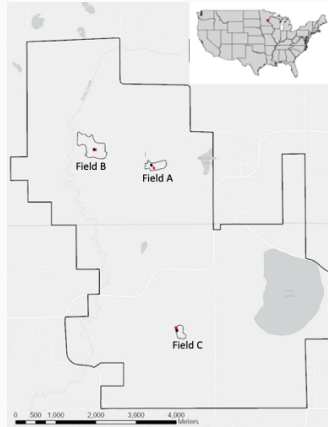
Research questions:

- 1) How has aboveground biomass changed as a function of disturbance (disking) and nutrient addition?

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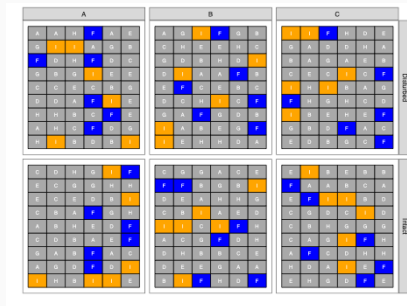
- 1) How has aboveground biomass changed as a function of disturbance (disking) and nutrient addition?
- 2) How are these effects mediated by diversity?

Introduction to the dataset



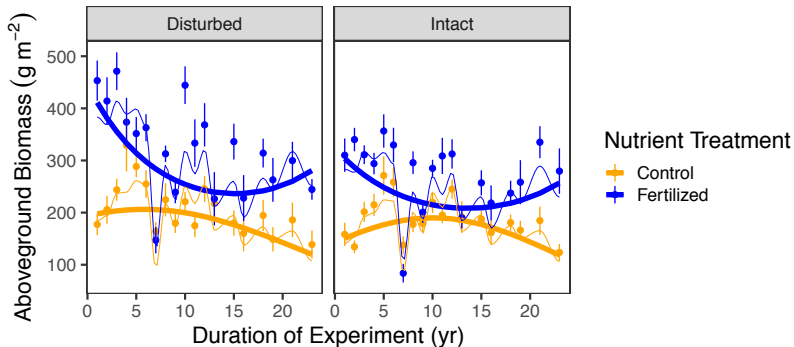
Location of the study site (Cedar Creek Ecosystem Science Reserve), the location of the three study fields within the reserve, and location of the 35 x 55 m intact (black) and disturbed (red) plots within each field.

Introduction to the dataset



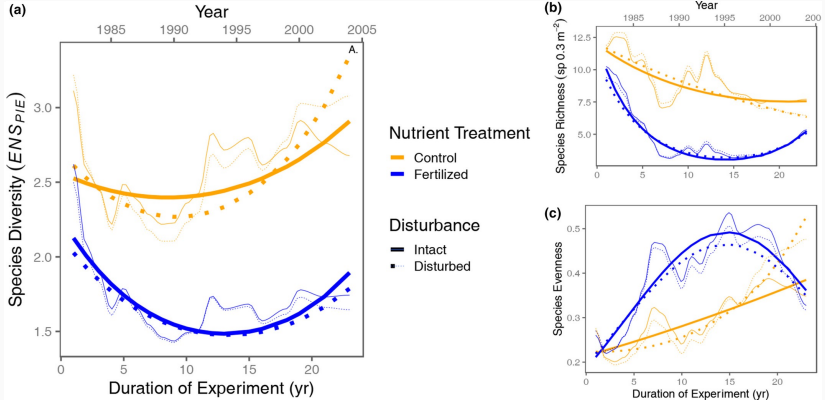
Location of the 4 x 4 m nutrient treatment plots within each 35 x 55 m Intact or Disturbed plot within each of three fields (A, B, and C). Letters indicate the nutrient treatments.

Introduction to the dataset



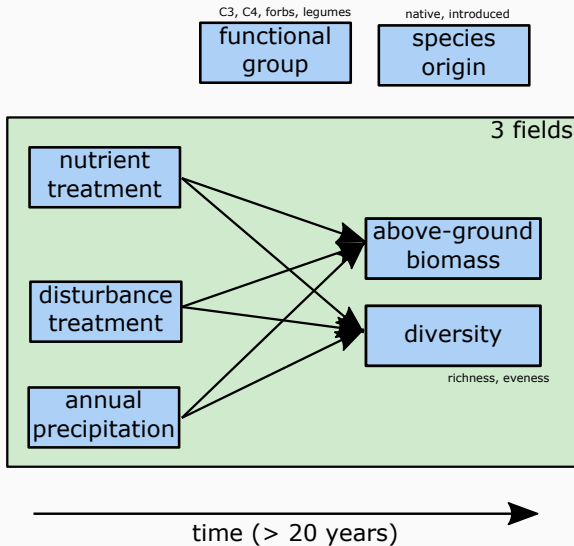
Effect of soil disturbance (disking) and nutrient enrichment on live, aboveground plant biomass. Colors indicate nutrient addition treatment: Control and NPK+ (all nutrients plus $9.5 \text{ g N m}^{-2} \text{ yr}^{-1}$).

Introduction to the dataset

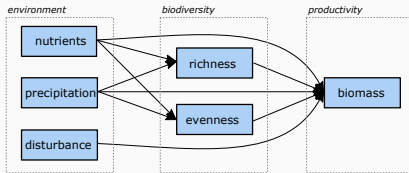


Effect of soil disturbance (disking) and nutrient enrichment on (a) diversity (ENS_{PIE}), (b) richness (S , species $0.3\ m^{-2}$), and (c) evenness ($ENS_{PIE}\ S^{-1}$).

Question of interest: what is the effect of richness on biomass?



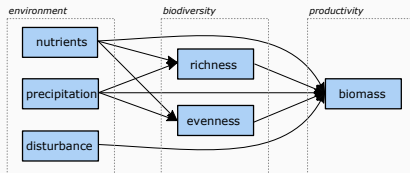
Meta-model



Meta-model are conceptual models that allow to link data with theory.

- 1) Productivity (biomass) is directly influenced by the environment (nutrients, disturbance and precipitation)

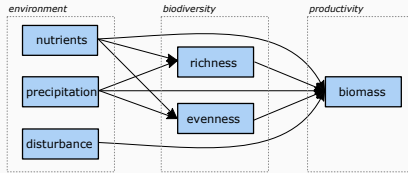
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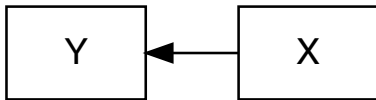
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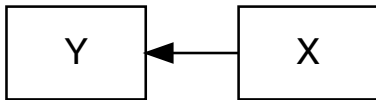
- 1) Productivity (biomass) is directly influenced by the environment (nutrients, disturbance and precipitation)
- 2) Productivity (biomass) is directly influenced by biodiversity (richness and evenness).
- 3) The environment also influences biodiversity and thus, have an indirect effect on productivity via biodiversity.

Graphical models:



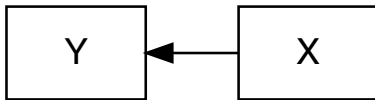
- Directed acyclic graphs (no loops).

Graphical models:



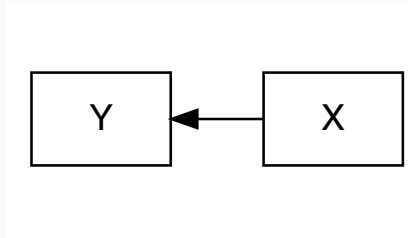
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- Variables are nodes (boxes).
- Edges (one-headed arrows) are causal relationships such as X affects Y.
- Edges (two-headed arrows) are non-causal relationships such as X and Y are correlated.

Exercise:

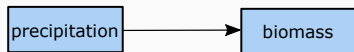
Exercise:

- Draw a meta-model of the dataset you will work with on Thursday.

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- Make a table with putative causal relationships.

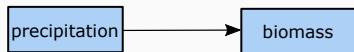
A simple bivariate model.



```
lm(biomass ~ precipitation)
```

- Linear regression

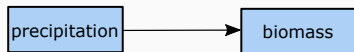
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- Linear regression
- Regression coefficient quantifies the strength of relationship

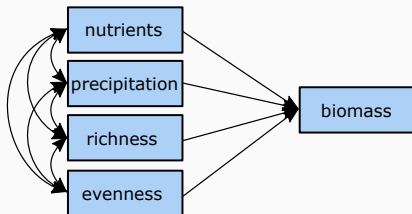
A simple bivariate model.



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- Linear regression
- Regression coefficient quantifies the strength of relationship
- Change in Y for one unit change in X

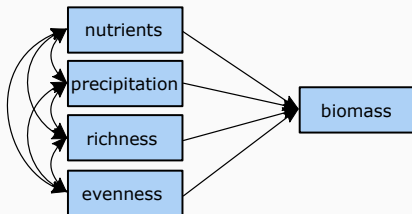
Multiple independent variables



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lm(biomass ~ precipitation + nutrients + ...)
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- Often more than one independent variable important → multiple regression

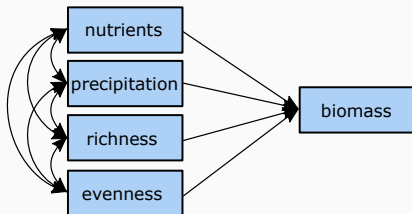
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- Estimates partial regression coefficients (i.e. effect of precipitation on biomass when nutrient addition is fixed)

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- Often more than one independent variable important → multiple regression
- Estimates partial regression coefficients (i.e. effect of precipitation on biomass when nutrient addition is fixed)
- Only direct effects.

From regression models to SEM

SEM: variance-covariance matrix

```
##           [,1]      [,2]      [,3]
## [1,] 1875.3209  429.8712  462.4775
## [2,]  429.8712 1306.9817 -262.8231
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```

- Comparison of specified SEM to observed variance-covariance matrix

SEM: variance-covariance matrix

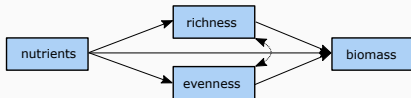
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```

- Comparison of specified SEM to observed variance-covariance matrix
- The variances appear along the diagonal and covariances appear in the off-diagonal elements

```
##           [,1]      [,2]      [,3]
## [1,] 1.0000000 0.2745780 0.3885349
## [2,] 0.2745780 1.0000000 -0.2644881
## [3,] 0.3885349 -0.2644881 1.0000000
```

- Correlation matrix is standardized variance-covariance matrix

Multiple independent variables



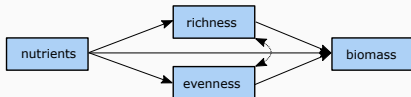
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lm(evenness ~ nutrients)
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lm(biomass ~ richness + evenness)
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- Indirect effect is the effect of an independent variable on a dependent variable through one or more intervening or mediating variables.

Multiple independent variables



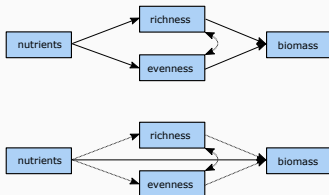
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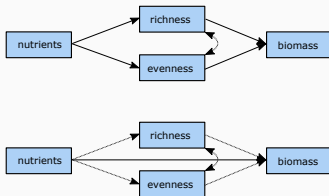
- Indirect effect is the effect of an independent variable on a dependent variable through one or more intervening or mediating variables.
- Indirect effects can be quantified by the product of the compound path.

Mediation



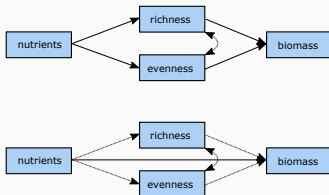
- Tests whether a particular variable has a mediating effect on a path.

Mediation



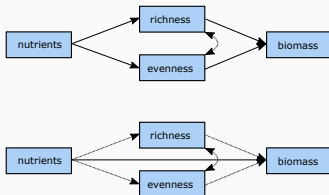
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Mediation



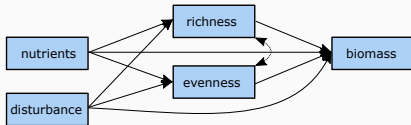
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Mediation



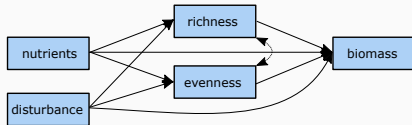
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- Possibilities: complete mediation, partial mediation, no mediation.

System level approach



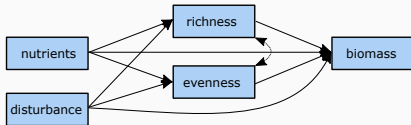
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System level approach



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System level approach



- **Exogenous** variables only have paths emanating from them (i.e., do not have arrows going into them).
- **Endogenous** variables have paths directed into them.
- An endogenous variable can also have arrows directing out of it, but the sole condition is that they must be predicted.

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Grace's 8 rules of path coefficients

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- 4) When variables are connected by more than one pathway, each pathway is the 'partial' regression coefficient.

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- 6) Unanalyzed (residual) correlations among two endogenous variables are their partial correlations.
- 7) The total effect one variable has on another is the sum of its direct and indirect effects.
- 8) The total effect (including undirected paths) is equivalent to the total correlation.

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- *Root-mean squared error of approximation (RMSEA)*: statistic penalizes models based on sample size. A value < 0.10 is acceptable, and anything < 0.08 is good.
- *Standardized root-mean squared residual (SRMR)*: the standardized difference between the observed and predicted correlations. A value < 0.08 is considered good.

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- Global estimation based on variance-covariance matrix¹

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“t-rule” to quickly gauge whether a model is under-, just, or overidentified:

$$t \leq \frac{n(n+1)}{2}$$

t = number of unknowns (parameters to be estimated, i.e. variances & covariances)

n = number of knowns (observed variables).

The LHS is how many pieces of information we want to know.

RHS: information we have (number of unique cells in the observed variance-covariance matrix).

- Replication should be at least 5x the number of estimated coefficients (not error variances or other correlations).

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- Ideally, replication is 5-20x the number of estimated parameters.
- The larger the sample size, the more precise (unbiased) the estimates.

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- 10) Interpret and present results visually

Lavaan syntax

Define model:

```
simple <-  
"mass.above ~ nadd + rich + even + precip.mm + disk  
rich ~ nadd + precip.mm  
even ~ nadd + precip.mm"
```

Fit model:

```
fit.simple <- sem(simple, data = seabloom)
```

Formula type	R	Meaning	Example
regression	~	is regressed on	$y \sim x$
correlation	~~	correlate errors for	$y1 \sim\sim y2$
latent	=~	set reflective indicators	$\text{Height} =\sim y1 + y2 + y3$
composite	<~	set formative indicators	$\text{Comp1} <\sim 1*x1 + x2 + x3$
intercept	~ 1	estimate mean for y	$y \sim 1$
labelling	*	name coefficients	$y \sim b1*x1 + b2*x2$
defining	:=	define quantity	$\text{Total} := b1*b3 + b2$

Questions?

Live coding session

Your turn: working with the
Seabloom dataset

Exercise 1

- Exploration of dataset (variables and treatments)

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 - Multiple regression (direct effects of predictors on AGB)

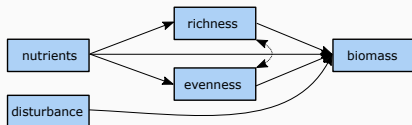
Exercise 1

- Exploration of dataset (variables and treatments)
- Check collinearity and normality
- Fitting linear models to estimate coefficients
 - Multiple regression (direct effects of predictors on AGB)
 - Multiple regression (indirect effects on richness and evenness)

Exercise 1

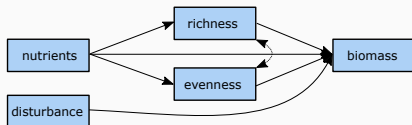
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 - What can you conclude?

Exercise 2



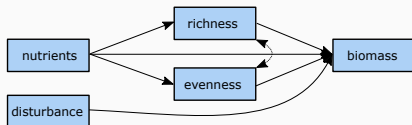
- Fitting of above SEM to Seabloom data:

Exercise 2



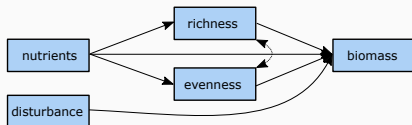
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 - Assess model goodness of fit.

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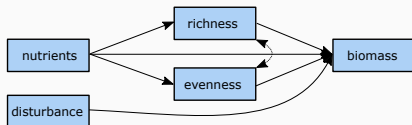
- Fitting of above SEM to Seabloom data:
 - Assess model goodness of fit.
 - Investigate the modification indices. Are there paths to add that are reasonable?

Exercise 2



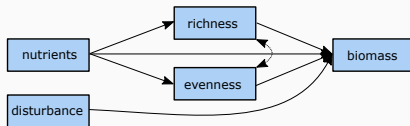
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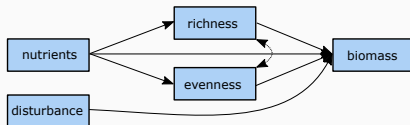
Exercise 3



After finding a model with good fit:

- Model analysis:

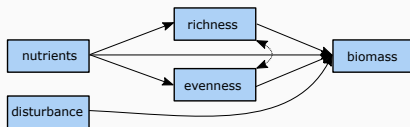
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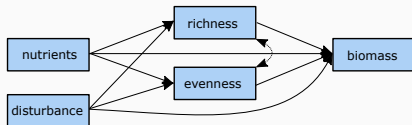
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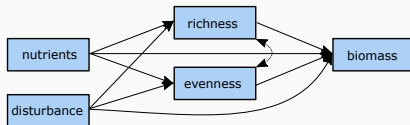
- Model analysis:
 - Calculate the standardized coefficients.
 - Add derived quantities (direct and indirect effects of nutrients and disturbance).

Exercise 4



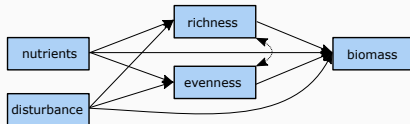
- Saturated model:

Exercise 4



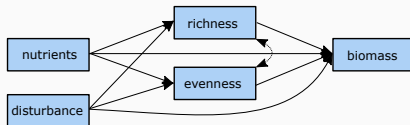
- Saturated model:
 - Model comparison with simpler models used previously.

Exercise 4



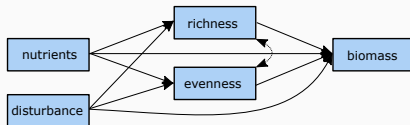
- Saturated model:
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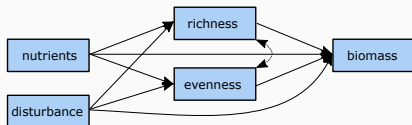
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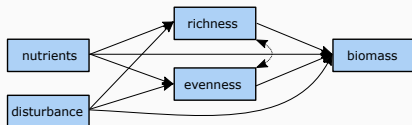
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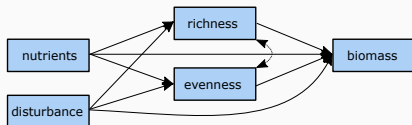
- Perform mediation analysis:

Exercise 5



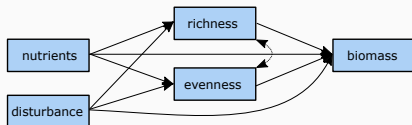
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- Perform mediation analysis:
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 - Add paths from disturbance to richness and evenness, remove the direct paths to AGB (both nutrients and disturbance).

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- Perform mediation analysis:
 - Is the effect of disturbance mediated via its effect on richness and evenness, rather than directly on biomass?
 - Add paths from disturbance to richness and evenness, remove the direct paths to AGB (both nutrients and disturbance).
 - Compare model fit to simple model. What do you conclude?