

Evaluation de la dynamique des reseaux trophiques avec RCaN

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The historical team behind RCaN

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Sam Subbey



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Benjamin Planque



Ulf Lindstrøm



Elliot Sivel



Food-web assessment using RCaN

- I. Chance and necessity
- II. Making sense of different sources of data
- III. Principles of CaN modelling
- IV. A one species example
- V. A food-web example
- VI. RCaN and RCaN Constructor

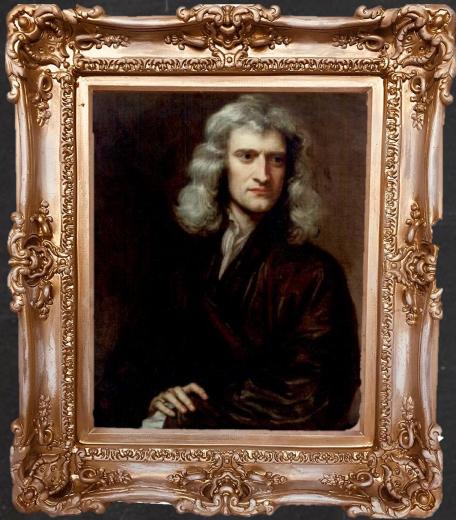


I

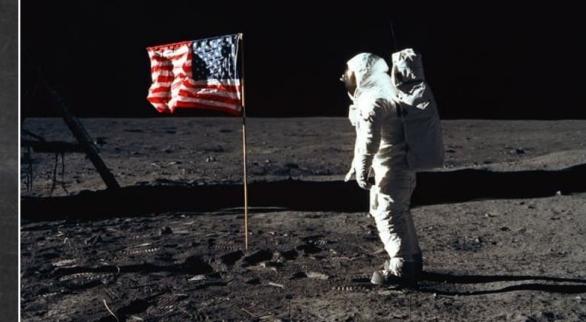
Chance and necessity



Rocket science is easy for modellers



Newton

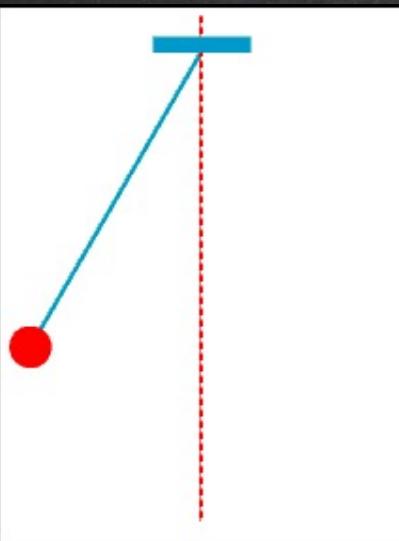


Apollo 11 - 1969



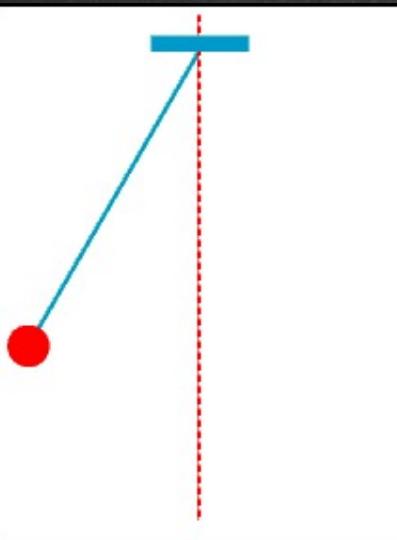
Simple physics is generally ...simple

pendulum

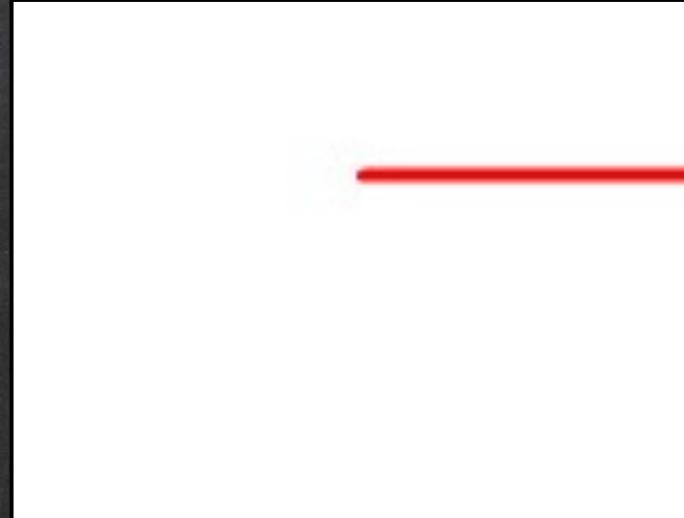


Simple physics can sometimes be complex

pendulum



Double pendulum



Deterministic chaos
=
We know the equations
but
we cannot predict

A Knight in fuzzy armour



Frank Knight (1885-1972)

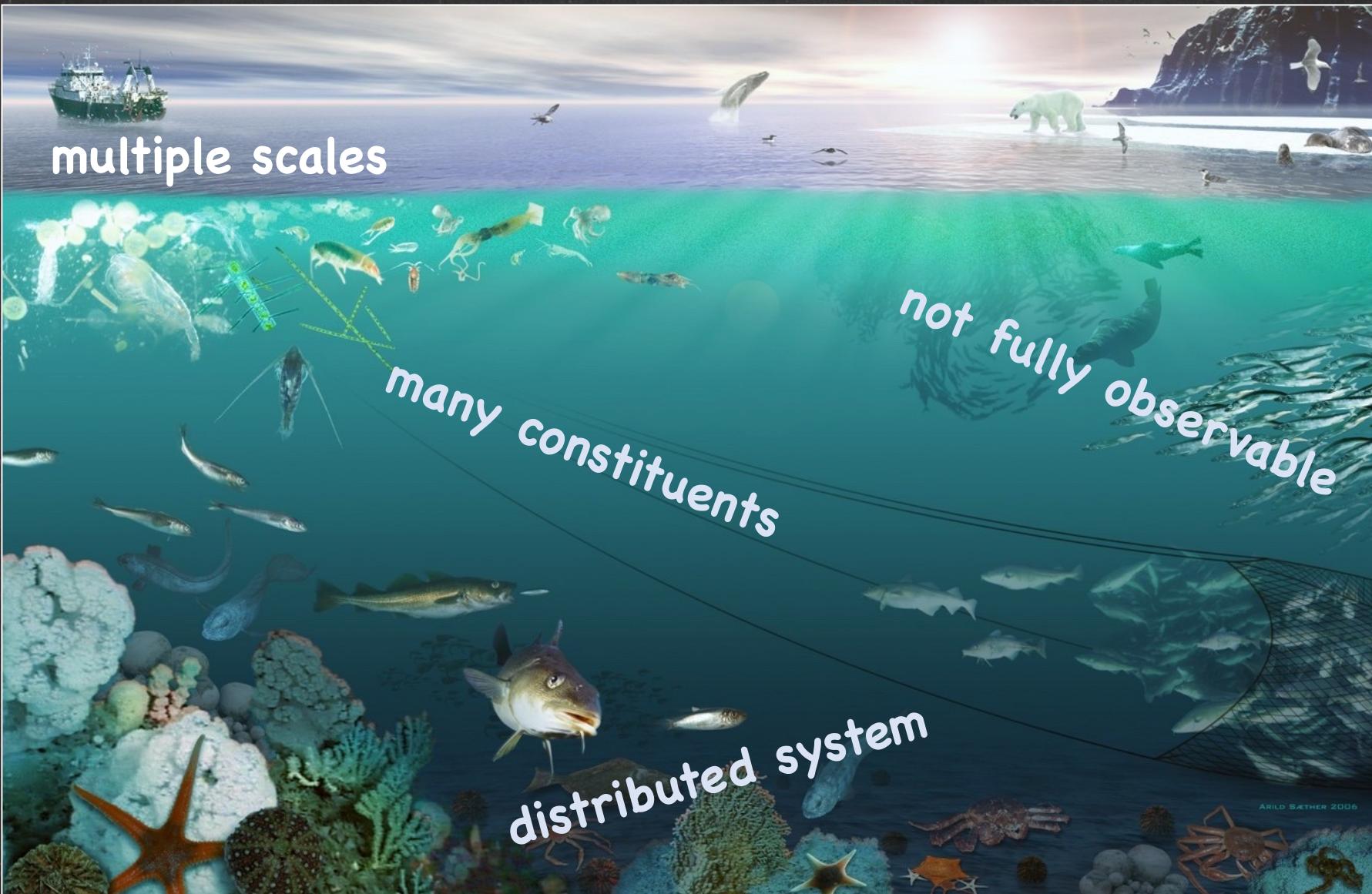
University of Chicago Centennial Exhibition Catalogues

ur
Knightian uncertainty
=
Unquantifiable risk

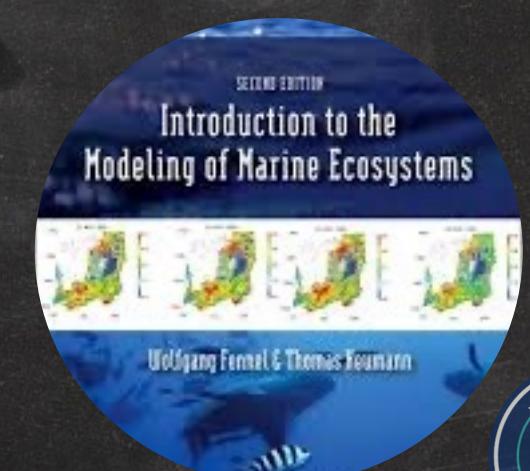
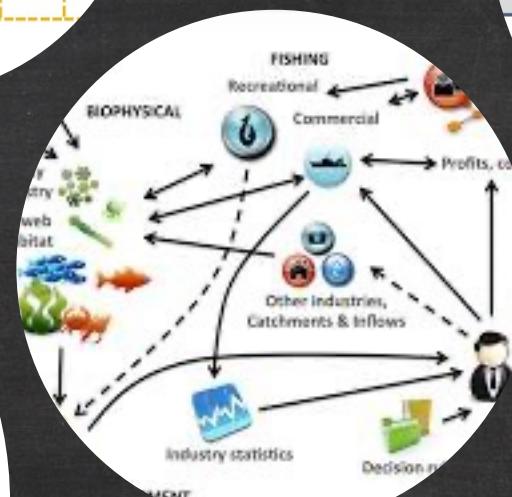
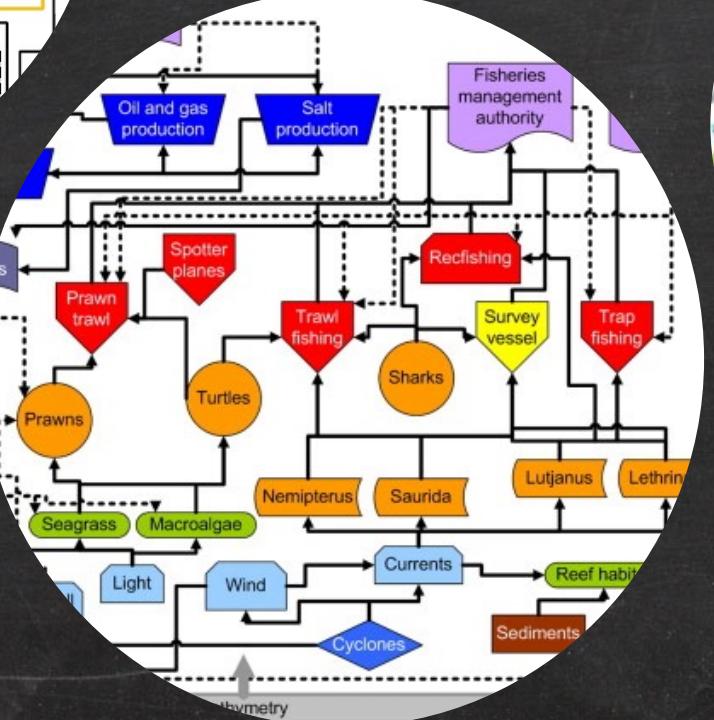
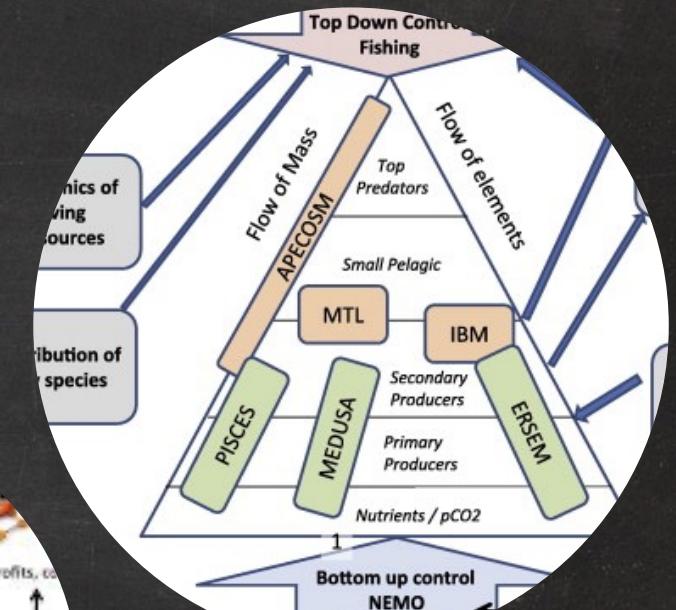
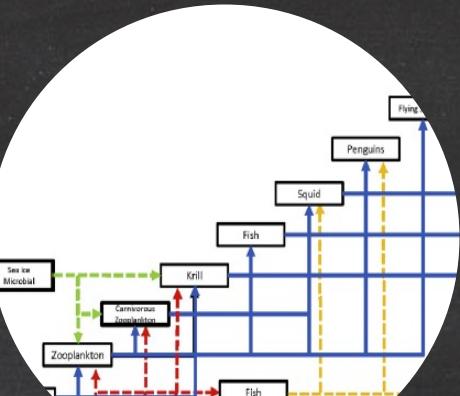
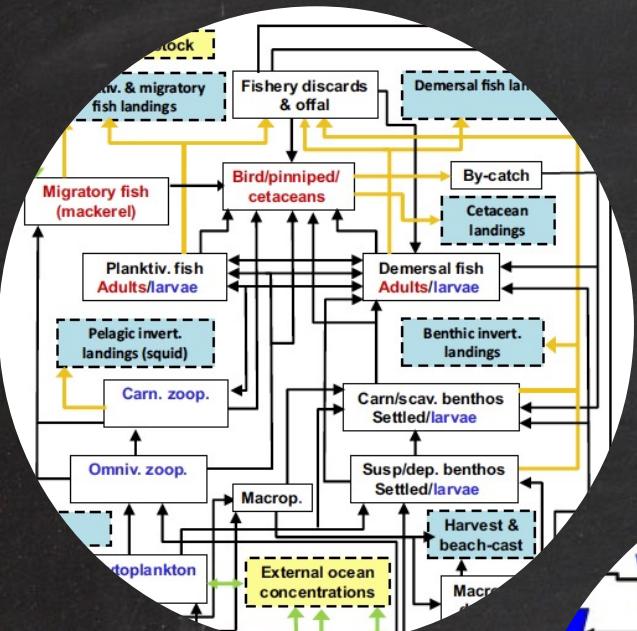
Orrin H. Pilkey & Linda Pilkey-Jarvis



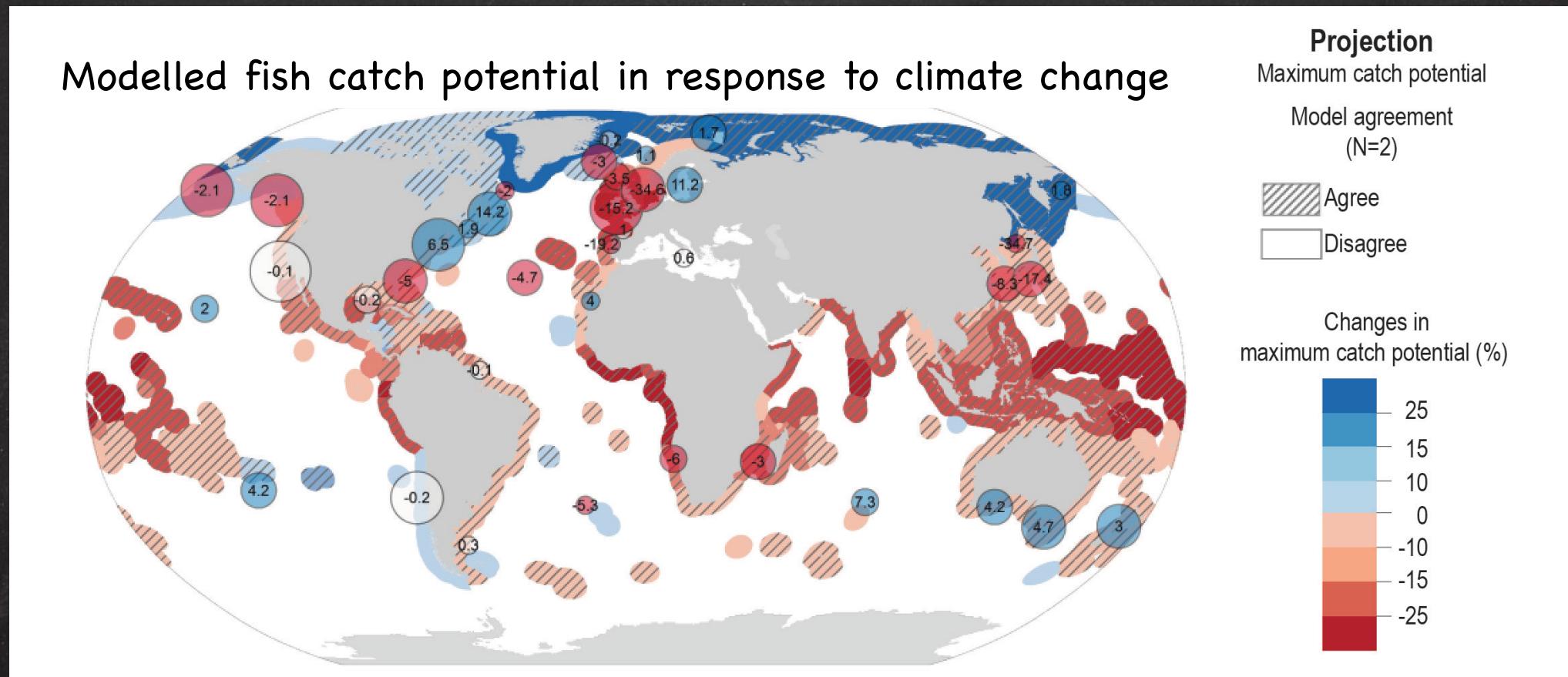
Ecosystems are complicated and complex



Modellers can model ecosystems



Modellers can model ecosystems



IPCC, 2019: Summary for Policymakers. In: IPCC Special Report on the Ocean and Cryosphere in a Changing Climate



Why should one trust ecosystem models?

Ecosystems are too complex to be modelled

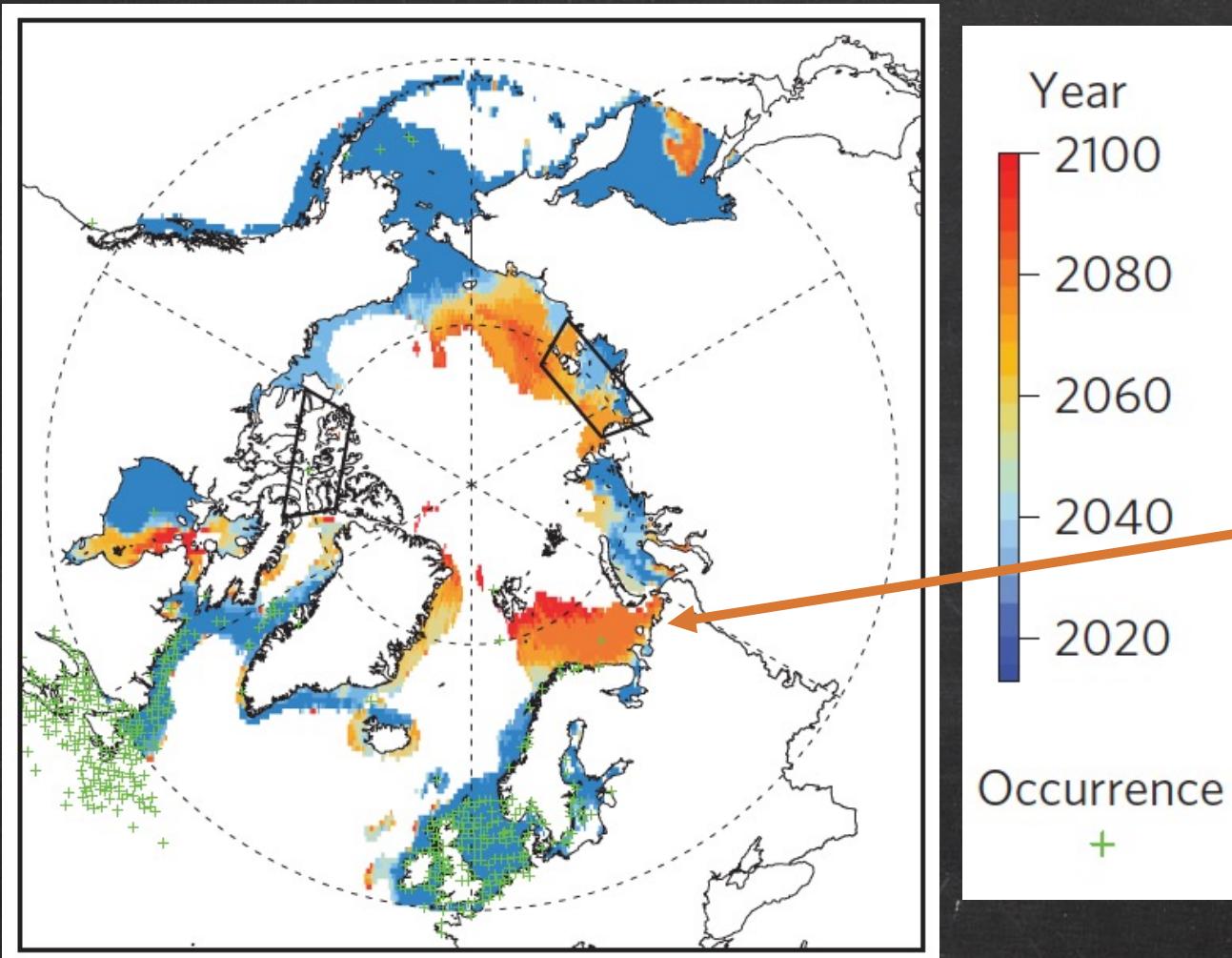
No theory

Many assumptions

Different models give different results

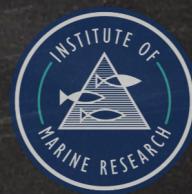


Why should one trust ecosystem models?



predicted year in which the conditions become suitable for Atlantic cod
(*Gadus morhua*). Wisz et al. 2015

Cod will enter the
Barents Sea by 2070-
2100



Reconciling models and practitioners

There's enough we know to tell something about the natural system



Optimist

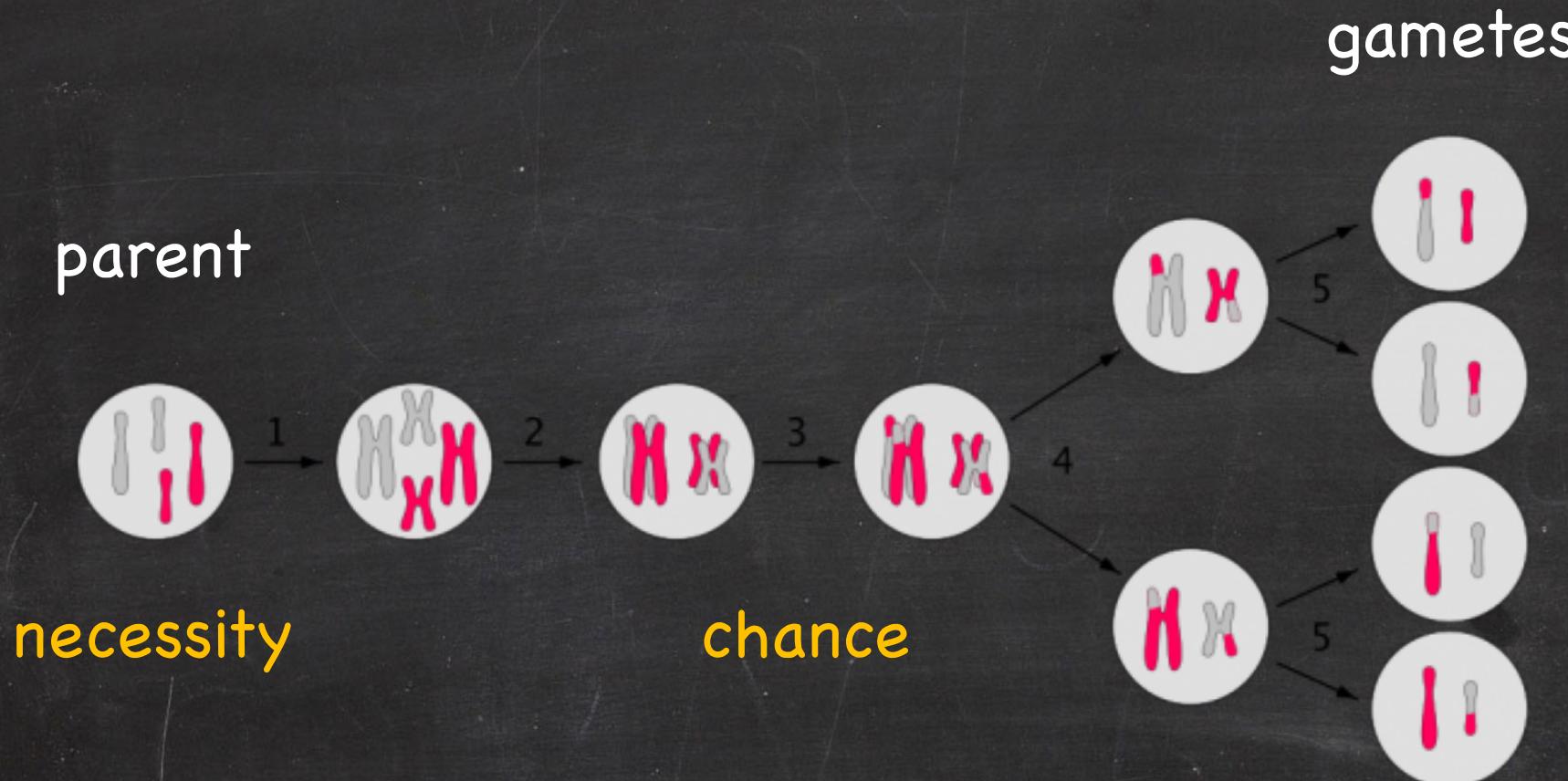


Pessimist

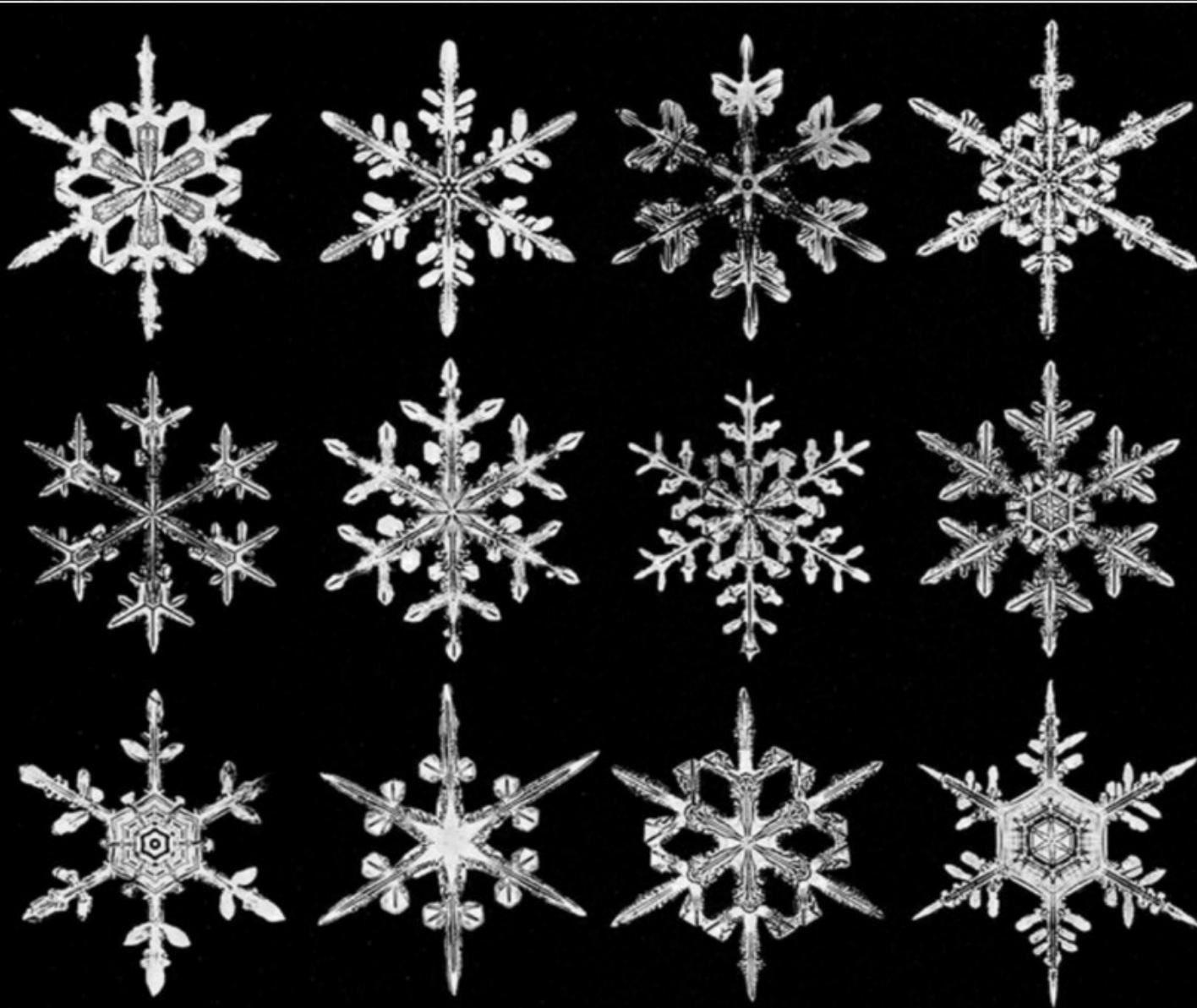
There's not enough we know to tell something about the system

Chance and necessity modelling (CaN):
Model what we know and leave the rest to chance

Chance and Necessity: sexual reproduction



Chance and Necessity: snow flakes

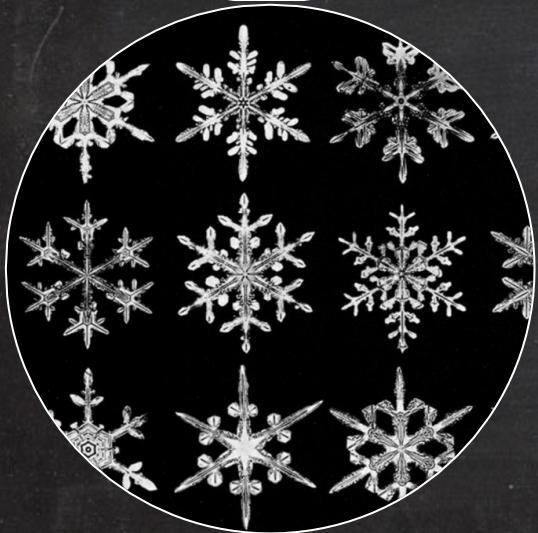
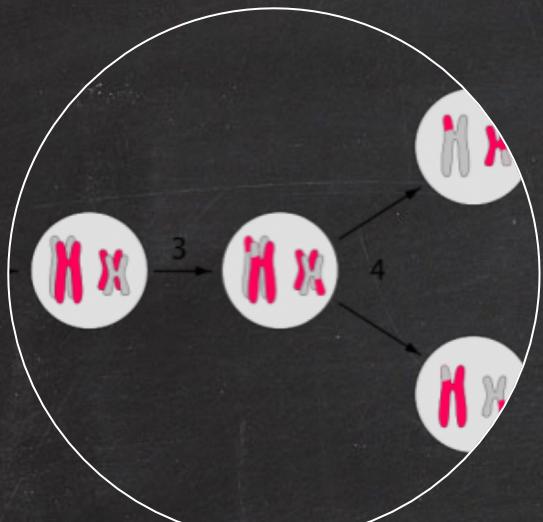


Wilson Bentley 1865-1931

Necessity: humidity & cold
Chance...gives the shape



There is no 'most likely' outcome



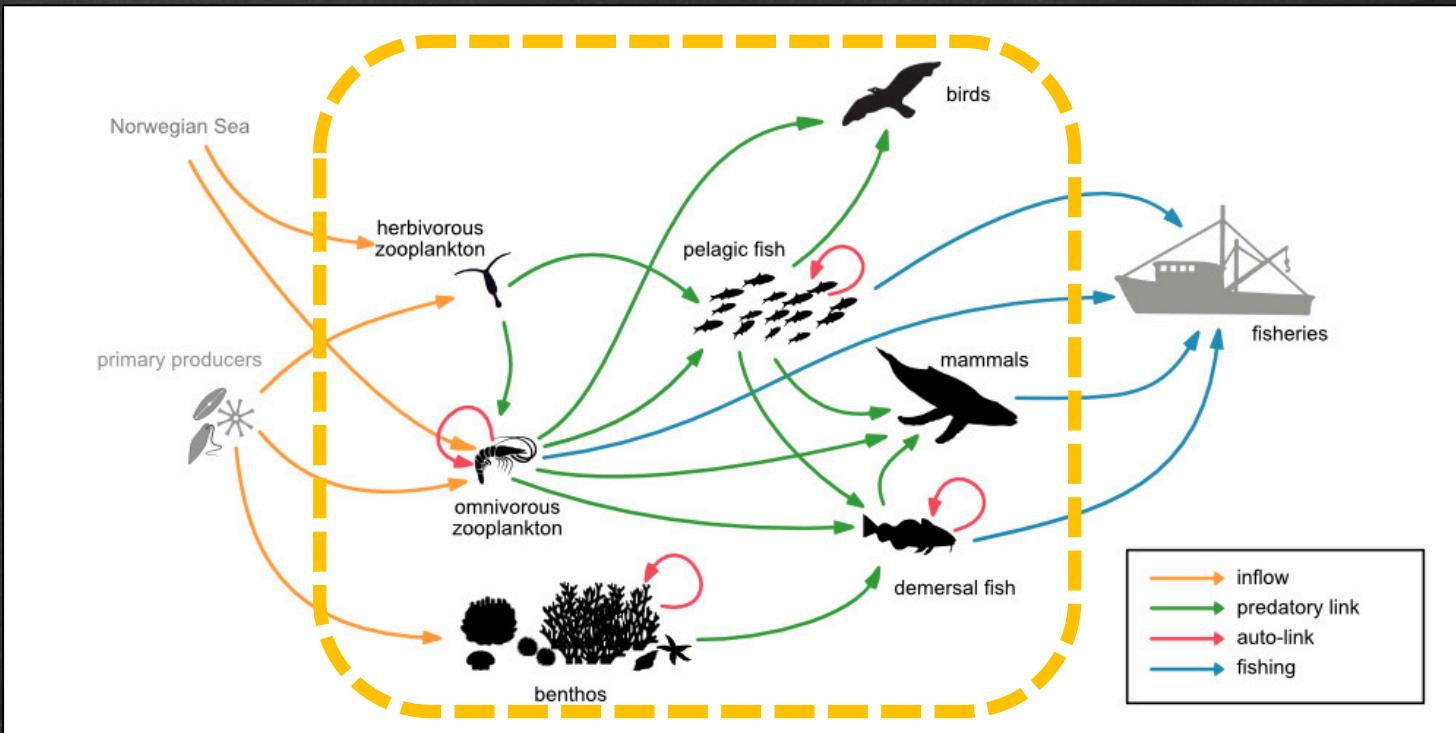
Contribution to the Themed Section: 'Science in support of a nonlinear non-equilibrium world'

Food for Thought

Modelling chance and necessity in natural systems

Benjamin Planque  ^{1*} and Christian Mullon  ²

Explore the diversity of the
Barents Sea food web dynamics



II

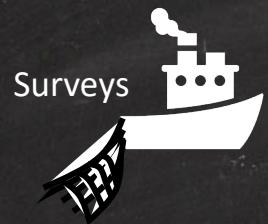
Making sense of different sources of data



Information

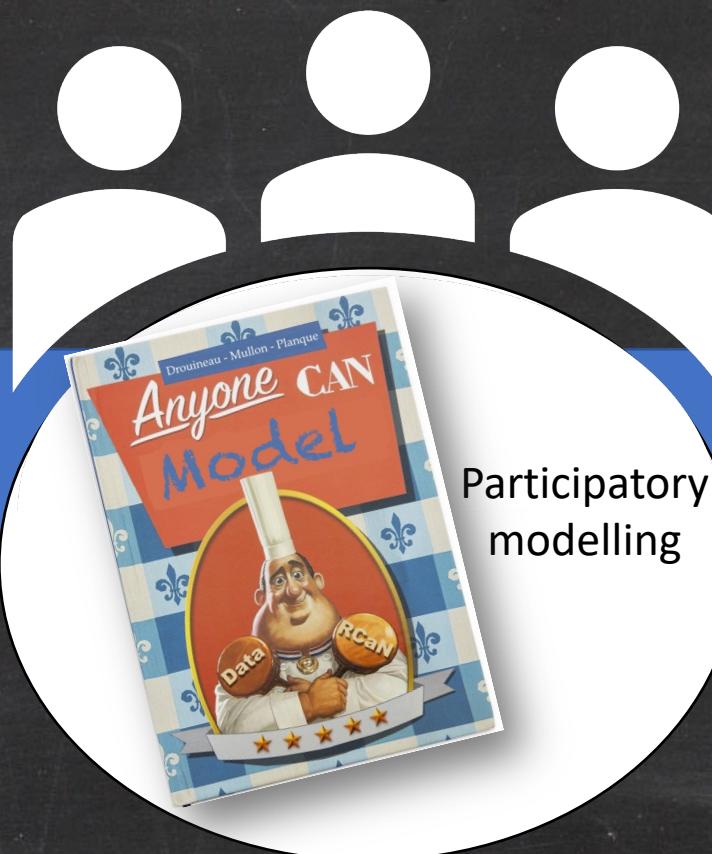
CaN

IEA



hypotheses
uncertainties
& unknowns

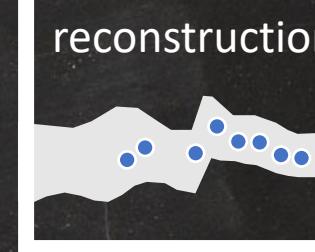
Chance and Necessity
modelling



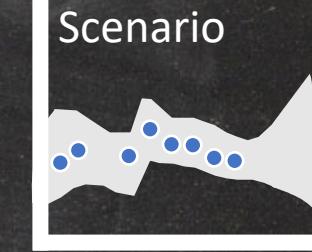
Participatory
modelling

Are the assumptions,
information and data consistent?

Historical
reconstructions



Short term forecasts
Scenario

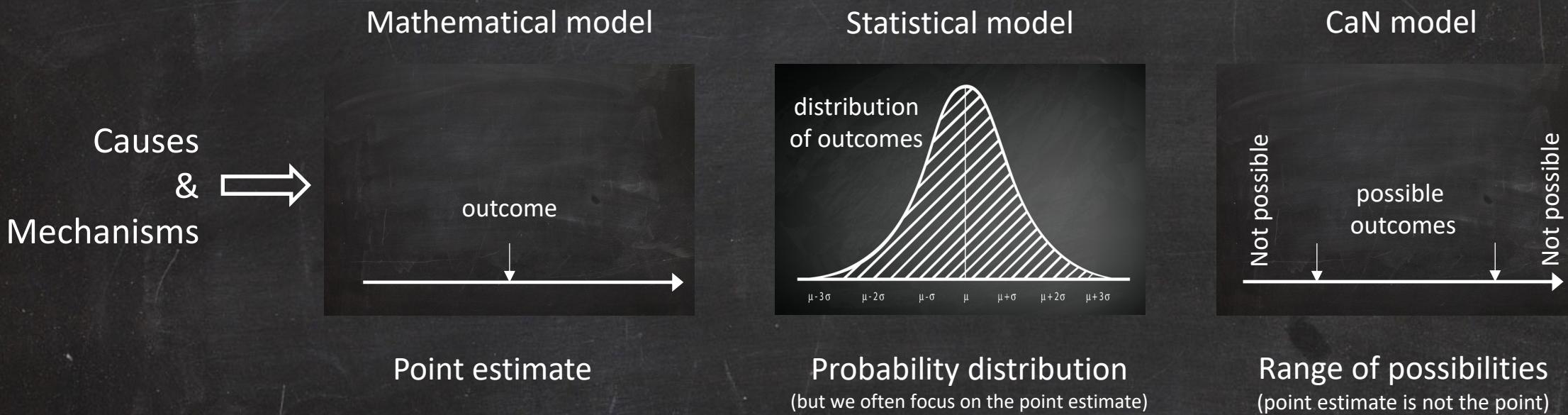


III

Principles of CaN modelling



CaN modelling is about sampling “possible” outcomes



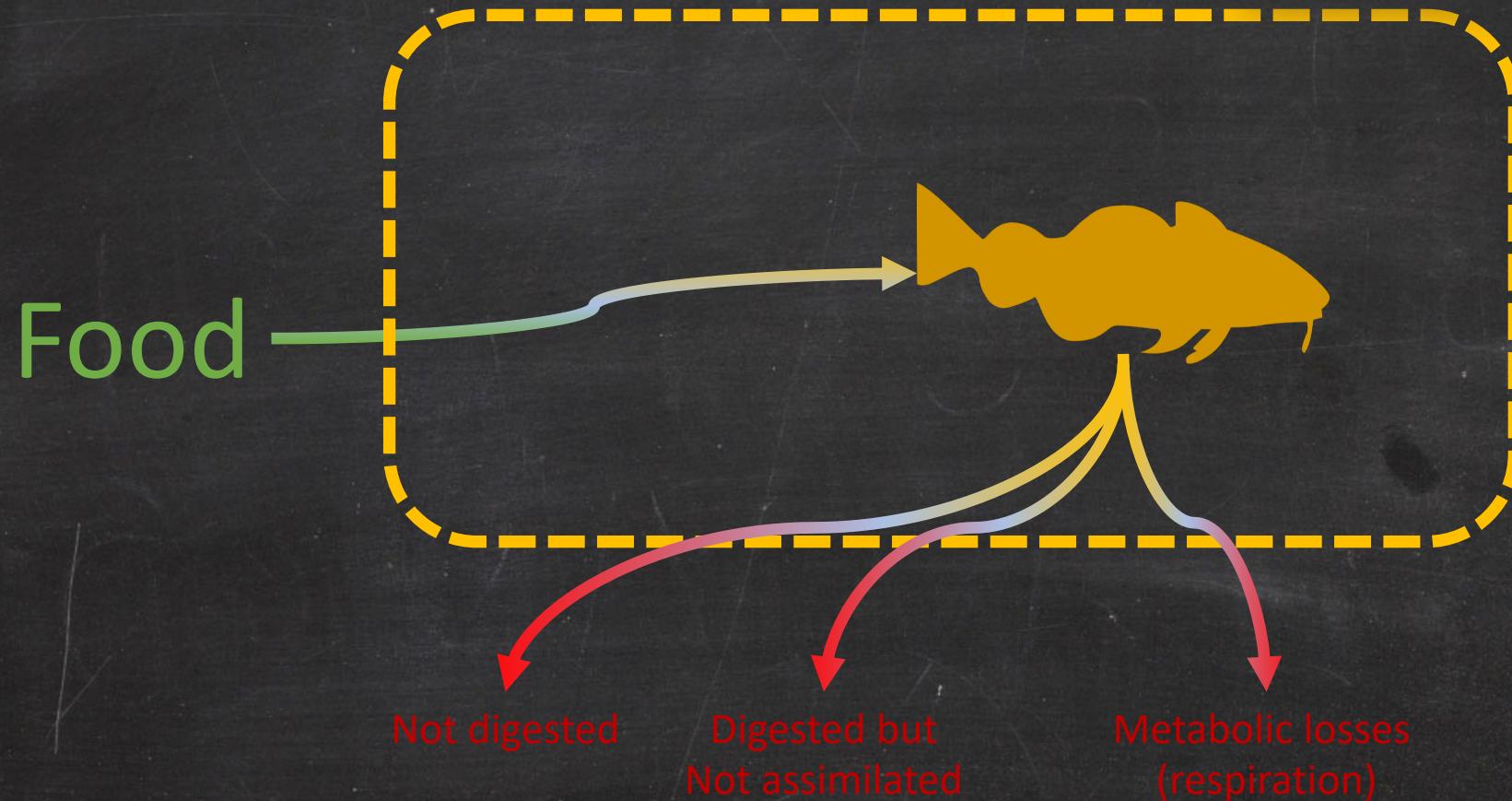
IV

A one species example

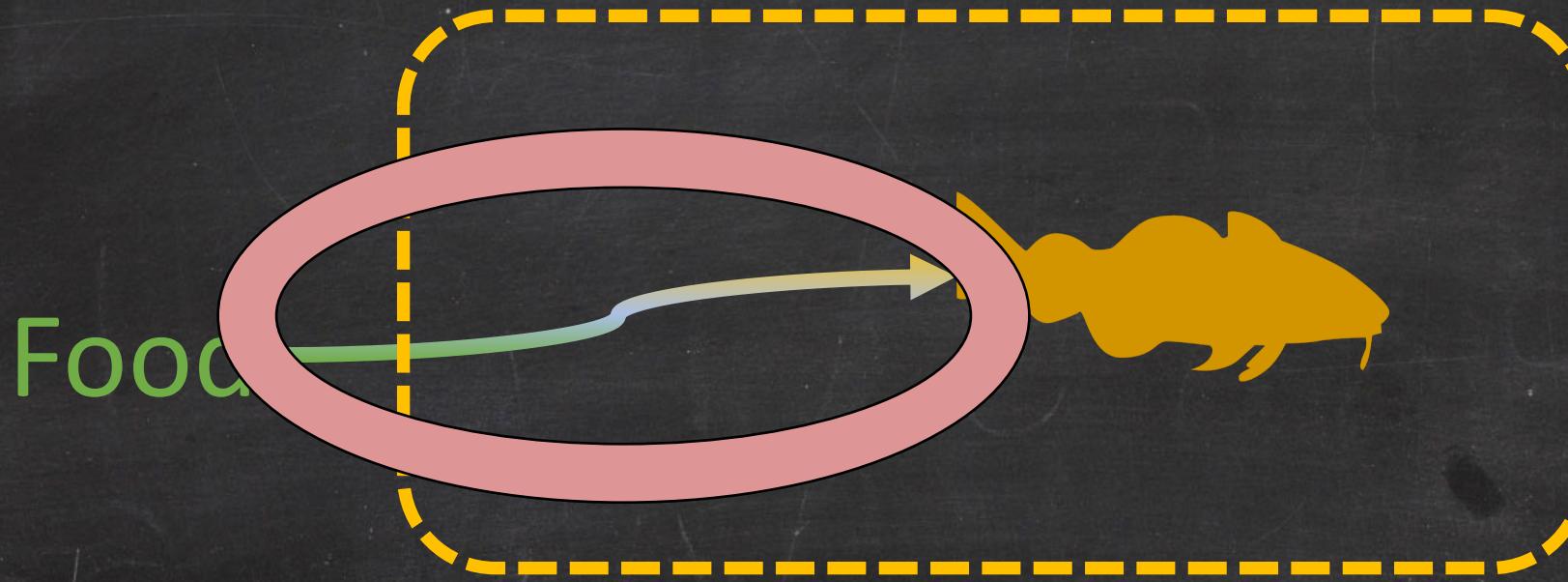


Part of the ingested food is ‘lost’ and the rest contributes to population growth

$$B_{t+1} = (1 - H) B_t + \sum_f N_f F_f$$

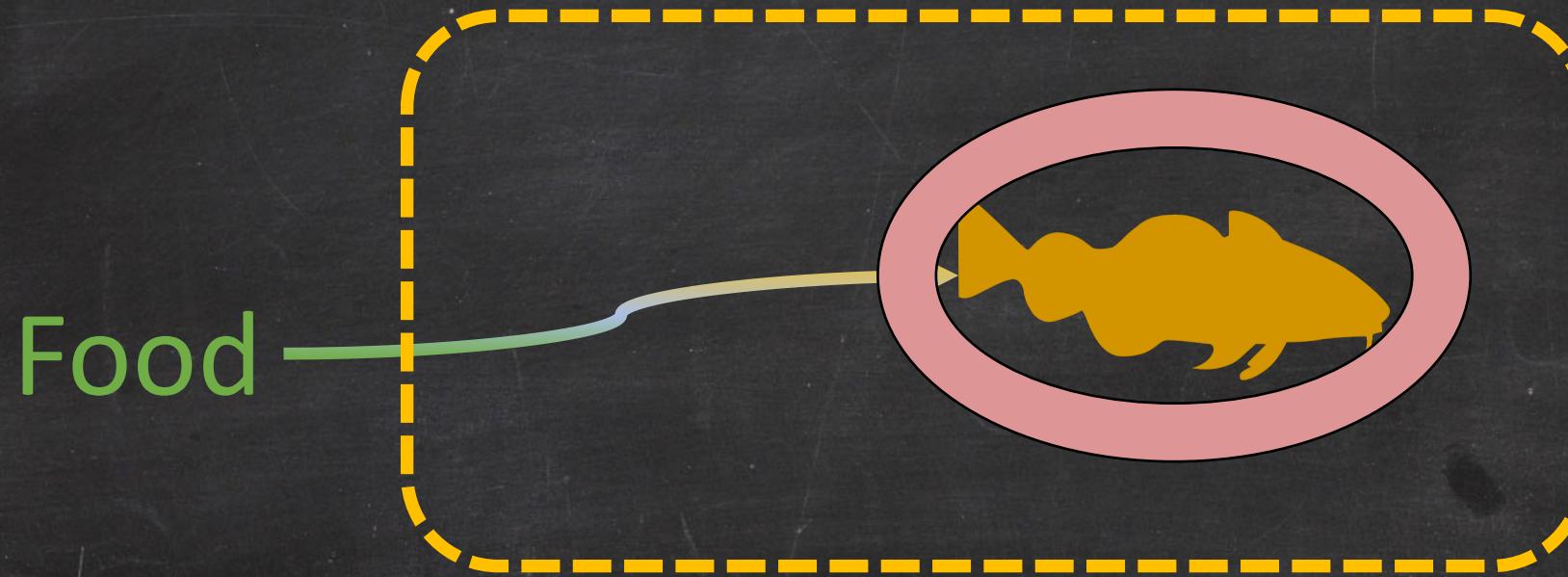


The quantity ingested is indeterminate (chance) but constrained (necessity)



Flow (food \rightarrow fish) \leq Biomass.of.fish \times satiation
Flow (food \rightarrow fish) \leq food available

The change in biomass from one year to the next is indeterminate (chance) but constrained (necessity)



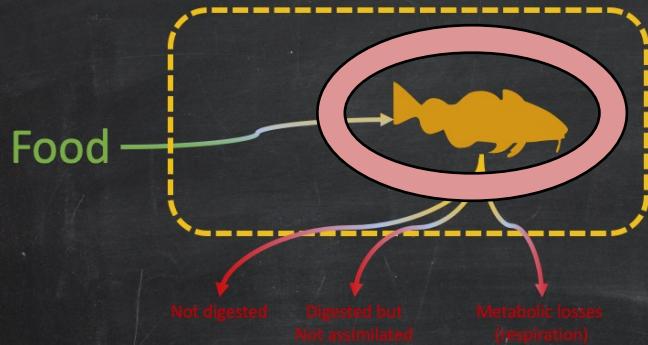
$$\text{Biomass.fish} \times \exp(-\rho) \leq \Delta(\text{Biomass.of.fish}) \leq \text{Biomass.of.fish} \times \exp(+\rho)$$

Sampling model solutions

Flow (food \rightarrow fish) \leq Biomass.of.fish x satiation

Flow (food \rightarrow fish) \leq food available (Carrying Capacity)

Biomass.fish x exp(-rho) \leq Delta(Biomass.of.fish) \leq Biomass.of.fish x exp(+rho)



Model input parameters

Satiation = 5.5

Inertia = 0.25

Metabolic losses = 1.65

Digestibility (food) = 0.65

Assimilation Efficiency = 0.93

Refuge Biomass = 0.01

Constraints

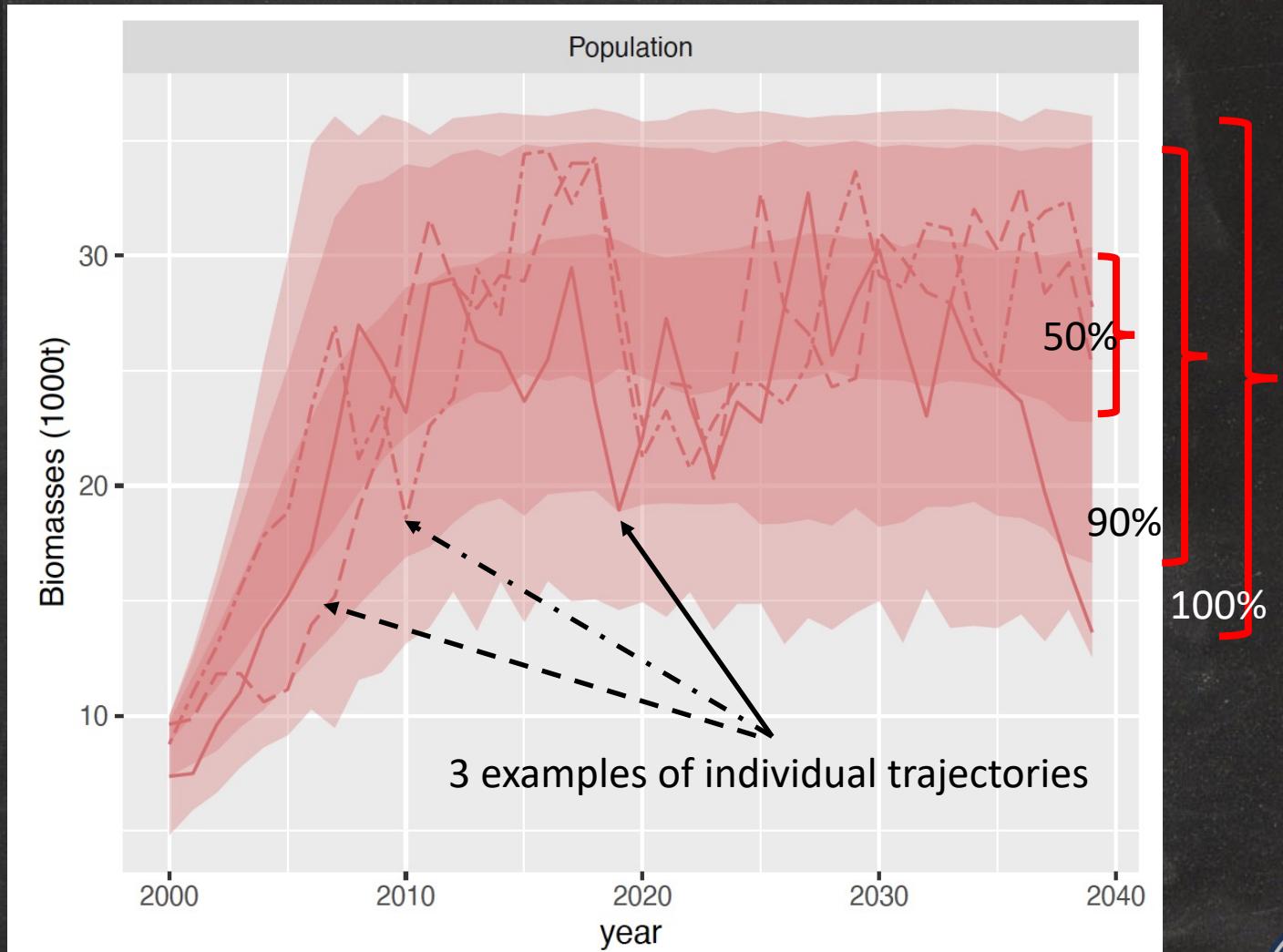
Satiation

Inertia

Max consumption (<100)

Starting value (<10)

Biomass

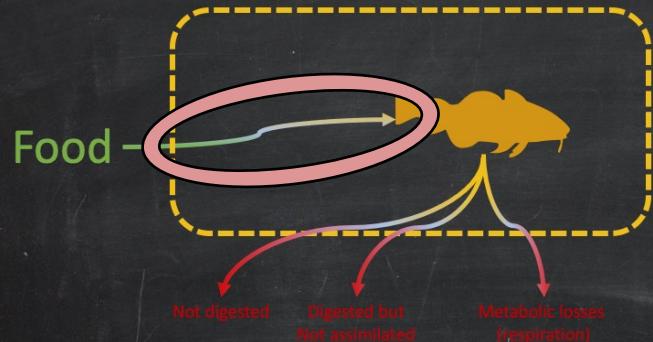


Sampling model solutions

Flow (food \rightarrow fish) \leq Biomass.of.fish x satiation

Flow (food \rightarrow fish) \leq food available (Carrying Capacity)

Biomass.fish x exp(-rho) \leq Delta(Biomass.of.fish) \leq Biomass.of.fish x exp(+rho)



Model input parameters

Satiation = 5.5

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Constraints

Satiation

Inertia

Carrying capacity (<100)

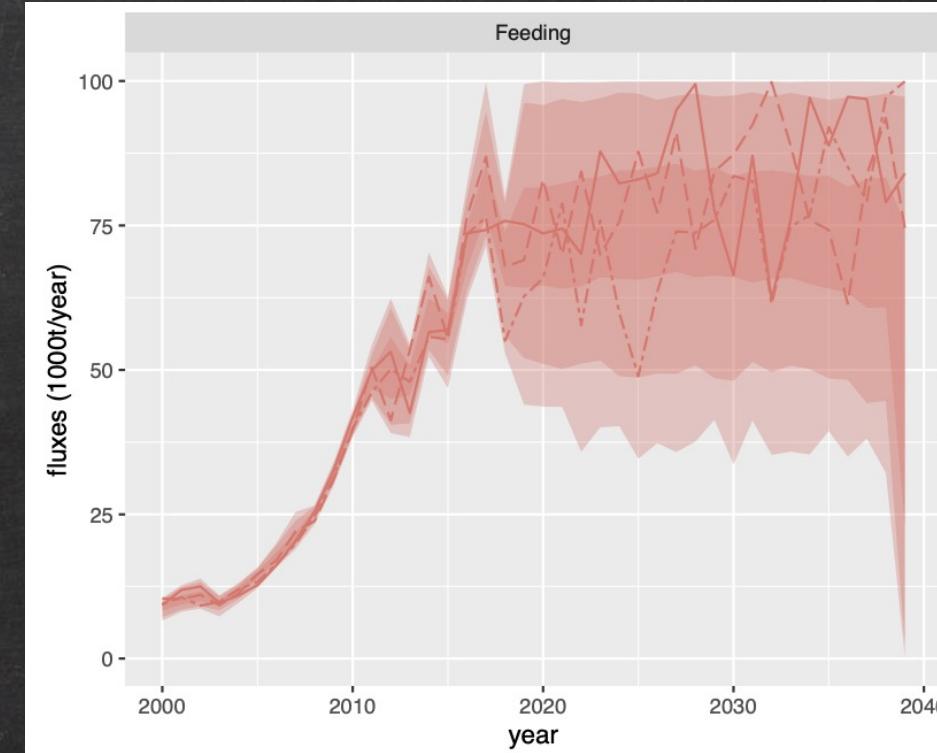
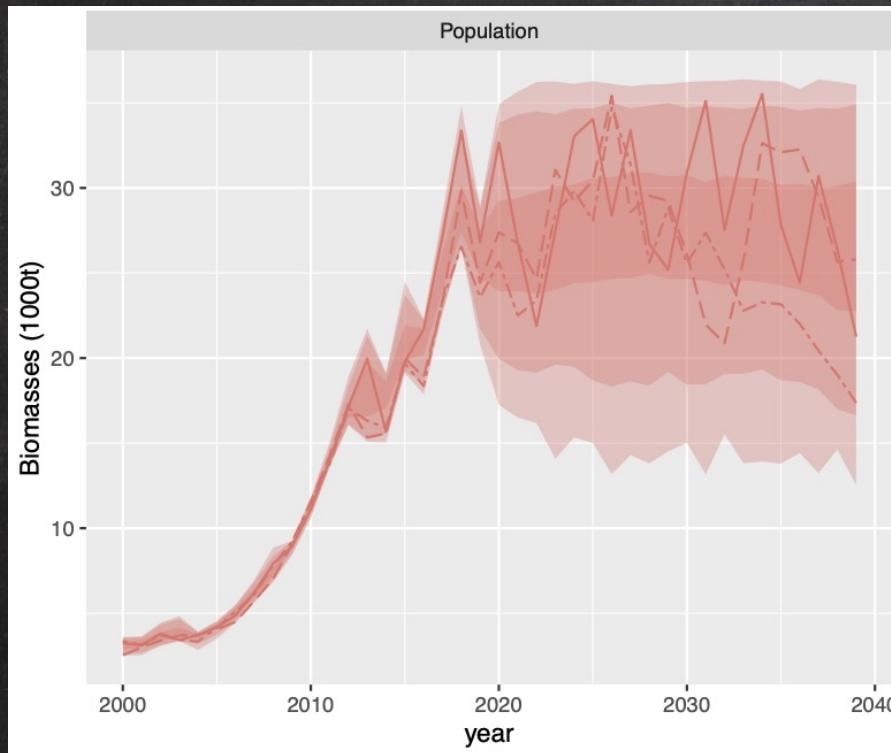
Starting value (<10)

Trophic flow

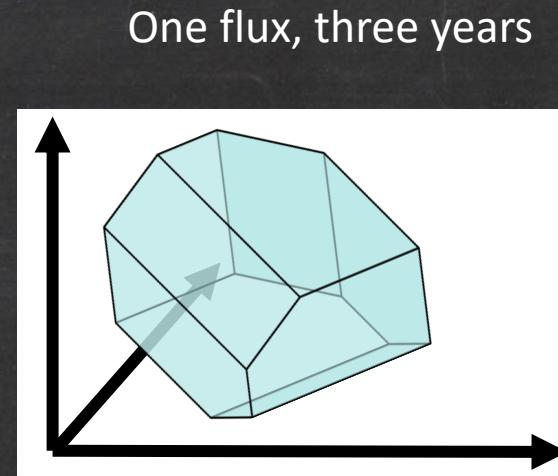
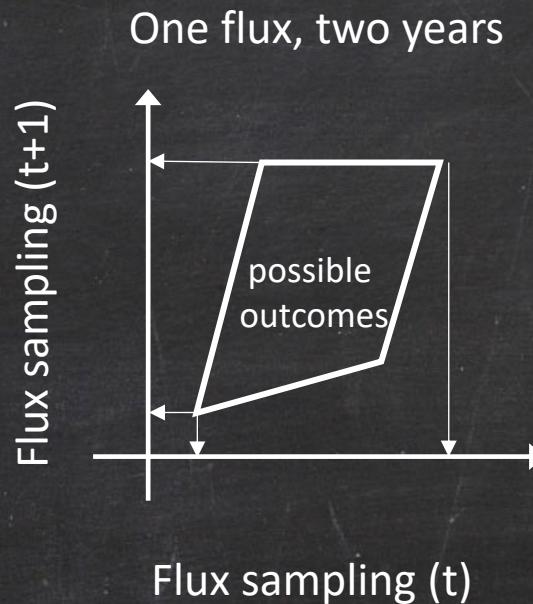
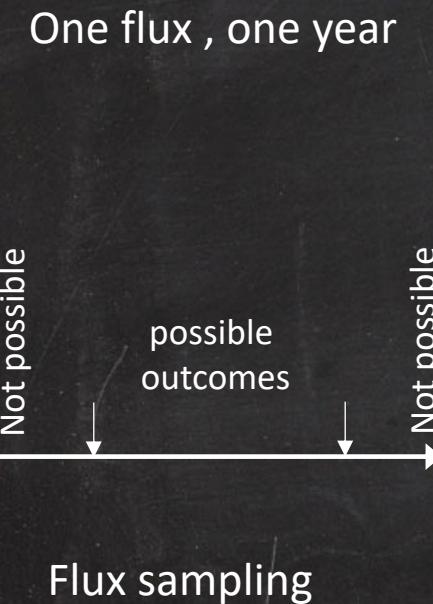


What if we have observations about the population?

- 1. We already had one observation: the starting max value
- 2. We add biomass estimates, with uncertainties, for the period 2000-2019

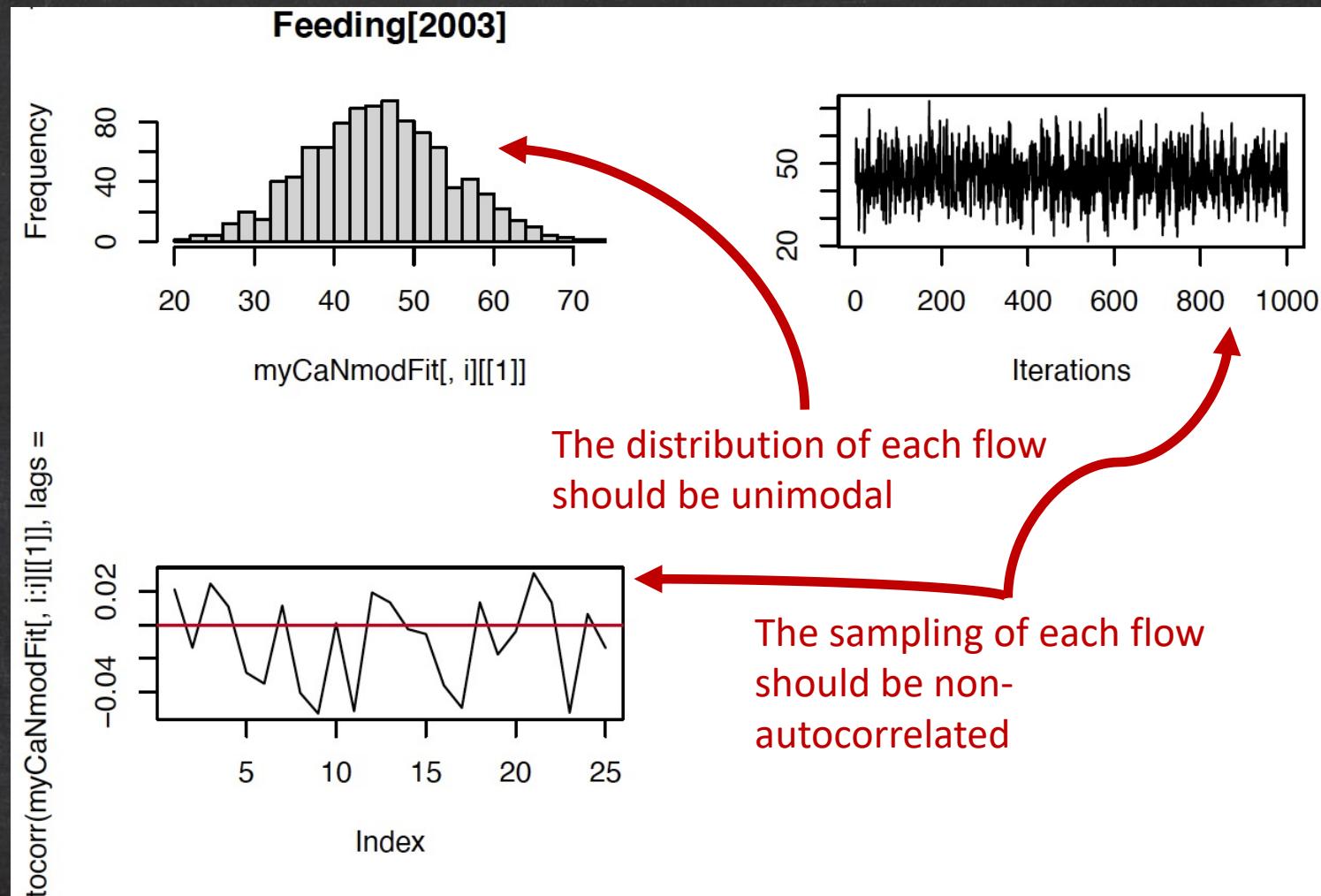
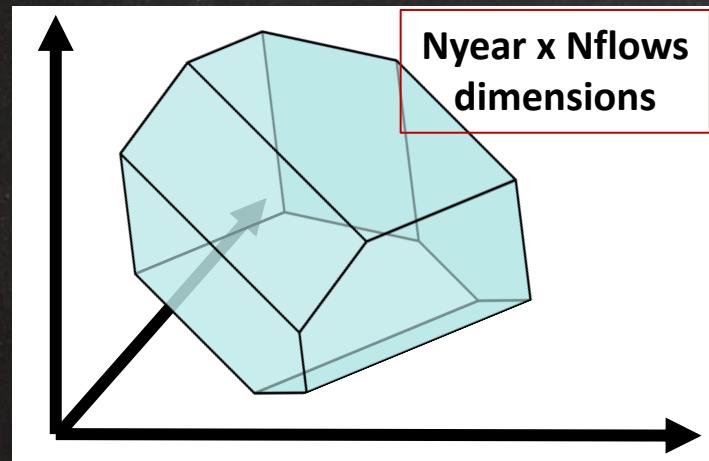


A short digression on polytope sampling



Many dimensions
-
complex polytope

A short digression on polytope sampling

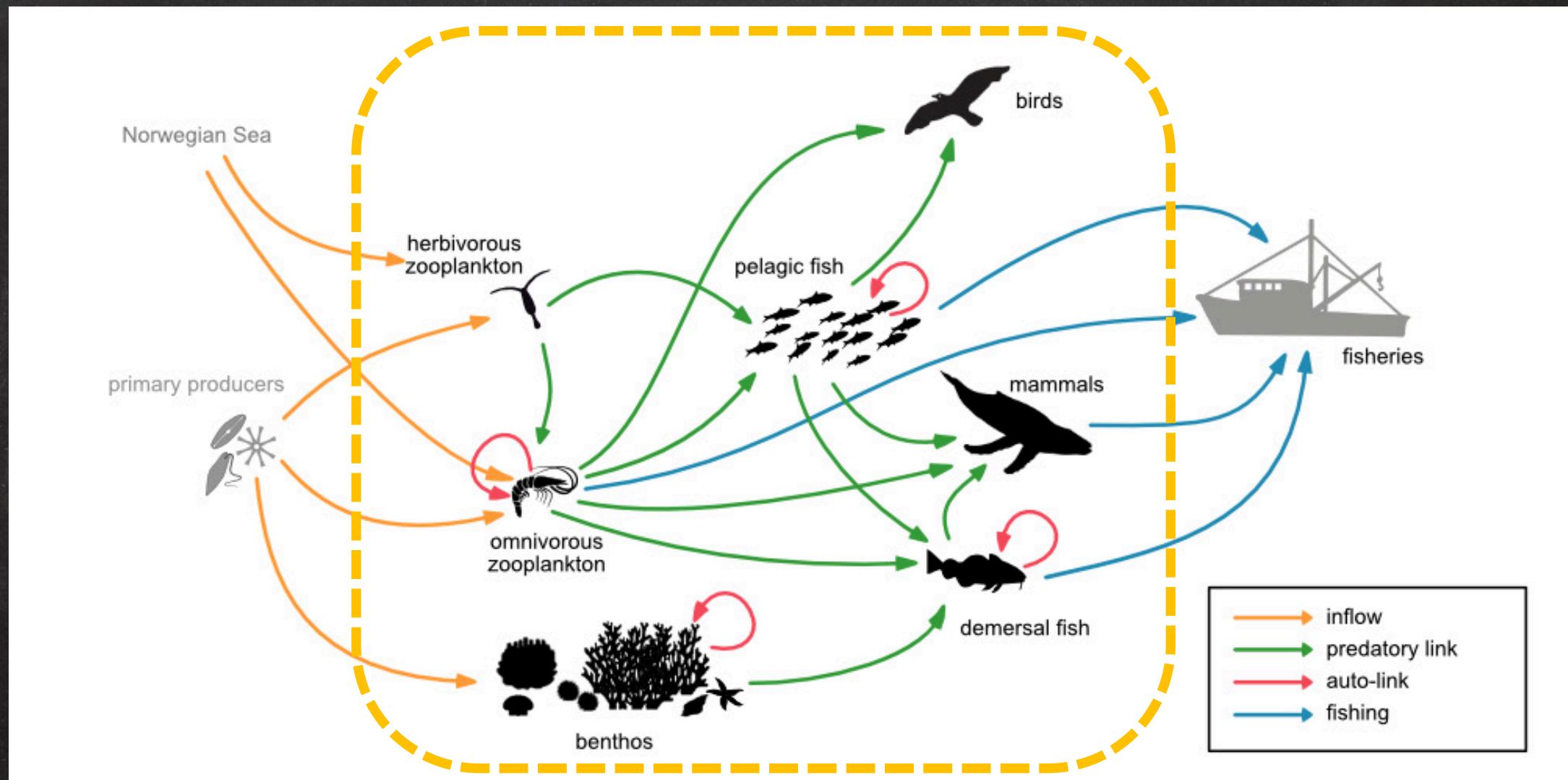


V

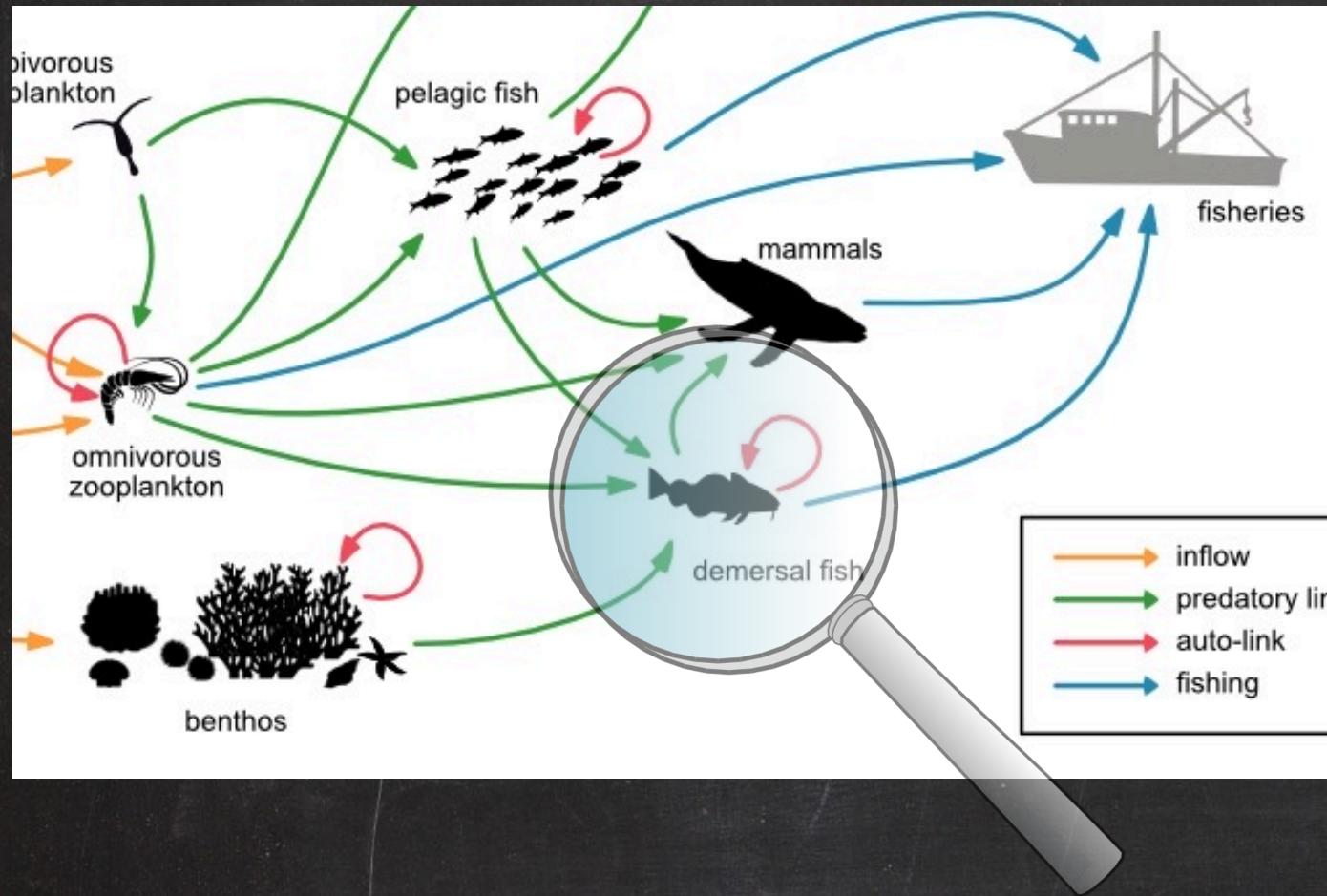
A food-web example



Simplified food-web for the Barents Sea



Simplified food-web for the Barents Sea: focus on demersals



General processes:

Variation in demersals biomass is the result of gains (feeding) minus losses (predation, fishing, non-assimilates/digested fraction and metabolic losses)

'General' constraints:

Satiation: max feeding by demersals depends on their total biomass

Inertia: the change in biomass of demersals from one year to the next is limited

Refuge Biomass: the biomass of demersals cannot go below a minimal level.

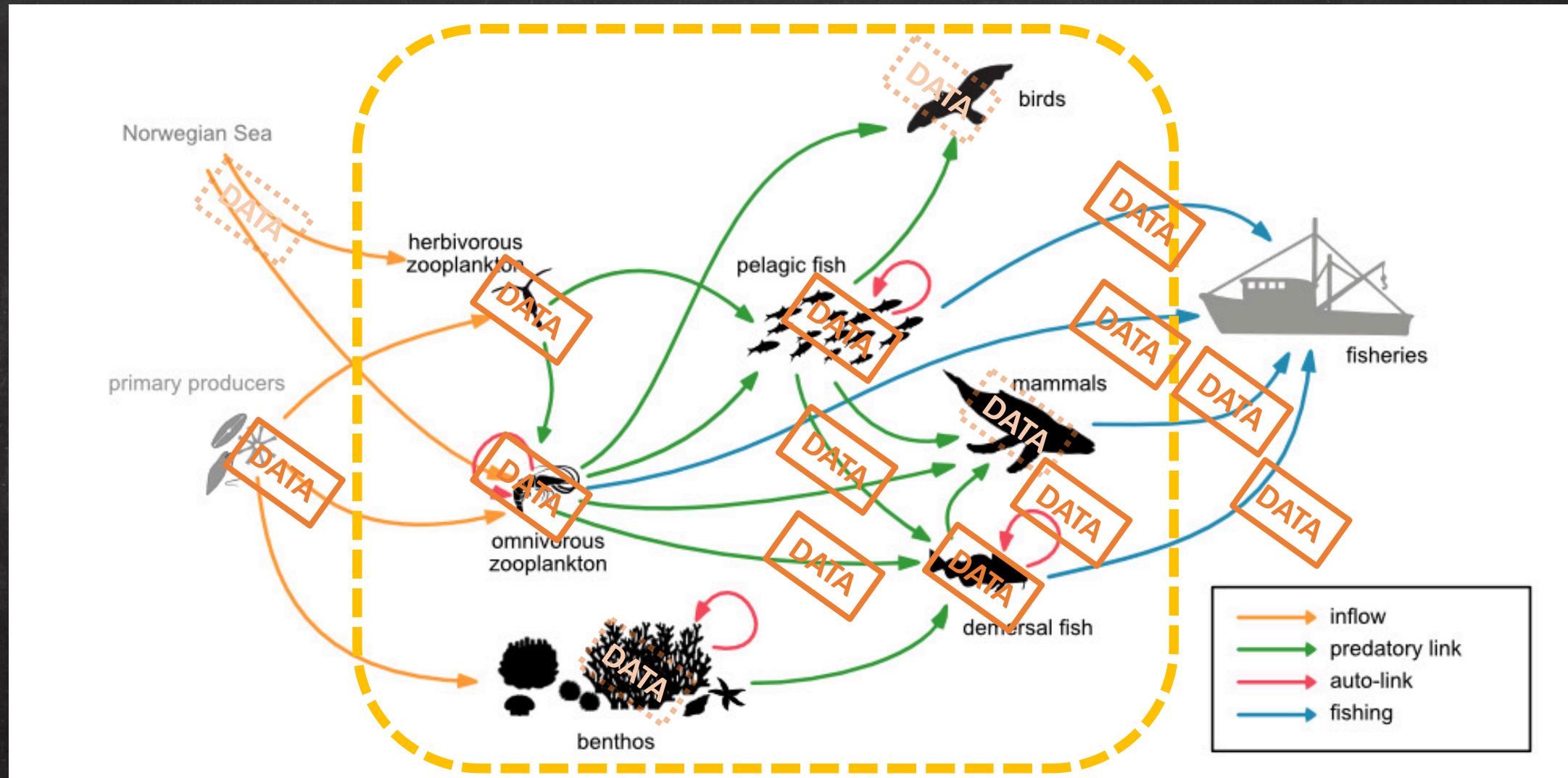
'Observational' constraints:

Biomass variations follow observations

Trophic flows follow observations

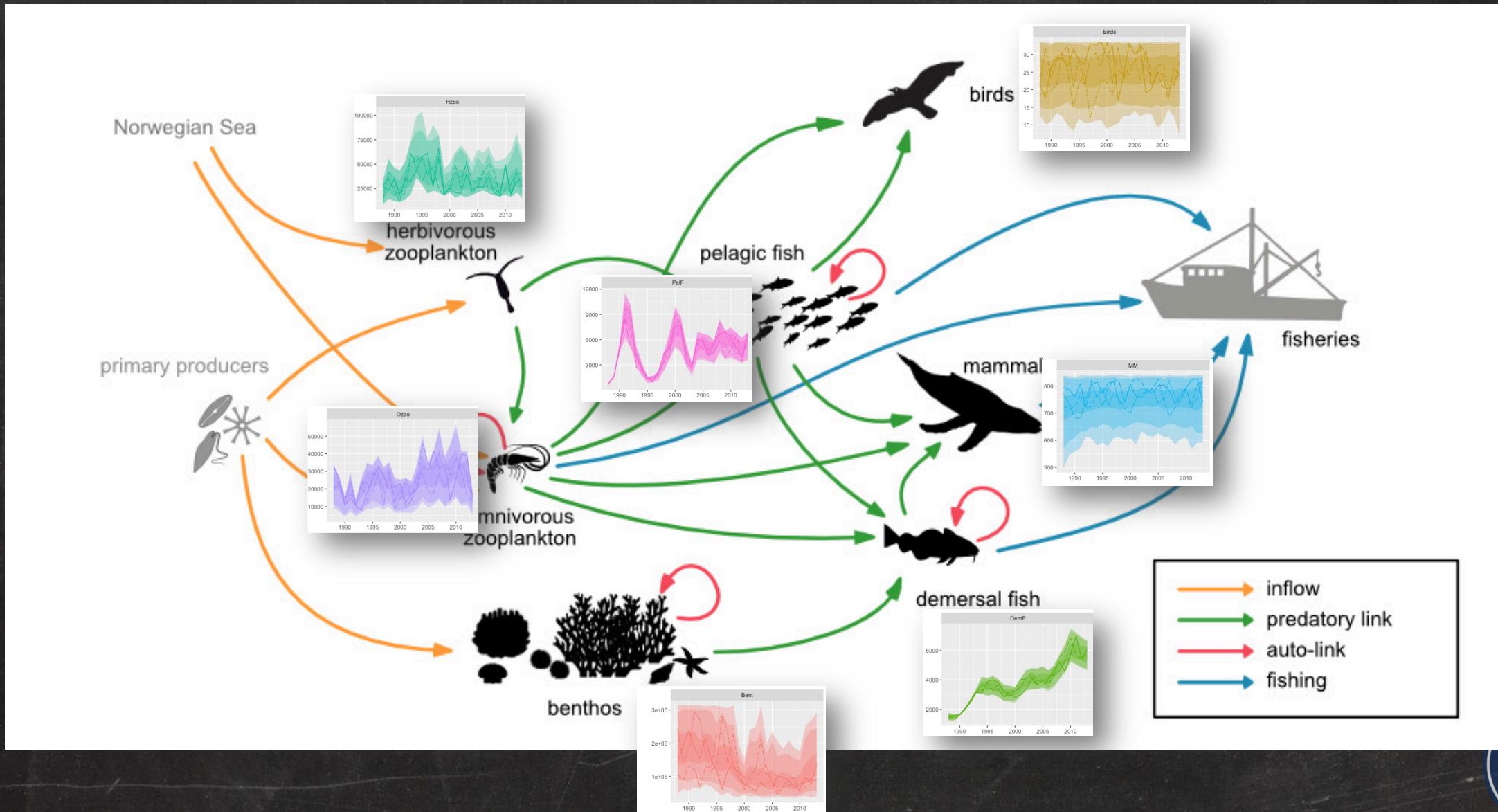
Fisheries catch follow observations

A lot of data...but not everywhere and not of equal quality

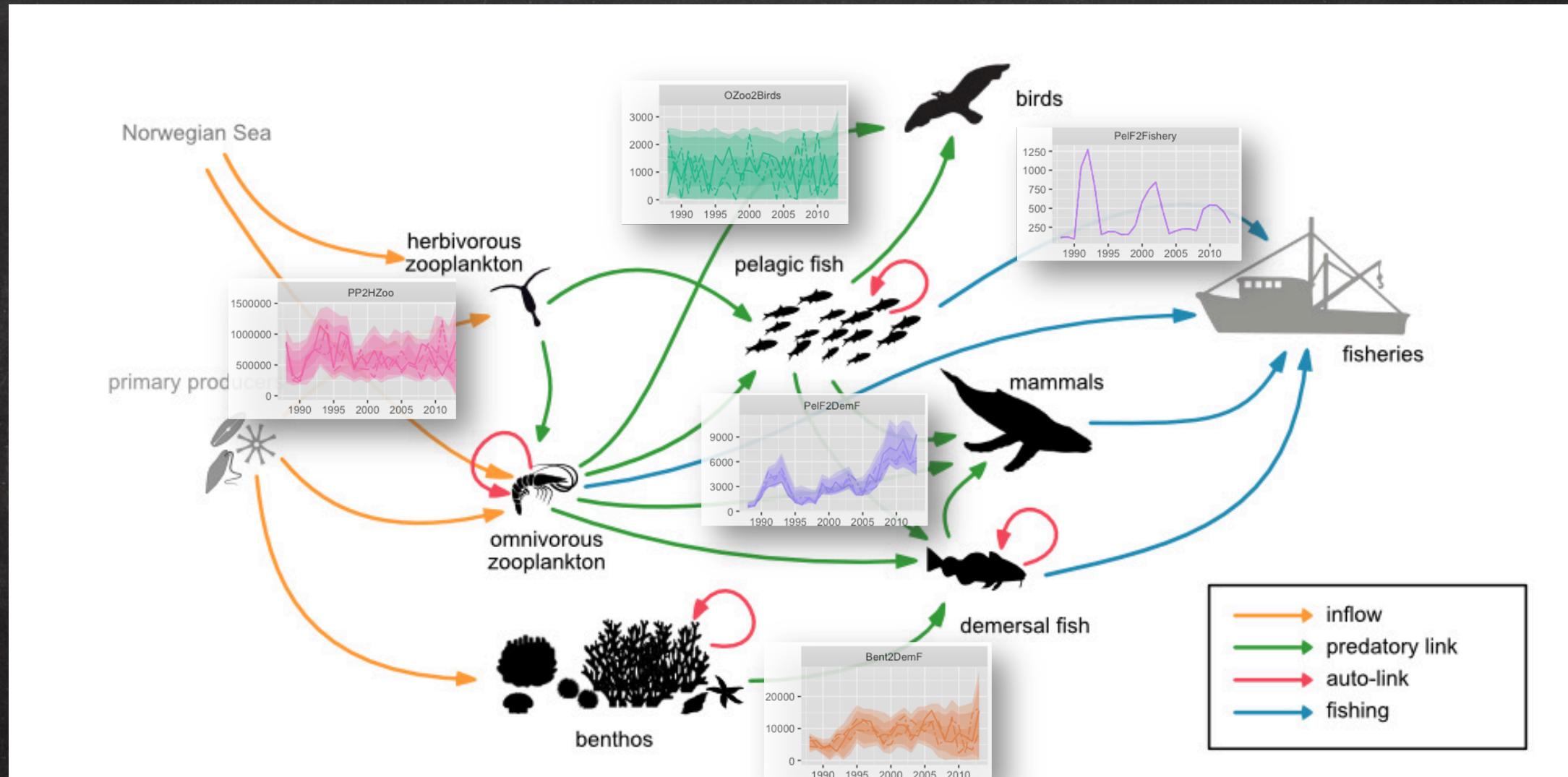


Multiple data types: absolute estimates, relative indices, ranges of values, diet fractions, ...

Reconstructed biomasses



Reconstructed flows (trophic and non-trophic)



Information

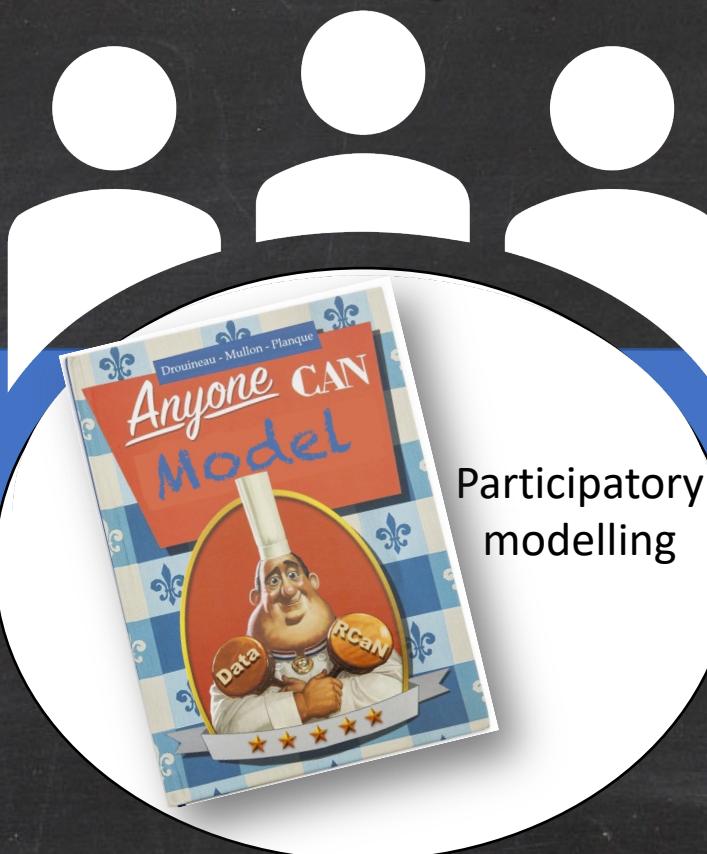
CaN

IEA



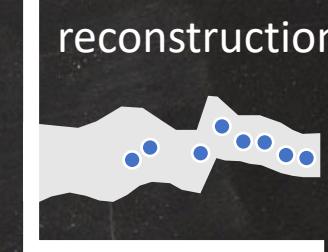
Integrate
uncertainties
& unknowns

Chance and Necessity
modelling

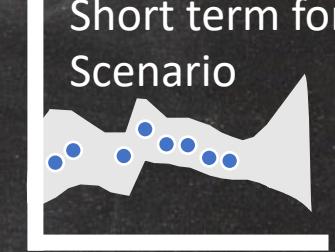


Participatory
modelling

Historical
reconstructions



Short term forecasts
Scenario



A summary of all the ‘ingredients’ of an RCaN model

1. Components: trophospecies, fishing fleets, food sources
2. Input biological parameters: Assimilation efficiency, digestibility, other losses, inertia, satiation
3. Fluxes: trophic, import/export, fishing
4. Observational series
5. Constraints



VI

RCaN and RCaN-Constructor



Excel template, RCaN & RCaN constructor

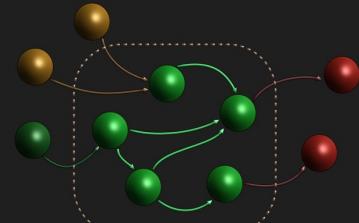


RCaN
Model constructor

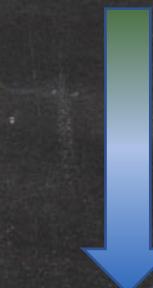
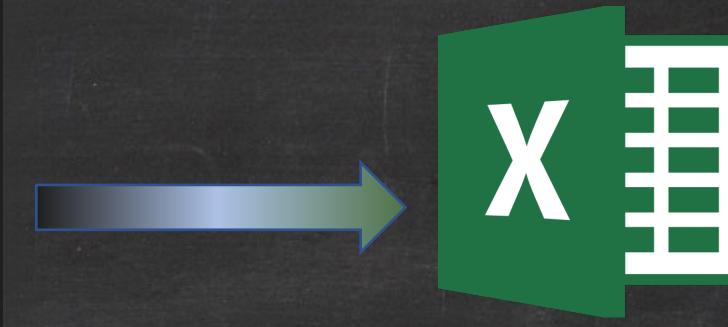
Christian Mullon
Hilaire Drouineau
Benjamin Planque

2020 - INRAE (France) / IMR (Norway)

$B_{t+1} = (1-H)B_t + \sum_f N_f F_f$

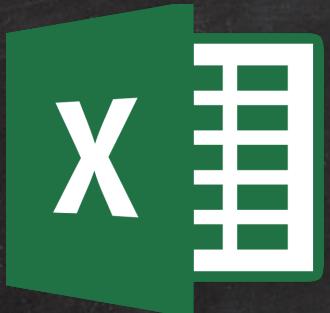


RCaN
Excel template



The RCaN template

RCaN
Excel template



- Document in xlsx format
- Compulsory worksheets:
 1. Components & 2. Input parameters
 3. Fluxes
 4. Input time-series
 5. Constraints
- Additional worksheets:
 6. INFO
 7. InputFiles

Worksheet #1. Components & input parameters

C3 X ✓ f_x | 0.93

1	A	B	C	D	E	F	G	H	I
1	Component	Inside	Assimilation	Digestibility	OtherLosses	Inertia	Satiation	RefugeBiomass	
2	Food		0		0.65				
3	Population		1	0.93	0.85	1.65	0.25	5.5	0.01
4									
5									
6									

INFO Components & input parameter Fluxes Constraints Input time-series InputFiles + 150%



Worksheet #2. Fluxes

	A	B	C	D	E	F	G
1	Flux	From	To	Trophic			
2	Feeding	Food	Population	1			
3							
4							
5							
6							

INFO Components & input parameters Fluxes Constraints Input time-series InputFiles +

Ready 176%

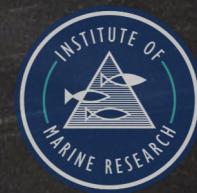


Worksheet #4. Input time-series

1	A	B	C	D	E	F	G	H
1	Year	ObservedBiomass						
2	2000	3						
3	2001	3.03						
4	2002	3.6966						
5	2003	4.029294						
6	2004	3.26372814						
7	2005	3.78592464						
8	2006	4.58096882						
9	2007	6.77983385						
10	2008	8.20359896						
11	2009	7.71138302						
12	2010	11.335733						
13	2011	16.3234556						
14	2012	19.2616776						
15	2013	18.1059769						

Components & input parameter Fluxes Constraints **Input time-series**

Ready



Worksheet #3. Constraints

	B	C	D	E
1	Constraint	Time-range	Active	Information
2	Population<=10	2000	1	Constraint on biomass
3	Feeding<=100	2000:2039	1	Constraint on flux. Carrying Capacity
4	Population<=ObservedBiomass*1.2	2000:2019	0	Constraint biomass on observations (upper bound)
5	Population>=ObservedBiomass/1.2	2000:2019	0	Constraint biomass on observations (lower bound)
6				
-				

INFO Components & input parameter Fluxes Constraints Input time-series InputFiles +

Ready 162%



Worksheet #1, #2, #3, #4: Connections

Components

Component	in_out	AssimilationE	Digestibility	OtherLosses	Inertia	Satiation	RefugeBiomass
Food	Out		0.65				
Population	In	0.93	0.85	1.65	0.25	5.5	0.01

Constraint	Time-range	Active	Information
Population<10	2000	2000	1 Constraint on biomass
Feeding<=100	2000:2039	2000	1 Constraint on flux. Carrying Capacity
Population< ObservedBiomass*1.2	2000:2019	0	0 Constraint biomass on observations (upper bound)
Population> ObservedBiomass/1.2	2000:2019	0	0 Constraint biomass on observations (lower bound)

Fluxes

A	B	C	D	E	F	G
1 Flux	From	To	Trophic			
3 Feeding	Food	Population	1			
4						
5						
6						

A	B	C	D	E	F	G	H
1 Year	ObservedBiomass						
2 2000	3						
3 2001	3.03						
4 2002	3.6966						
5 2003	4.029294						
6 2004	3.26372814						
7 2005	3.78592464						
8 2006	4.58096882						
9 2007	6.77983385						
10 2008	8.20359896						
11 2009	7.71120202						

Constraints

Data



Worksheet #5. INFO

- This contains all the metadata about the model
 - Authors, date-time, objectives, data sources, parameter calculations, literature, working group, etc...



Worksheet #6. InputFiles

- The list of CSV data files from which observations are taken
- ...this is only produced when building the template with RCaN constructor



More on constraints:

Default constraints

- *Satiation, inertia, refuge biomass, positive flows*
!!! These apply to all years

Model specific constraints

- Constant absolute limits (e.g. max PP)
- Varying absolute limits (e.g. time series on biomass, flows, catches)
- Varying relative limits (e.g. time series on survey indices)
- Ratios (e.g. diet ratios, biomass ratios)
- Inequalities & equalities can be used
!!! These can apply to selected years

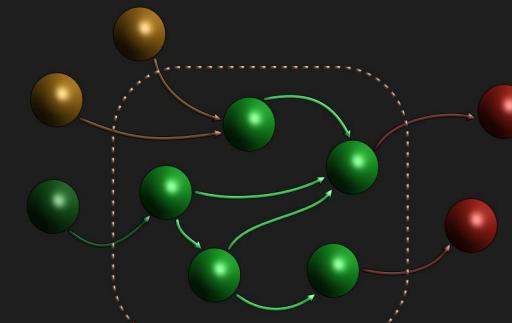


RCaN constructor

RCaN

Model constructor

Christian Mullon
Hilaire Drouineau
Benjamin Planque



$$B_{t+1} = (1 - H)B_t + \sum_f N_f F_f$$

2020 - INRAE (France) / IMR (Norway)

A graphical user interface (GUI) to support the construction of RCaN templates



PAUSE

Questions?



Derivation of biological parameters

Lindstrøm et al., 2017 Ecosystems

<https://doi.org/10.1007/s10021-016-0022-y>



Satiation

- Satiation (sigma): the maximum consumption rate per unit biomass of the predator. The coefficient is strictly positive. A coefficient of 10 indicates that a species group can eat up to 10 times its body weight in the course of 1 time unit (a year, this is equivalent to a maximum daily consumption rate of 2.7%).

$$\sigma = \frac{1}{\gamma K} f_J a_J M^{-0.25}$$

Based on body size and species specific physiology

(Q/B)

Borrowed from an existing Ecopath model



Inertia

- Inertia (rho): a parameter that bounds biomass variations between a maximum growth rate and a maximum mortality rate. The coefficient is strictly positive. The upper bound of biomass variation is given by $\exp(\rho)$ and the minimum bound rate is given by $\exp(-\rho)$. A value of 0.25 indicates that over 1 time unit (a year) the biomass of a species can reach at most $\exp(0.25) \approx 128\%$ and at least $\exp(-0.25) \approx 78\%$ of the original biomass.

$$r_{\max} = e^{a_s} e^{-\frac{b_s}{kT}} M^{-0.25}$$

$$\rho = r_{\max} \cdot d$$

Based on body size
and temperature
(metabolic theory of
ecology)

$$\rho = e^{a_l} age^{b_l} \lambda$$

Based on longevity
(life history correlates)

$$(\rho' = \bar{\rho} / \sqrt{NS}).$$

Correction for
multiple species in a
trophospecies group



Metabolic losses

- Other losses (μ): a coefficient that account for losses, i.e. metabolic losses and other mortality not explicitly accounted for in the model. The losses are expressed in the form $\exp(-\mu)$. The coefficient is strictly positive. A value of 2 indicates that over 1 time unit (a year) metabolic losses and other mortality losses will represent $1-\exp(-2) \approx 86\%$ of the original biomass (equivalent to $1-\exp(2/365) = 0.55\%$ daily losses).

$$\mu_{Gillooy} = 3 \left(e^{a_{bmr}} e^{-\frac{b_{bmr}}{T} 1000} M^{0.75} \right)$$

$BMR_{Gillooy}$

$$\mu_{Yodzis} = a_T M^{-0.25}$$

$$\mu_{BS} = 3 \left(BMR_{Mak.} 2^{\frac{(T_{BS}-T_M)}{10}} \right)$$

Based on body size and temperature
(metabolic theory of ecology,
experimental results)



Digestibility

- Digestibility correction factor (kappa): a correction factor that accounts for variations in energy content of a prey. The value of the parameter is between zero (no energy content) and one. The product of the potential assimilation efficiency by the digestibility correction factor is the absorption efficiency (the proportion of prey biomass digested and absorbed).



Assimilation Efficiency

- Potential assimilation efficiency (gamma): the proportion of the biomass ingested by a predator that is effectively assimilated. The value of the parameter is between zero (no assimilation) and one (complete assimilation). The product of the potential assimilation efficiency by the digestibility correction factor is the absorption efficiency (the proportion of prey biomass digested and absorbed).



Relevant papers

2009



A minimal model of the variability of marine ecosystems

Christian Mullon¹, Pierre Fréon¹, Philippe Cury², Lynne Shannon² & Claude Roy³

OPEN ACCESS Freely available online

Non-Deterministic Modelling of Food-Web Dynamics

Benjamin Planque^{1*}, Ulf Lindstrøm¹, Sam Subbey²

¹Institute of Marine Research, Tromsø, Norway, ²Institute of Marine Research, Bergen, Norway

2014



J. Math. Biol. (2016) 73:575–595
DOI 10.1007/s00285-015-0959-z

ICES Journal of Marine Science

ICES Journal of Marine Science (2020), 77(4), 1573–1588. doi:10.1093/icesjms/fsz173



Contribution to the Themed Section: 'Science in support of a nonlinear non-equilibrium world'

Food for Thought

Modelling chance and necessity in natural systems

Benjamin Planque ^{1*} and Christian Mullon ²

2020

New Results

RCaN : a software for Chance and Necessity modelling

Hilaire Drouineau, Benjamin Planque, Christian Mullon
doi: <https://doi.org/10.1101/2021.06.09.447734>

This article is a preprint and has not been certified by peer review [what does this mean?].

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2014

Non-Deterministic Modelling of Food-Web Dynamics

Benjamin Planque^{1*}, Ulf Lindstrøm¹, Sam Subbey²
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PLOS ONE

J. Math. Biol. (2016) 73:575–595
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2016

Mathematical Biology

Exploring stochasticity and imprecise knowledge based on linear inequality constraints

Sam Subbey^{1,2} · Benjamin Planque^{2,3} · Ulf Lindstrøm³

Ecosystems (2017) 20: 163–182
DOI: 10.1007/s10021-016-0022-y

2017

Multiple Patterns of Food Web Dynamics Revealed by a Minimal Non-deterministic Model

Ulf Lindstrøm,¹ Benjamin Planque,^{1,2*} and Sam Subbey^{2,3}

ECOSYSTEMS

zenodo

November 18, 2021

Quantification of trophic interactions in the Norwegian Sea pelagic food-web over multiple decades.

Benjamin Planque; Aurélien Favreau; Bérengère Husson; Erik Mousing; Cecilie Hansen; Cecilie Broms; Ulf Lindstrøm; Elliot Sivel

<https://github.com/inrae/RCaN>

+packages:

symengine, ggplot2, ggraph, coda, DT, tidyverse



<https://doi.org/10.1101/2021.06.09.447734>

New Results

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