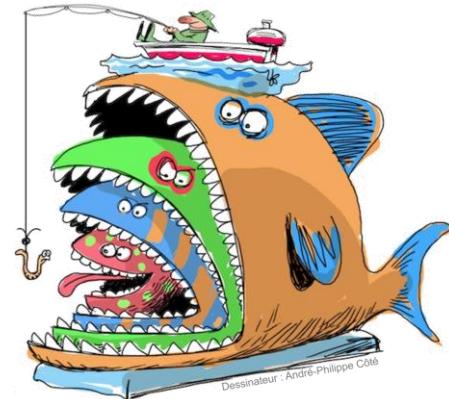


Fonctionnement trophique de l'écosystème mer Celtique : impact de la pêche et scénarios de gestion écosystémique



Pierre-Yves Hernvann

CONTEXTE



Dessinateur : André-Philippe Côté

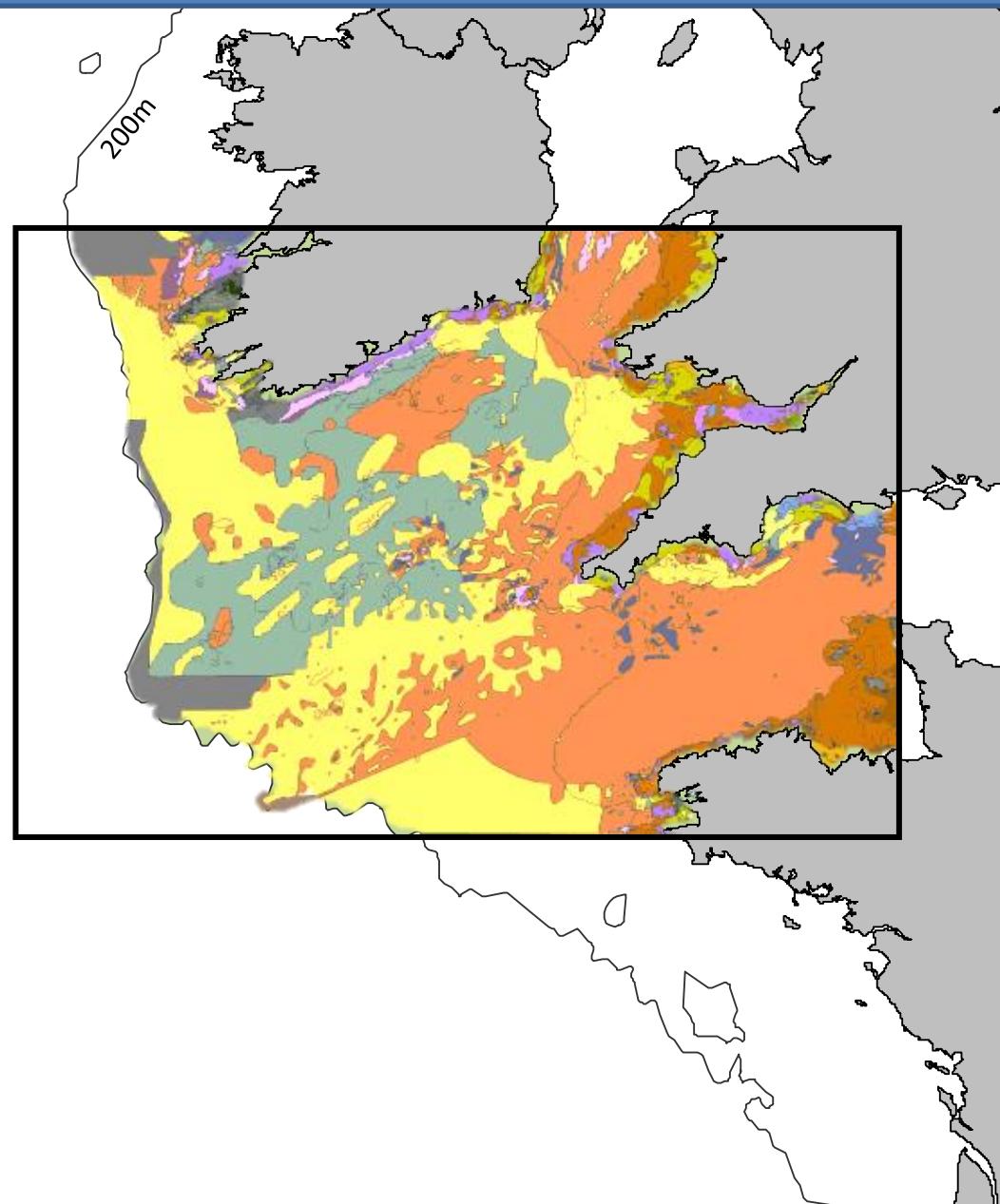
A CASE STUDY: THE CELTIC SEA

- North-West European ecosystem



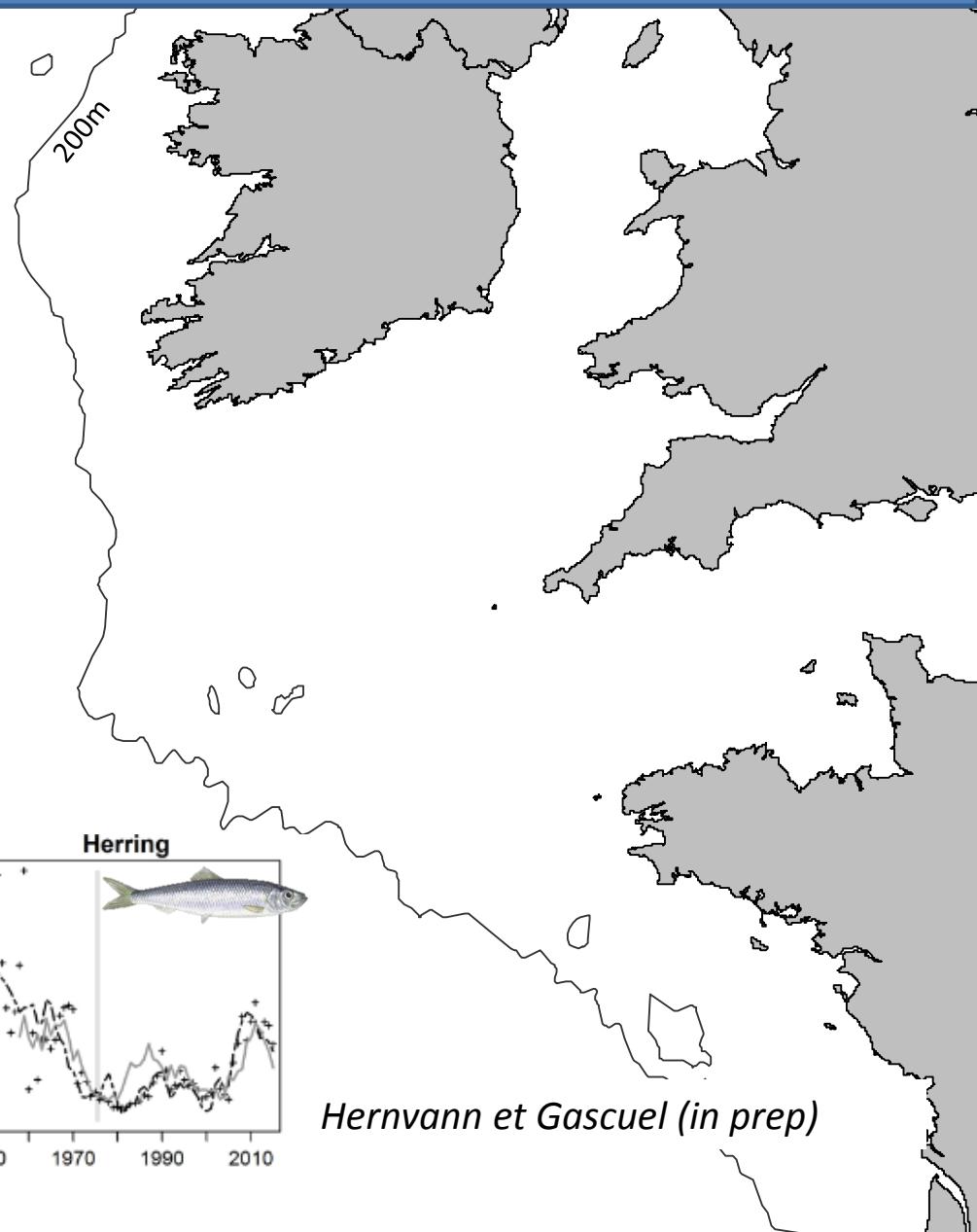
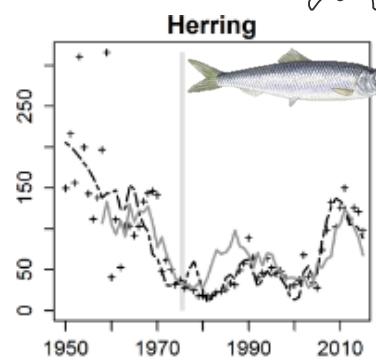
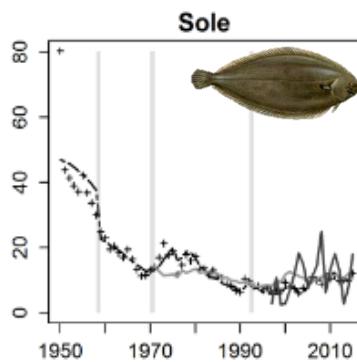
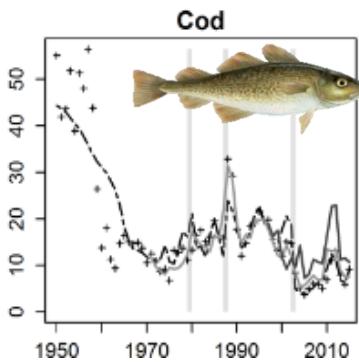
A CASE STUDY: THE CELTIC SEA

- North-West European ecosystem
- Great variety of habitats
→ diversity of species



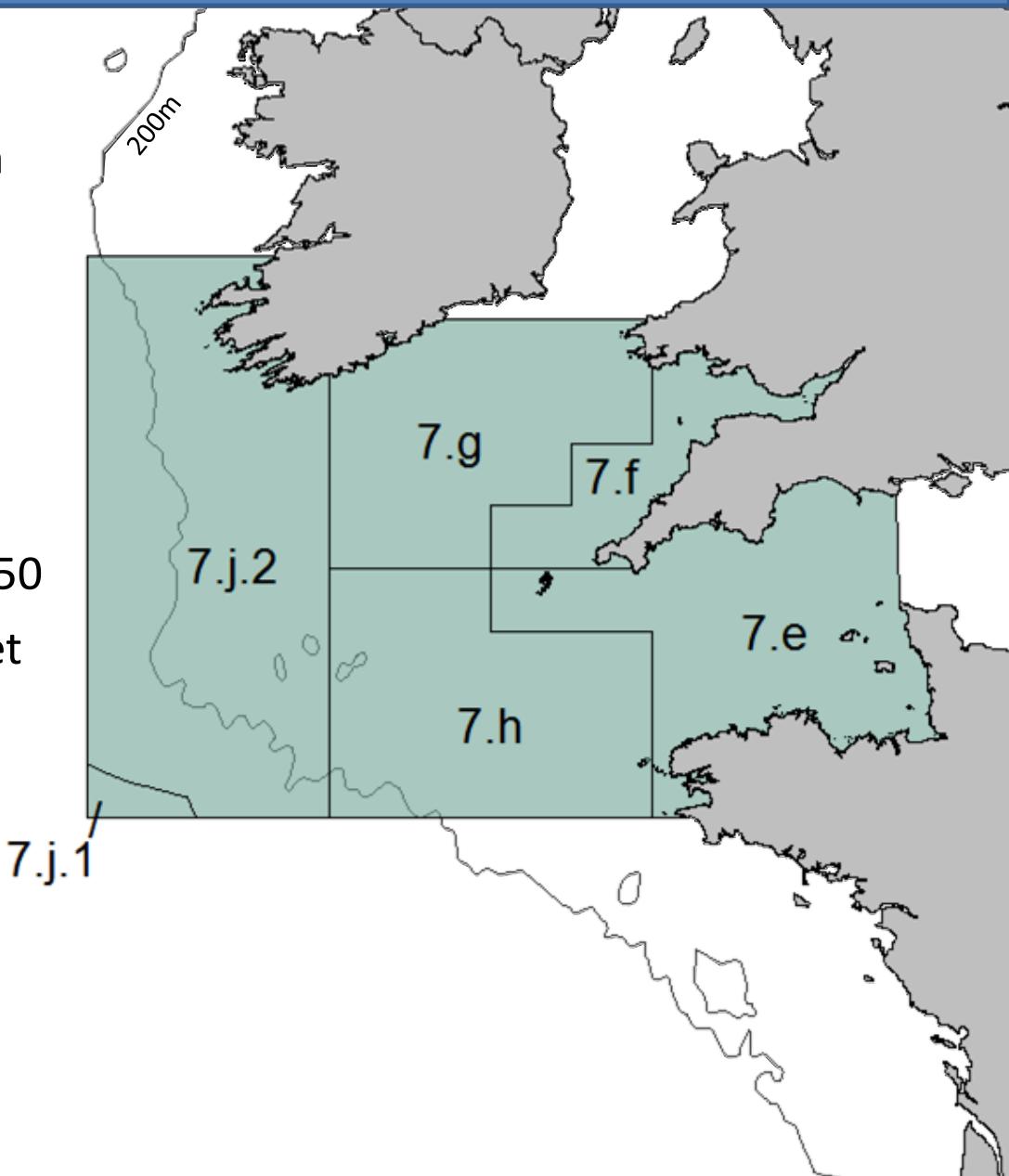
A CASE STUDY: THE CELTIC SEA

- North-West European ecosystem
- Great variety of habitats
→ diversity of species
- Intensive fishing since the XXth century and particularly after 1950
→ 75% reduction of main target species biomass in 1980



A CASE STUDY: THE CELTIC SEA

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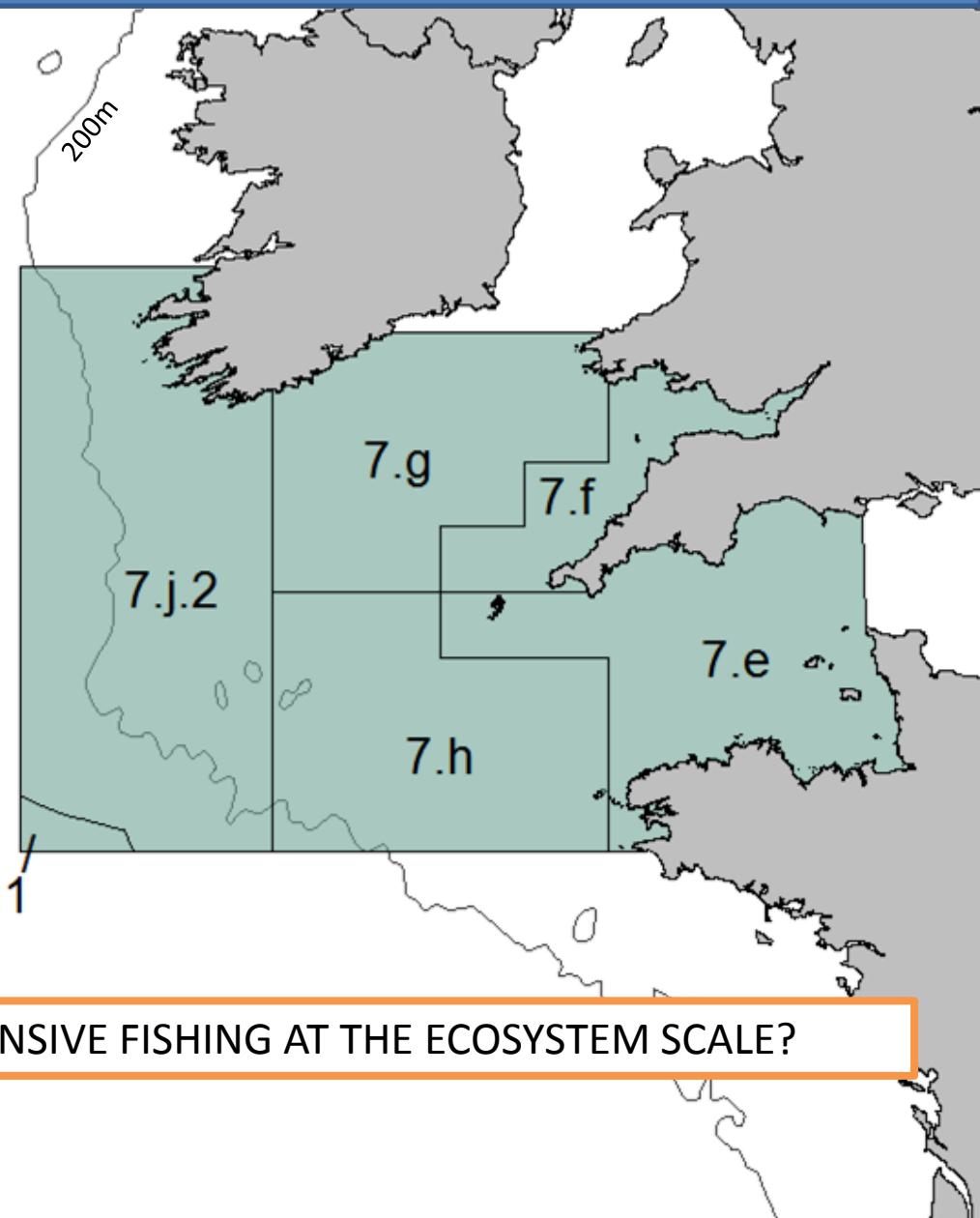
$$\text{[Flag 1]} + \text{[Flag 2]} + \text{[Flag 3]} + \text{[Flag 4]} = 340,000 \text{ t}$$

13.5% French catch in FAO27

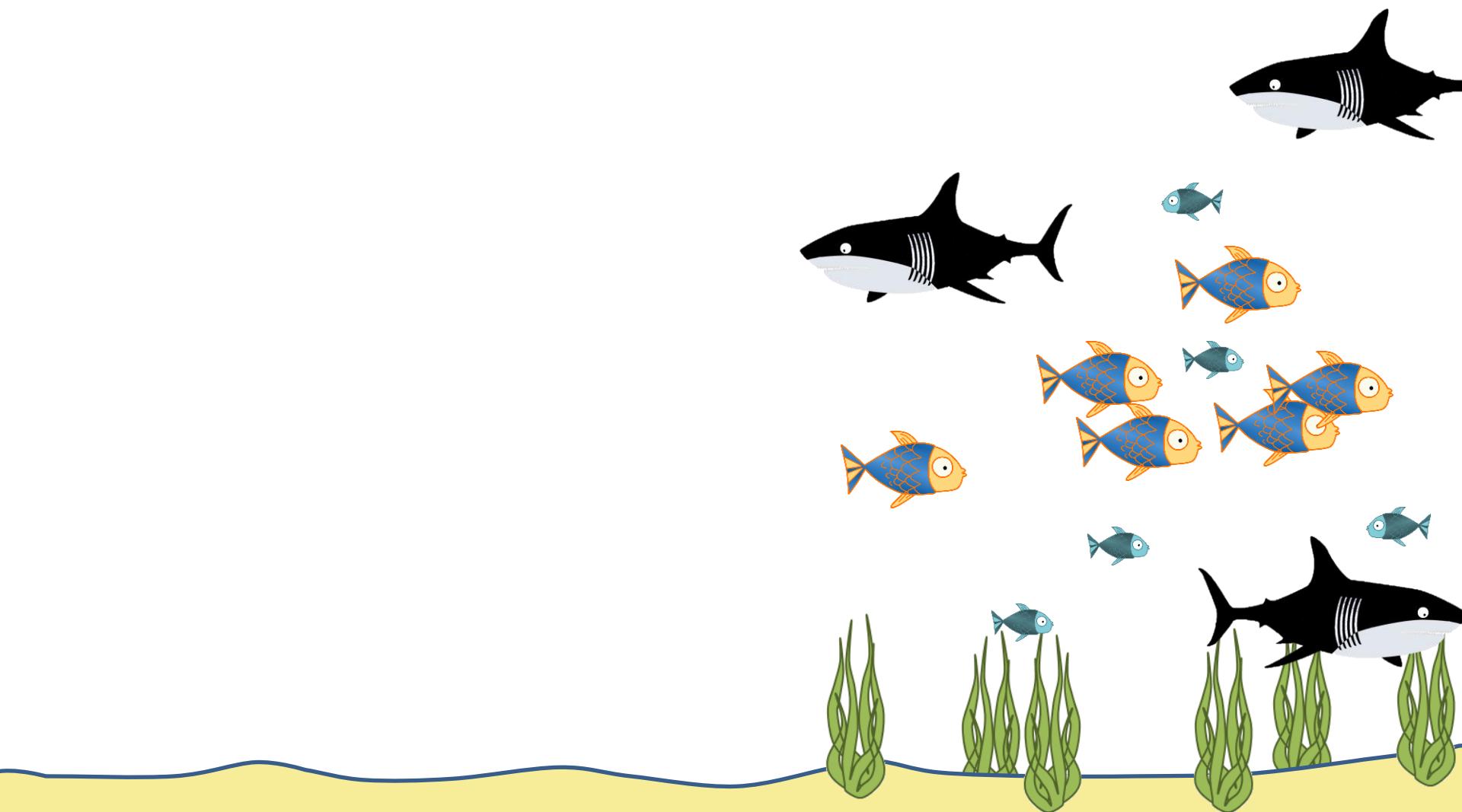
A CASE STUDY: THE CELTIC SEA

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WHAT IS THE IMPACT OF SUCH INTENSIVE FISHING AT THE ECOSYSTEM SCALE?

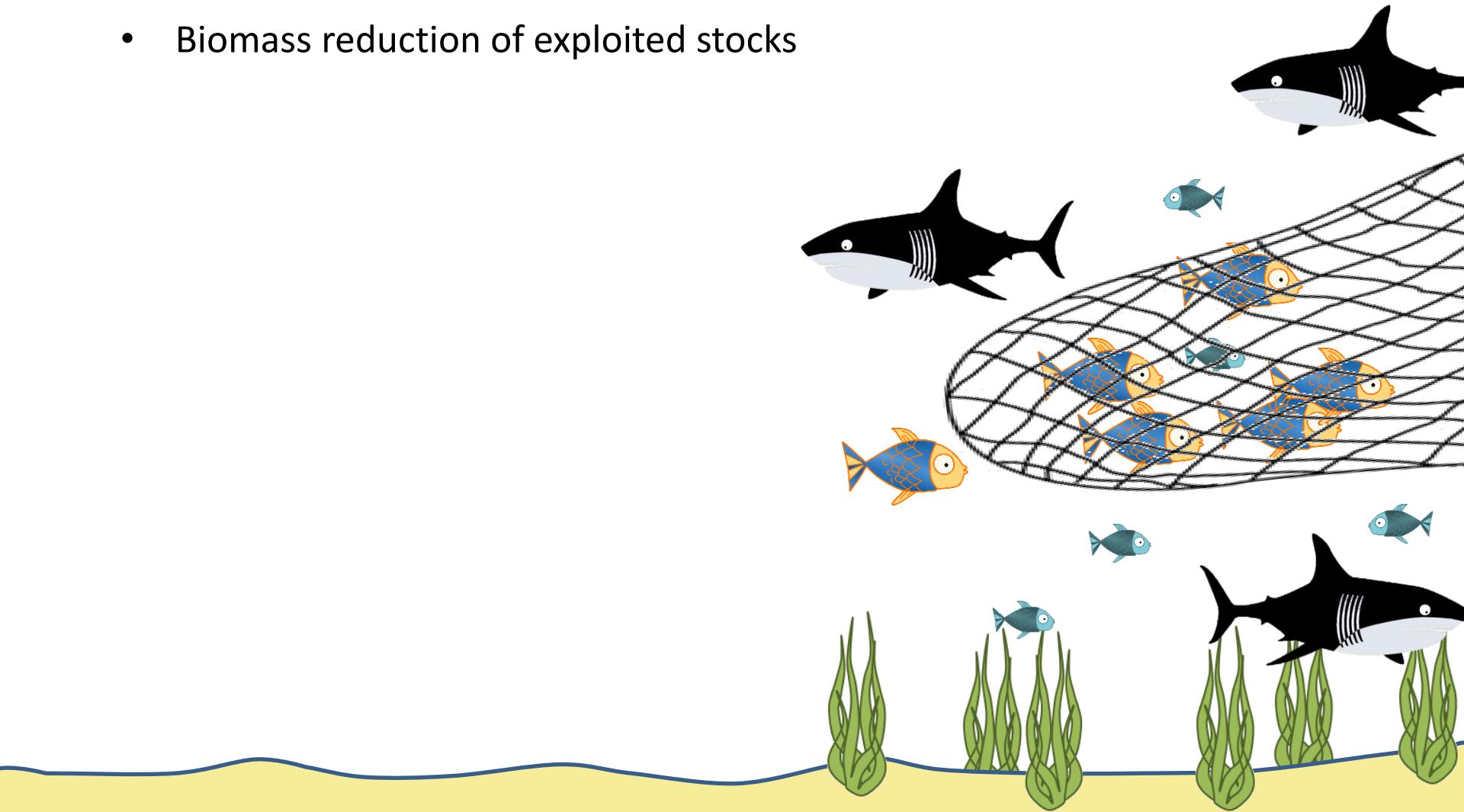


FISHING IMPACT ON ECOSYSTEMS – The trophic side



FISHING IMPACT ON ECOSYSTEMS

- Biomass reduction of exploited stocks



FISHING IMPACT ON ECOSYSTEMS – The trophic side

- Biomass reduction of exploited stocks
- Changes of species composition

Predators

Preys

Competitors



FISHING IMPACT ON ECOSYSTEMS – A TROPHIC VIEW

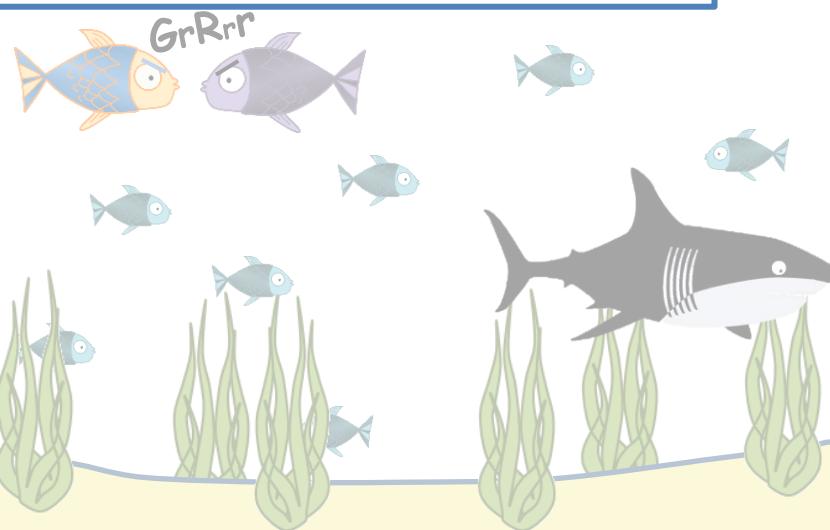
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MOVE TOWARD AN ECOSYSTEM-BASED FISHERIES MANAGEMENT

Preys

Competitors

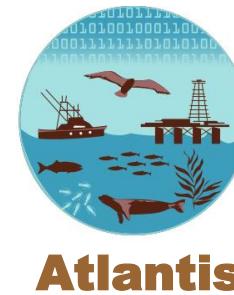


ABOUT ECOSYSTEM MODELLING

- Need of an ecosystem-based fisheries management (Pikitch et al., 2014; Garcia et al., 2003 ...)

Scientific Community + Policy Makers (CFP)

- One tool to support an EBFM is ecosystem modeling, especially trophic modeling (Plaganyi, 2007)

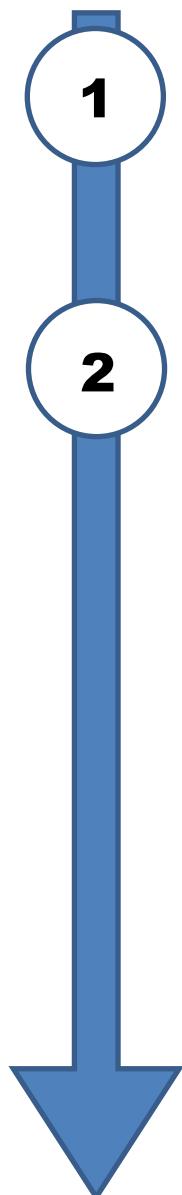


etc.

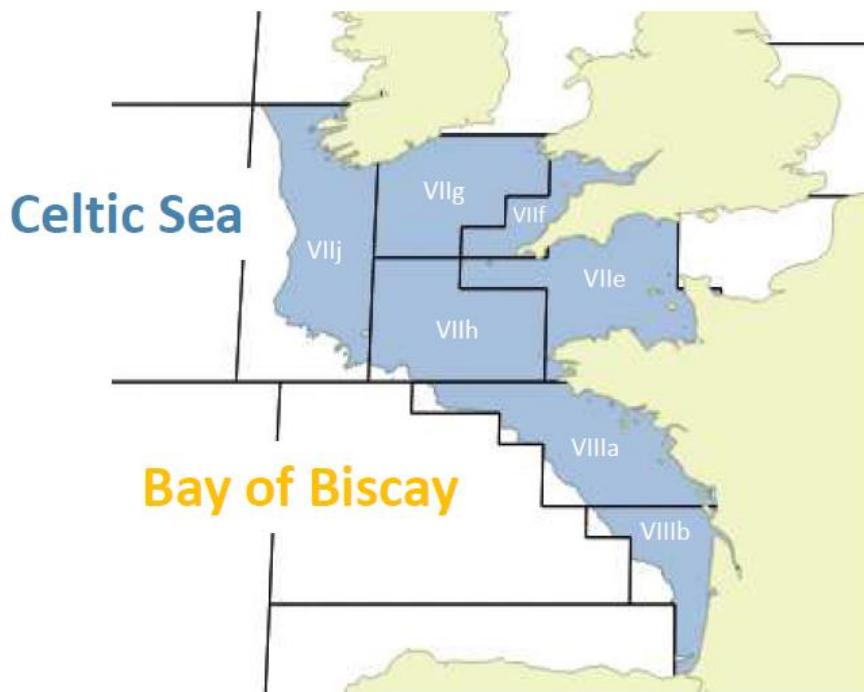
PRE-EXISTING MODELS

Guénette & Gascuel (2009)

Bentorcha et al (2017)



1980 - 2012



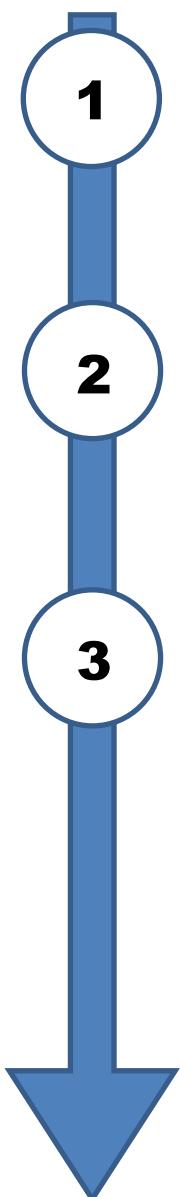
+ EBFM designed
+ Actualisation

PRE-EXISTING MODELS

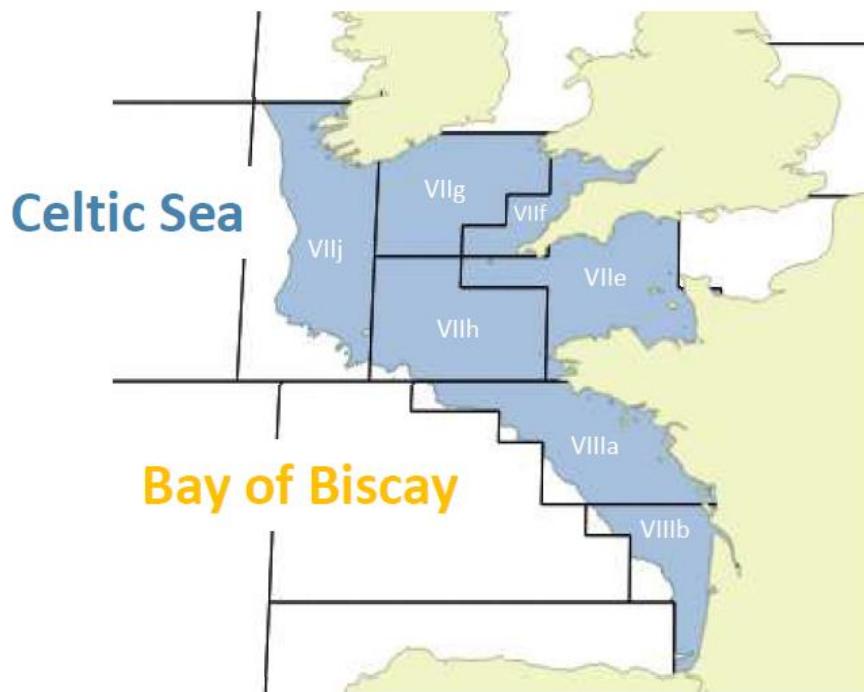
Guénette & Gascuel (2009)

Bentorcha et al (2017)

Moullec et al (2017)

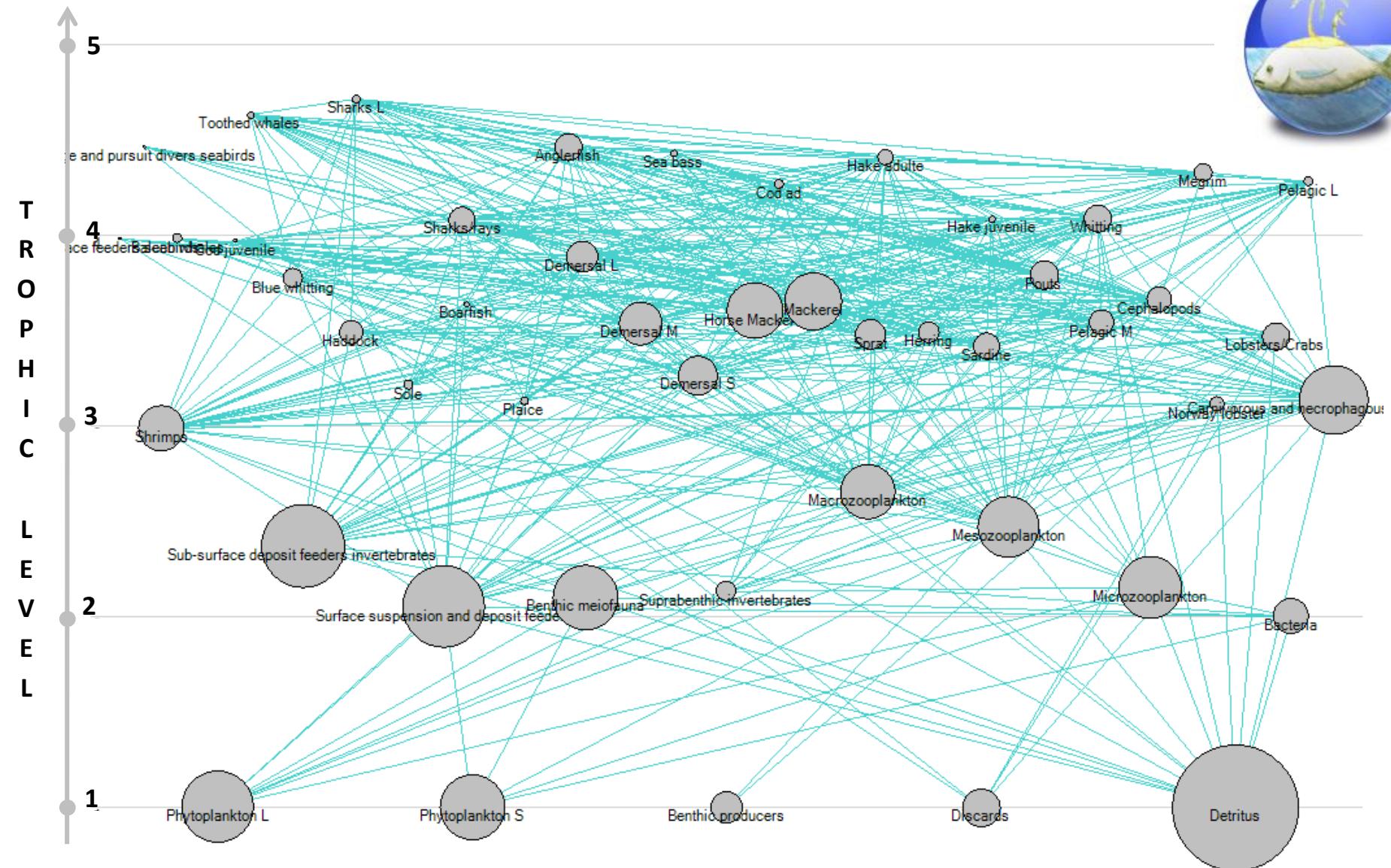


1980 - 2013



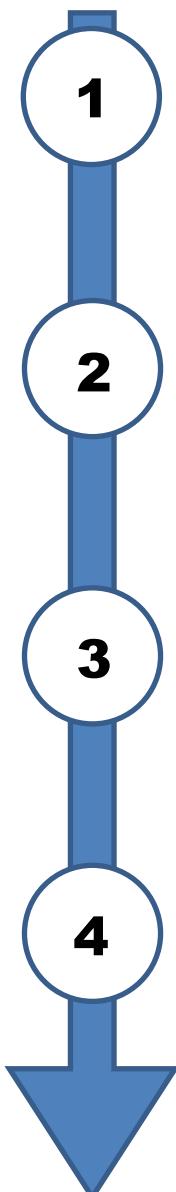
+ EBFM designed
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PRE-EXISTING MODELS – Celtic Sea model (Moullec et al. 2015)



PRE-EXISTING MODELS

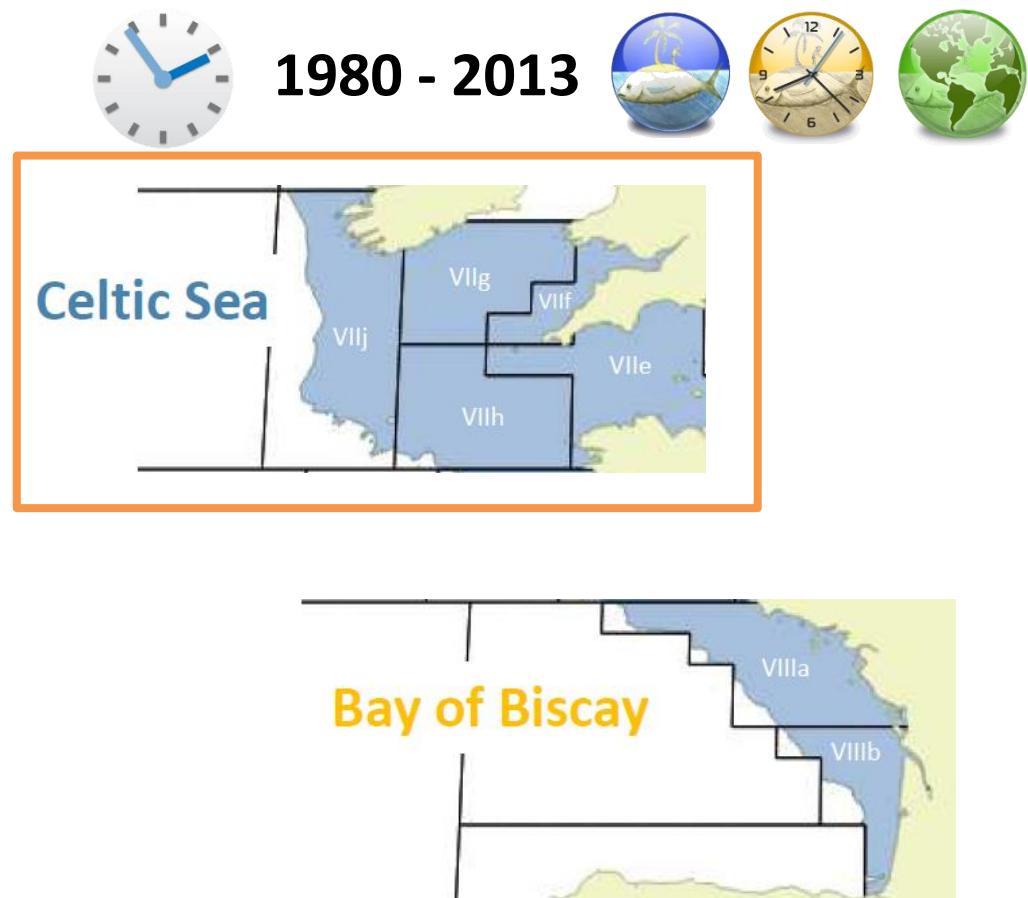
Guénette & Gascuel (2009)



Bentorcha et al (2017)

Moullec et al (2017)

Me!



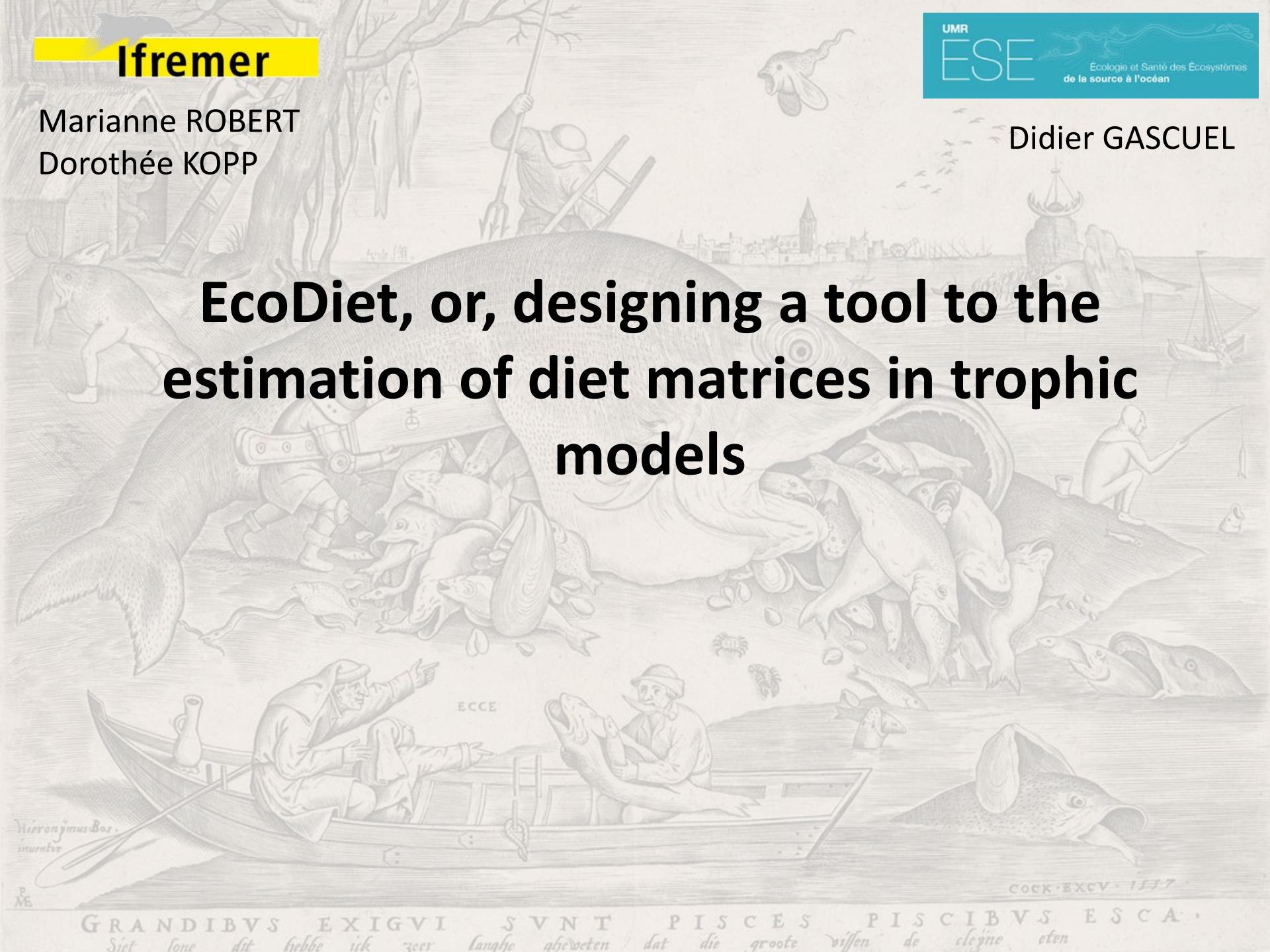
AU MENU

- What are the fisheries and environmental determinants of the Celtic Sea ecosystem dynamics?
What would be the impacts of various management measures and how can trophic models contribute to the identification of fisheries in order to minimize the impacts on ecosystems?
- In practical terms: Improve the predictive capacity of the Celtic model and Test realistic ecosystem-based fisheries management scenarios
- 3 main steps:
 - 1 Develop a generic method to estimate diet matrices in Ecopath models and apply the method to the Celtic Sea
 - 2 Spatialize the trophic model through Ecospace
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AU MENU

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EcoDiet, or, designing a tool to the estimation of diet matrices in trophic models



GRANDIBVS EXIGVI SVNT PISCES PISCIBVS ESCA.
Siet sou dit hebbe ik zeer langhe ghevachten dat die grote vissen de cleyne eten

~~CONTEXTE~~

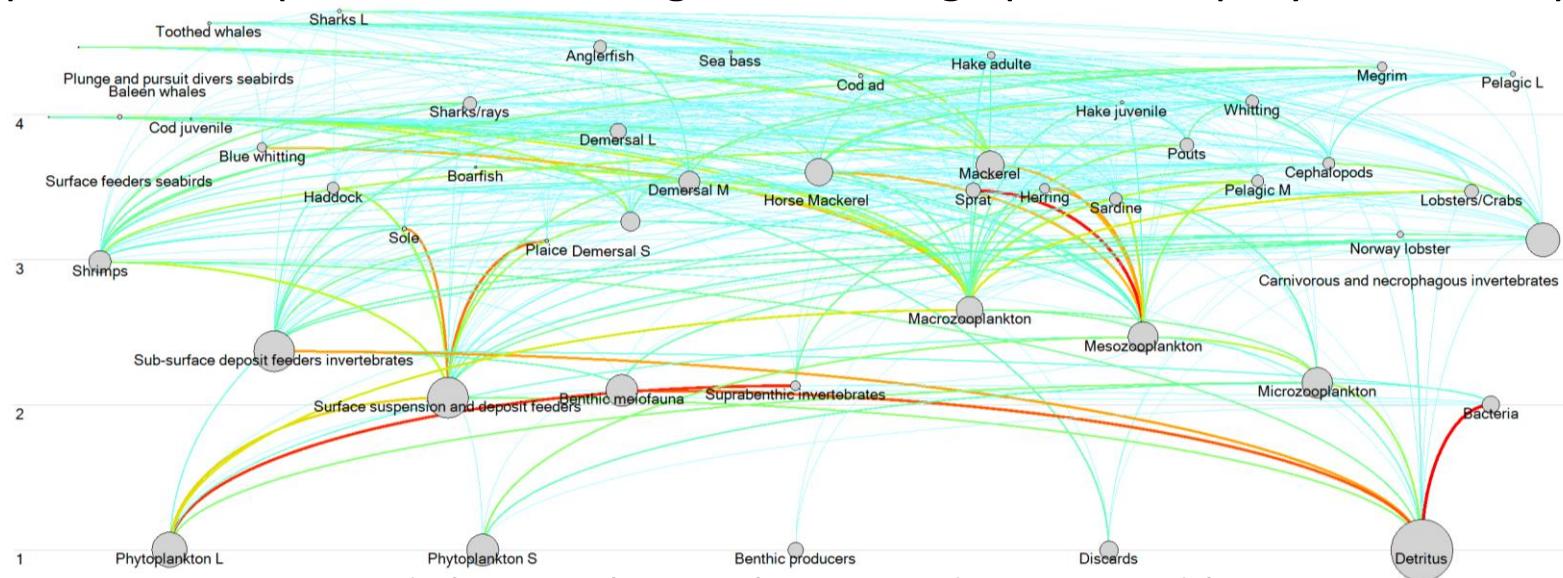
INTRO



Dessinateur : André-Philippe Côté

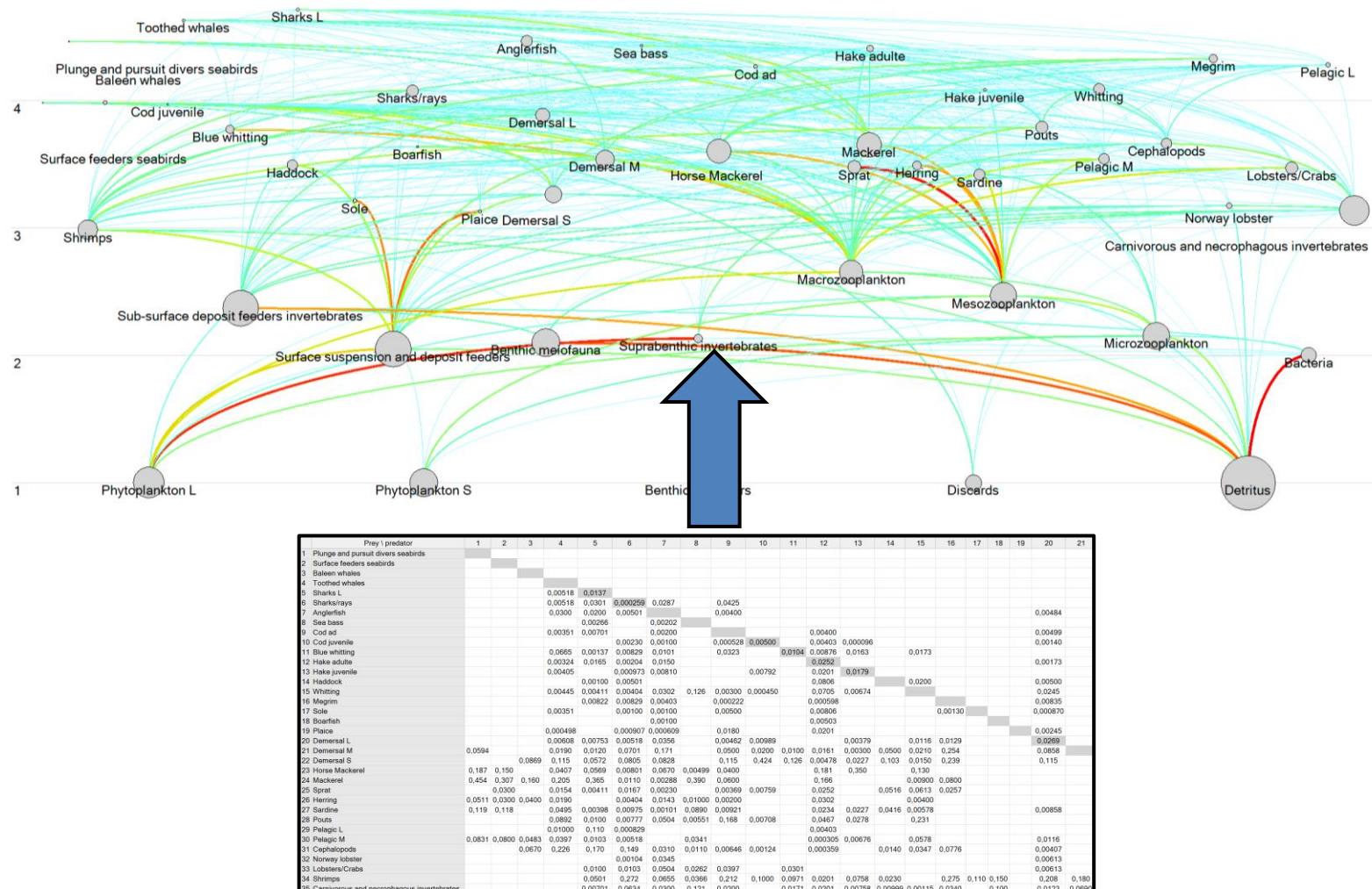
THE DIET MATRIX

Species or trophic connected together through predator/prey relationships



...and these relationships are determined by...

THE DIET MATRIX



...the DIET MATRIX !



THE DIET MATRIX

Predators (=consumers)

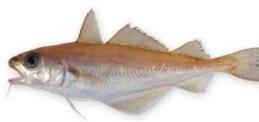


THE DIET MATRIX

Predators (=consumers)



THE DIET MATRIX

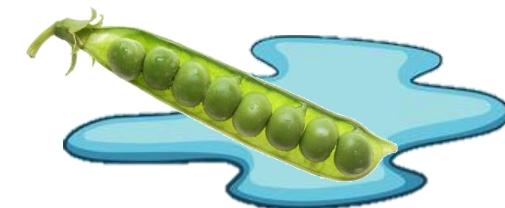


	Cod
Sharks/rays	0,04
Blue whiting	0,03
Sole	0,01
Plaice	0,02
Demersal M	0,05
Demersal S	0,12
Horse mack.	0,04
Mackerel	0,06
Sardine	0,01
Pouts	0,17
Cephalopods	0,01
Lobsters/Crabs	0,04
Shrimps	0,21
Carn/necro	0,02
SSDF	0,07
Macrozoo	0,05
Mesozoo	0,04
Sum	1,00

The « cod case »

« je mange ***% de bidule,
 ***% de truc,
 ***% de machin »

Unité =



THE DIET MATRIX

- Generally, diet matrix are based on information from the literature

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- In the case of the current Celtic Sea model, it comes from



THE DIET MATRIX

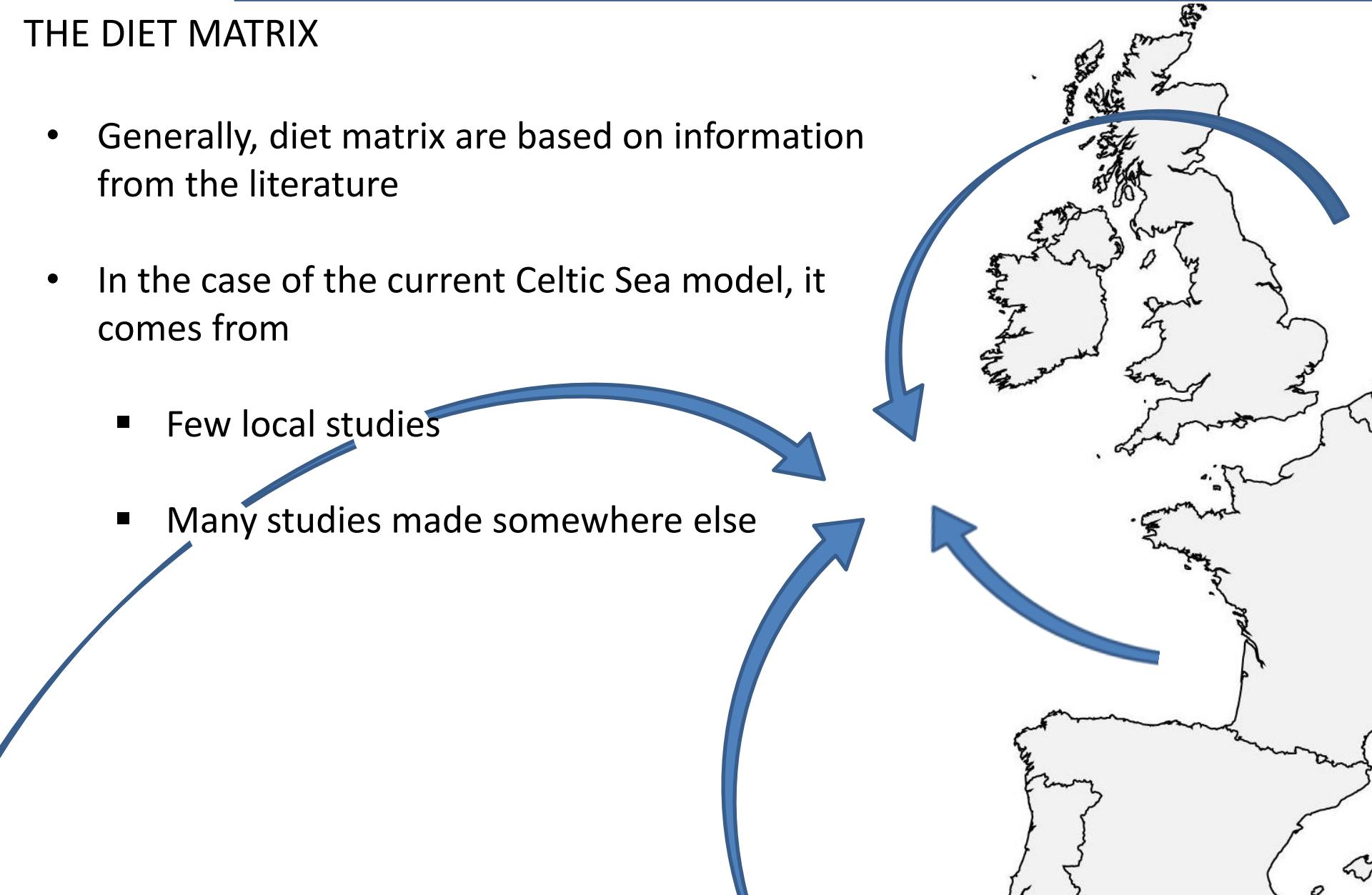
- Generally, diet matrix are based on information from the literature
- In the case of the current Celtic Sea model, it comes from
 - Few local studies



THE DIET MATRIX

- Generally, diet matrix are based on information from the literature
- In the case of the current Celtic Sea model, it comes from

- Few local studies
- Many studies made somewhere else



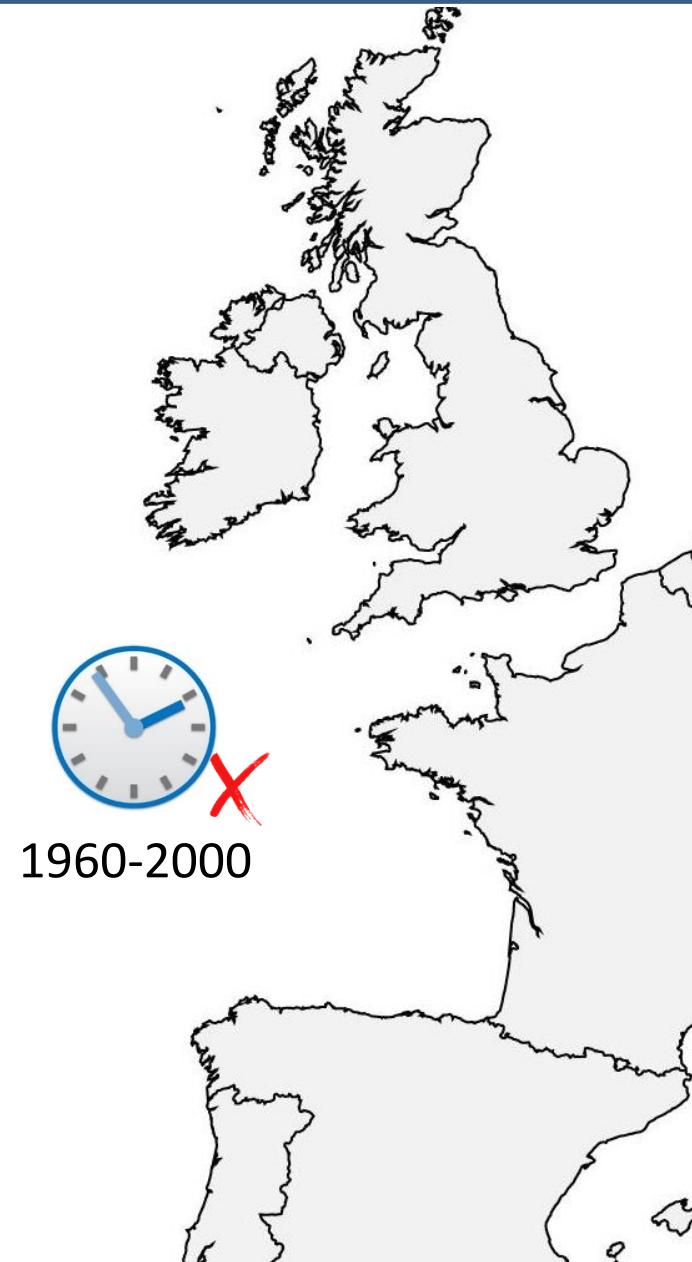
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→ NEED OF BETTER INFORMATION
(local and relevant)
ON SPECIES INTERACTIONS

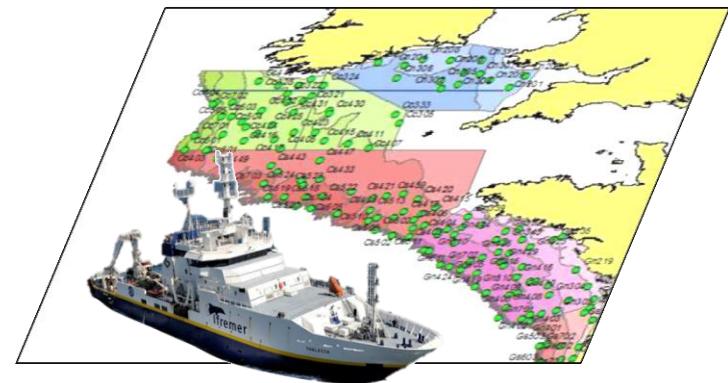


A GREAT OPPORTUNITY FOR TROPHIC MODELING IN THE CELTIC SEA

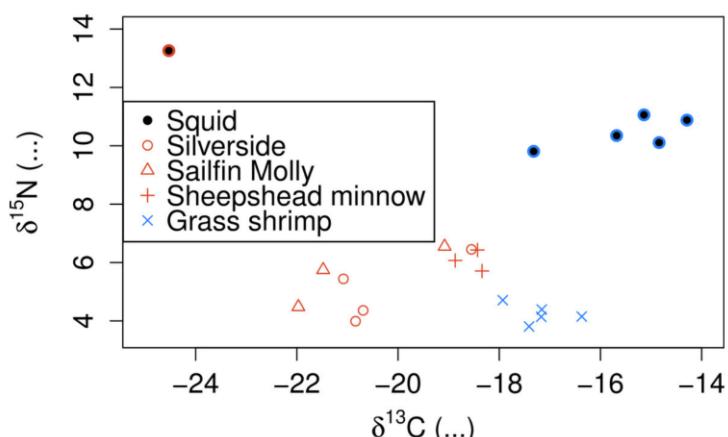
- EATME Project – IFREMER Lorient



- EVHOE survey 2014-2016 – Celtic Sea
- More than 2000 samples



Stable isotops analysis (SIA)



Stomach contents analysis (SCA)



METHODS



Dessinateur : André-Philippe Côté

CONSIDERATIONS ABOUT THE DATA:

- HOW TO USE EACH TYPE OF DATA?
- EVENTUAL DATA PREPARATION

STOMACH CONTENT ANALYSIS

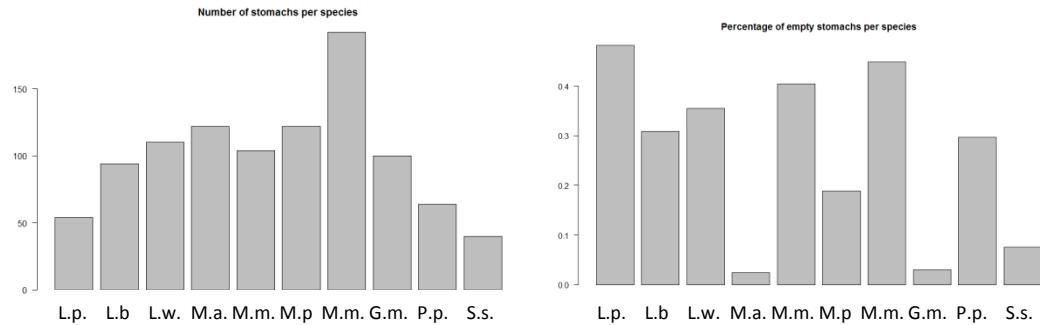


- 10 main commercial species



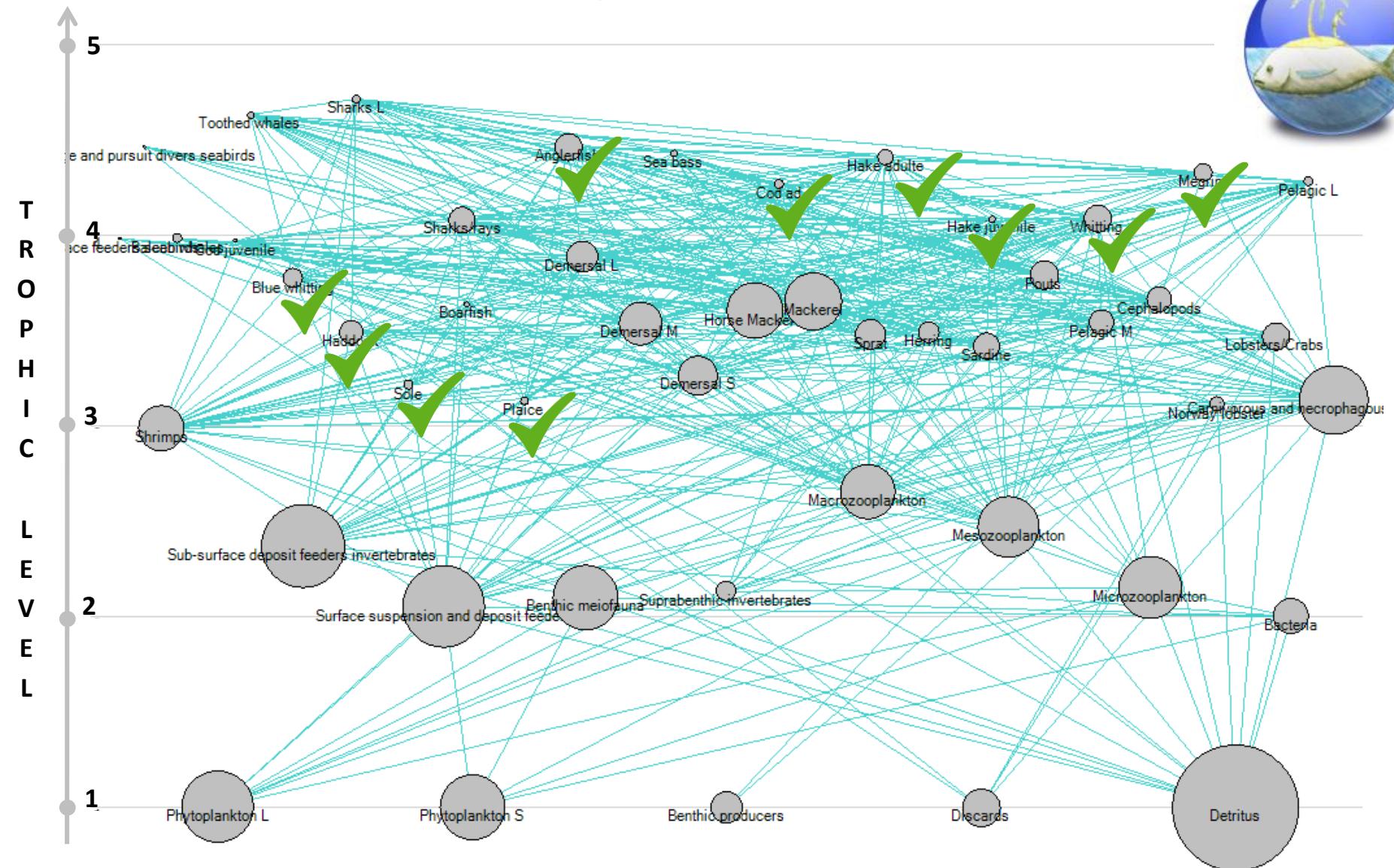
- > 1000 stomachs analyzed

75 % of non empty



- Identification as precise as possible of taxa - 206 different taxa
- Occurrence, numerical, ~~gravimetric, volumetric~~ ()

STOMACH CONTENT ANALYSIS



STOMACH CONTENT ANALYSIS



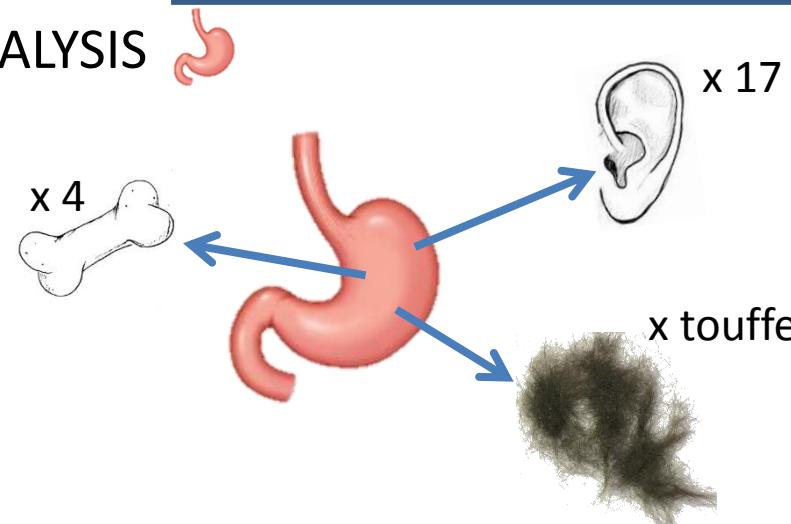
detail

Consumer_id	Prey_id	Nb_items	Finest taxon																					
Nom.latin	code_ind	Mass	Mass	No	Pa	Code_Proie	Prey_Ph	Prey_S	Prey_St	Prey_Cl	Prey_Sut	Prey_Si	Prey_Oi	Prey_Su	Prey_In	Prey_Si	Prey_F	Prey_G	Prey_Nb	Remarks	Prey_Taxon	Prey_T		
Gadus morhua	GM-Z1-1	25.11	19.77	5.34	1	0	GM-Z1-1-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	NA	NA	NA	NA	NA	NA	NA	1	1 paire d'yeu	Eumalacostri		
Gadus morhua	GM-Z1-1	25.11	19.77	5.34	1	0	GM-Z1-1-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1	1 paire d'yeu	Brachyura		
Gadus morhua	GM-Z1-10	30.37	18.63	11.74	1	0	GM-Z1-10-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	NA	NA	NA	NA	NA	NA	3	3 paires d'yeu	Eucarida		
Gadus morhua	GM-Z1-10	30.37	18.63	11.74	1	0	GM-Z1-10-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1	carapace san	Brachyura		
Gadus morhua	GM-Z1-10	30.37	18.63	11.74	1	0	GM-Z1-10-3	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1		Liocarcinus		
Gadus morhua	GM-Z1-11	38.13	14.68	23.45	1	0	GM-Z1-11-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	Portun	Polybii	Liocarc	NA				
Gadus morhua	GM-Z1-11	38.13	14.68	23.45	1	0	GM-Z1-11-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Caridea	Pandal	Pandal	NA	3	3 queues, 3 t	Pandalidae		
Gadus morhua	GM-Z1-11	38.13	14.68	23.45	1	0	GM-Z1-11-3	Echinode	Astero	NA	Ophiuro	NA	NA	Ophiuri	Ophiurir	Chiloph	NA	Ophiur	Ophiur	NA	1	1 disque san	Ophiura	
Gadus morhua	GM-Z1-11	38.13	14.68	23.45	1	0	GM-Z1-11-4	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Anomur	NA	NA	NA	2	3 P1 (2 gauch	Anomura		
Gadus morhua	GM-Z1-12	21.7	16.02	5.68	1	1	GM-Z1-12-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Caridea	Proces	Proces	NA	1	rostre tronq	Processidae		
Gadus morhua	GM-Z1-13	81.61	56.56	25.05	1	1	GM-Z1-13-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1	périopodes	Brachyura		
Gadus morhua	GM-Z1-13	81.61	56.56	25.05	1	1	GM-Z1-13-2	Chordata	Verteb	Gnathio	Actinop	NA	NA	NA	NA	NA	NA	NA	NA	1	chair, arêtes	Actinopteri		
Gadus morhua	GM-Z1-14	26.89	24.49	2.4	0	1	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA			
Gadus morhua	GM-Z1-15	22.52	20.14	2.38	1	1	GM-Z1-15-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Caridea	Crango	Crango	Crango	1	entièr	Crangon all		
Gadus morhua	GM-Z1-15	22.52	20.14	2.38	1	1	GM-Z1-15-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Caridea	Crango	NA	NA	1	1 pince subc	Crangonoide		
Gadus morhua	GM-Z1-16	20.66	15.5	5.16	1	0	GM-Z1-16-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	21	5 têtes sans	Eumalacostri		
Gadus morhua	GM-Z1-16	20.66	15.5	5.16	1	0	GM-Z1-16-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1	1 œil, 1 pinc	Brachyura		
Gadus morhua	GM-Z1-17	67.62	56.88	10.74	1	1	GM-Z1-17-1	Mollusca	NA	NA	Bivalvia	Pteriomc	NA	Pectino	NA	Pectinc	Pectini	Palliol	Pallio	2	1 bivalve, 1 v	Palliolium tig		
Gadus morhua	GM-Z1-17	67.62	56.88	10.74	1	1	GM-Z1-17-2	Chordata	Verteb	Gnathio	Actinop	NA	NA	NA	NA	NA	NA	NA	NA	1	arc branchial	Actinopteri		
Gadus morhua	GM-Z1-18	26.97	15.83	11.14	1	0	GM-Z1-18-1	Mollusca	NA	NA	Bivalvia	Pteriomc	NA	Pectino	NA	Pectinc	Pectini	Palliol	Pallio	1	1 bivalve	Palliolium tig		
Gadus morhua	GM-Z1-18	26.97	15.83	11.14	1	0	GM-Z1-18-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	NA	NA	NA	NA	NA	NA	8	8 paires d'yeu	Eucarida		
Gadus morhua	GM-Z1-18	26.97	15.83	11.14	1	0	GM-Z1-18-3	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1	1 paire d'yeu	Brachyura		
Gadus morhua	GM-Z1-18	26.97	15.83	11.14	1	0	GM-Z1-18-4	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	Portun	Polybii	Liocarc	2	2 pinces, 1 co	Liocarcinus v		
Gadus morhua	GM-Z1-18	26.97	15.83	11.14	1	0	GM-Z1-18-5	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	Xanthic	Xanthic	NA	1	1 corps enti	Xanthidae		
Gadus morhua	GM-Z1-18	26.97	15.83	11.14	1	0	GM-Z1-18-6	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	Cancroi	Atelecy	Atelecy	1	1 corps enti	Atelecyclus i		
Gadus morhua	GM-Z1-19	34.65	17.63	17.02	1	0	GM-Z1-19-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	Portun	Polybii	Macrac	1	entier	Macropipus i		
Gadus morhua	GM-Z1-19	34.65	17.63	17.02	1	0	GM-Z1-19-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	5	1 chair carap	Brachyura		
Gadus morhua	GM-Z1-19	34.65	17.63	17.02	1	0	GM-Z1-19-3	Mollusca	NA	NA	Bivalvia	Heterodc	NA	Cardito	NA	NA	Crassat	Astarti	Astarti	1	entièr	Astarte sulca		
Gadus morhua	GM-Z1-19	34.65	17.63	17.02	1	0	GM-Z1-19-4	Echinode	Echino	NA	Echinoic	NA	NA	NA	NA	NA	NA	NA	NA	1	fragments di	Echinoidea		
Gadus morhua	GM-Z1-2	31.58	24.28	7.3	0	0	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA			
Gadus morhua	GM-Z1-20	10.58	8.4	2.18	1	0	GM-Z1-20-1	Annelida	NA	NA	Polychaet	NA	NA	NA	NA	NA	NA	NA	NA	NA	1	tube membr	Polychaeta	
Gadus morhua	GM-Z1-20	10.58	8.4	2.18	1	0	GM-Z1-20-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Caridea	Pandal	Pandal	NA	1	céphalo rost	Pandalidae		
Gadus morhua	GM-Z1-20	10.58	8.4	2.18	1	0	GM-Z1-20-3	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	1	tête sans ros	Anomura		
Gadus morhua	GM-Z1-20	10.58	8.4	2.18	1	0	GM-Z1-20-4	Mollusca	NA	NA	Gastrop	Caenoga	NA	Littorini	NA	Naticoi	Naticid	NA	1	columelle tr	Naticidae			
Gadus morhua	GM-Z1-21	15.27	10.6	4.67	1	0	GM-Z1-21-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	NA	NA	NA	NA	NA	NA	NA	3	3 têtes	Eumalacostri		
Gadus morhua	GM-Z1-21	15.27	10.6	4.67	1	0	GM-Z1-21-2	Mollusca	NA	NA	Bivalvia	Pteriomc	NA	Pectino	NA	Pectinc	Pectini	Palliol	Pallio	1	1 bivalve	Palliolum tig		
Gadus morhua	GM-Z1-21	15.27	10.6	4.67	1	0	GM-Z1-21-3	Chordata	Verteb	Gnathio	Actinop	NA	NA	NA	NA	NA	NA	NA	NA	1	otolithes tr	Actinopteri		
Gadus morhua	GM-Z1-22	46.97	27.47	19.5	1	0	GM-Z1-22-1	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	Leucos	Leucos	Ebalia	1	carapace ave	Ebalia tuberc		
Gadus morhua	GM-Z1-22	46.97	27.47	19.5	1	0	GM-Z1-22-2	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Anomur	Paguro	Paguric	NA	1	corps avec	Paguridae		
Gadus morhua	GM-Z1-22	46.97	27.47	19.5	1	0	GM-Z1-22-3	Arthropo	Crustac	Multicr	Malacos	Eumalacc	Eucarid	Decapo	Pleocym	Brachyu	NA	NA	NA	2	1 pince	Brachyura		



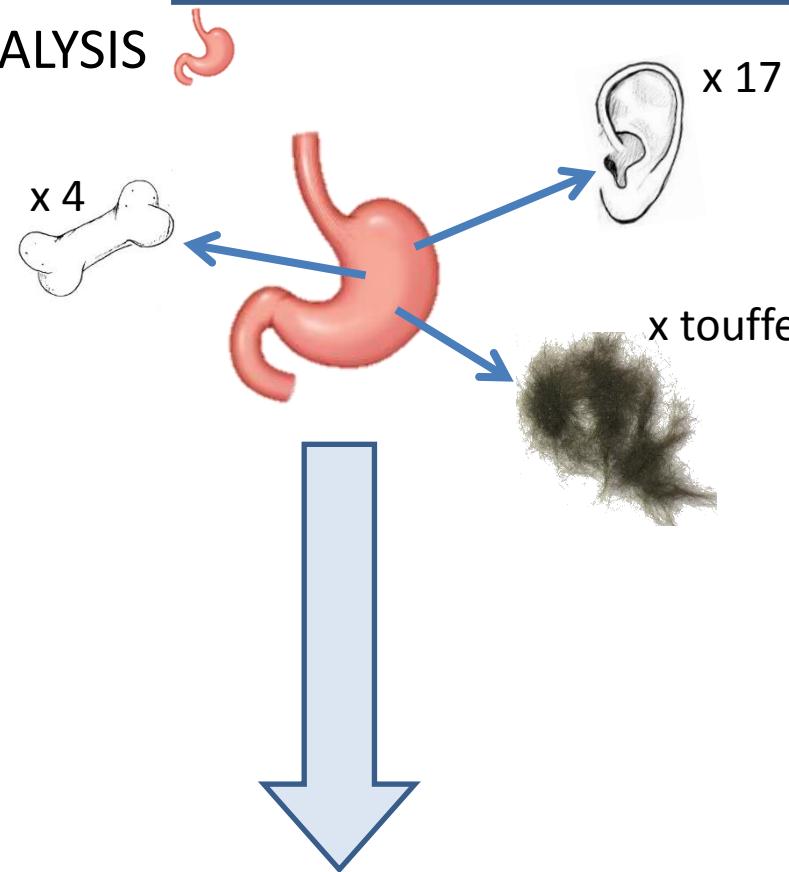
STOMACH CONTENT ANALYSIS

Number of prey
items per stomach

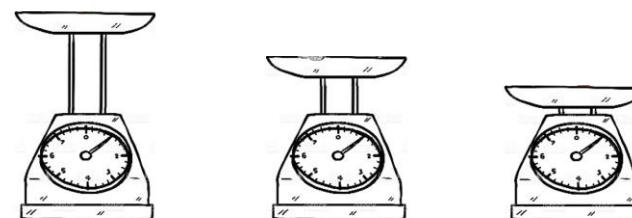


STOMACH CONTENT ANALYSIS

Number of prey items per stomach



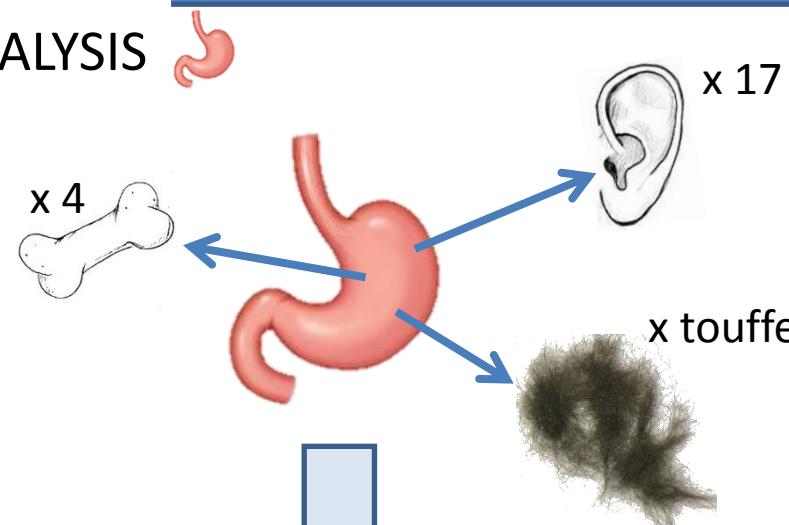
Eaten quantity of each prey type



STOMACH CONTENT ANALYSIS

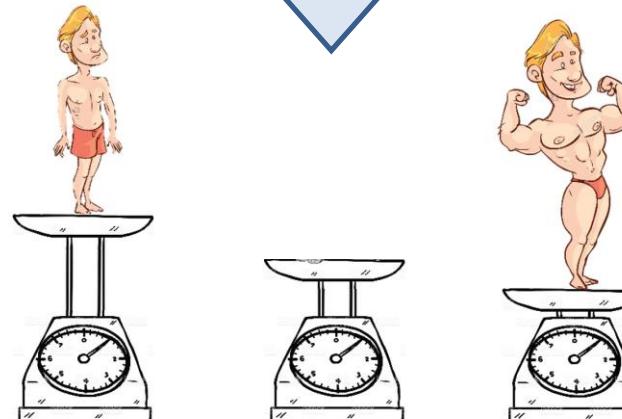
Number of prey items per stomach

- Number → Weight?
- Question of digestibility



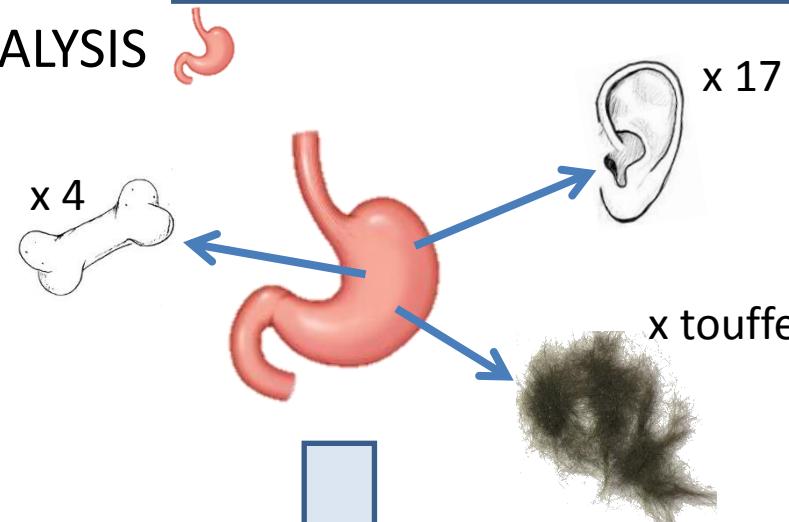
PREY ITEMS = ESTIMATORS
OF MATTER FLOWS?

Eaten quantity of each prey type



STOMACH CONTENT ANALYSIS

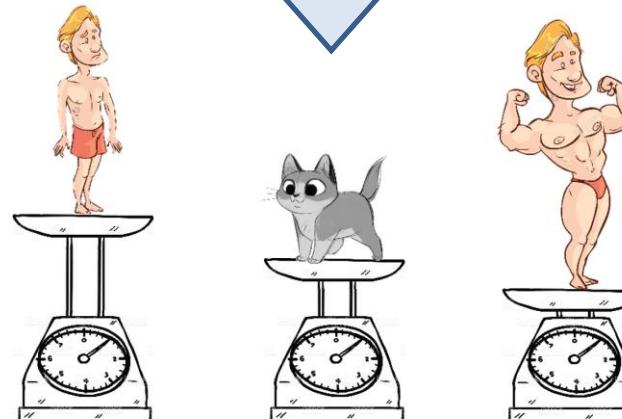
Number of prey items per stomach



- Number → Weight?
- Question of digestibility
- Compare species...

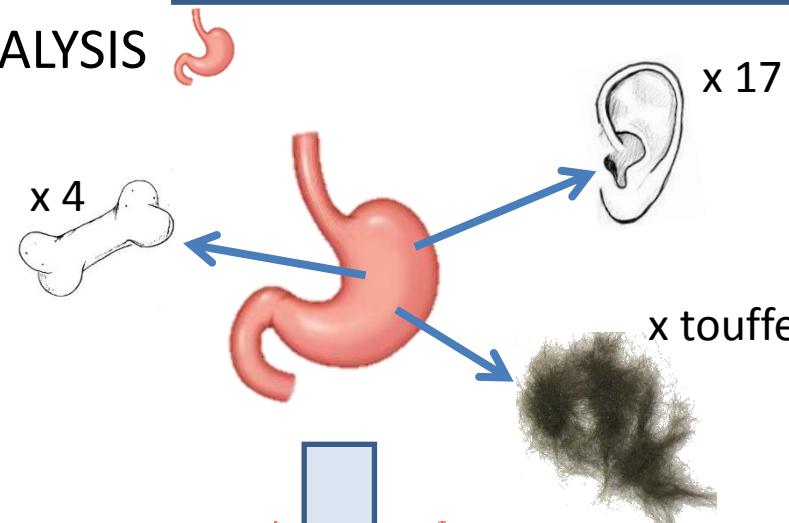
PREY ITEMS = ESTIMATORS
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Eaten quantity of each prey type



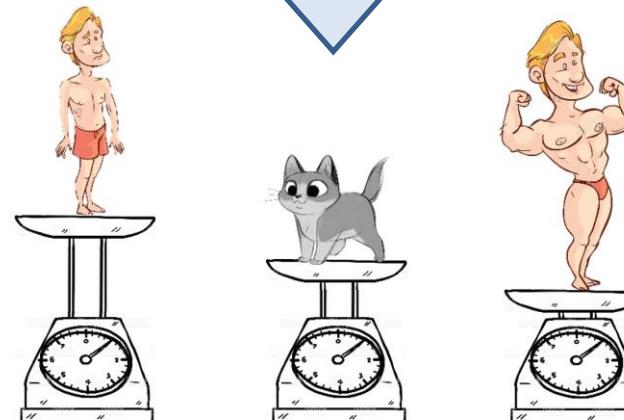
STOMACH CONTENT ANALYSIS

Number of prey items per stomach



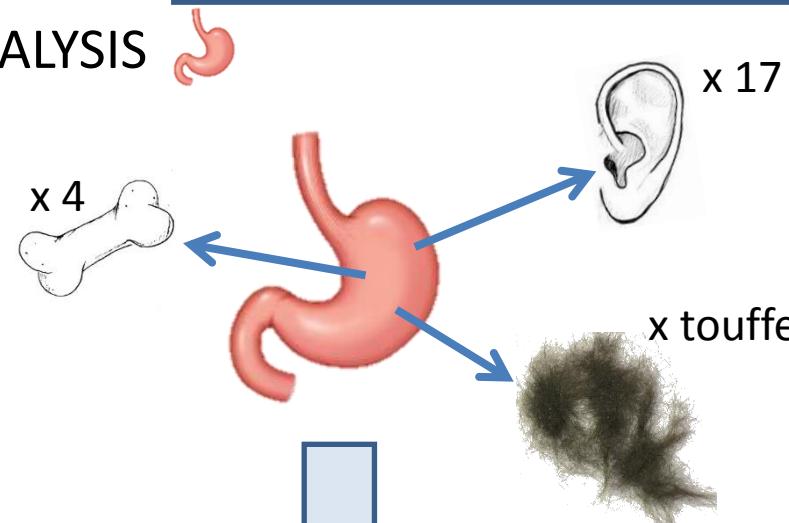
PREY ITEMS ~~ESTIMATORS OF MATTER FLOWS?~~

Eaten quantity of each prey type

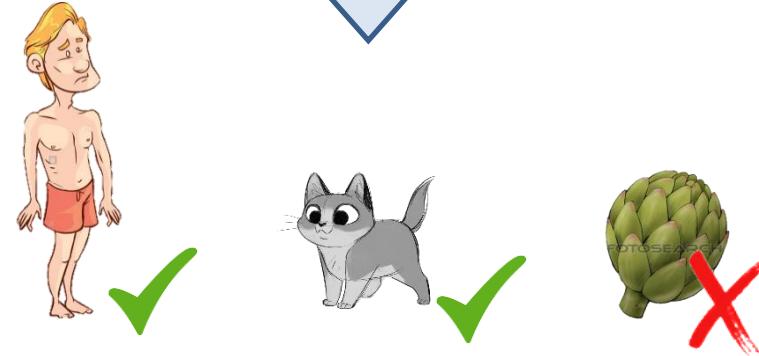


STOMACH CONTENT ANALYSIS

Number of prey items per stomach

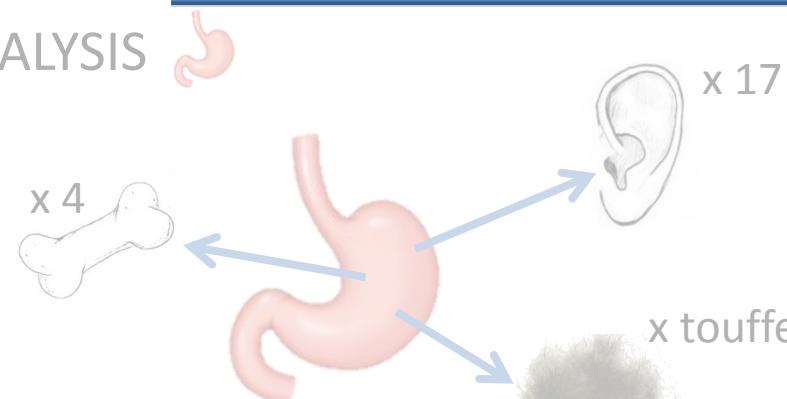


Frequencies of observation



STOMACH CONTENT ANALYSIS

Number of prey
items per stomach



**SCA will give information on the
links to be considered in the diet
matrix**

Frequencies of
observation



STABLE ISOTOPES ANALYSIS



- ~ all functionnal groups in the ecosystem (the 10 commercial species and their potential preys)



- > 1000 samples
- Both carbon and nitrogen stable isotopes

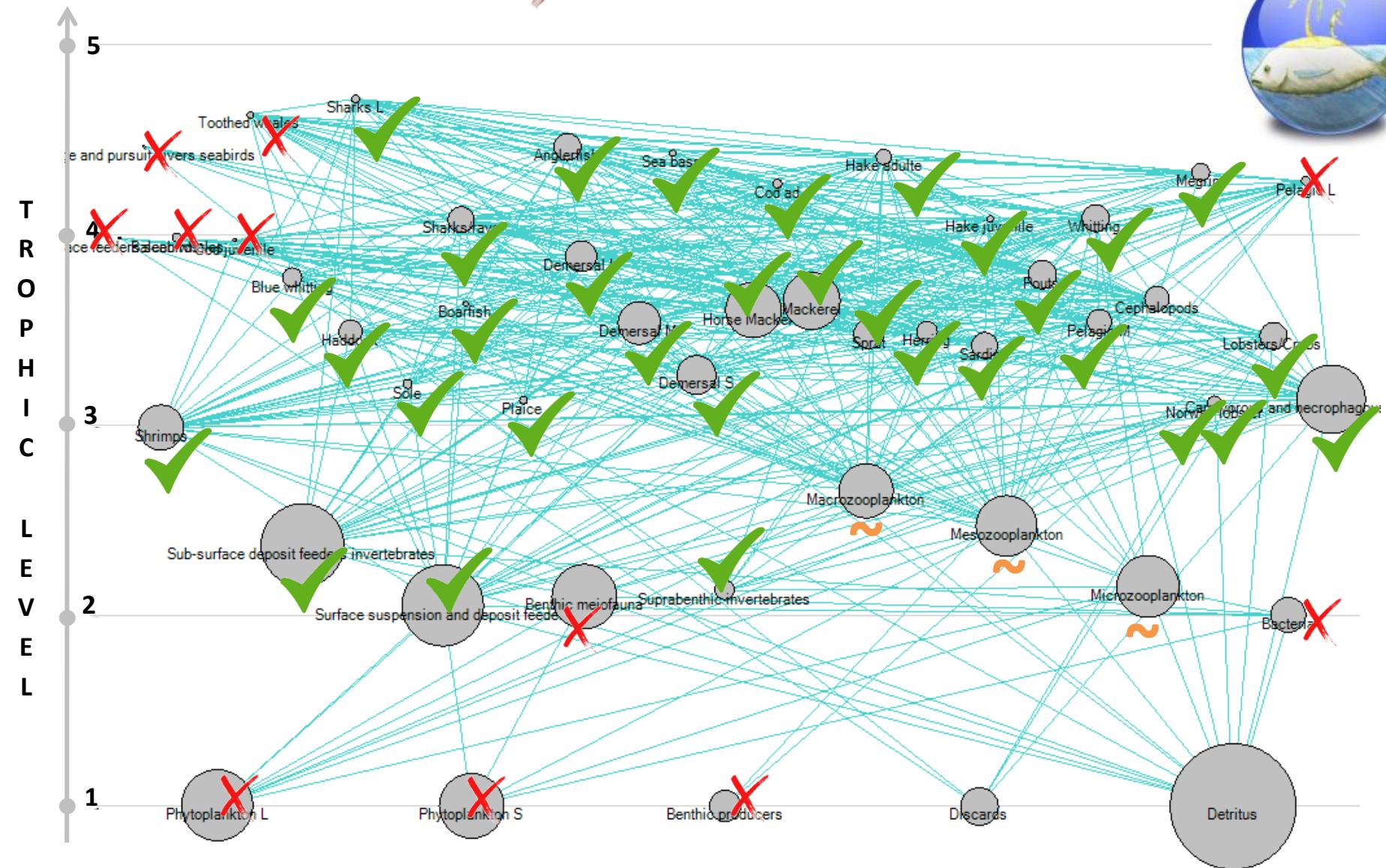
 $\delta^{13}\text{C}$

Identification of the
primary source

 $\delta^{15}\text{N}$

Identification of the
trophic position

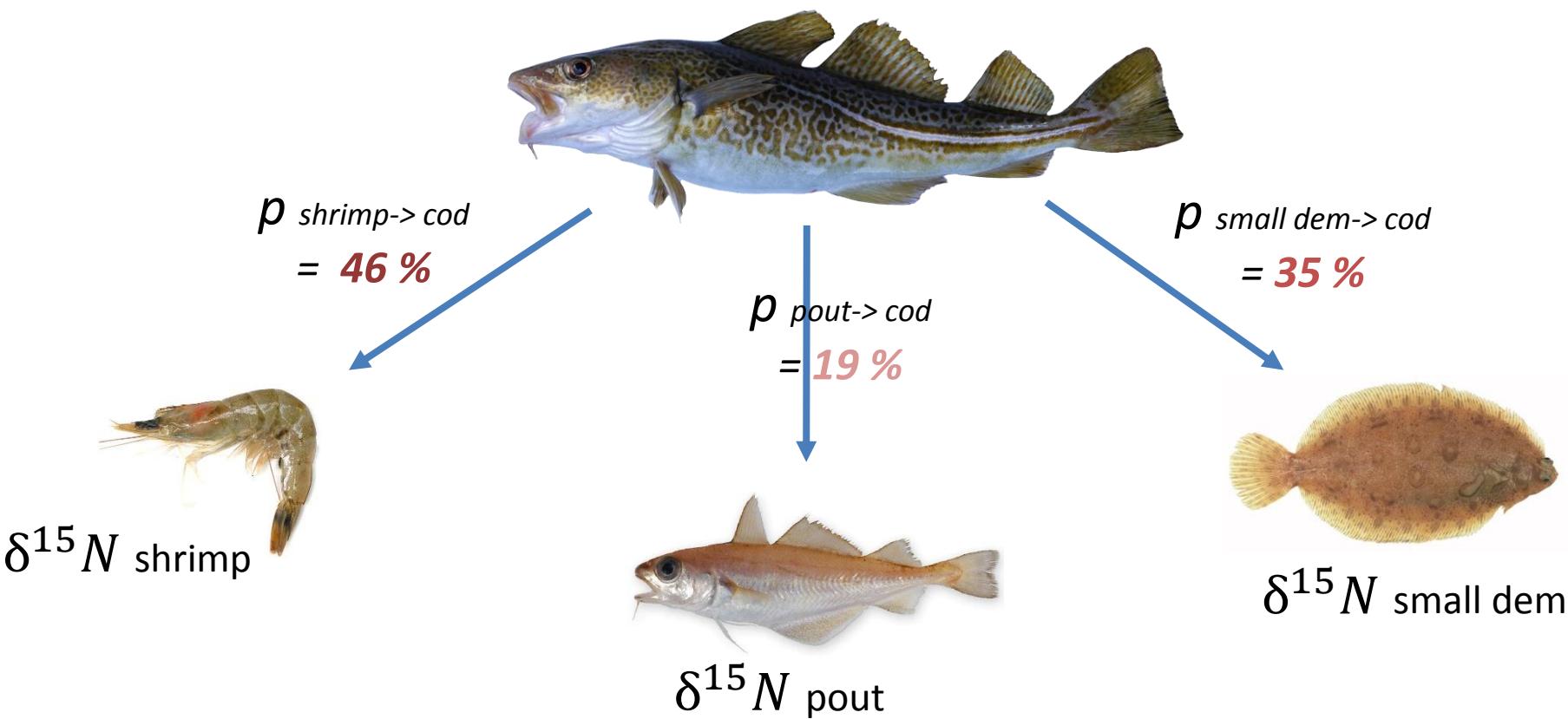
STABLE ISOTOPES ANALYSIS



TROPHIC ENRICHMENT CONCEPT

« You are what you eat »

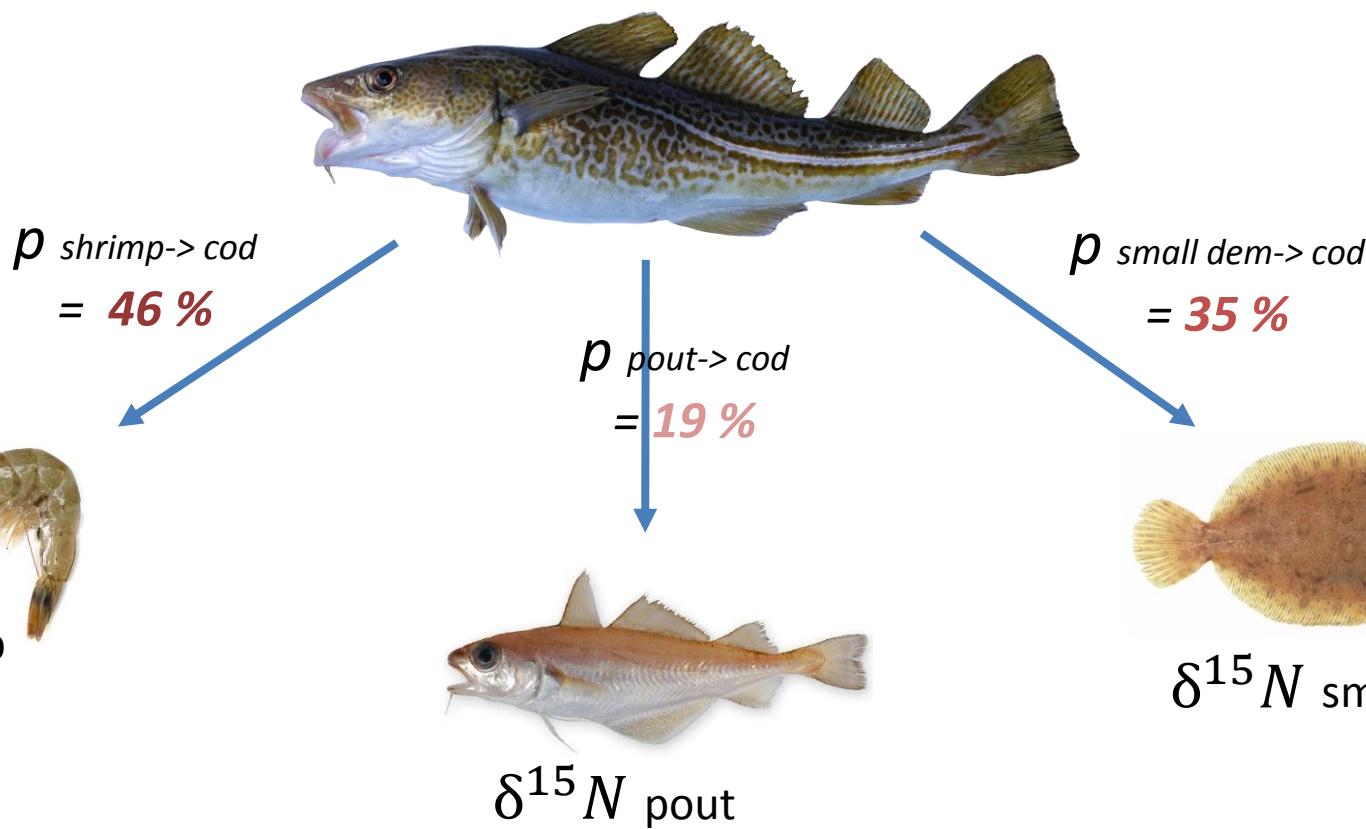
Fry 2006 etc



TROPHIC ENRICHMENT CONCEPT



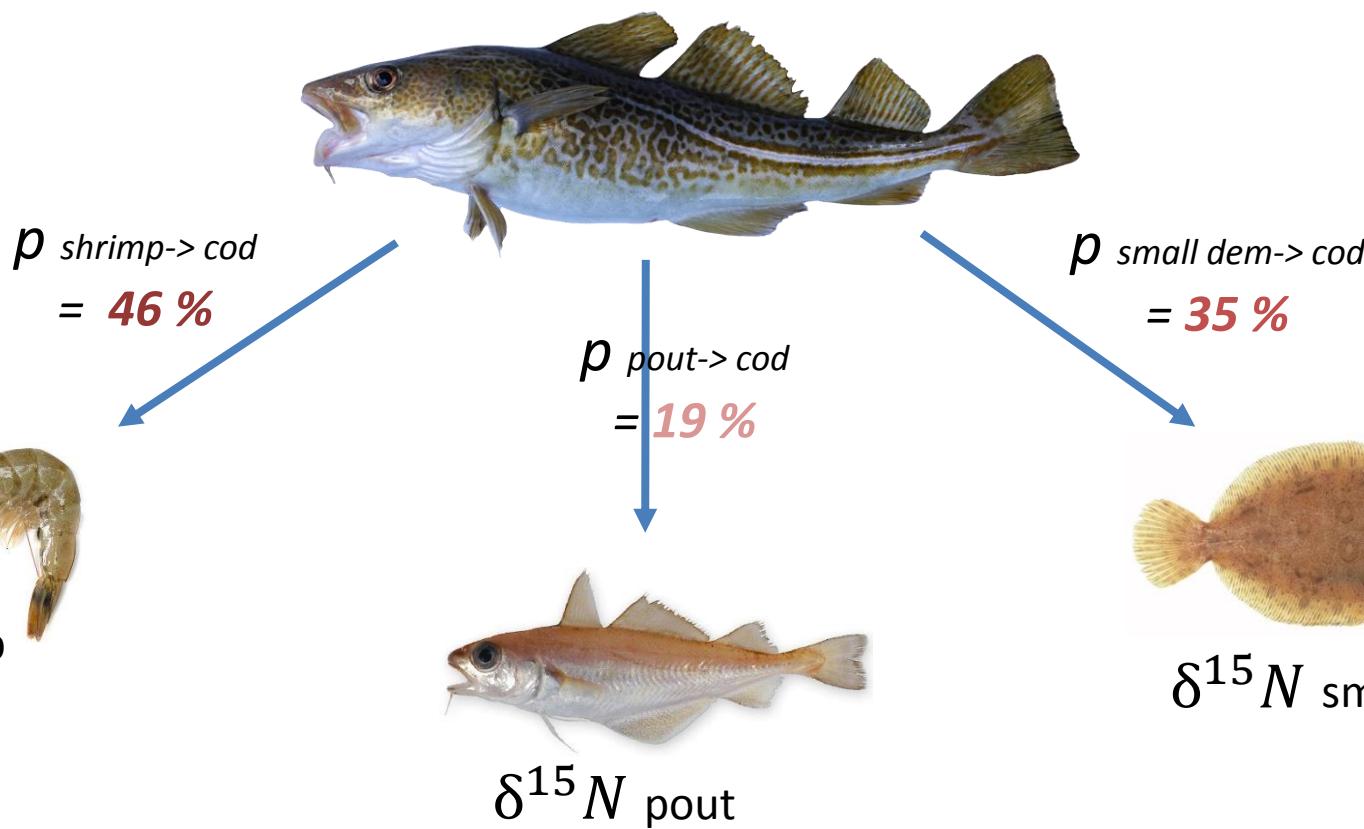
$$\delta^{15}N_{cod} = \frac{0.46 * q_{N,shrimp} * (\delta^{15}N_{shrimp} + \Delta) + 0.19 * q_{N,pout} * (\delta^{15}N_{pout} + \Delta) + 0.35 * q_{N,sm\ dem} * (\delta^{15}N_{small\ dem} + \Delta)}{0.46 * q_{N,shrimp} + 0.19 * q_{N,pout} + 0.35 * q_{N,sm\ dem}}$$



TROPHIC ENRICHMENT CONCEPT



$$\delta^{15}N_{\text{cod}} \sim 0.46 * (\delta^{15}N_{\text{shrimp}} + \Delta) + 0.19 * (\delta^{15}N_{\text{pout}} + \Delta) + 0.35 * (\delta^{15}N_{\text{small dem}} + \Delta)$$

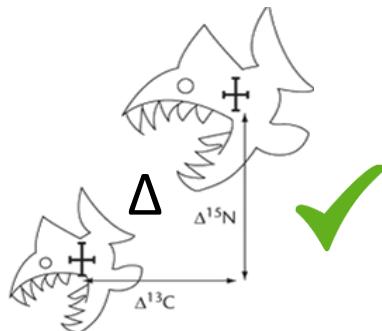


TROPHIC ENRICHMENT CONCEPT



→ ISOTOPIC SIGNATURE of CONSUMER and SOURCES + TROPHIC ENRICHMENT

= DETERMINATION OF THE DIET



$p_{shrimp \rightarrow cod}$

= ? %

$p_{small\ dem \rightarrow cod}$

= ? %

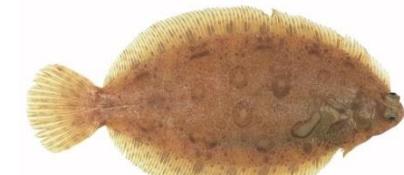


$\delta^{15}N_{shrimp}$

A green checkmark is located to the right of the shrimp.

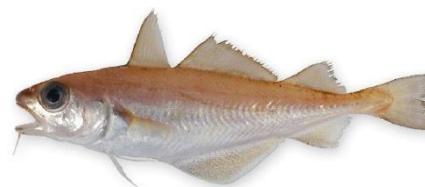
$p_{pout \rightarrow cod}$

= ? %



$\delta^{15}N_{small\ dem}$

A green checkmark is located to the right of the small demersal fish.

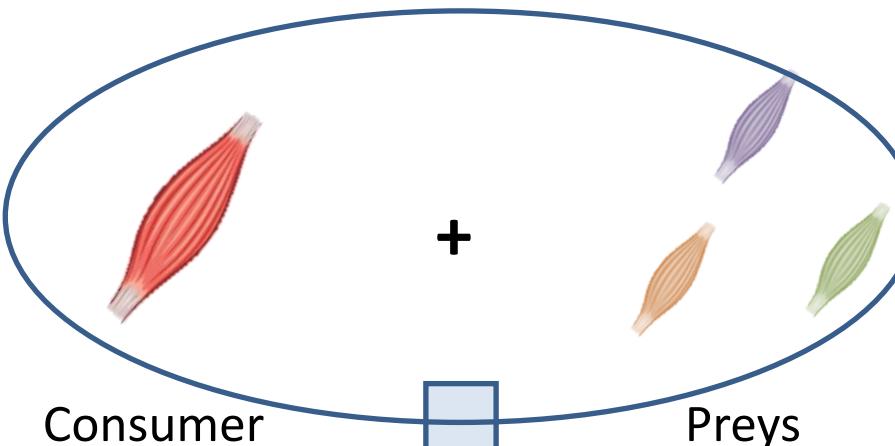


$\delta^{15}N_{pout}$

A green checkmark is located to the right of the pout.

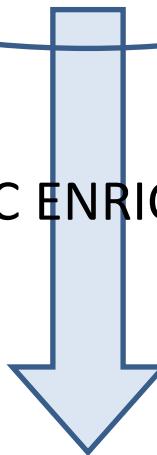
STABLE ISOTOPES ANALYSIS

Stable isotopes concentrations

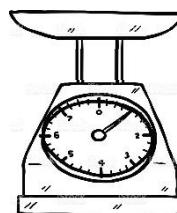
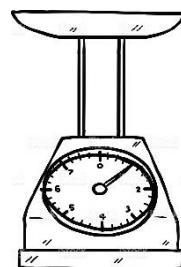


$\delta^{13}\text{C}$
&
 $\delta^{15}\text{N}$

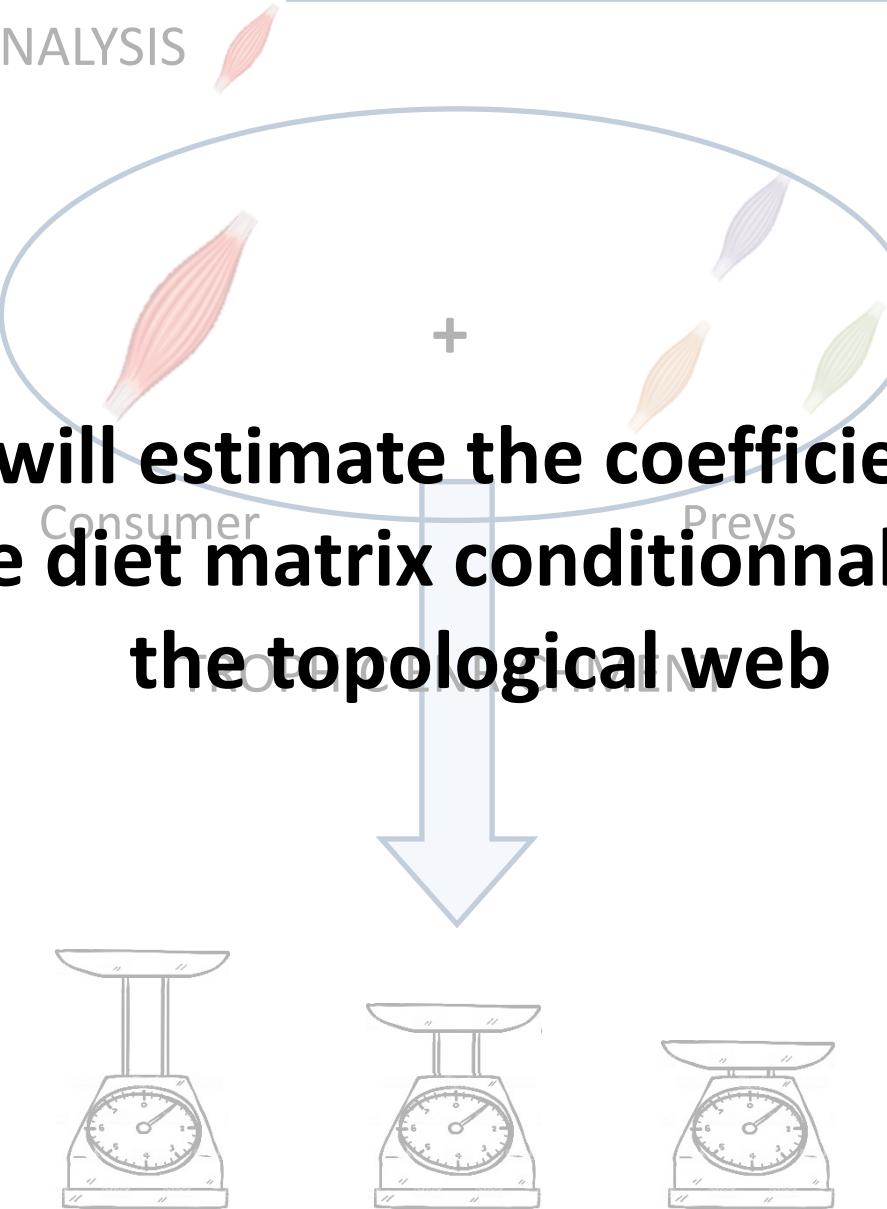
TROPHIC ENRICHMENT



Eaten quantity of
each prey type



STABLE ISOTOPES ANALYSIS

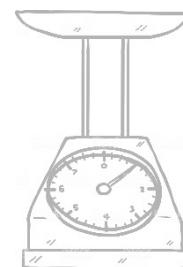


Stable isotopes concentrations

$\delta^{13}\text{C}$
&
 $\delta^{15}\text{N}$

**SIA will estimate the coefficients of
the diet matrix conditionnally to
the topological web**

Eaten quantity of
each prey type



LITERATURE...because it is still useful!



- Does the predator X eat the prey Y?
 - If we miss something!
 - Because of the temporal coverage of our SCA and SIA data
- Are there estimates of the contributions?

Qualitative or Quantitative studies?
- What about data quality?
 - Local vs Foreign
 - Recent vs Old

LITERATURE...because it is still useful!



- Does the predator X eat the prey Y?
 - If we miss something!
 - Because of the temporal coverage of our SCA and SIA data
- Are there estimates of the contributions?

**Literature can inform on both
topological web and contributions**

Qualitative or Quantitative studies?
- What about data quality?
 - Local vs Foreign
 - Recent vs Old

OK, SO KNOW WE CAN START MODELLING: Building EcoDiet

Diversity of data = SIA & SCA
to couple

+

Integrate « a priori » knowledge from the literature

+

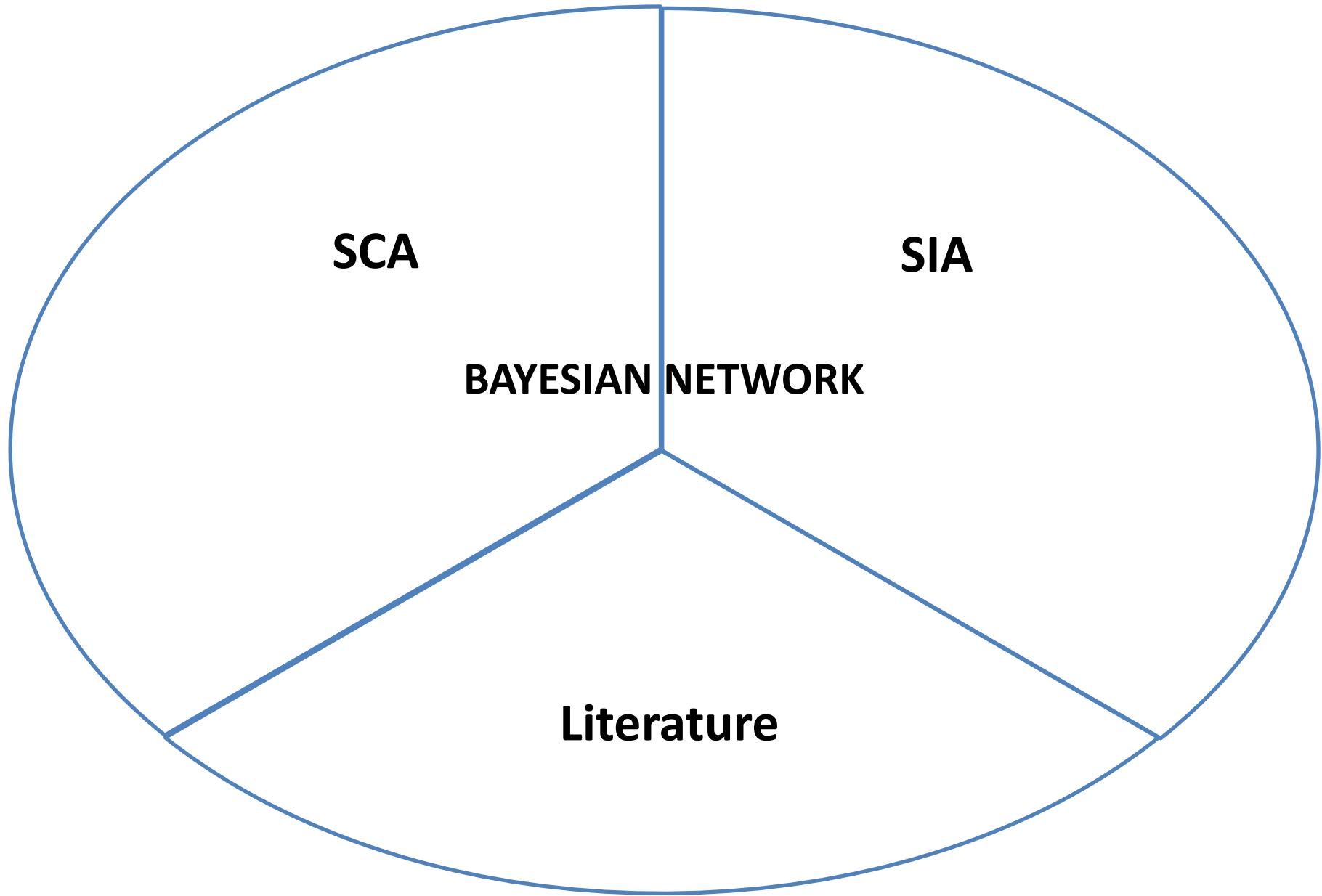
Need of taking into account uncertainty related to diet estimates



INTEGRATED BAYESIAN MODEL

=

EcoDiet



STOMACH CONTENT ANALYSIS



For each consumer: Found vs Not found

b

N stomachs = N « trials »



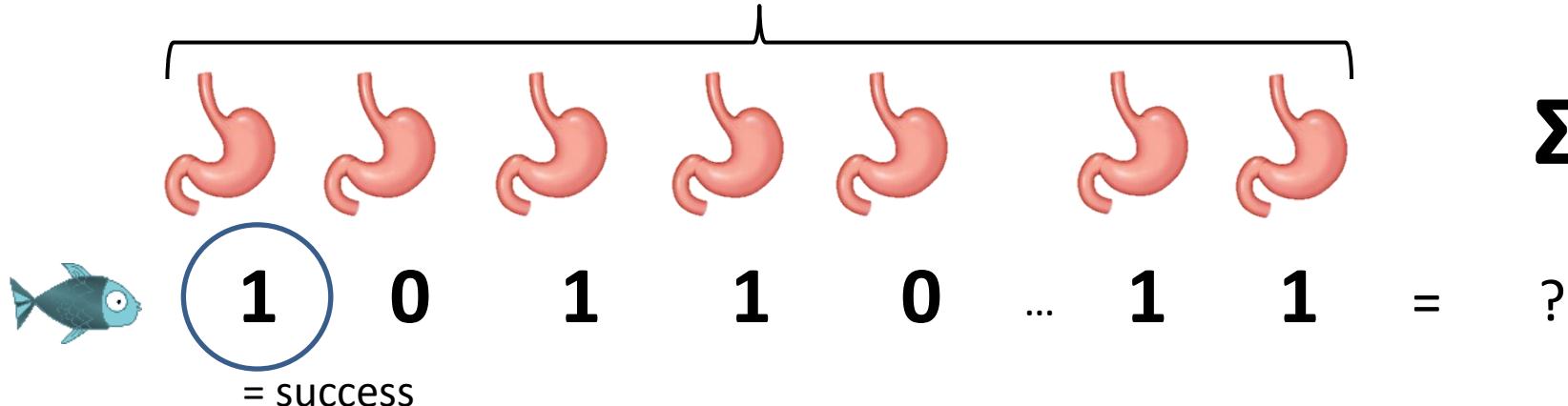
STOMACH CONTENT ANALYSIS



For each consumer: Found vs Not found

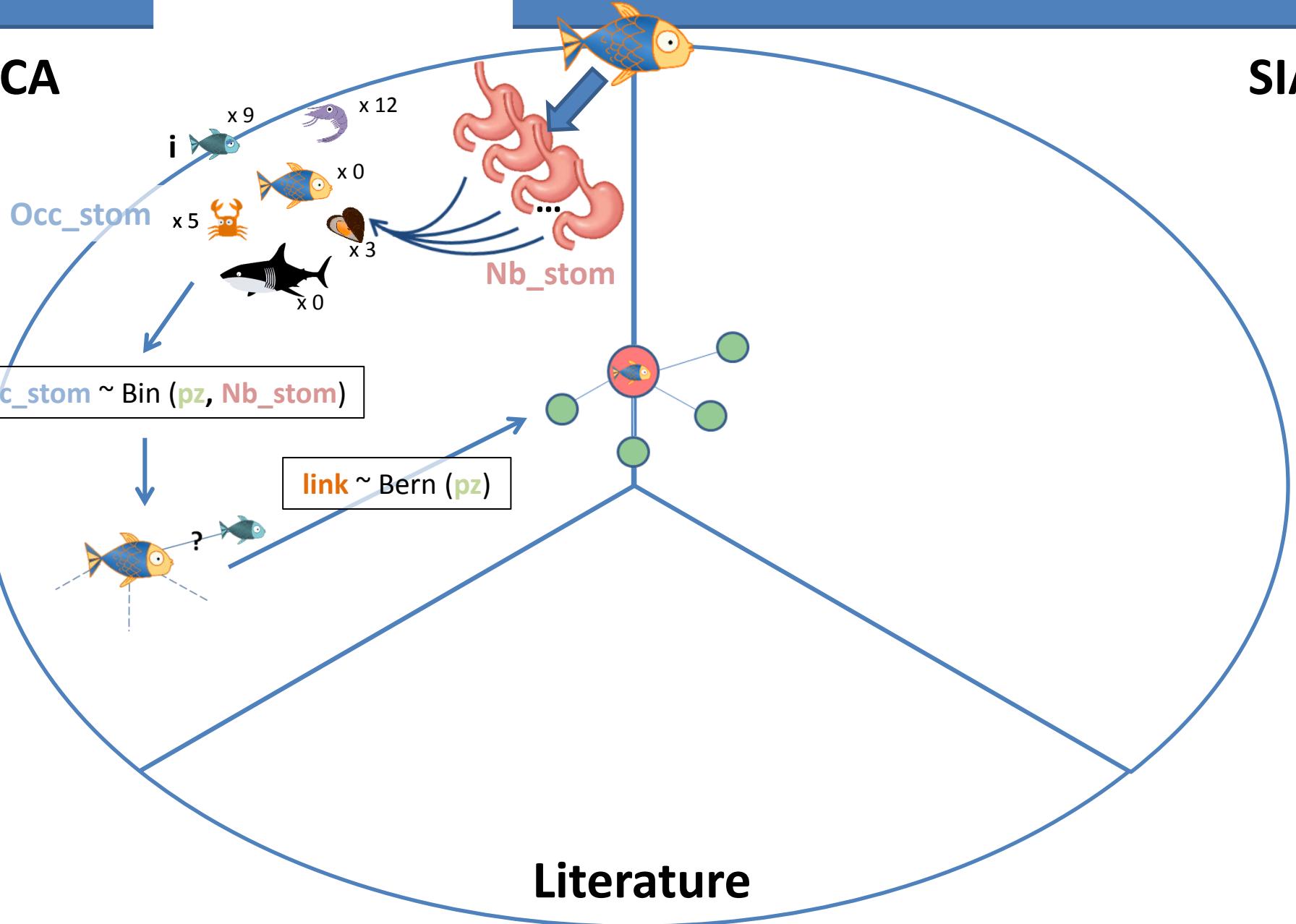
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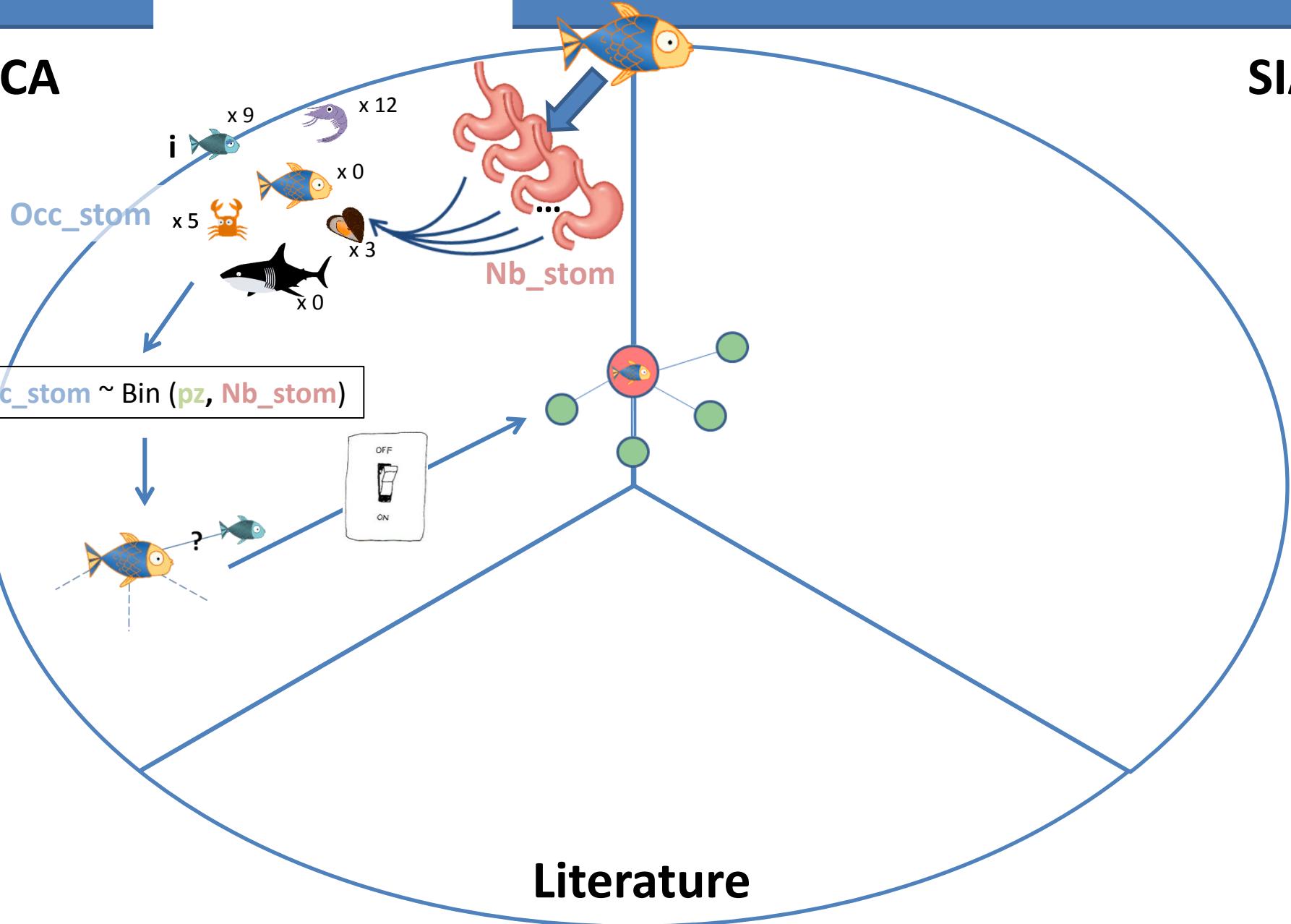
SCA

SIA



SCA

SIA



STABLE ISOTOPES ANALYSIS



- SIA module based on a pre-existing global mixing model

IsoWeb: A Bayesian Isotope Mixing Model for Diet Analysis of the Whole Food Web



Taku Kadoya, Yutaka Osada, Gaku Takimoto

- Main equations

$$X_{ij} \sim N(s_{ij}, \sigma_{ij}^2)$$

$$c_{j\mathbf{k}_i[m]} \sim N(\Lambda_j, \tau_{j\mathbf{k}_i[m]}^2)$$

$$s_{ij} = \frac{\sum_{m=1}^{M_i} p_{i\mathbf{k}_i[m]} Q_{j\mathbf{k}_i[m]} (s_{j\mathbf{k}_i[m]} + c_{j\mathbf{k}_i[m]})}{\sum_{m=1}^{M_i} p_{i\mathbf{k}_i[m]} Q_{j\mathbf{k}_i[m]}}$$

$$p_{i\mathbf{k}_i[1]}, \dots, p_{i\mathbf{k}_i[M_i]} \sim Dirichlet(\alpha_{i1}, \dots, \alpha_{iM_i})$$

Data:

X_{ij} = observed isotope ratio of element j in consumer i ; normally distributed with mean s_{ij} and variance σ_{ij}^2 .

$Q_{j\mathbf{k}_i[m]}$ = observed concentration of element j in resource $\mathbf{k}_i[m]$.

Estimated:

s_{ij} = mean of isotope ratio of element j in consumer i .

σ_{ij}^2 = residual variances of isotope ratio of element j in consumer i .

$c_{j\mathbf{k}_i[m]}$ = trophic enrichment factor of element j for the trophic link from resource $\mathbf{k}_i[m]$ to consumer i ($\mathbf{k}_i[m]$ is the m th resources of consumer i).

$p_{i\mathbf{k}_i[m]}$ = dietary proportion of resource $\mathbf{k}_i[m]$ of consumer i .

Priors:

Λ_j and $\tau_{j\mathbf{k}_i[m]}^2$ is the mean and variance of prior distribution
 $\alpha_{i1} = \dots = \alpha_{iM_i}$ are the parameters of a Dirichlet prior

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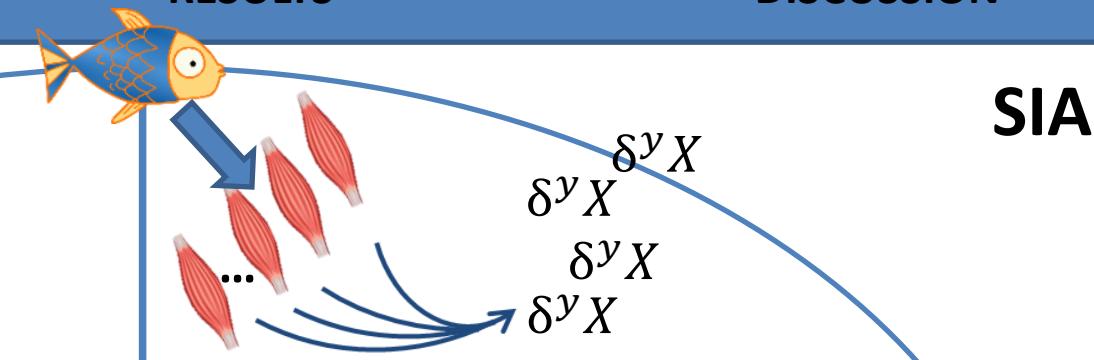
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SCA

SIA



$$\delta^y X \sim \text{Norm}(\mu, \sigma^2)$$

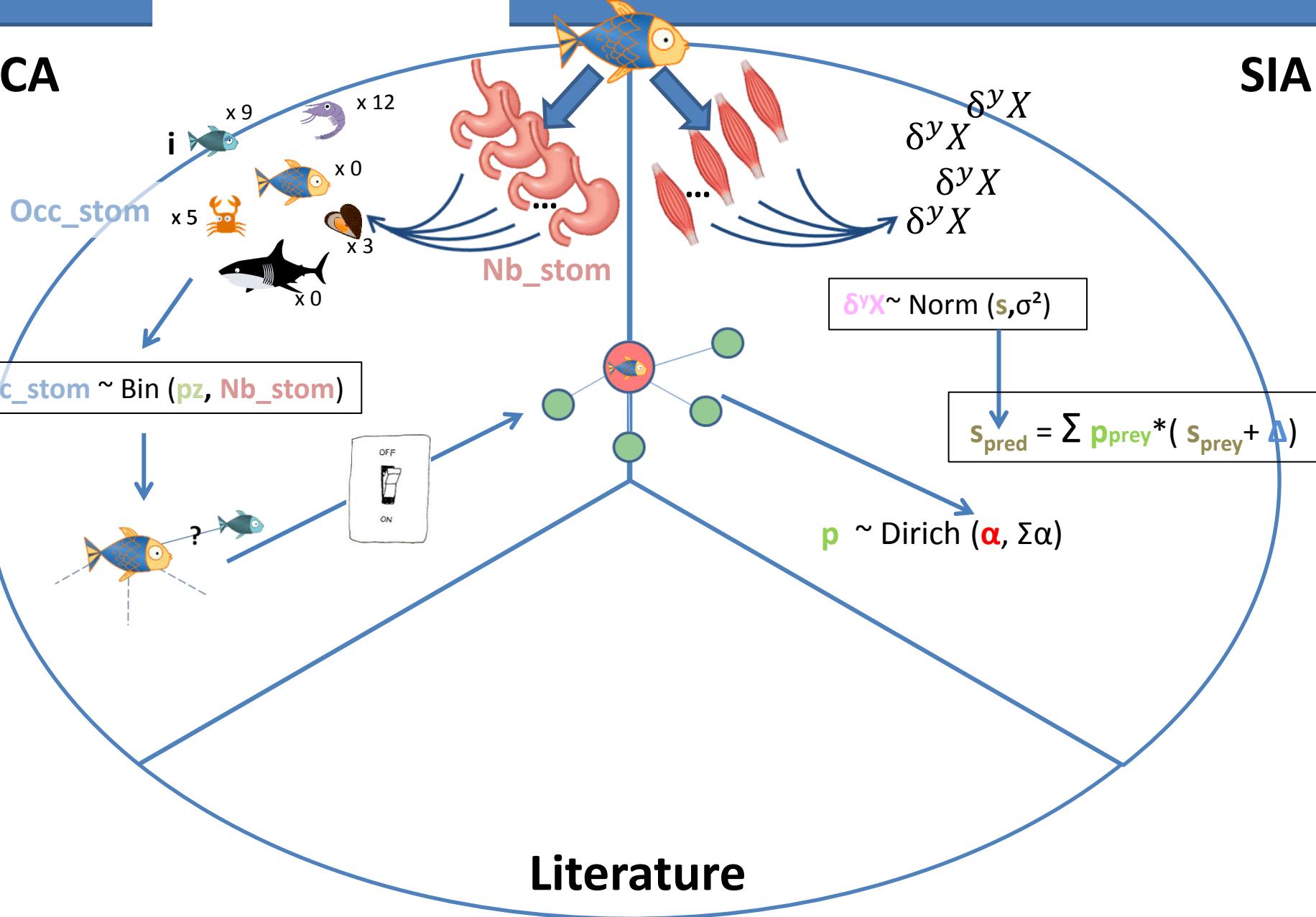
$$s_{\text{pred}} = \sum p_{\text{prey}} * (s_{\text{prey}} + \delta^y X)$$

$$p \sim \text{Dirich}(\alpha, \Sigma \alpha)$$

Literature

SCA

SIA



LITERATURE



- how to assess the quality of the data

- Consistency with EwE approach: using the Pedigree scores
- = A mark ($\in [0;1]$) that decreases when the data becomes more uncertain

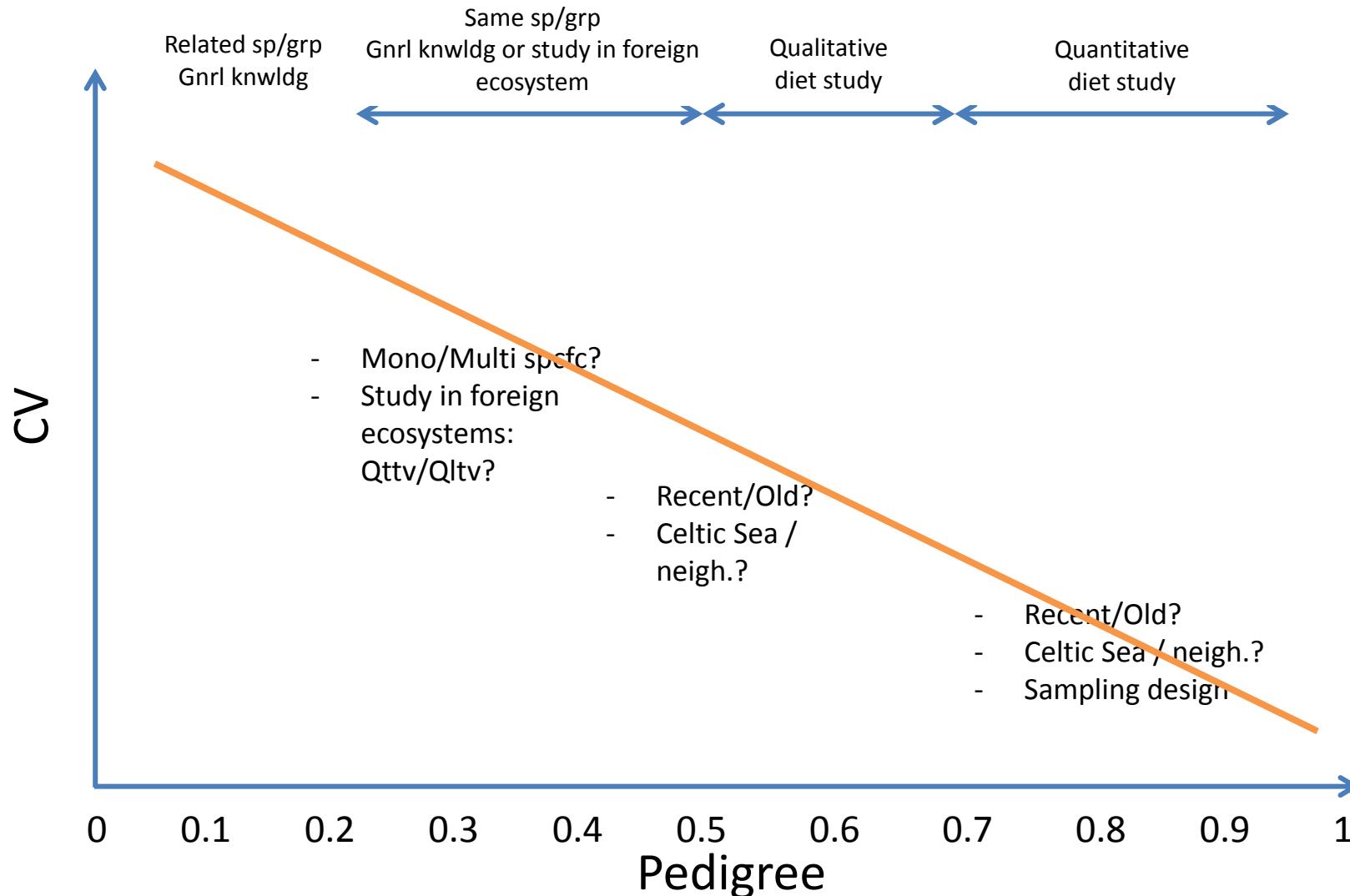
Parameter	Category	Description	Index	Default confidence interval (+/- %)
Biomass (<i>B</i>)	1	Missing parameter (estimated by Ecopath)	0.0	n.a.
	2	From other model	0.0	80
	3	Guesstimates	0.0	80
	4	Approximate or indirect method	0.4	50-80
	5	Sampling based, low precision	0.7	40
	6	Sampling based, high precision	1.0	10
<i>P/B</i> and <i>Q/B</i>	1	Missing parameter (estimated by Ecopath)	0.0	n.a.
	2	Guesstimates	0.1	90
	3	From other model	0.2	80
	4	Empirical relationships	0.5	50
	5	Similar group/species, similar system	0.6	40
	6	Similar group/species, same system	0.7	30
	7	Same group/species, similar system	0.8	20
	8	Same group/species, same system	1.0	10
Diets (<i>DC</i>)	1	General knowledge of related group/species	0.0	80
	2	From other model	0.0	80
	3	General knowledge for same group/species	0.2	80
	4	Qualitative diet composition study	0.5	50
	5	Quantitative but limited diet composition study	0.7	40
	6	Quantitative, detailed, diet composition study	1.0	30
Catches	1	Guesstimates	0.0	>80
	2	From other model	0.0	>80
	3	FAO statistics	0.2	80
	4	National statistics	0.5	50
	5	Local study, low precision/incomplete	0.7	30
	6	Local study, high precision/complete	1.0	10

+ C.V.

LITERATURE



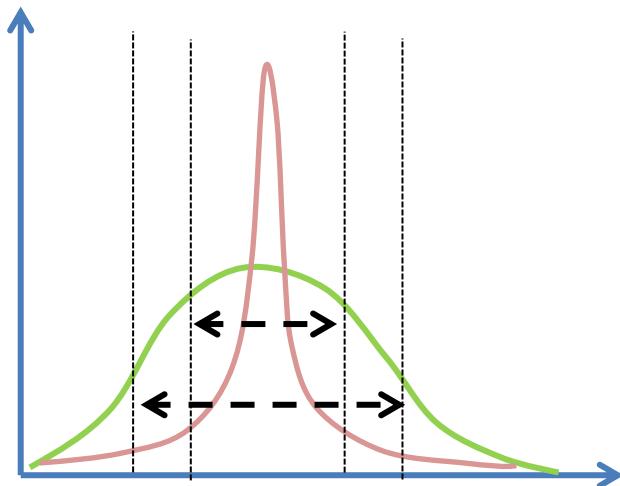
- Pedigree → a target C.V. by consumer



LITERATURE



- how to assess the quality of the data



non informative prior :

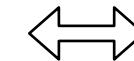
$$pz[i,j] \sim \text{Beta}(1,1)$$

\sum

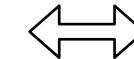


Weight Lit vs Stom:

Pedigree = 1



=



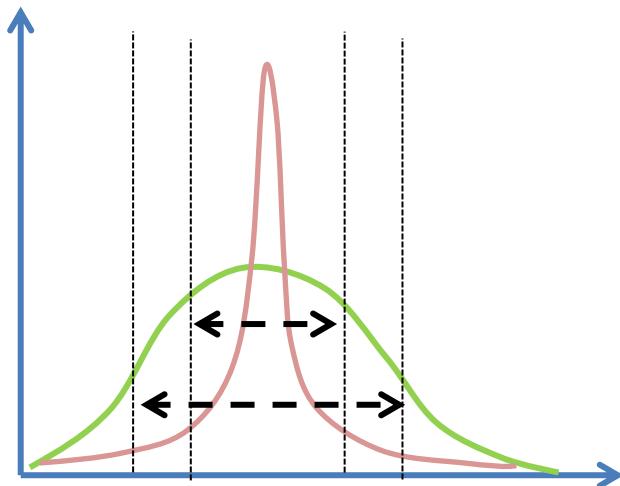
$\sum \sim \text{Sampling size of}$



LITERATURE



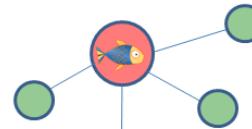
- how to assess the quality of the data



informative prior :

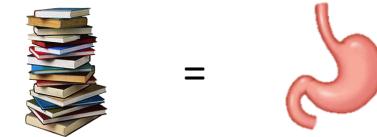
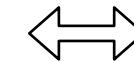
$$pz[i,j] \sim \text{Beta}(x,y)$$

(underbrace symbol)
 \sum



Weight Lit vs Stom:

Pedigree = 1

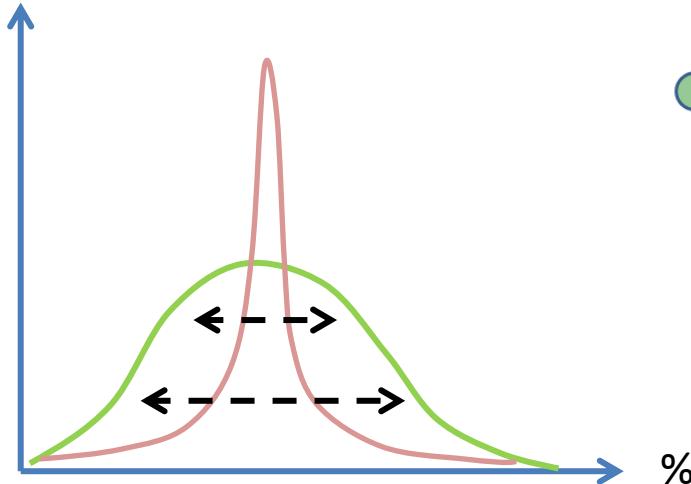


$\sum \sim \text{Sampling size of}$

LITERATURE

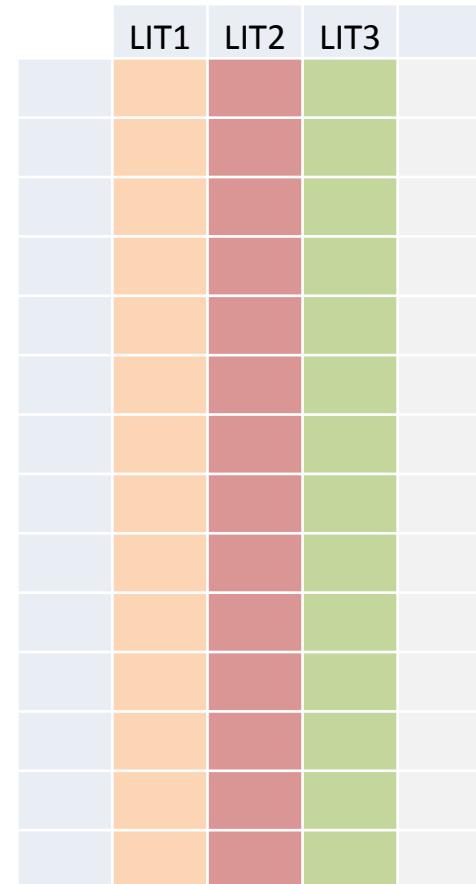
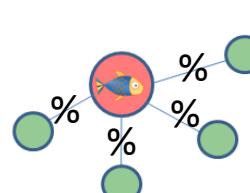


- how to assess the quality of the data



non informative prior :

$$p_{[i,j]} \sim \text{Beta}(p_{\text{lit}}, 1)$$

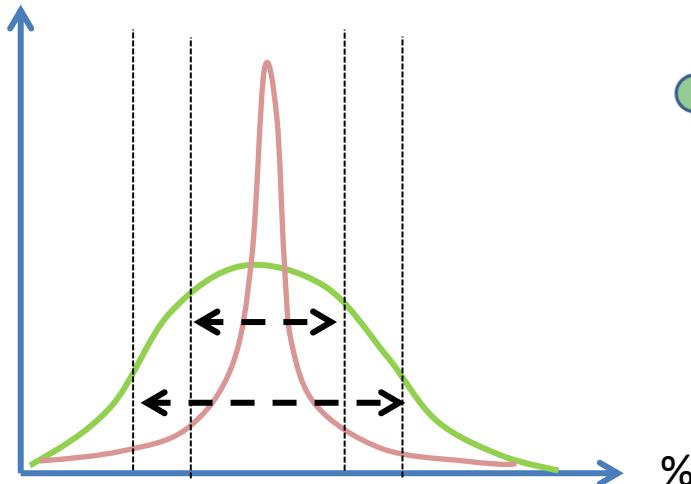


\sum = determined by $CV_{pedigree}$

LITERATURE

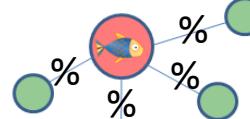


- how to assess the quality of the data



informative prior :

$$p_{[i,j]} \sim \text{Dirich}(\alpha_{[i,j]}, \sum \alpha_{[i,.]})$$



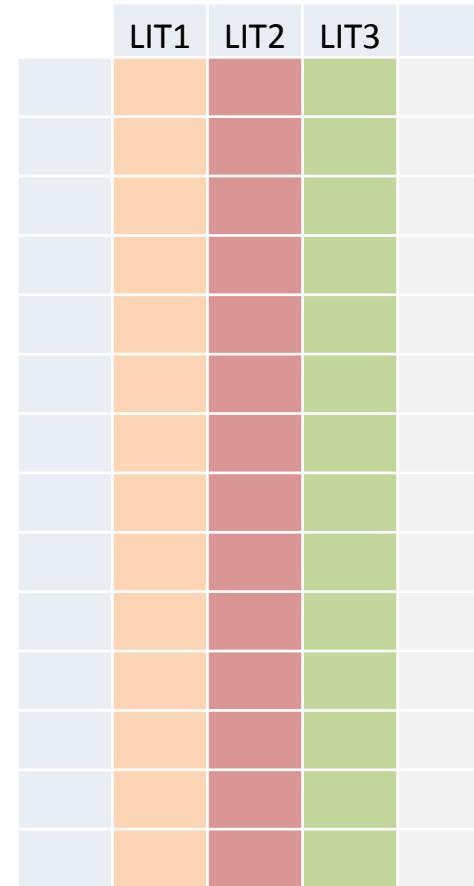
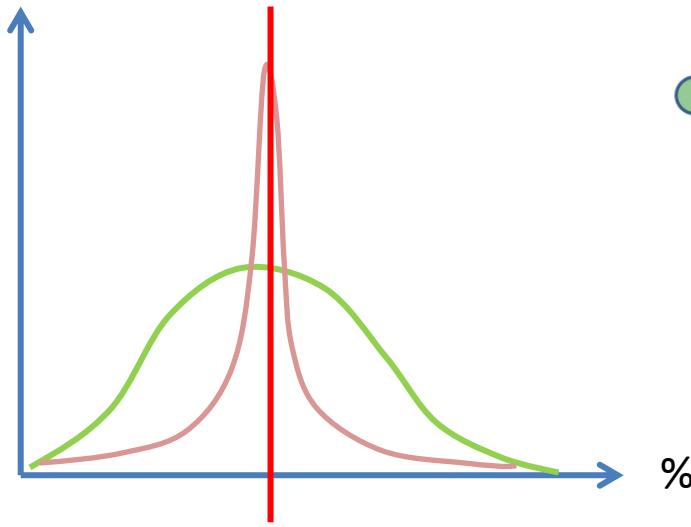
	LIT1	LIT2	LIT3	
1	Light Blue	Light Orange	Light Red	
2	Light Blue	Light Orange	Light Red	
3	Light Blue	Light Orange	Light Red	
4	Light Blue	Light Orange	Light Red	
5	Light Blue	Light Orange	Light Red	
6	Light Blue	Light Orange	Light Red	
7	Light Blue	Light Orange	Light Red	
8	Light Blue	Light Orange	Light Red	
9	Light Blue	Light Orange	Light Red	
10	Light Blue	Light Orange	Light Red	
11	Light Blue	Light Orange	Light Red	
12	Light Blue	Light Orange	Light Red	

\sum = determined by $CV_{pedigree}$

LITERATURE



- how to assess the quality of the data



informative prior :

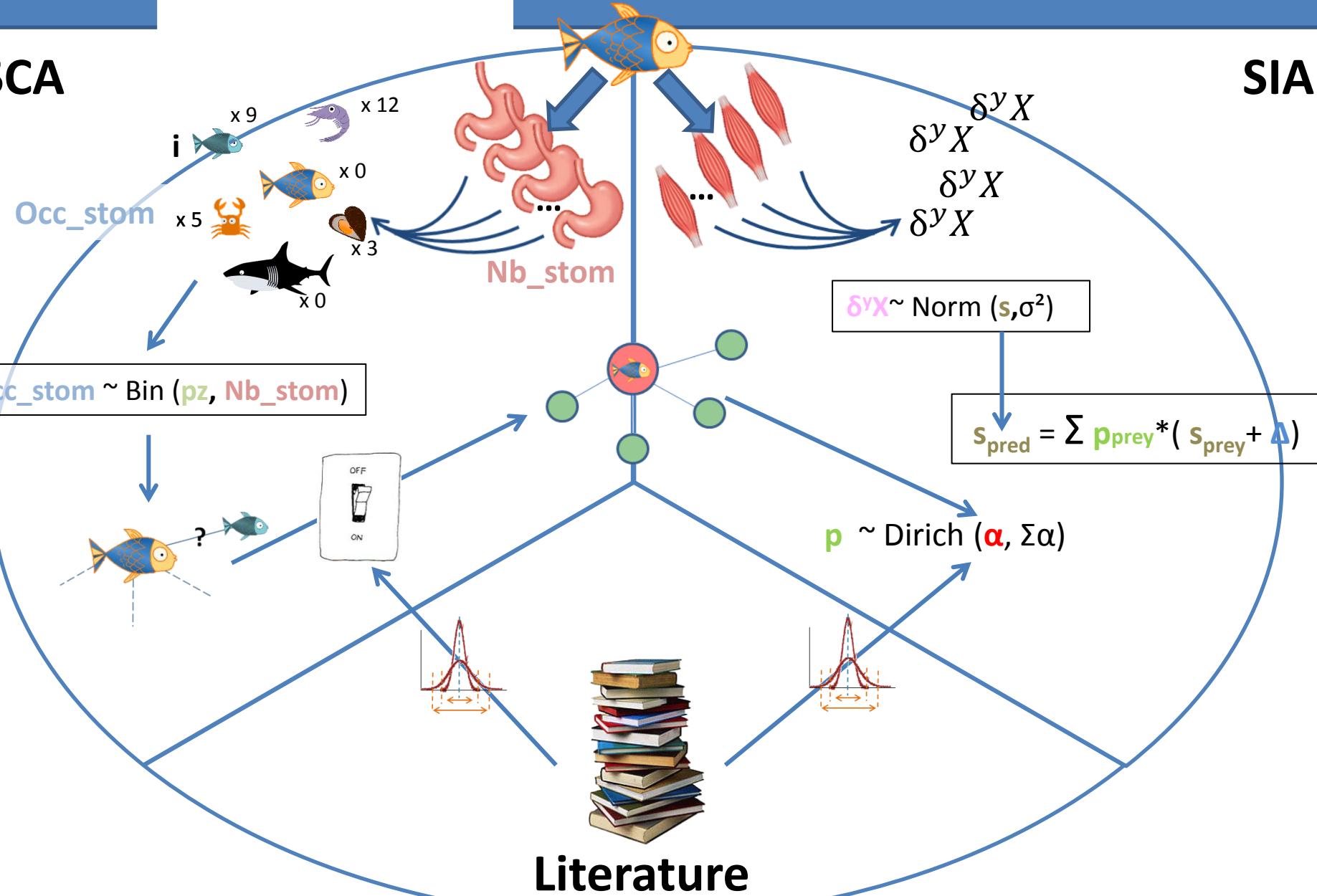
$$p[i,j] \sim \text{Beta}(\alpha[i,j], \sum \alpha[i,])$$

$$\alpha[i,j] / \sum \alpha[i,]$$

\sum = determined by $CV_{pedigree}$

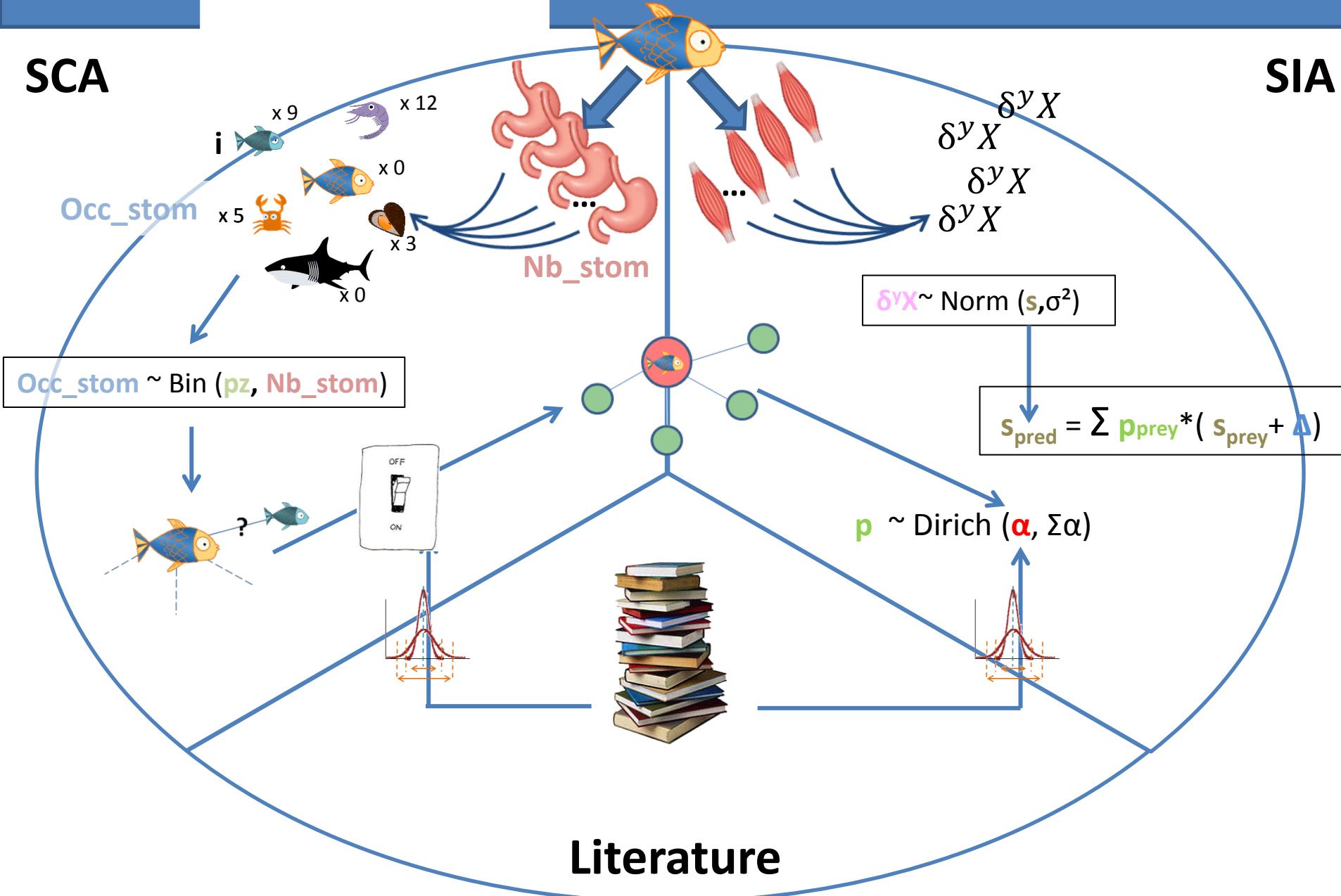
SCA

SIA



SCA

SIA



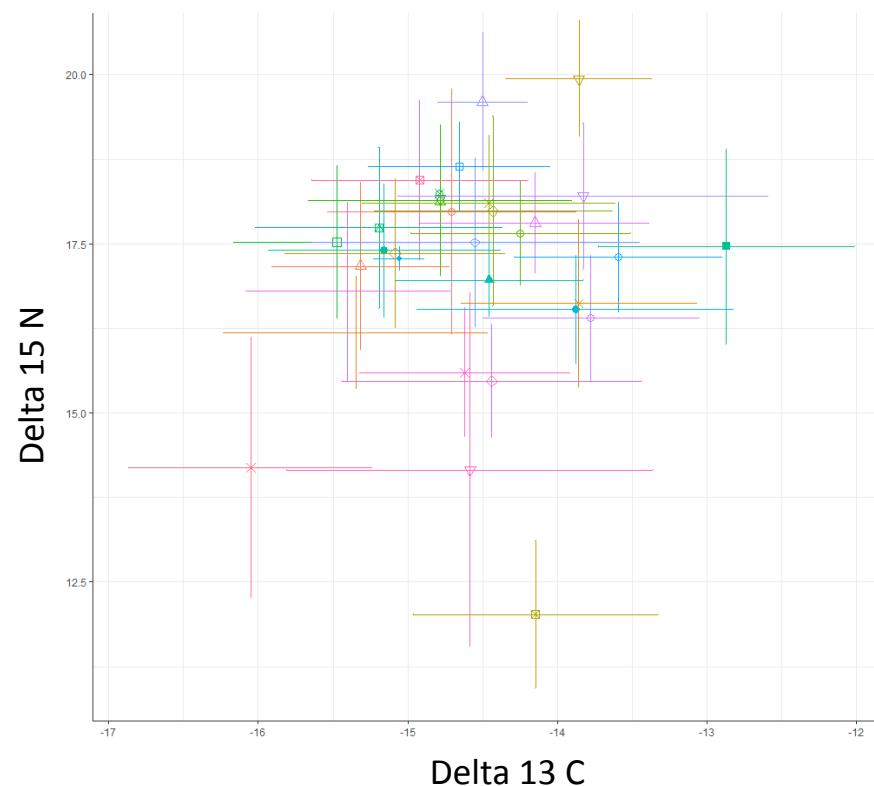
(first)
RESULTS



Dessinateur : André-Philippe Côté

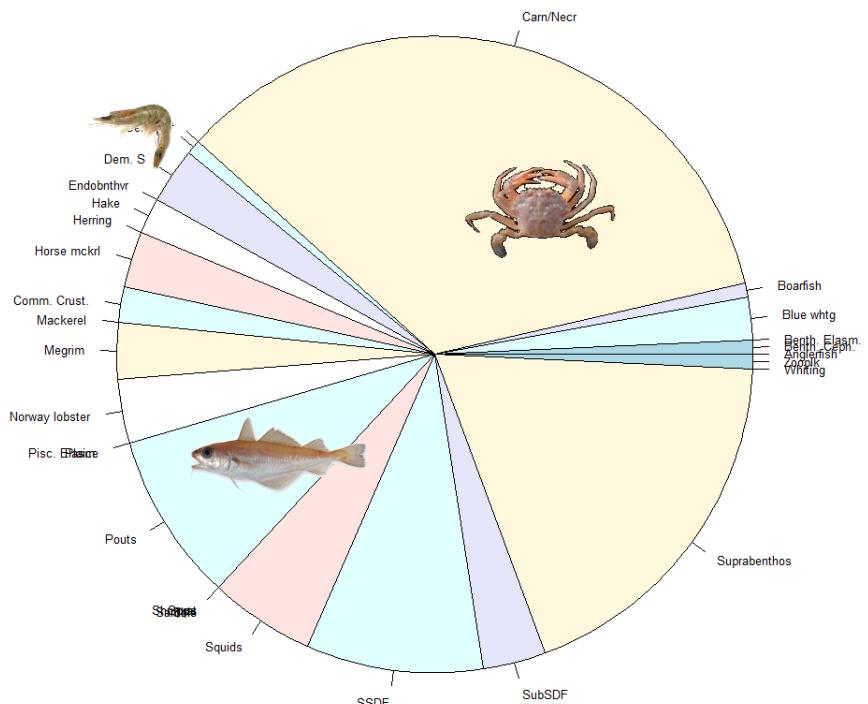
« EASY » RESULTS

SIA for all trophic groups



(Data prep.: baseline correction)

SCA for one species: cod



/!\ Occurrences of each prey in separated stomachs

(Data prep.: reassignment of undet. grp)

FIRST RUN OF EcoDiet ON BOTH SIA & SCA DATA IN THE CELTIC SEA

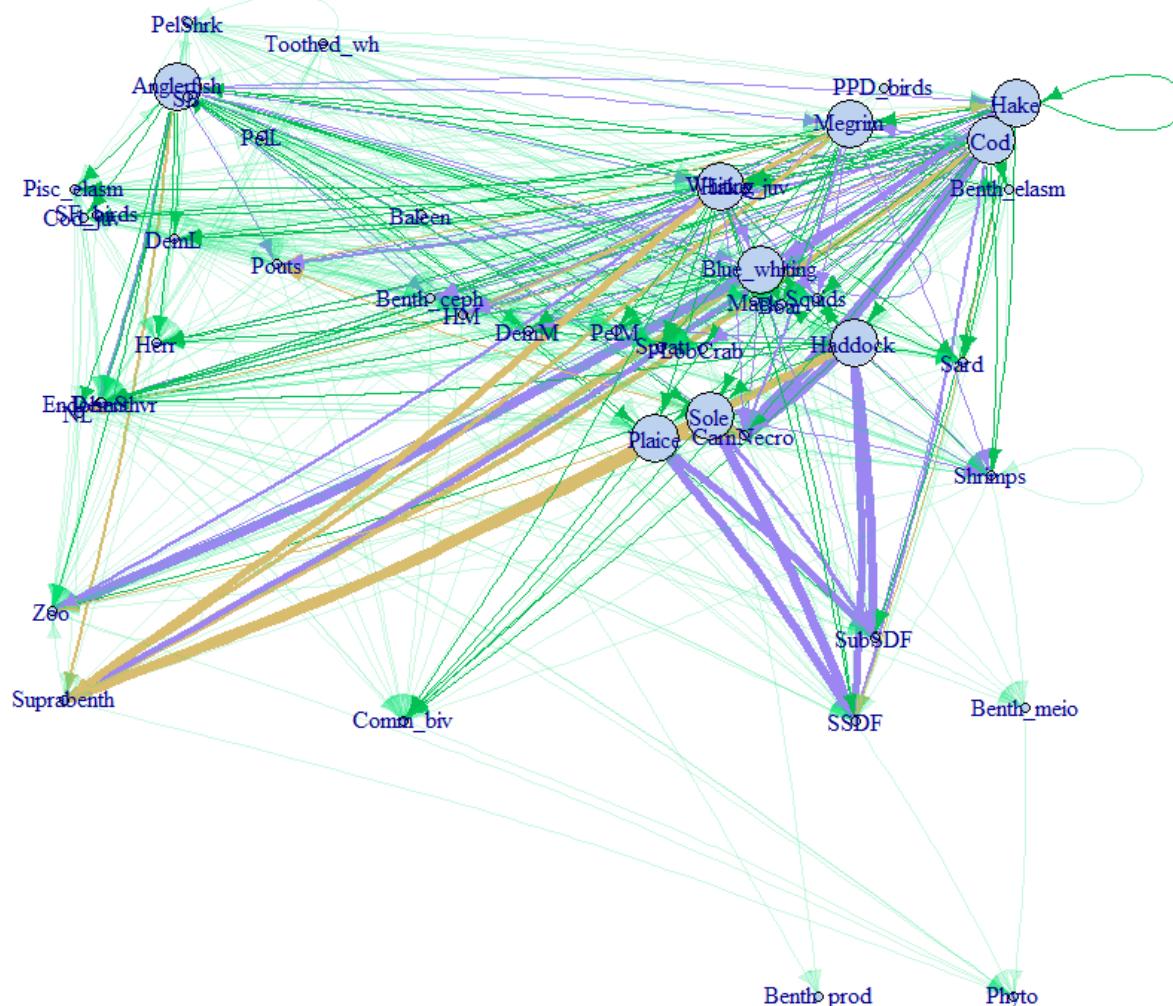
TOPOLOGICAL FOOD-WEB

only Stom
only lit.
Stom & lit.
only lit.

SCA available

SCA not available

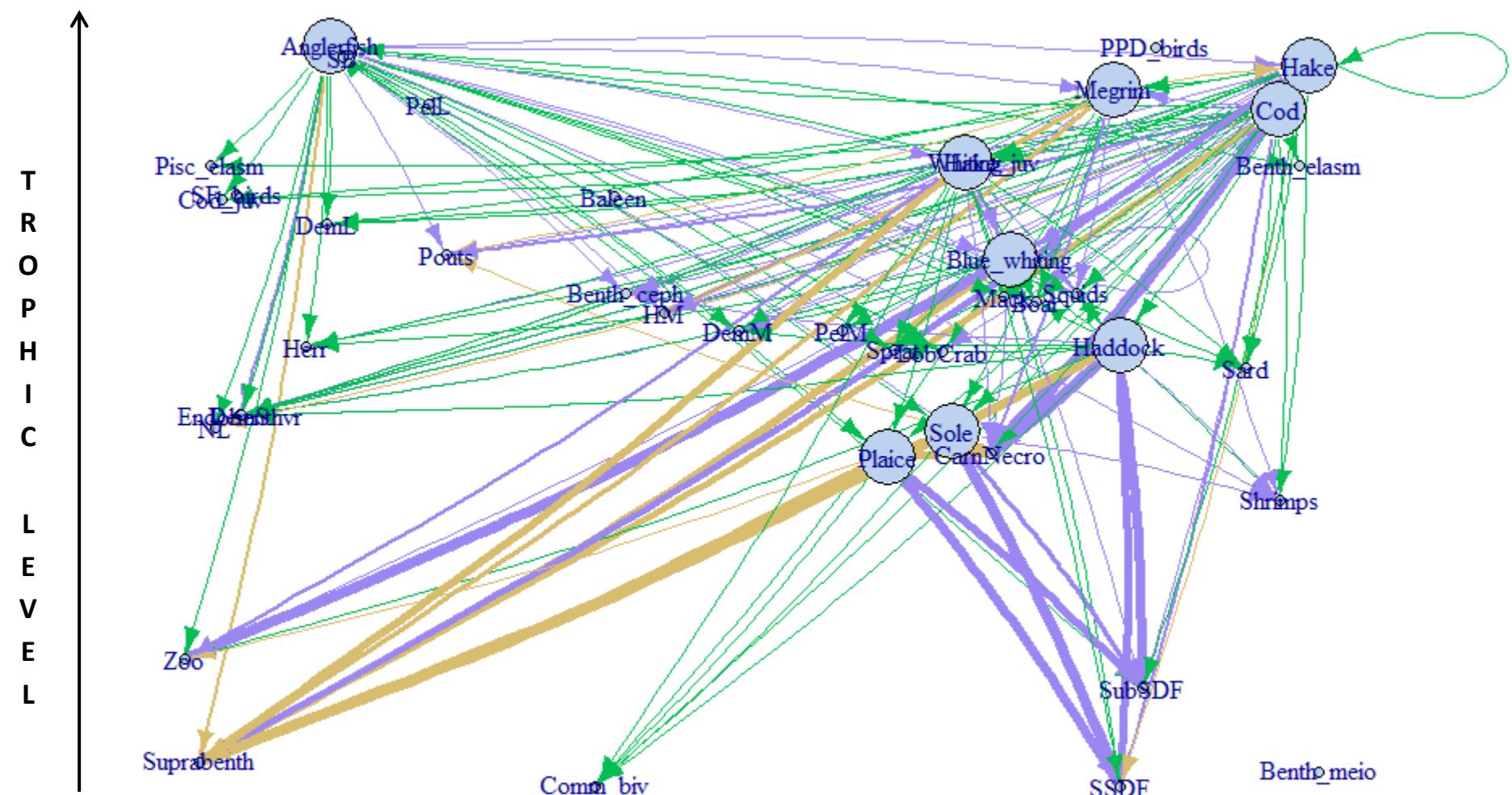
TROPHIC LEVEL



(Width is proportional to the probability that a link exist between two groups)

TOPOLOGICAL FOOD-WEB deduced by SCA

only Stom only lit. Stom & lit.

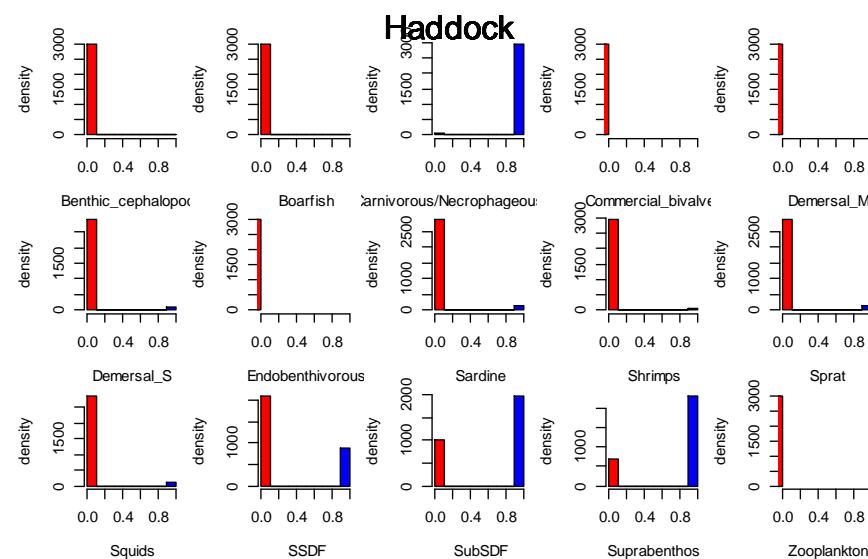
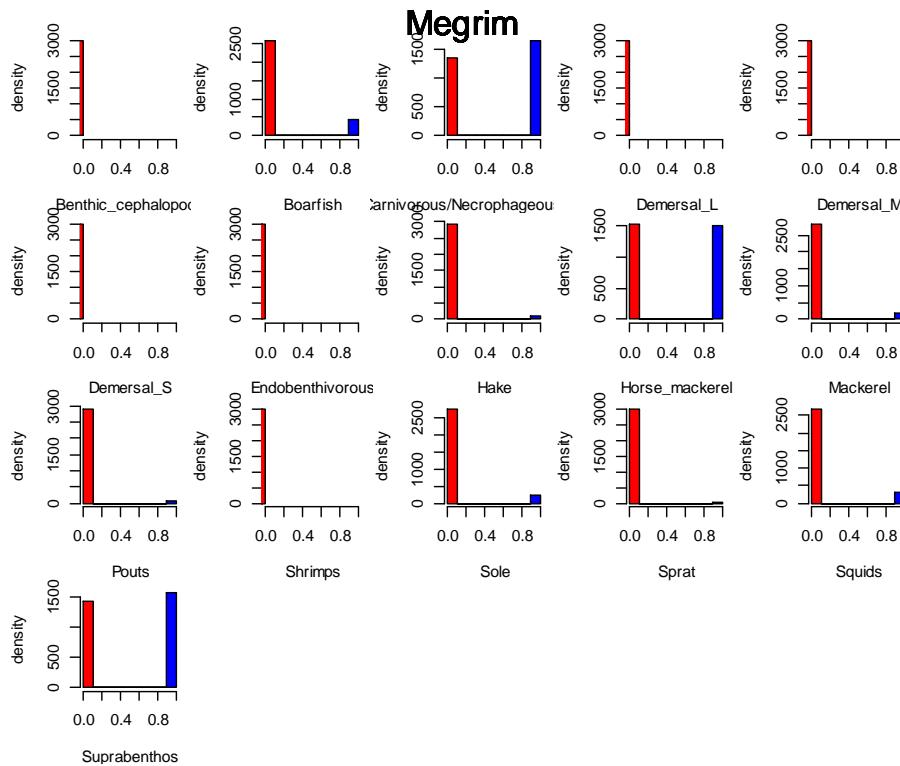


(Width is proportional to the probability that a link exist between two groups)

TOPOLOGICAL FOOD-WEB

Take back posteriors of the several parameters

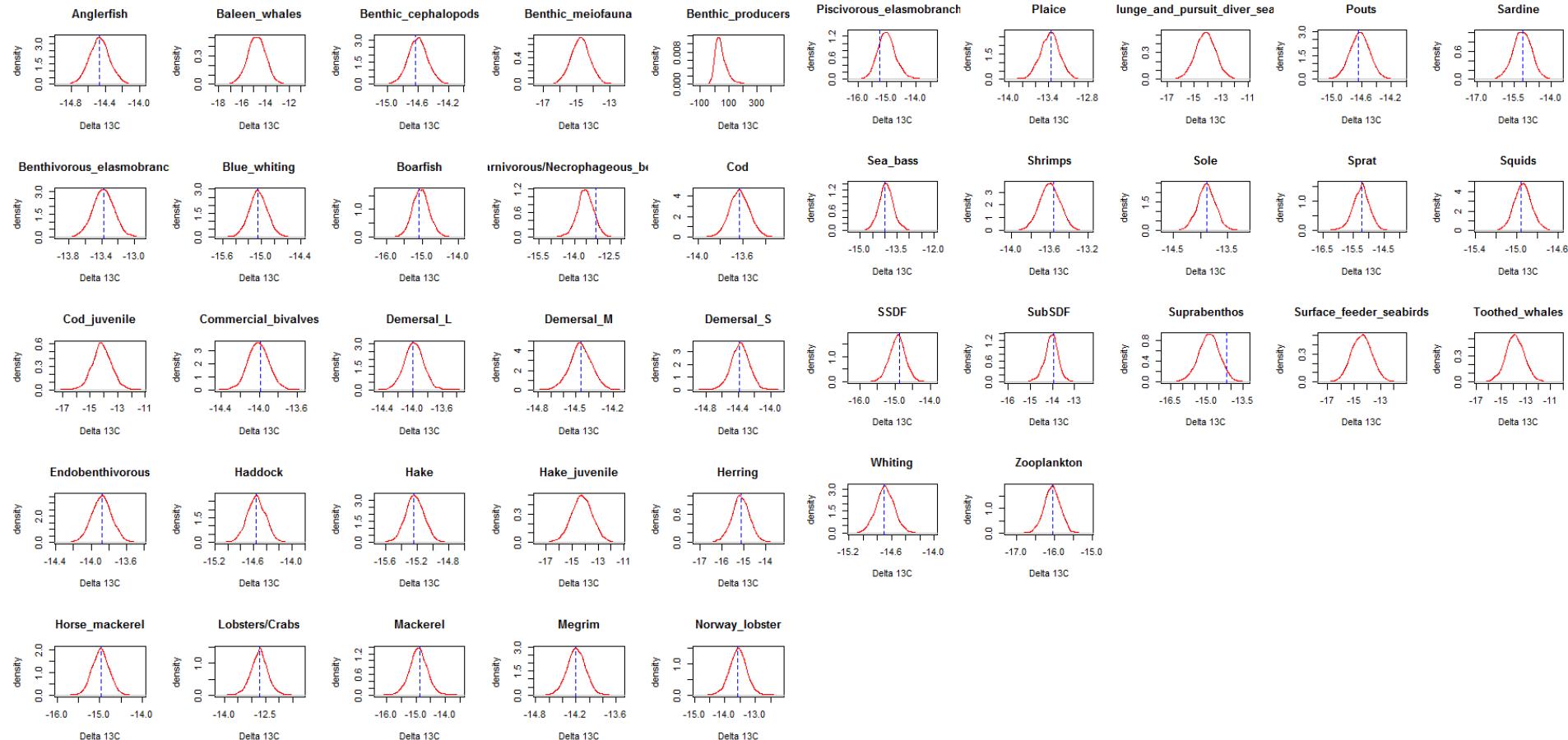
« SWITCHER »



STABLE ISOTOPES MODULE

Take back posteriors of the several parameters

ISOTOPIC SIGNATURES

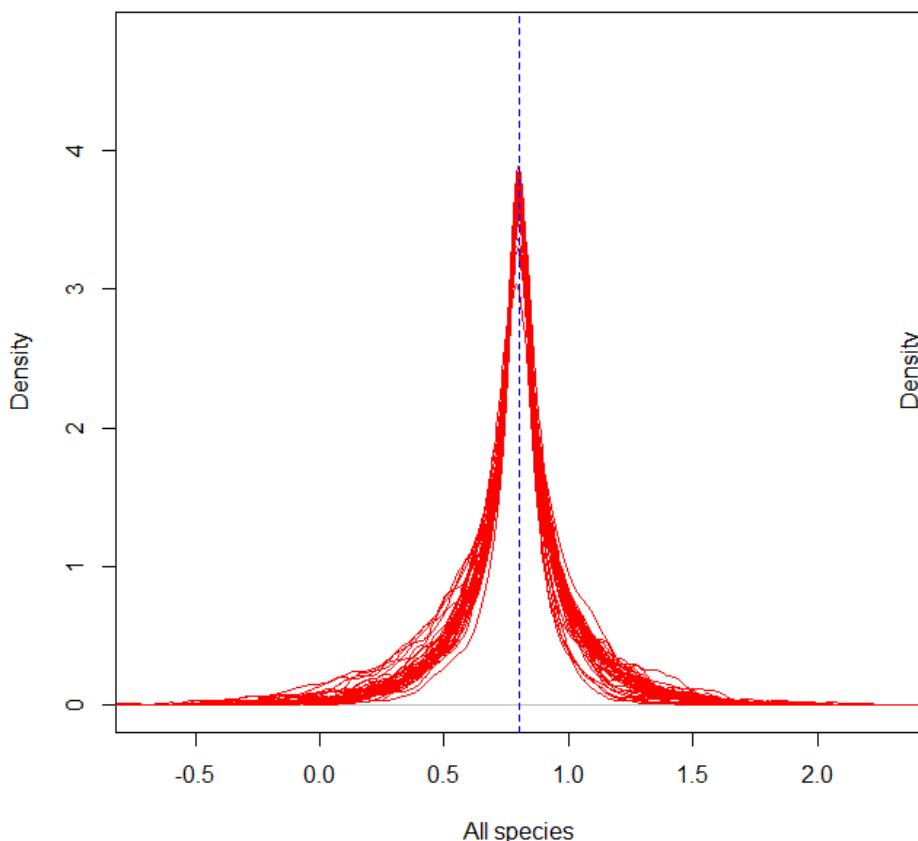


STABLE ISOTOPES MODULE

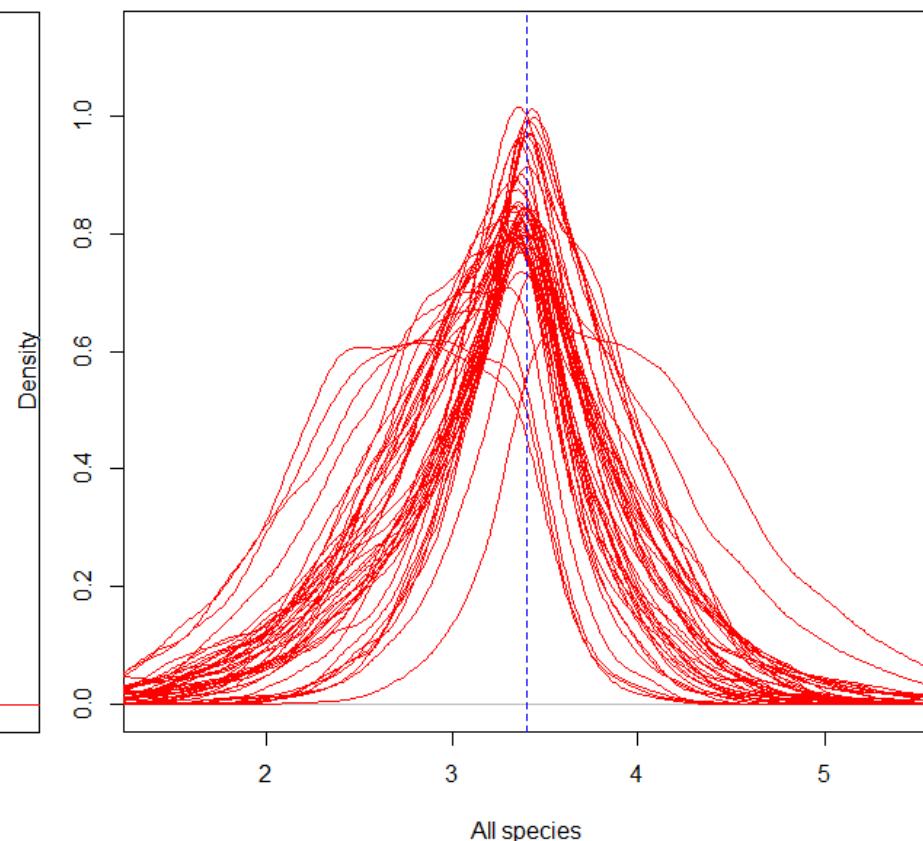
Take back posteriors of the several parameters

TROPHIC ENRICHMENT

Carbon trophic enrichment

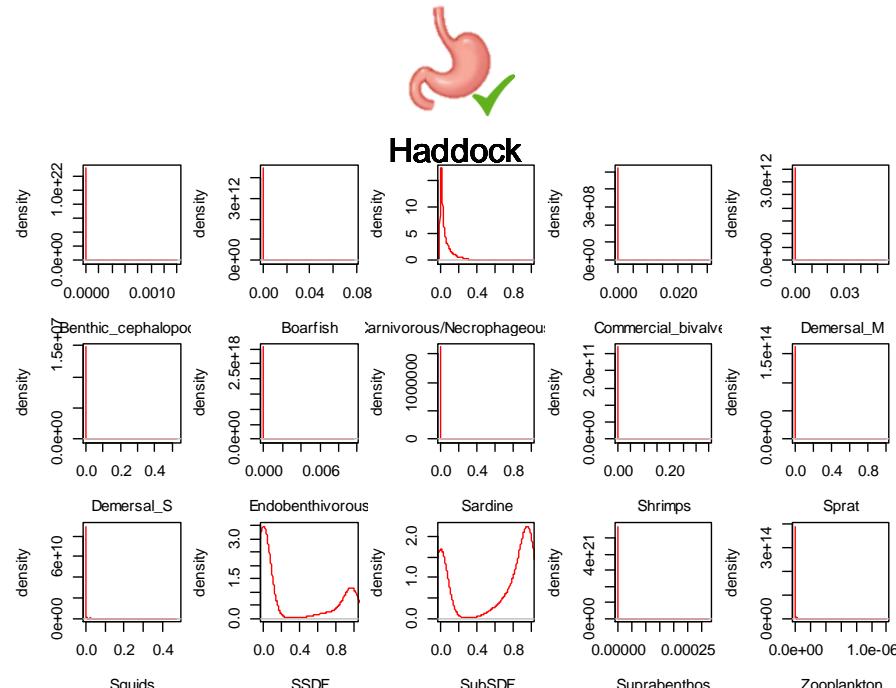


Nitrogen trophic enrichment



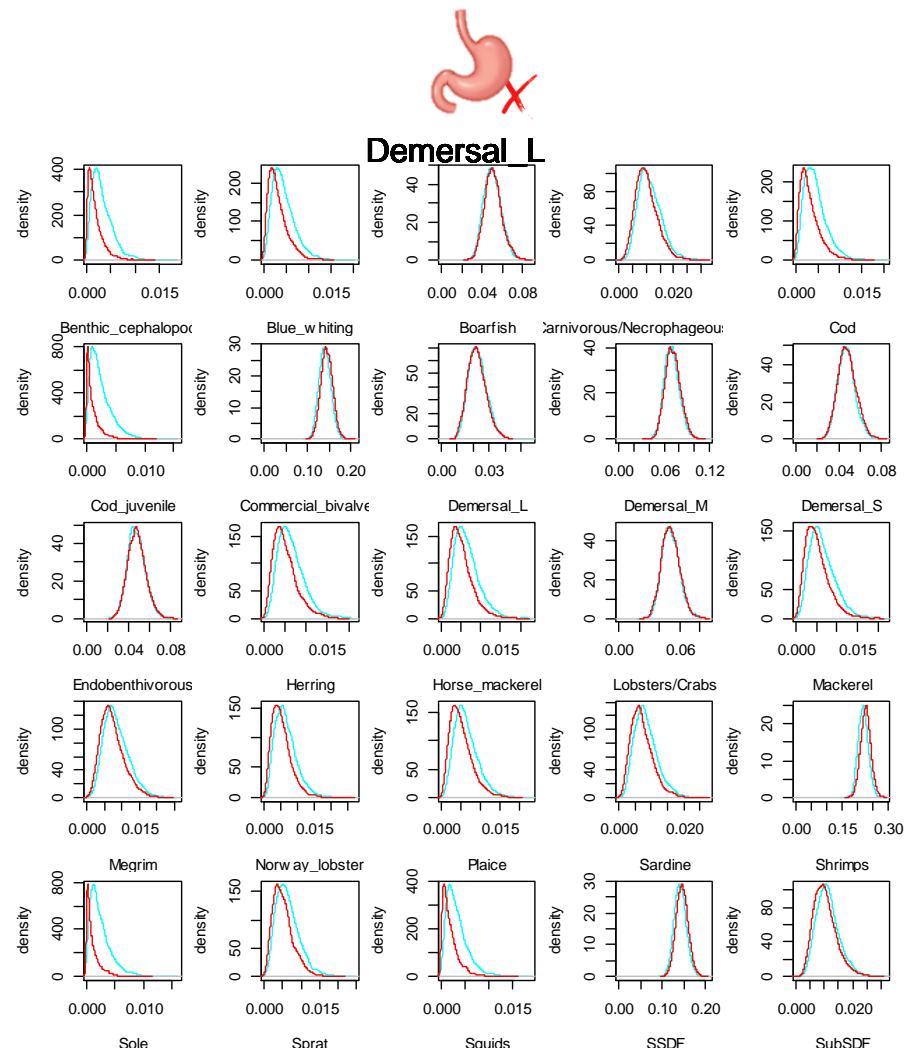
STABLE ISOTOPES MODULE

Take back posteriors of the several parameters



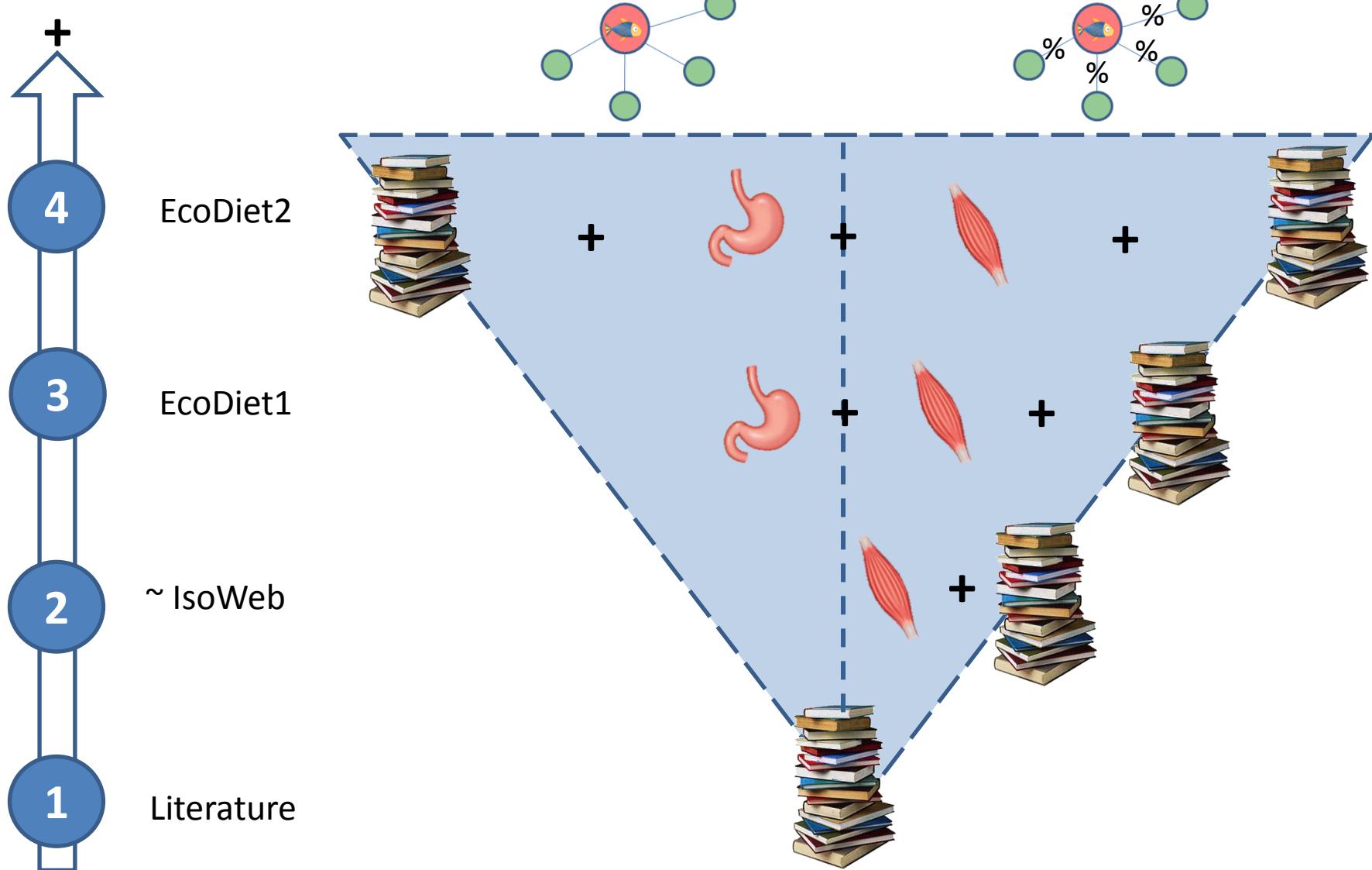
Haddock

CONTRIBUTIONS



Demersal L

COMPARING THE RESULTS - Strategic design



A NEW VISION OF DIETS IN THE CELTIC SEA – the cod case

Piscivorous_elasmobranchs

Mackerel

Plaice

Pouts

Sardine

Herring

Horse_mackerel

Lobsters/Crabs

Endobenthivorous

Demersal_S

Demersal_M

Demersal_L

Commercial_bivalves

Cod_juvenile

Carnivorous/Necrophageous_benthos

Boarfish

Blue_whiting

Benthivorous_elasmobranchs

Benthic_cephalopods

Anglefish

Zooplankton

SSDF

Squids

Sole

Sole

SSDF

Sole

Sole

SSDF

Sole

Sole

SSDF

Sole

SSDF</div

A NEW VISION OF DIETS IN THE CELTIC SEA – the cod case

1



3

2

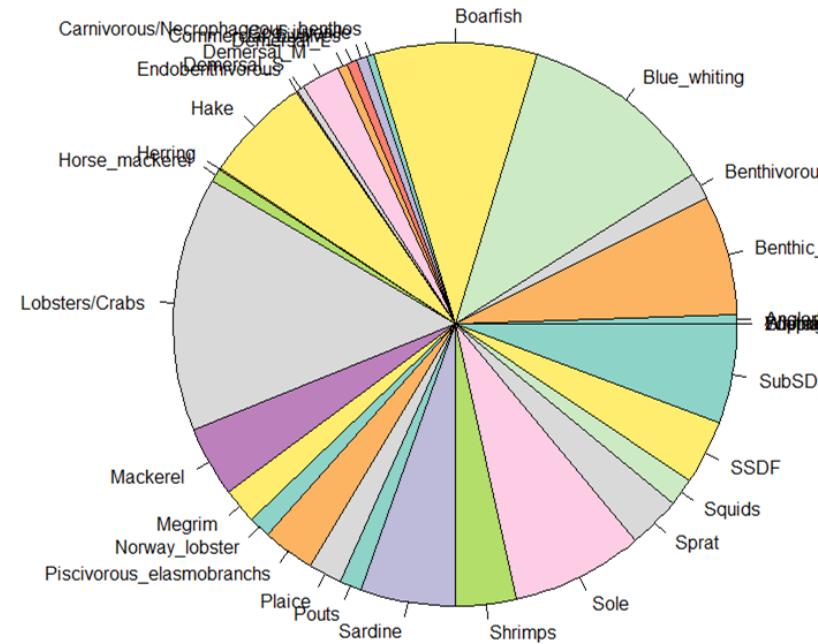
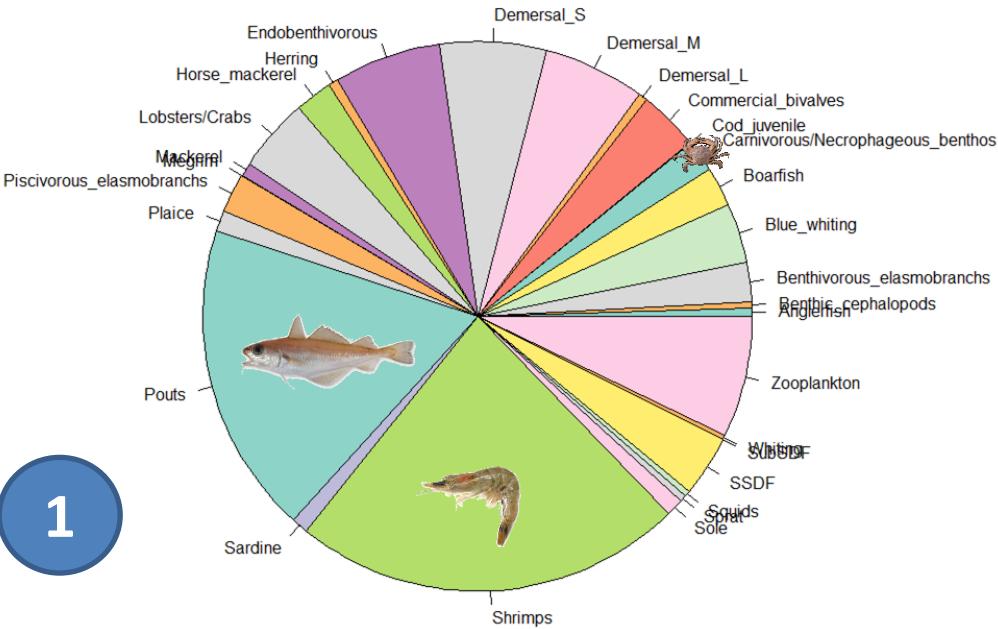


4

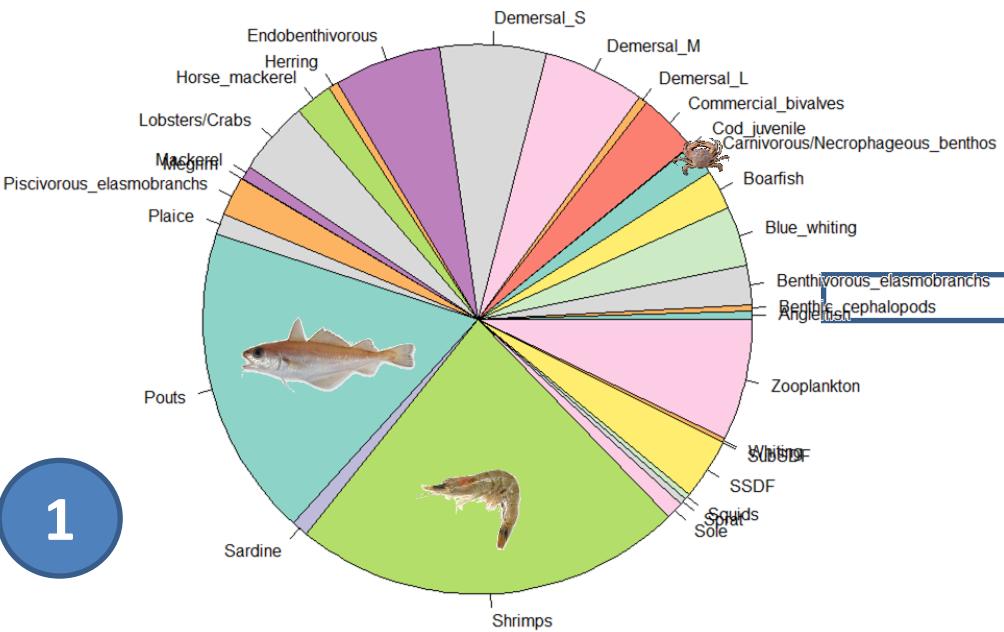


A NEW VISION OF DIETS IN THE CELTIC SEA – the cod case

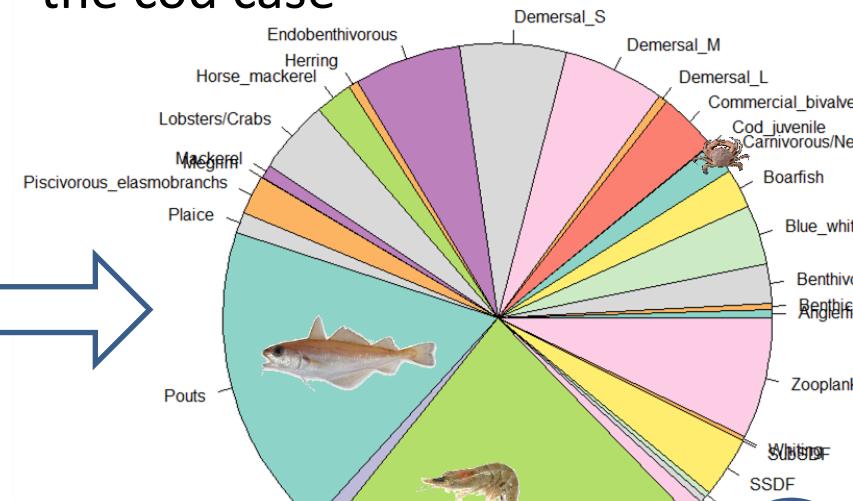
1



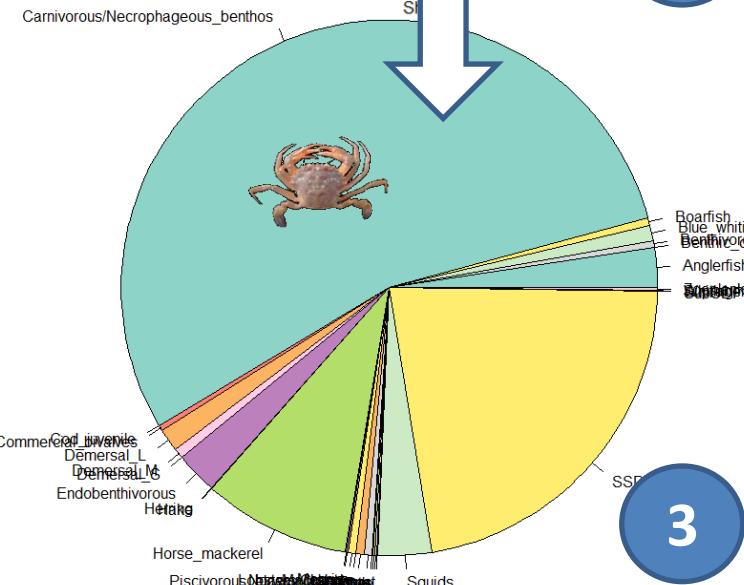
A NEW VISION OF DIETS IN THE CELTIC SEA – the cod case



1

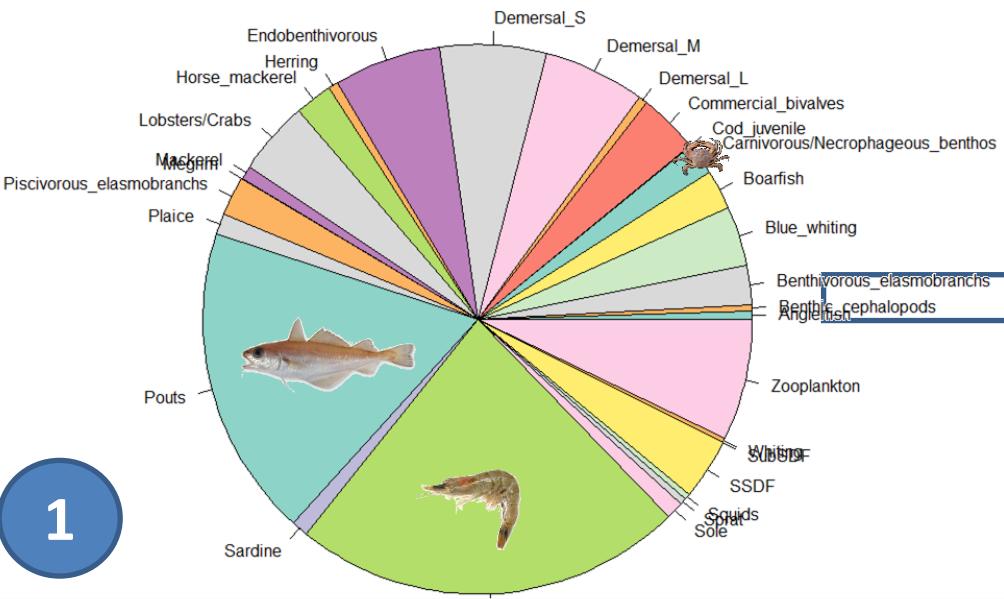


2

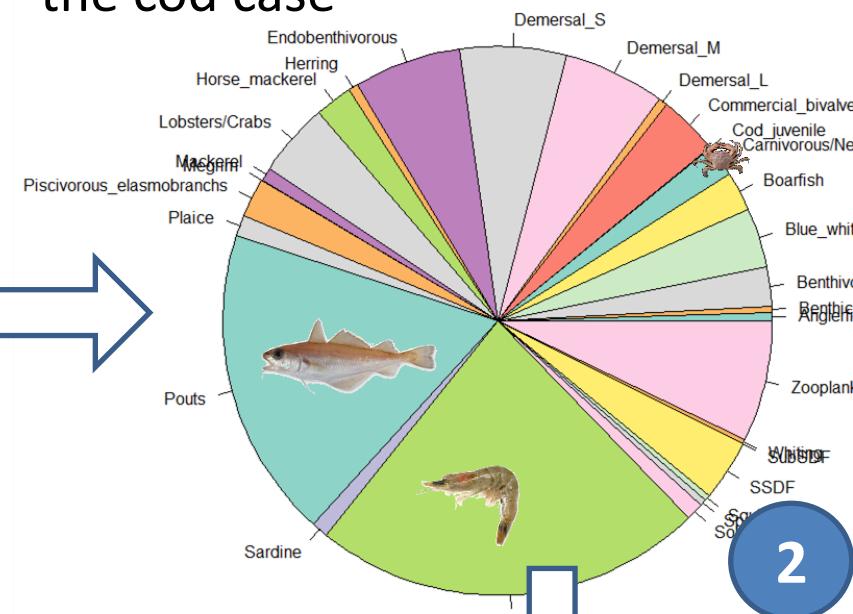


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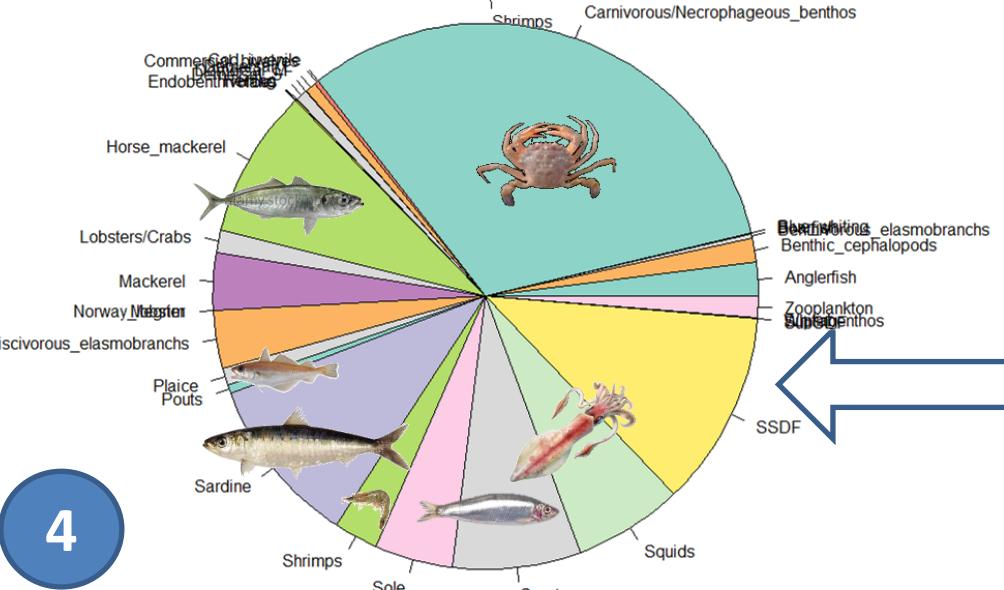
A NEW VISION OF DIETS IN THE CELTIC SEA – the cod case



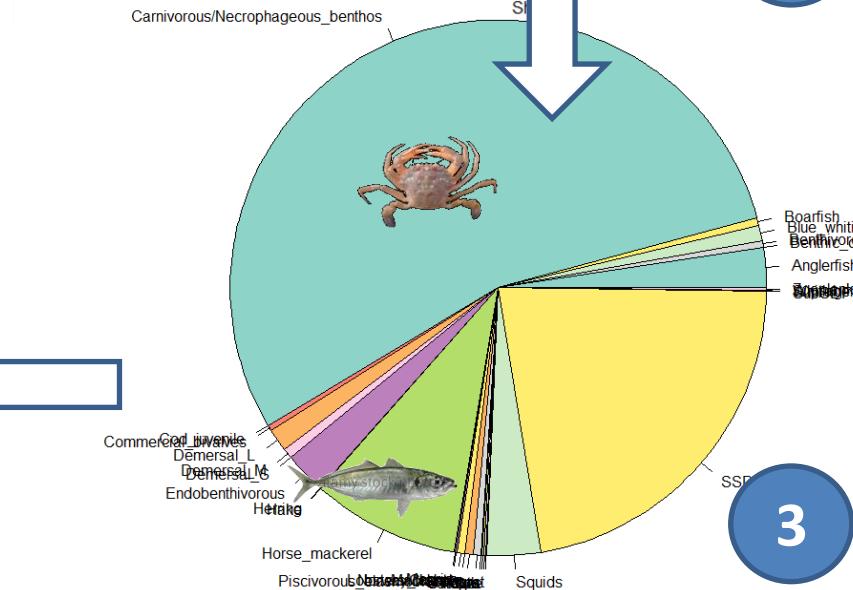
1



2



4



3

DISCUSSION



Dessinateur : André-Philippe Côté

STILL TO DO

- Inclusion of final a « Pedigreed » diet matrix as prior
- Remove variation for the trophic enrichment factor
- Sensitivity analysis on data pretreatment of SIA & SCA

ABOUT EcoDiet

- EcoDiet: an innovative integrated model to couple both SCA and SIA...
....and literature data
- Main strengths:
 - SCA as occurrences = less time-consuming
 - Distinction between preys with similar isotopic signatures
 - Better estimates of contributions since it artificially decrease the number of potential preys
- Promising results: new vision of trophic functionning in the Celtic Sea

Perspectives

