

Global change and aquatic ecosystems: An opportunity to combine fundamental and applied researches

Simon Blanchet
Station d'Ecologie Théorique et Expérimentale (UMR 5321)



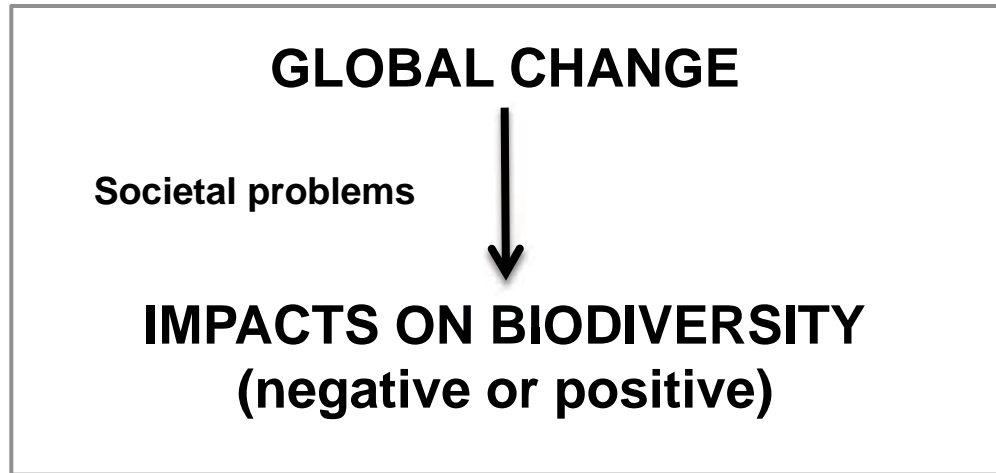
**AS MOST ECOSYSTEMS,
FRESHWATERS ARE IMPACTED BY MULTIPLE STRESSES**



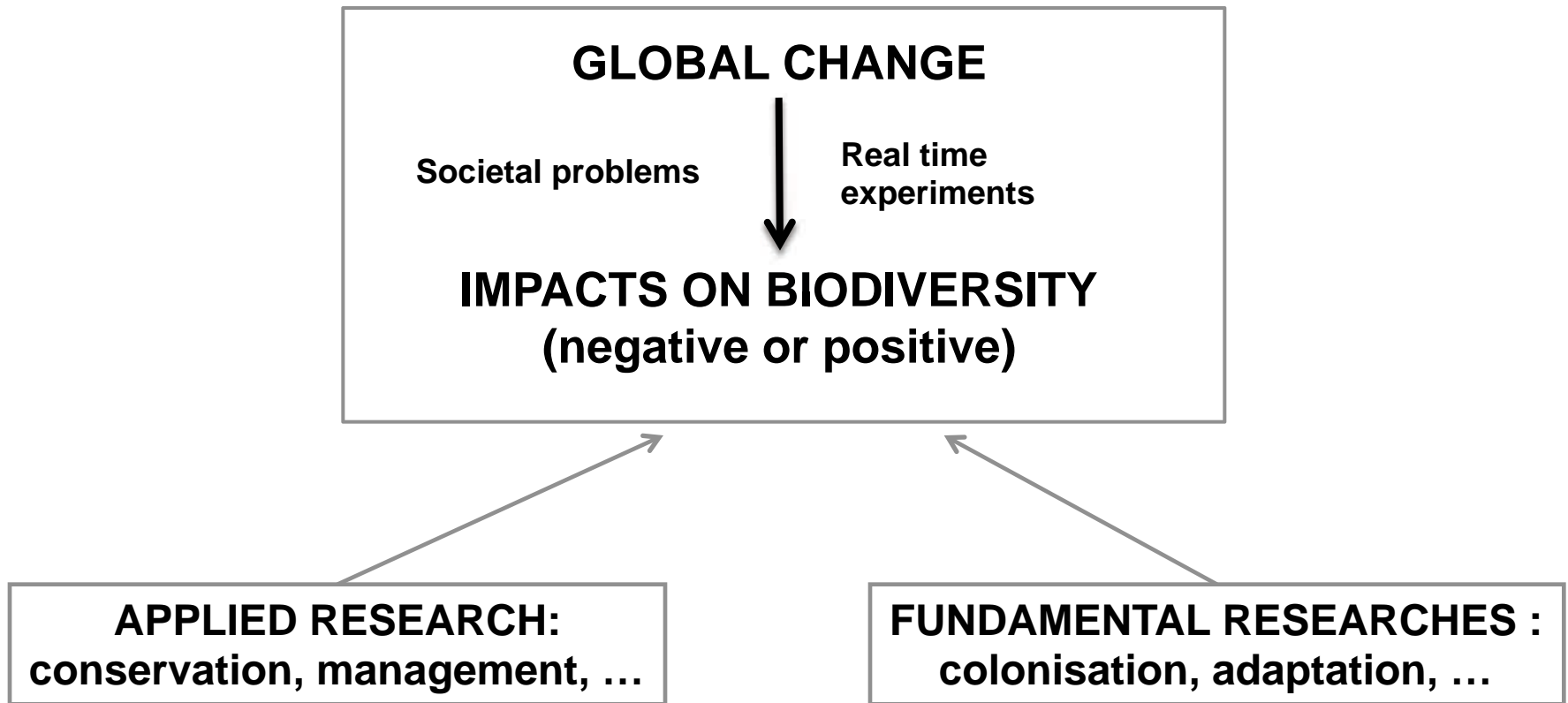
GLOBAL CHANGE

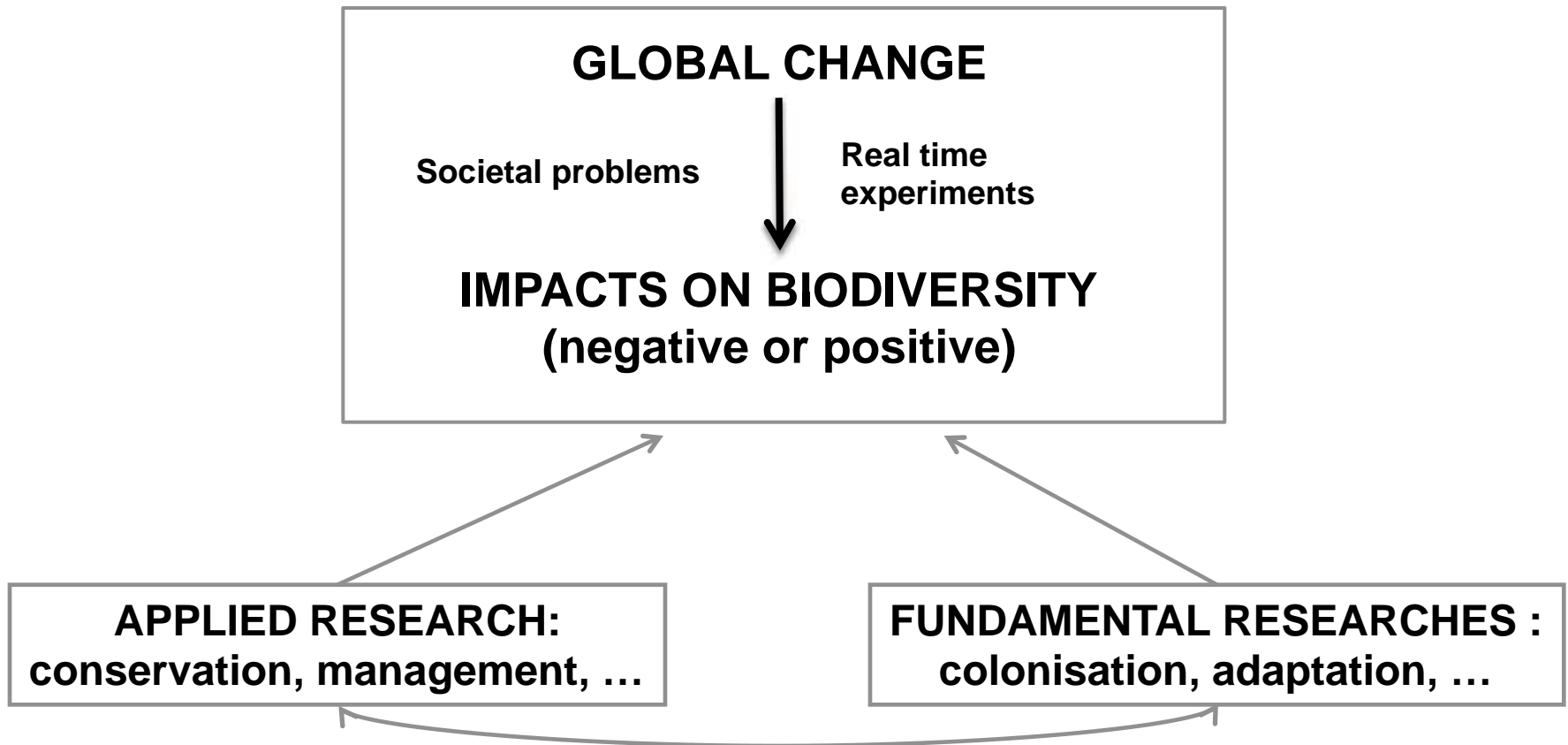


IMPACTS ON BIODIVERSITY
(negative or positive)



**APPLIED RESEARCH:
conservation, management,....**





A UNIQUE FRAMEWORK TO COMBINE APPLIED AND FUNDAMENTAL RESEARCHES



Two research axes

**Ecology and evolution of host-
parasite interactions**
Biological invasions

**Intraspecific diversity in river
networks**
*Conservation and habitat
fragmentation*

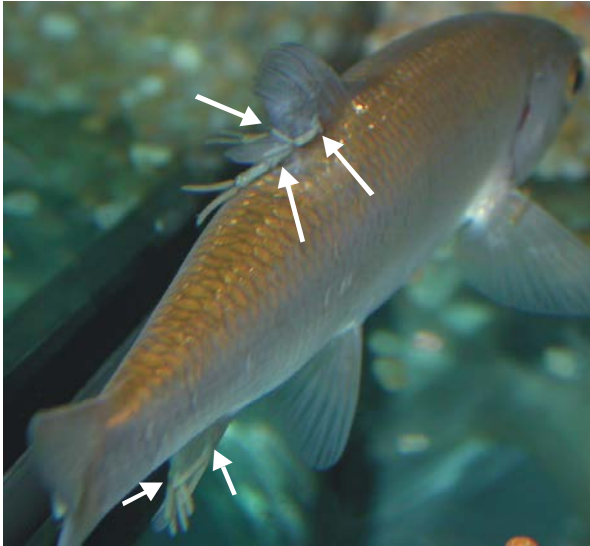
Both applied and fundamental questions
Molecular tools, empirical surveys, simulations, experiments



- 1) PATHWAYS OF INVASION FOR A NEGLECTED PARASITE**
- 2) ENVIRONMENTAL AND GENETIC BASES OF HOST DEFENSES**
- 3) GENOMIC BASES OF HOST DEFENSES**



THE TRACHELIASTES-DACE INTERACTIONS:



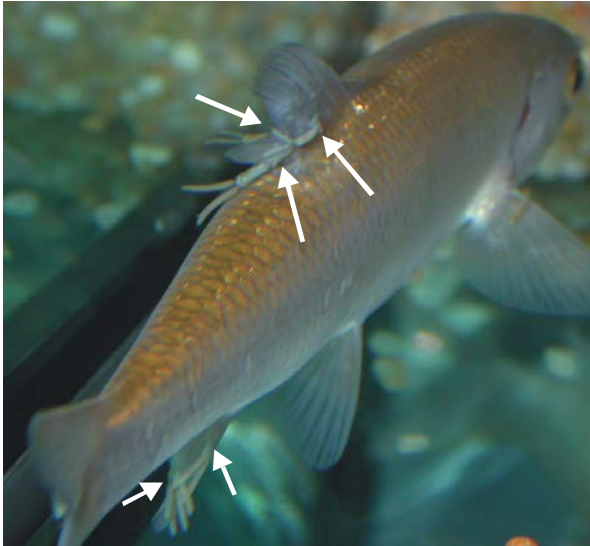
Dace (*Leuciscus sp*)



Copepod (*Tracheliaestes polycolpus*)



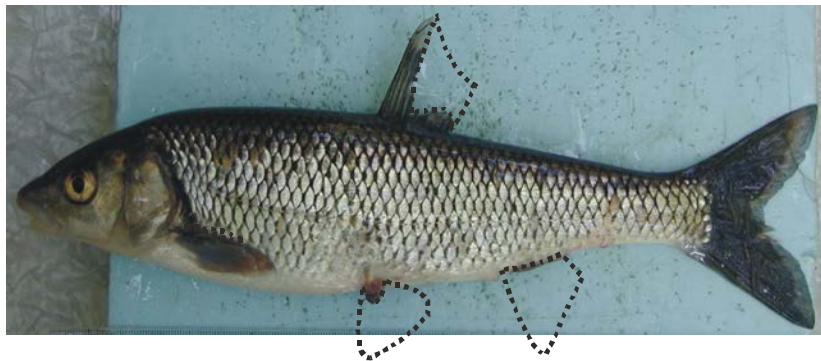
THE TRACHELIASTES-DACE INTERACTIONS:



Dace (*Leuciscus sp*)

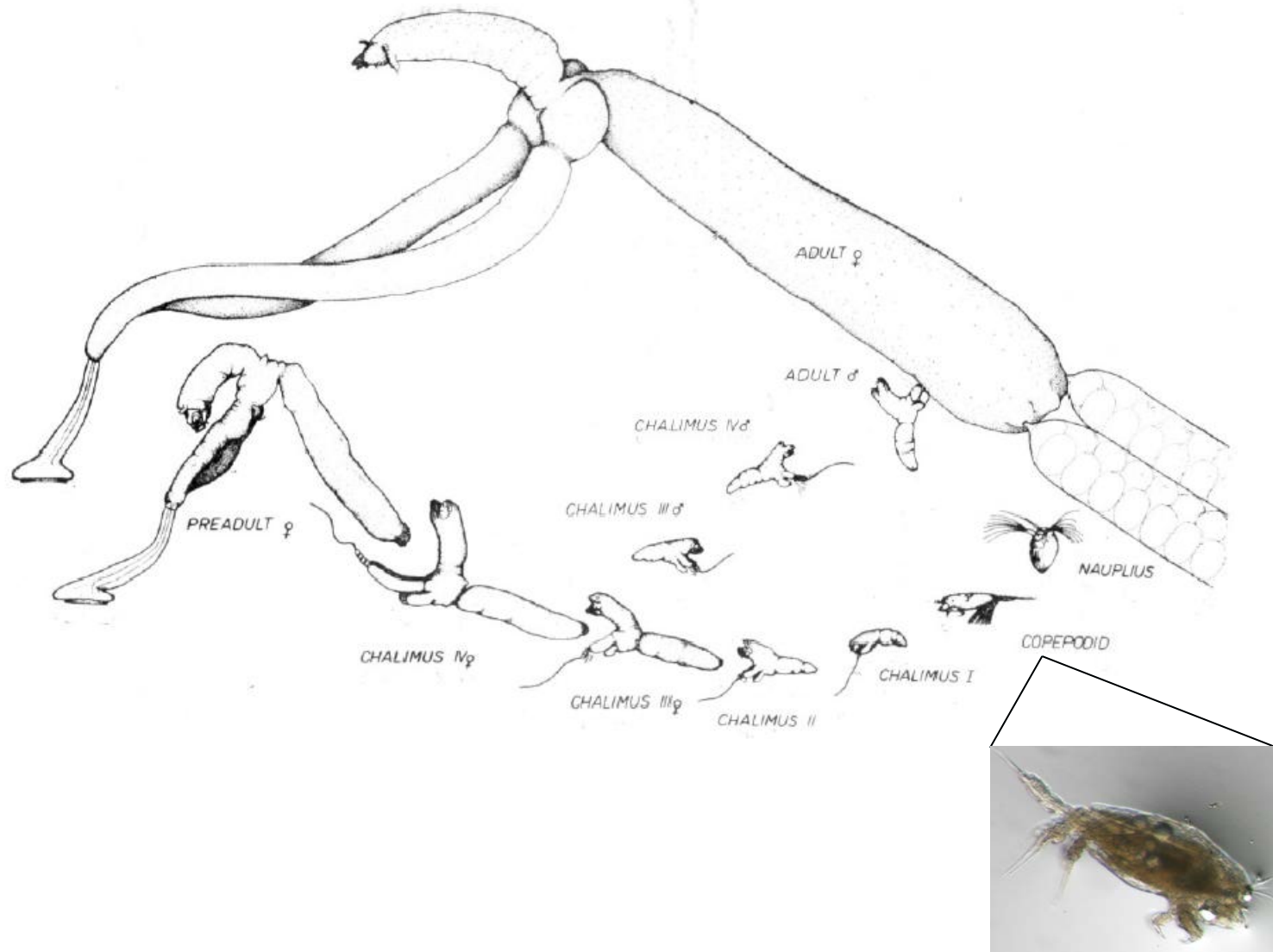


Copepoe (*Tracheliastes polycolpus*)





LIFE CYCLE (DIRECT CYCLE, A SINGLE HOST SPECIES):





(I) PATHWAYS OF INVASION

- Widespread in the North-Eastern part of the Palearctic
- Detected since the 1930' in the UK, the 1960' in France



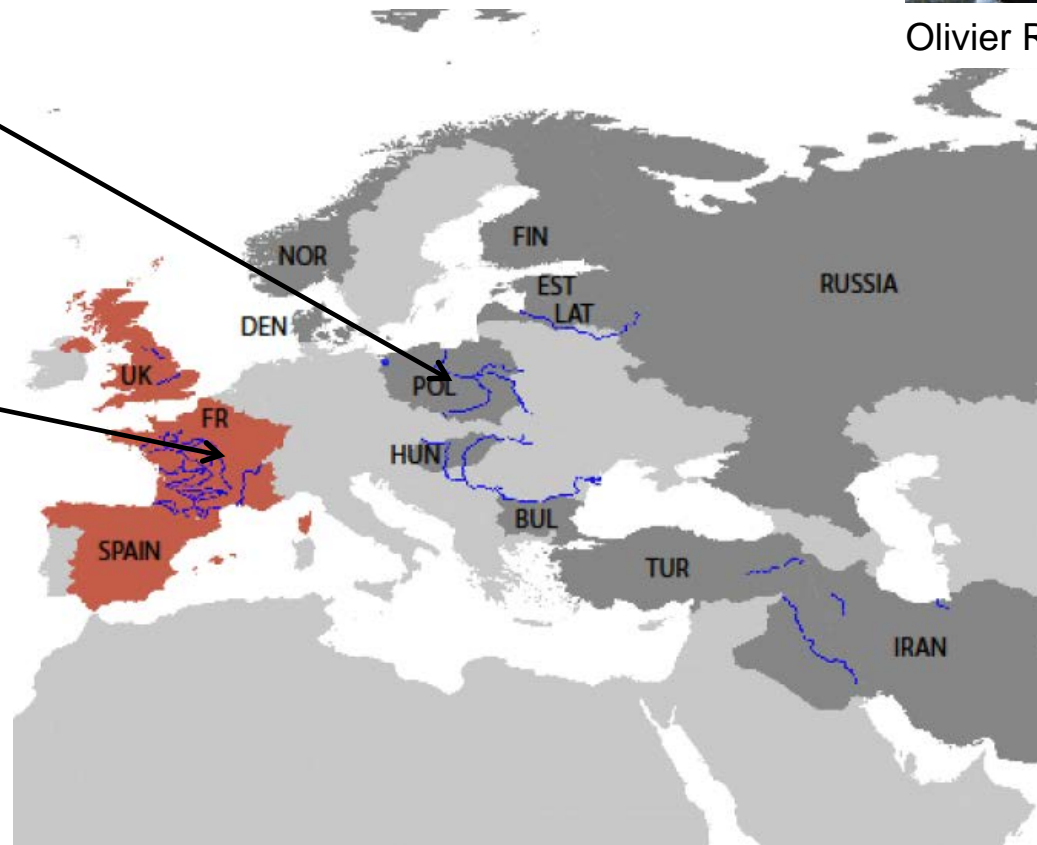
Olivier Rey



Lisa Fourtune

Historical presence

Recently detected

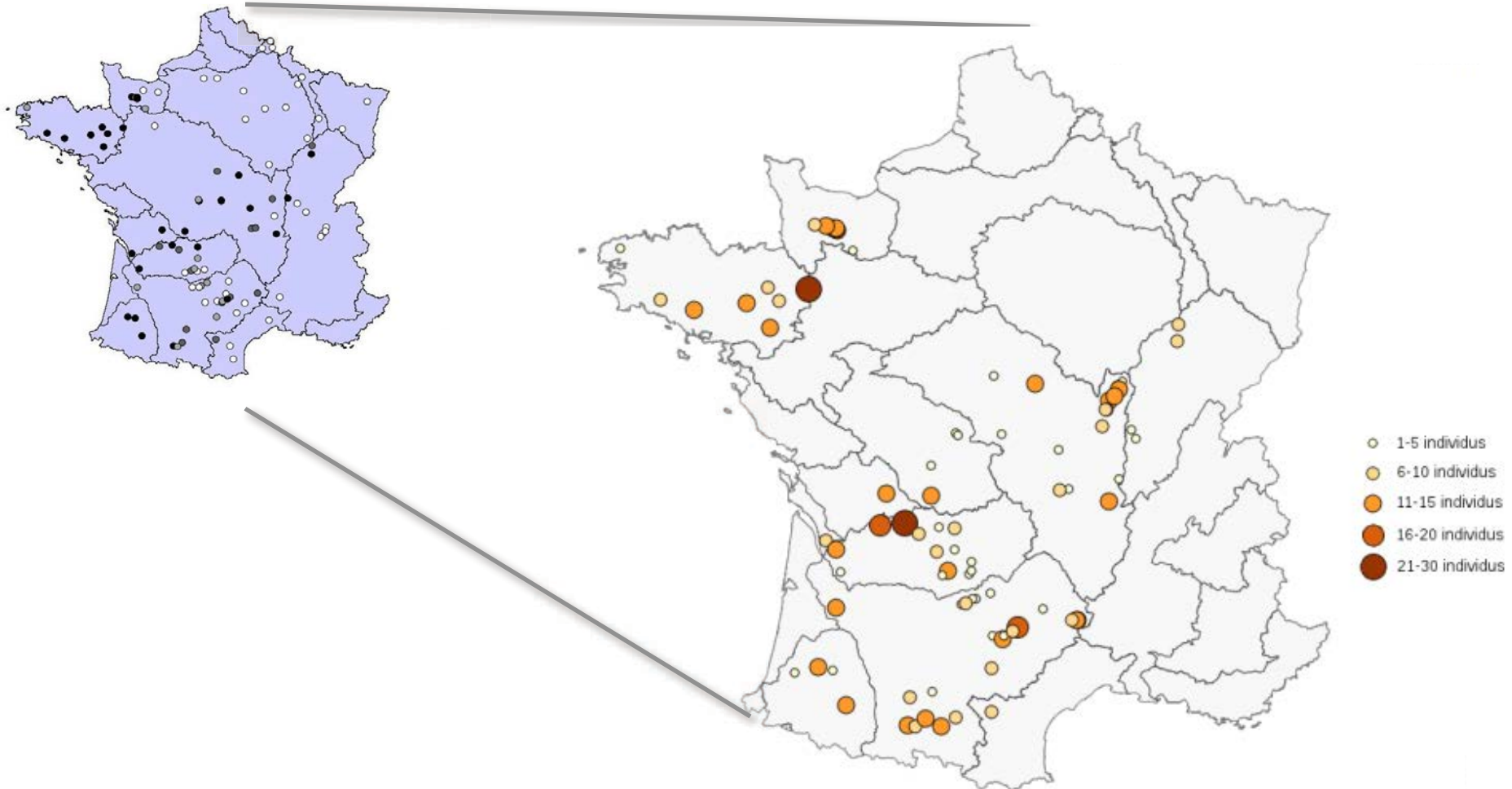


Is *T. polycolpus* a non-native species in France?
What is its invasion history? (potential sources are unknown)

(I) PATHWAYS OF INVASION

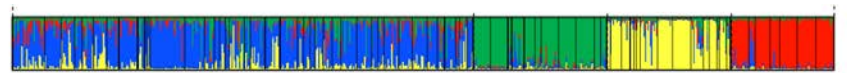
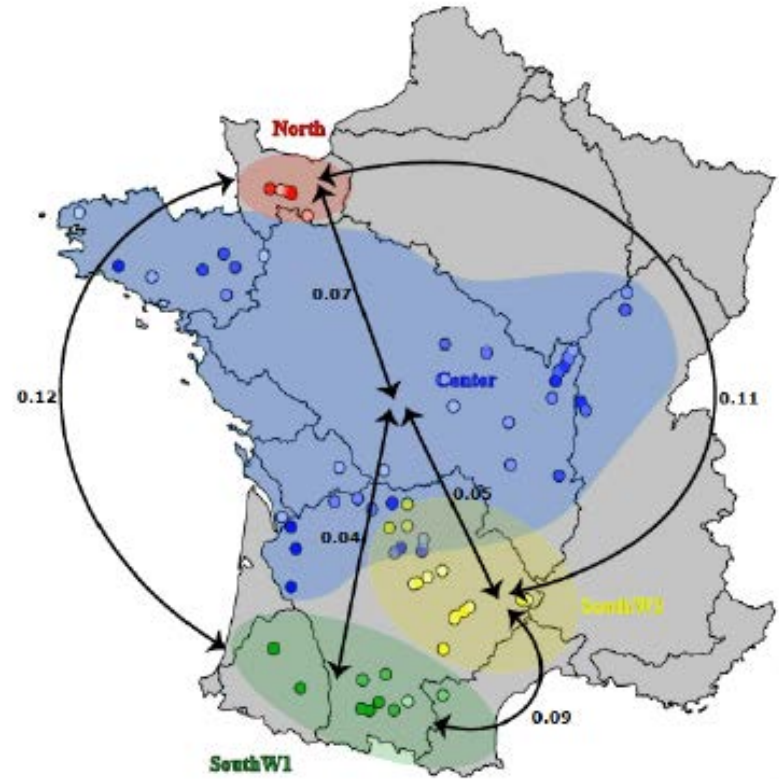


SAMPLING ACROSS FRANCE:
673 individuals, 86 sites, 61 rivers, 16 microsatellite loci





FOUR MAIN GENETIC CLUSTERS *(Structure, Pritchard et al. 2000)*



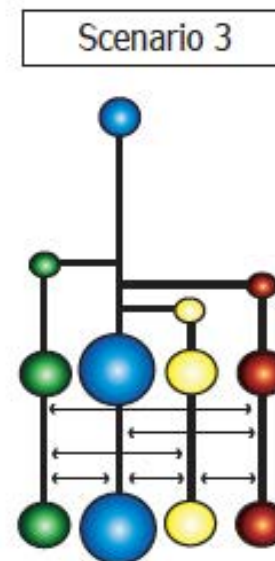
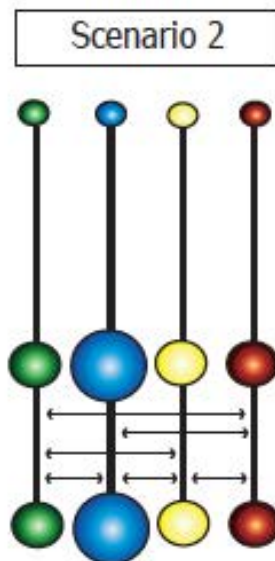
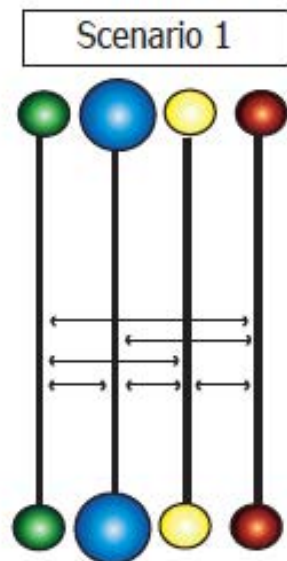
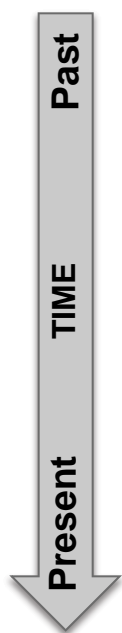
Two possible types of scenarios:

« Native » : historical presence in france, recent expansion

« Non-native » : introduction and recent expansion



Native scenarios: 3 models

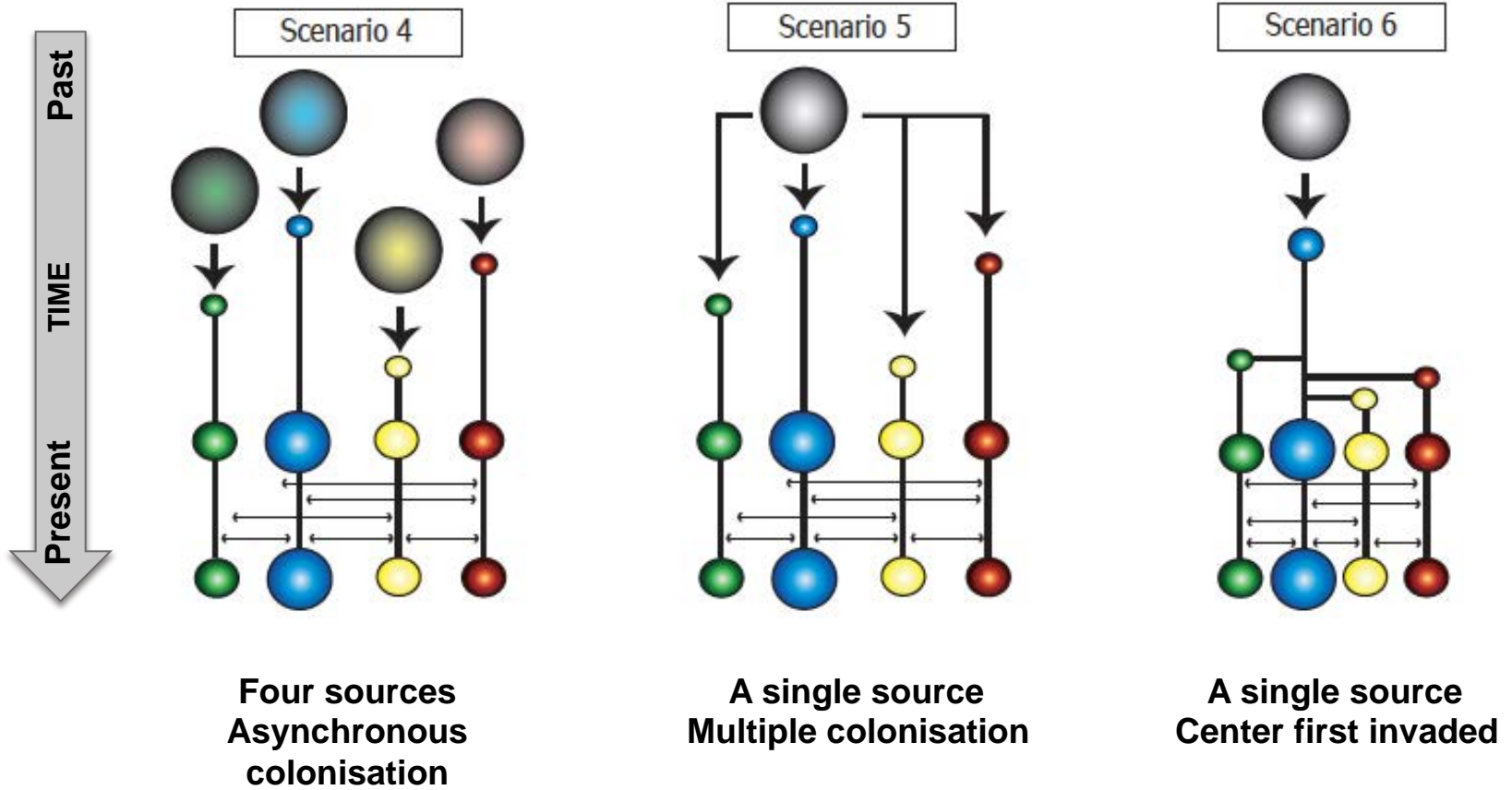


The four clusters historically exist
Tracheliastes discovered only recently
Stabilité (S1) vs. exponential growth (S2)

Only the Center cluster
historically exists
Recent colonisation of
new basins

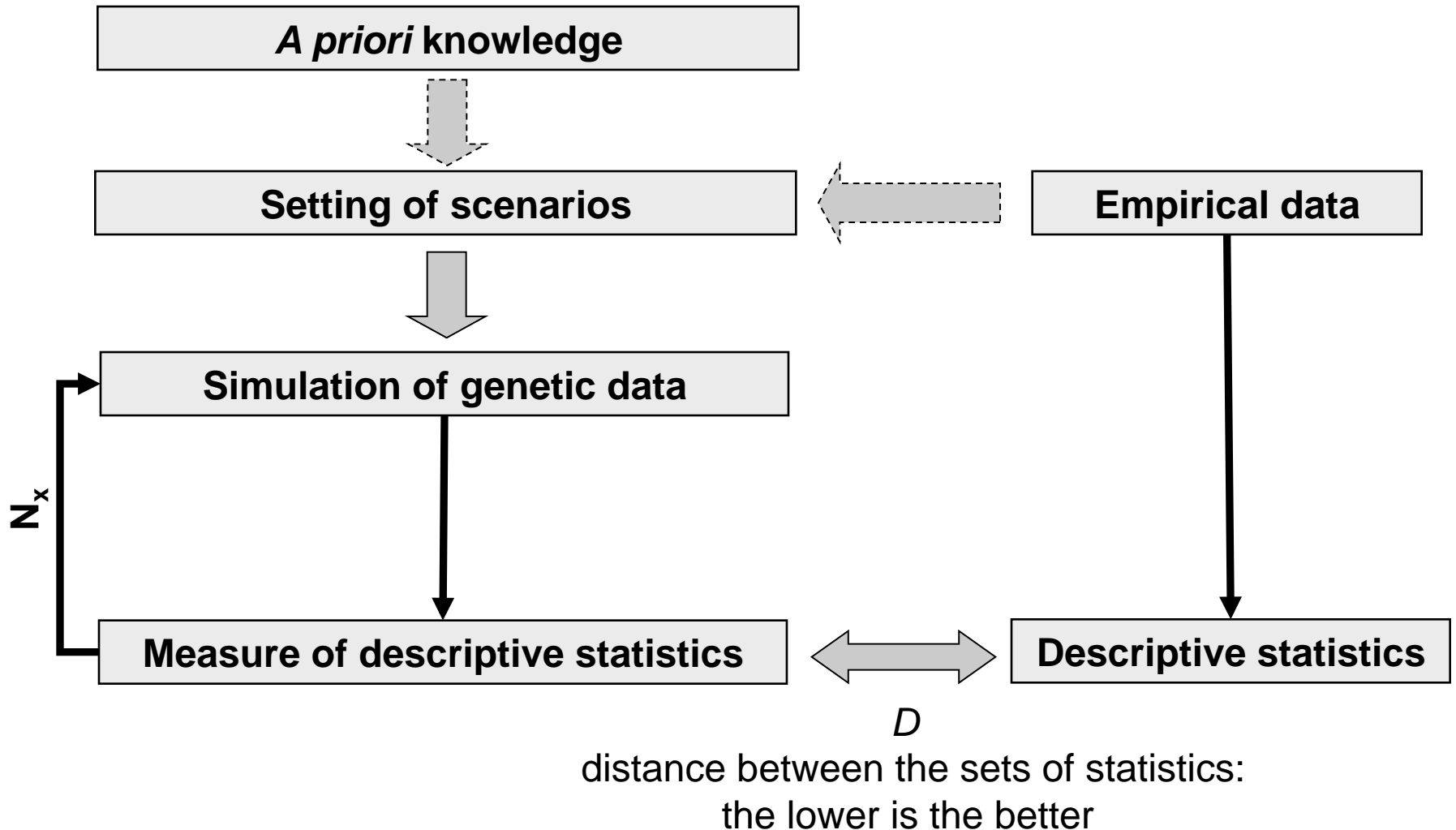


Non-native scenarios: 3 models





SIMULATION AND MODEL COMPARISON: « ABC » APPROACH

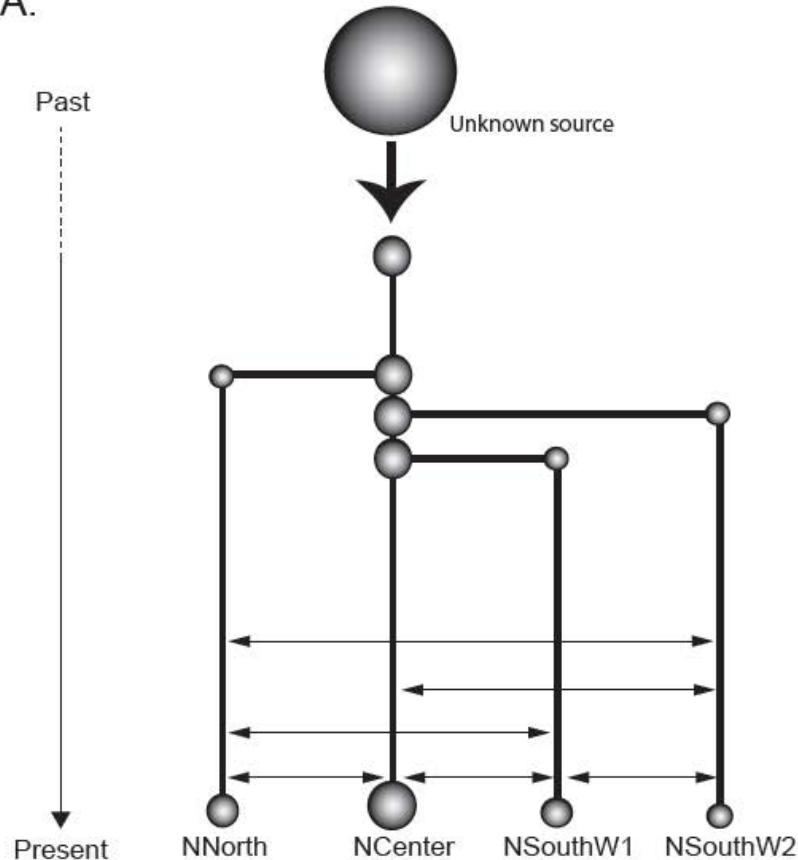


(I) PATHWAYS OF INVASION

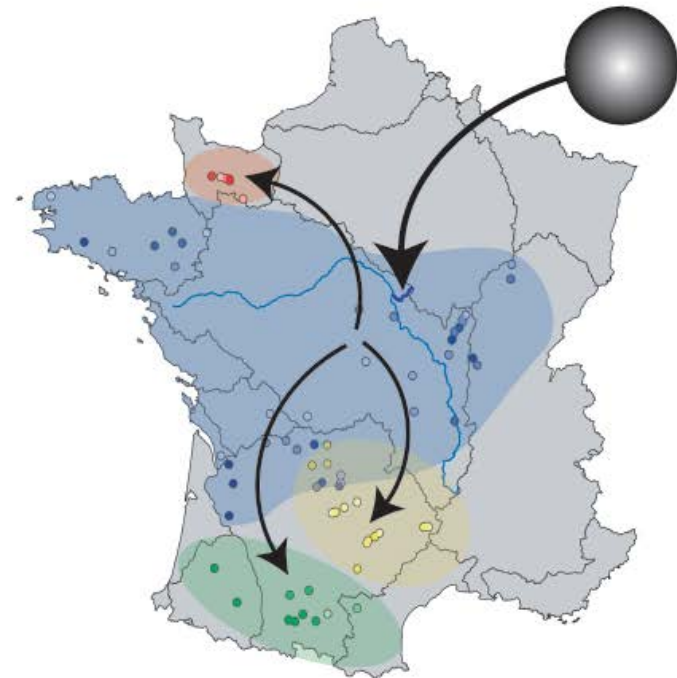


→ One of the « non-native » scenario is most likely:

A.



B.

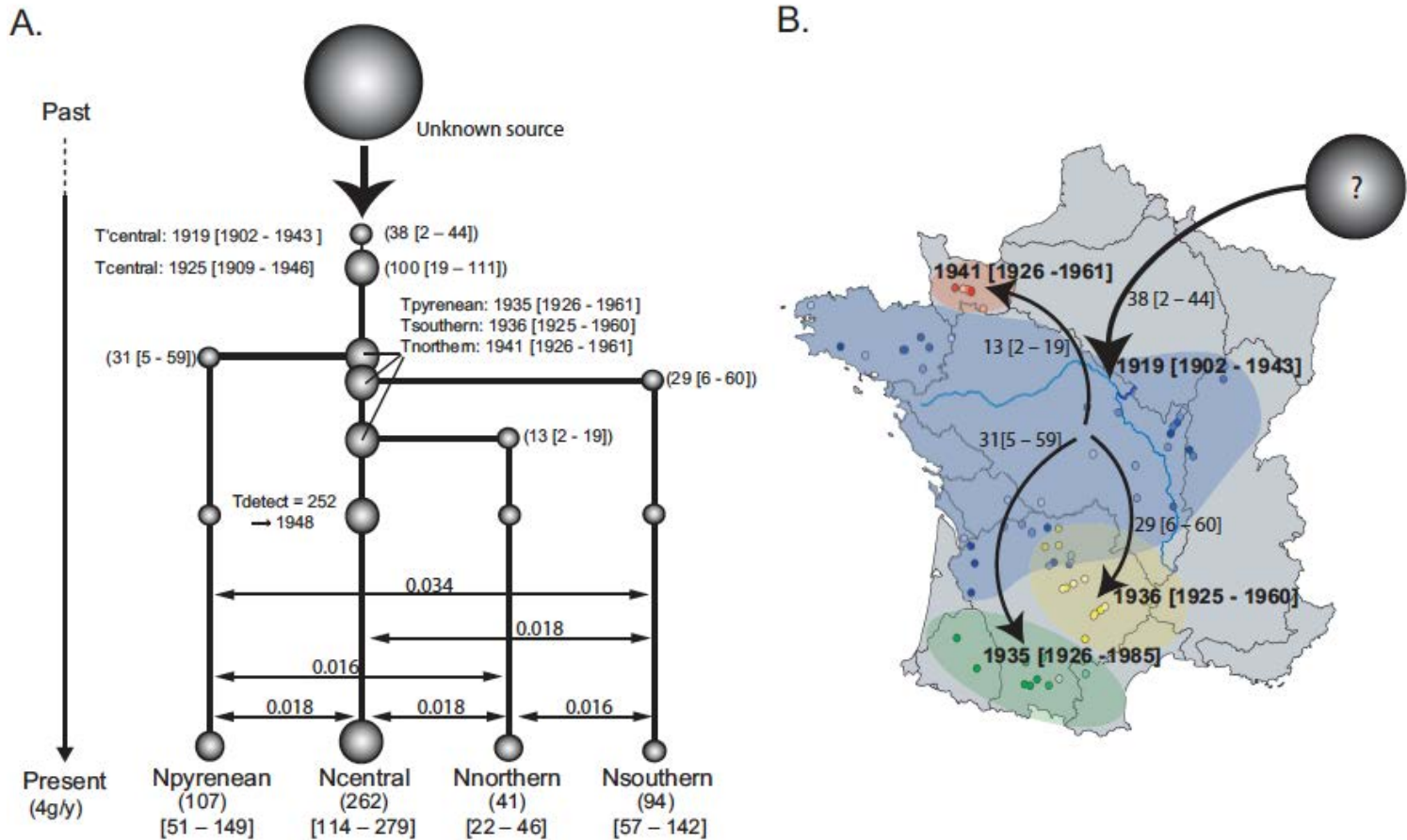


Tracheliastes has been introduced in Center France in the Loire river from a single unknown source, then spread to other river basins

(I) PATHWAYS OF INVASION

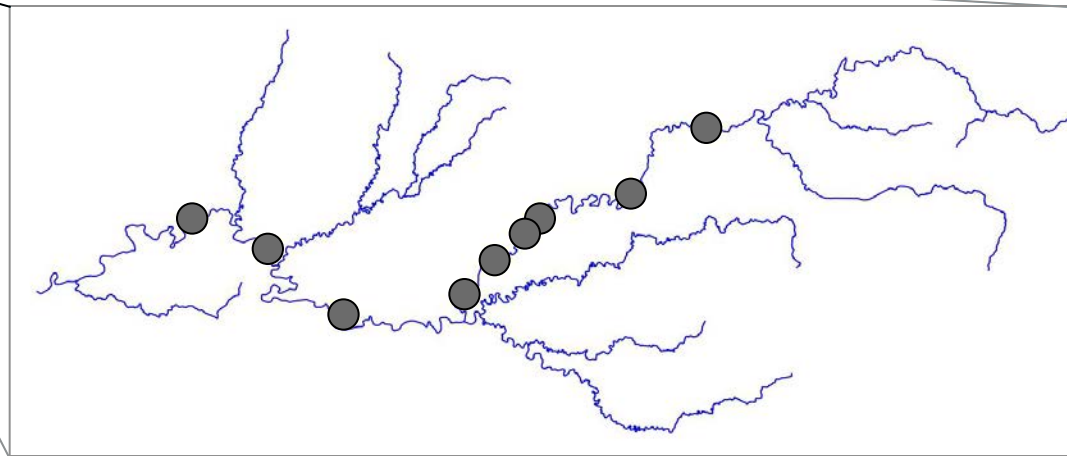
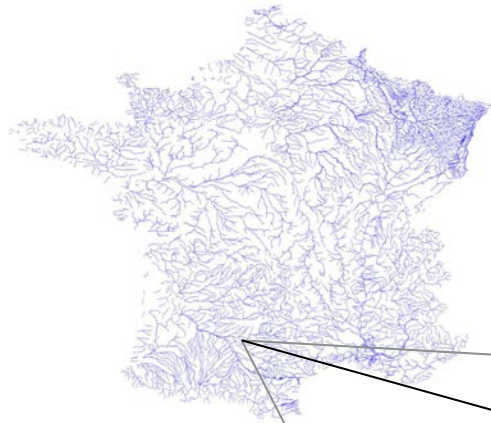


→ Further simulations are ran to infer key parameter values:



Tracheliastes first arrived in the 1920's (observed in the UK in 1930) from a few numbers of founders (20). It quickly colonised other river basins, i.e. 15 years later.

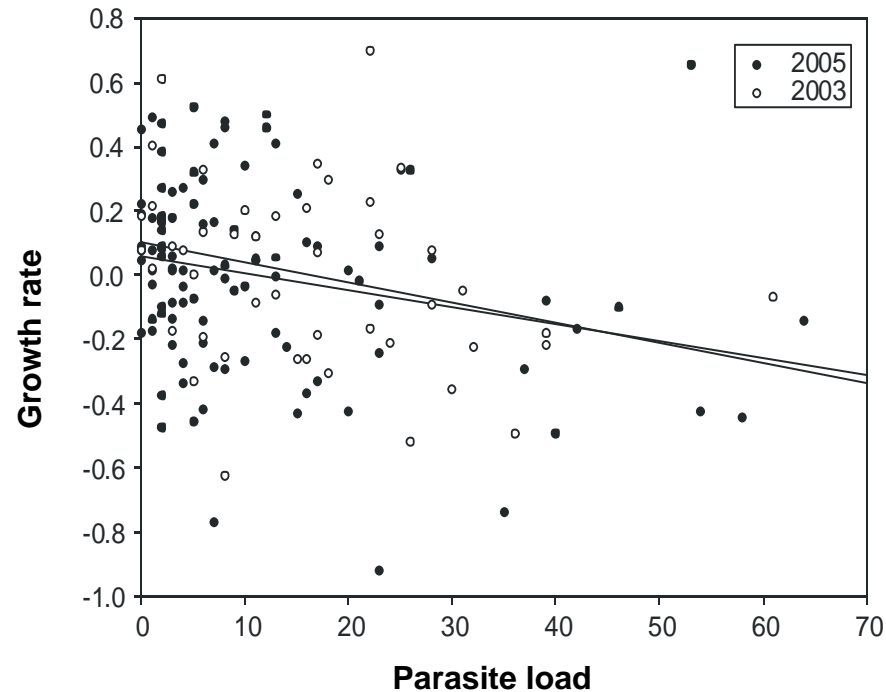
LONG-TERM SURVEY ON NATURAL POPULATIONS: VIAUR AND CÉLÉ RIVERS



**8 sampling sites from 2003 to 2018 x 2 rivers:
Tissue samples– biometry – parasitic indices**



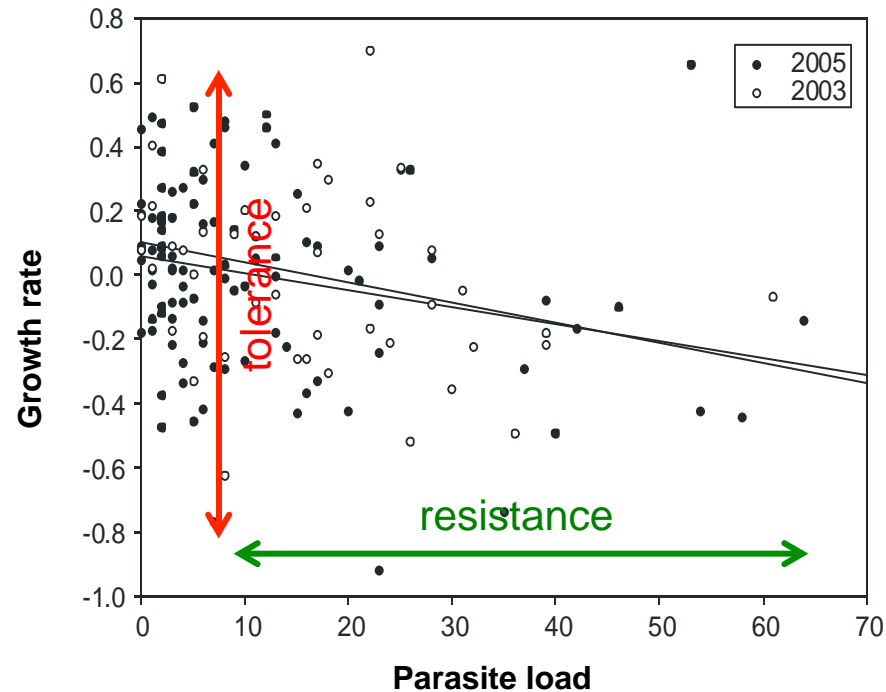
LONG-TERM SURVEY ON NATURAL POPULATIONS: VIAUR AND CÉLÉ RIVERS



- The parasite affects the dace individual fitness (evidences for population effects – selective mortality)



LONG-TERM SURVEY ON NATURAL POPULATIONS: VIAUR AND CÉLÉ RIVERS



- The parasite affects the dace individual fitness (evidences for population effects – selective mortality)
- Large inter-individual variance in terms of resistance and tolerance to the parasite



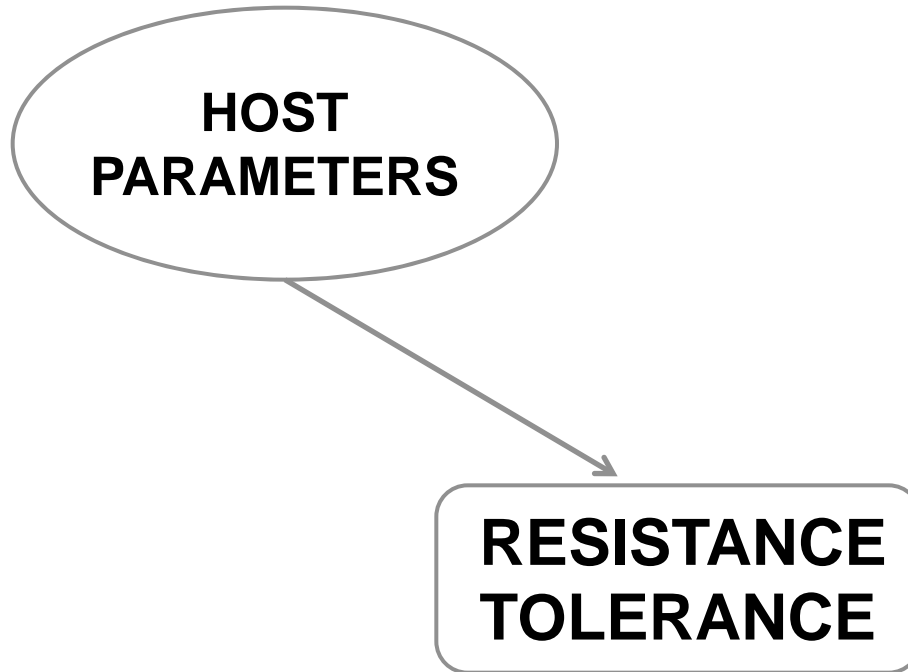
GENETIC AND ENVIRONMENTAL BASES



Maxime Cardon



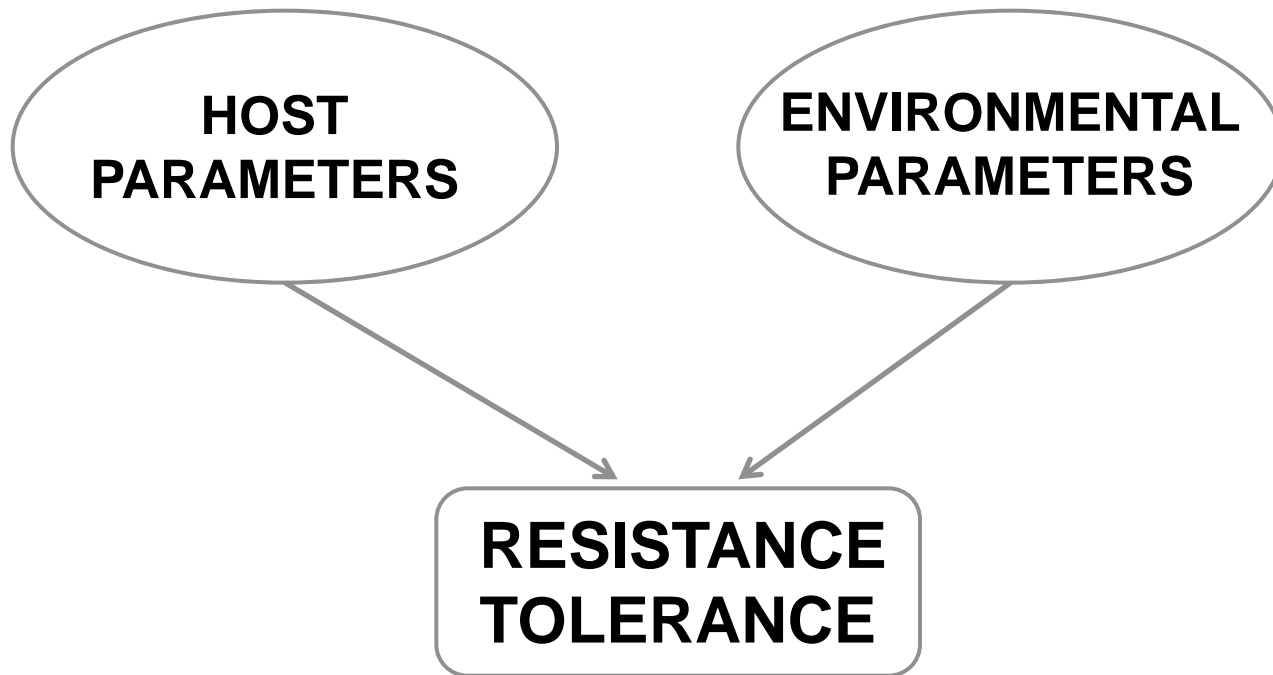
GENETIC AND ENVIRONMENTAL BASES



1) "HOST" MODEL



GENETIC AND ENVIRONMENTAL BASES

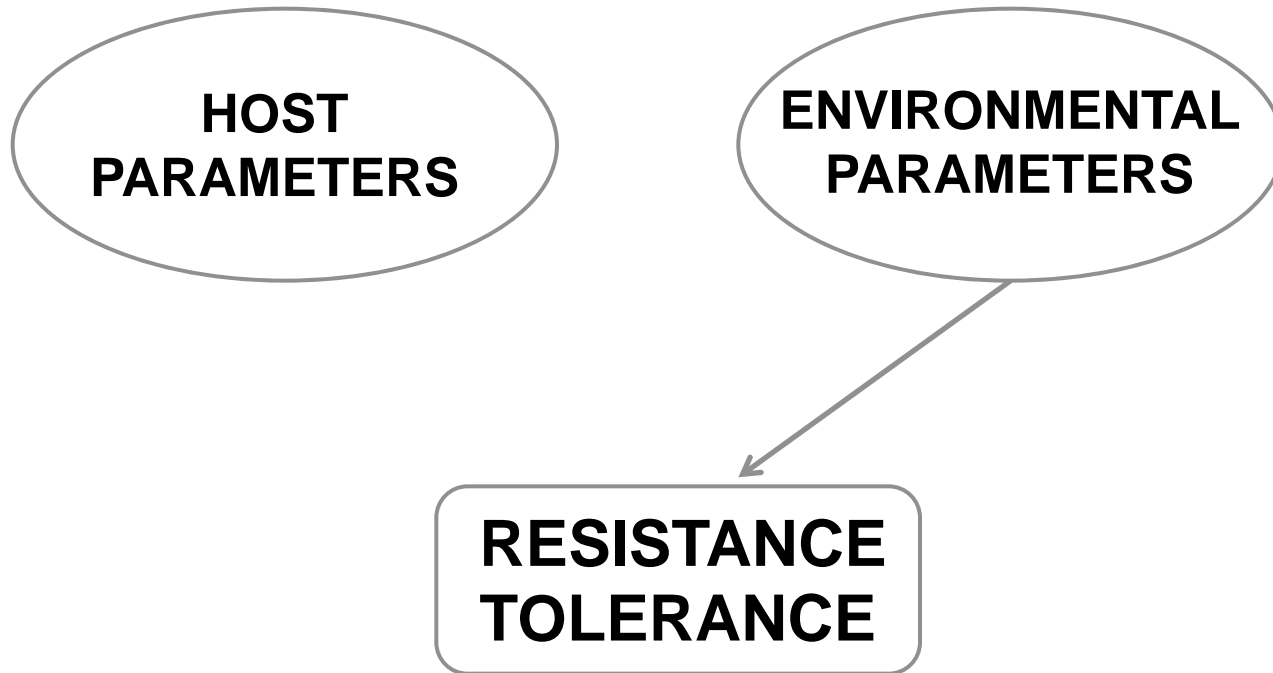


1) "HOST" MODEL

2) "HOST – ENVIRONMENT" MODEL



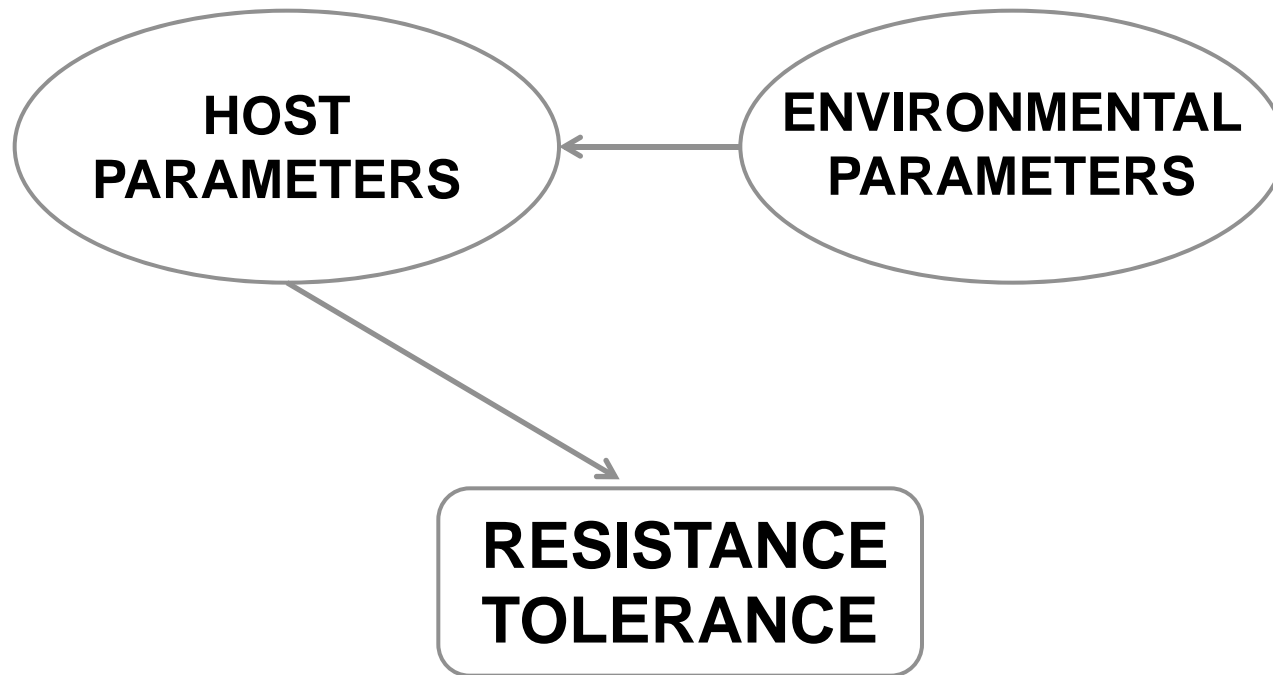
GENETIC AND ENVIRONMENTAL BASES



- 1) "HOST" MODEL
- 2) "HOST – ENVIRONMENT" MODEL
- 3) "DIRECT ENVIRONMENT" MODEL



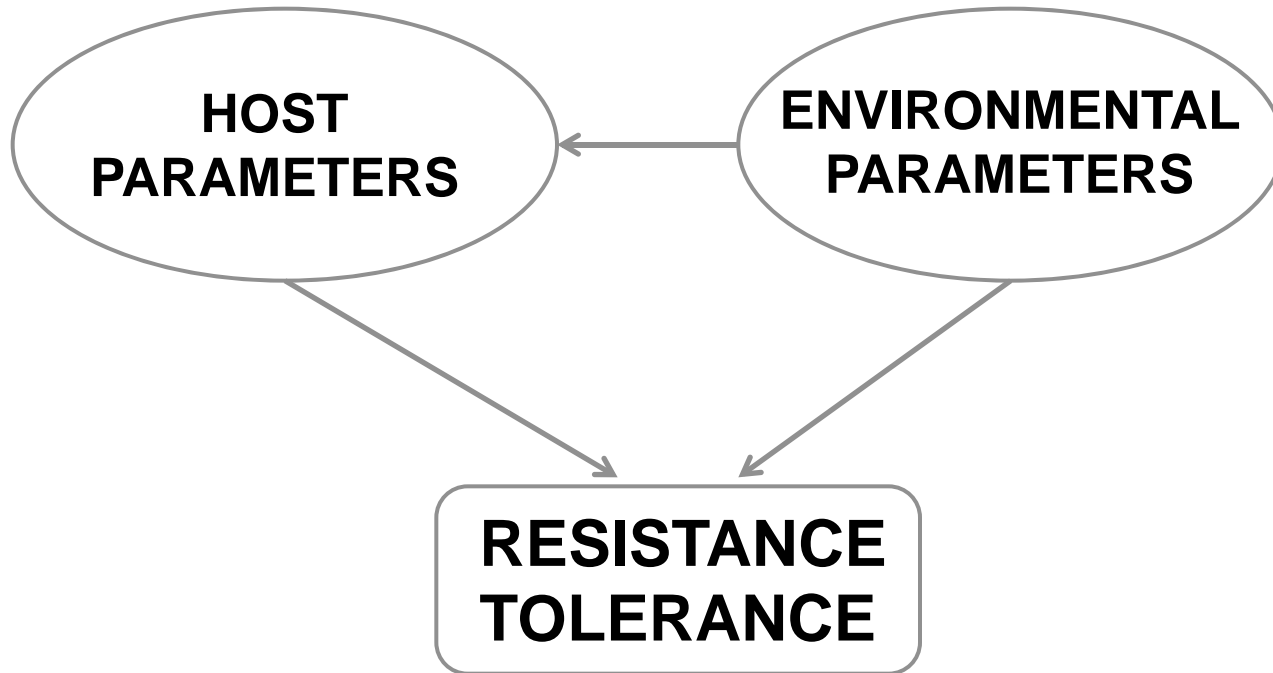
GENETIC AND ENVIRONMENTAL BASES



- 1) "HOST" MODEL
- 2) "HOST – ENVIRONMENT" MODEL
- 3) "DIRECT ENVIRONMENT" MODEL
- 4) "INDIRECT ENVIRONMENT" MODEL



GENETIC AND ENVIRONMENTAL BASES



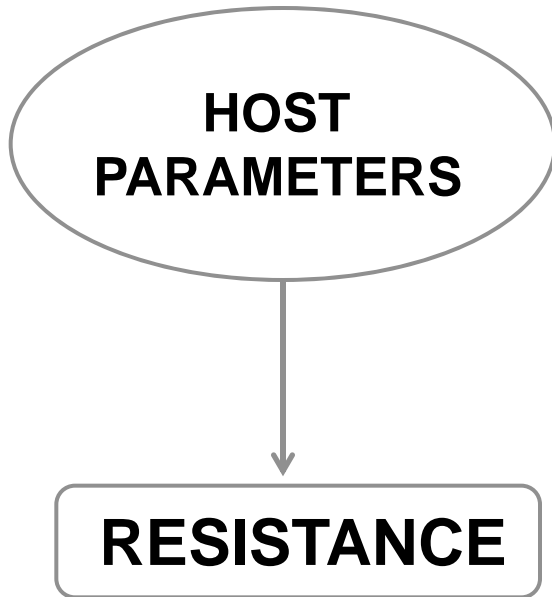
- 1) "HOST" MODEL
- 2) "HOST – ENVIRONMENT" MODEL
- 3) "DIRECT ENVIRONMENT" MODEL
- 4) "INDIRECT ENVIRONMENT" MODEL
- 5) "HOST – DIRECT + INDIRECT ENVIRONMENT" MODEL

SEM
+
AIC



GENETIC AND ENVIRONMENTAL BASES

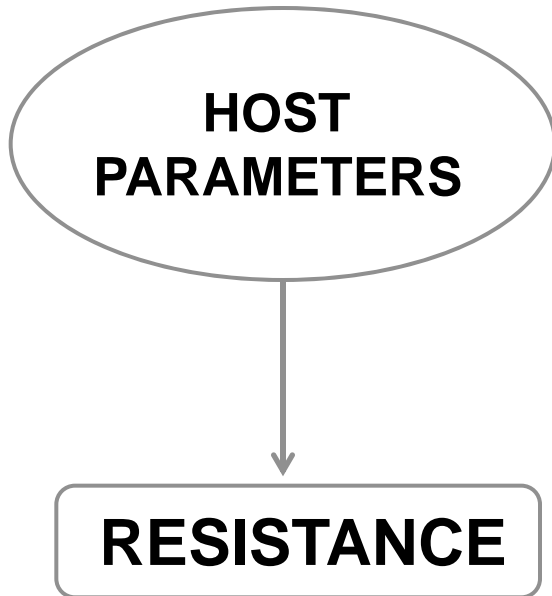
“HOST” MODELS



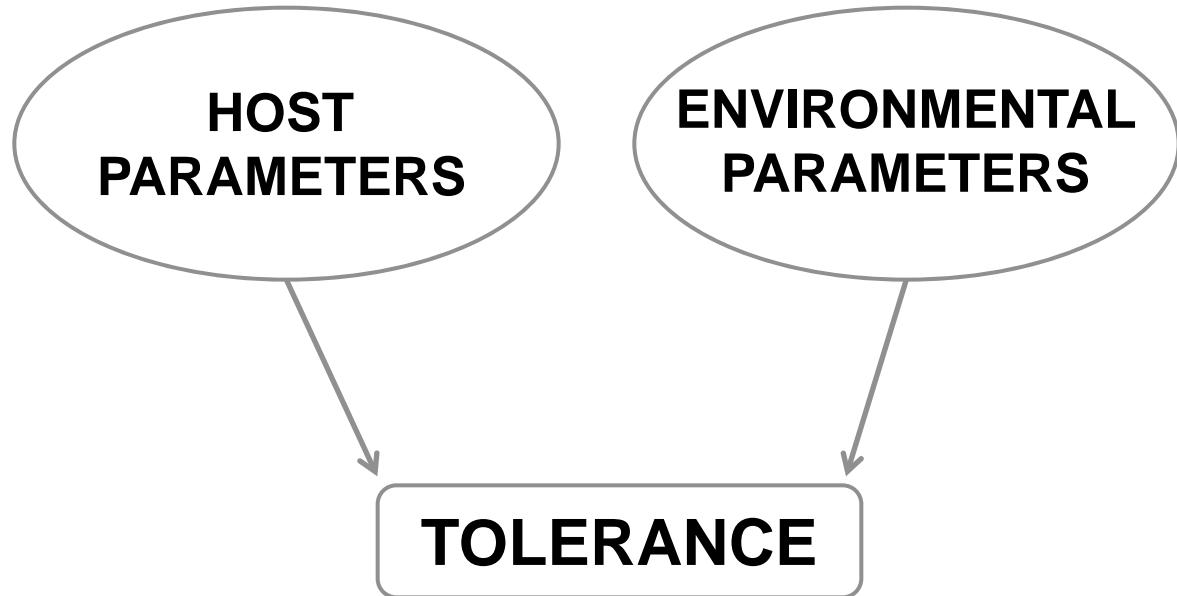


GENETIC AND ENVIRONMENTAL BASES

“HOST” MODELS



“HOST + ENVIRONMENT” MODELS



- ENVIRONMENTAL BASES FOR TOLERANCE (thermal regime)
- GENETIC BASES FOR BOTH RESISTANCE AND TOLERANCE?



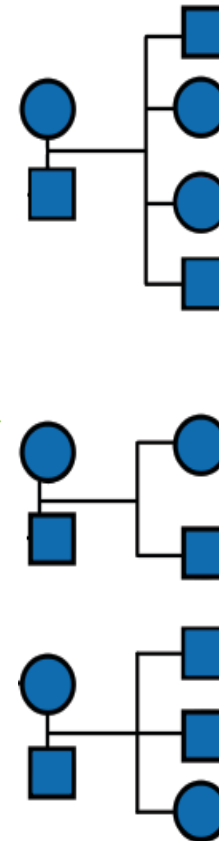
GENETIC BASES (HERITABLE)

Molecular data => inferring the parental links (pedigrees)



Elise Mazé

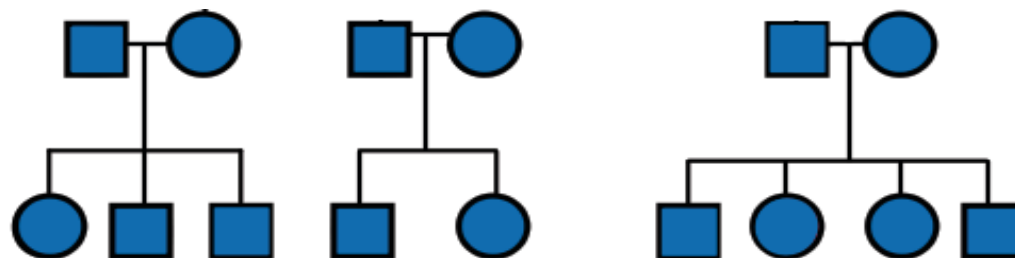
Job Name	Date	Status	Commands
2012-2-14 11 15 4720locus	Thu May 31 15:26:30 2012	100.0% done in 168. min	View details Delete
2012-2-17 15 15 0813marquenas	Thu May 31 15:26:31 2012	100.0% done in 162. min	View details Delete
2012-2-24 16 23 15globalbis	Thu May 31 15:26:32 2012	100.0% done in 5085. min	View details Delete
2012-2-24 16 26 110markeur	Thu May 31 15:26:48 2012	100.0% done in 989. min	View details Delete
2012-2-28 12 36 25global2loc	Thu May 31 15:26:50 2012	100.0% done in 329. min	View details Delete
2012-2-28 12 37 43	Thu May 31 15:26:54 2012	100.0% done in 327. min	View details Delete
2012-2-28 18 06 49globalrebis	Thu May 31 15:26:58 2012	100.0% done in 838. min	View details Delete
2012-2-8 12 52 54	Thu May 31 15:27:07 2012	100.0% done in 195. min	View details Delete
2012-3-14 10 12 23noFSC-W5	Thu May 31 15:27:09 2012	100.0% done (now in run 10. 0. min to go)	View details Delete
2012-3-15 13 00 44	Thu May 31 15:27:13 2012	100.0% done (now in run 10. 0. min to go)	View details Delete
2012-3-15 13 02 12restart-FSC vers noFSC	Thu May 31 15:27:17 2012	100.0% done (now in run 10. 0. min to go)	View details Delete
2012-3-1 10 32 56globalFSC	Thu May 31 15:27:22 2012	100.0% done in 1018. min	View details Delete
2012-3-1 10 37 01global-noFSC	Thu May 31 15:27:30 2012	100.0% done in 753. min	View details Delete
2012-3-1 17 58 39weigh	Thu May 31 15:27:35 2012	100.0% done in 1021. min	View details Delete
2012-3-2 16 09 58W5-FSC-restart-temp	Thu May 31 15:28:01 2012	100.0% done (now in run 10. 0. min to go)	View details Delete
2012-3-2 16 11 55W10-FSC-restart-temp	Thu May 31 15:28:09 2012	100.0% done (now in run 10. 0. min to go)	View details Delete



~ 250 FULL-SIB FAMILIES (2-4 INDIV / FAM)



GENETIC BASES (HERITABLE)



RESISTANCE

1 2 1 10 14

7 5 3 6

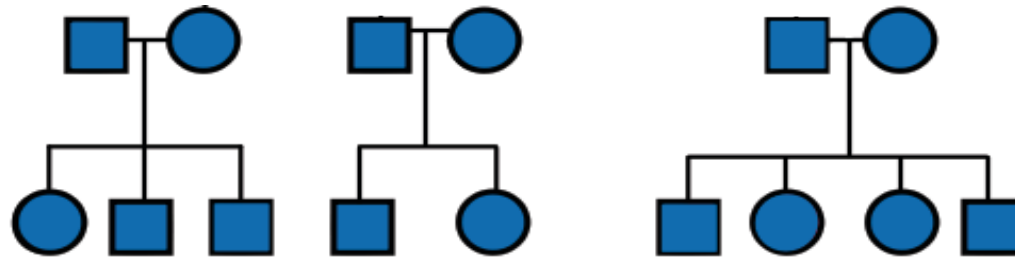
TOLERANCE

9 6 7 3 5

2 6 6 9



GENETIC BASES (HERITABLE)



RESISTANCE	1	2	1	10	14	7	5	3	6
TOLERANCE	9	6	7	3	5	2	6	6	9

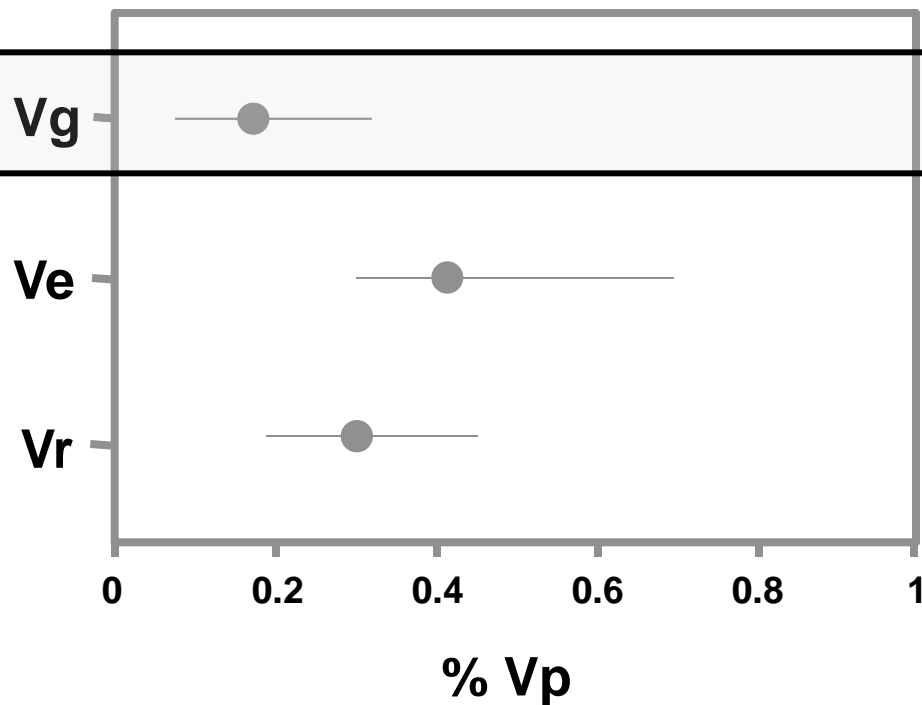
“ANIMAL MODELS” :

$$V_p = V_g + V_e + V_r$$

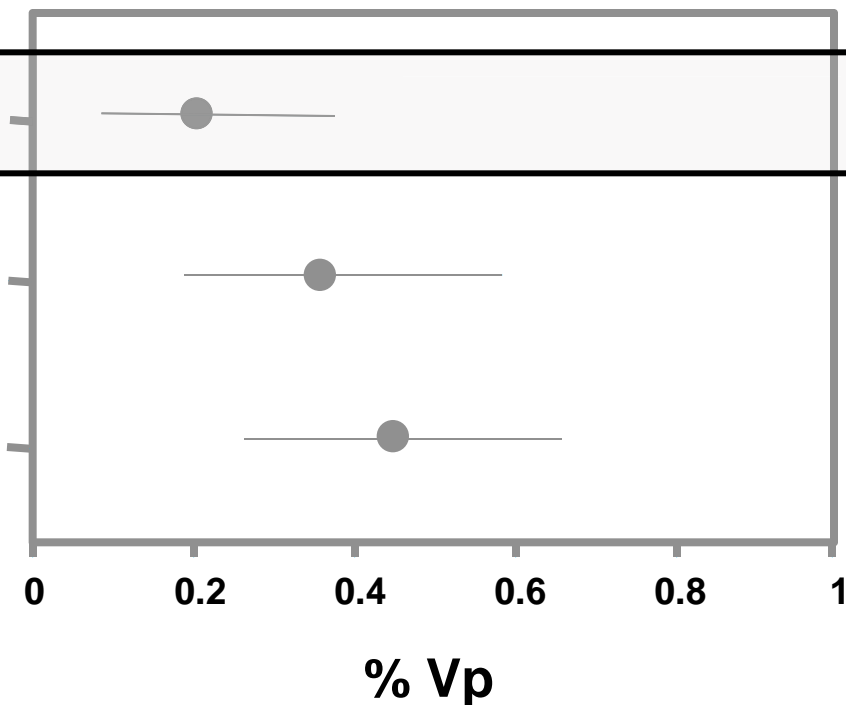
$$\Rightarrow H^2 = V_g / V_p$$



RESISTANCE



TOLERANCE



- SIGNIFICANT HERITABILITY VALUES FOR BOTH TRAITS
 - GENETIC BASES FOR BOTH DEFENSIVE TRAITS
- BASES FOR HOST ADAPTATION TO THIS NEW SELECTIVE PRESSURE



GOING INTO THE PHYSIOLOGICAL PATHWAYS OF RESISTANCE



Eglantine
Mathieu-Bégné



GOING INTO THE PHYSIOLOGICAL PATHWAYS OF RESISTANCE



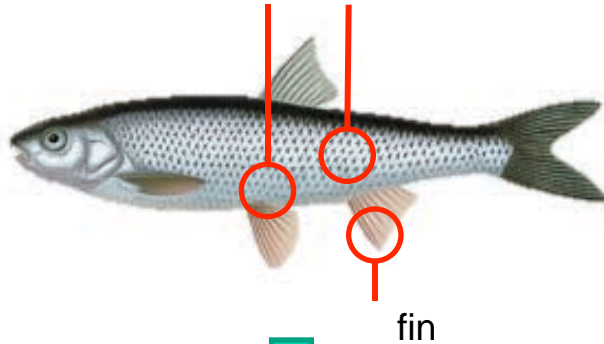
Eglantine
Mathieu-Bégné



4 healthy vs.
4 parasitized

kidney

spleen



fin

RNA-seq



Gene expression profile
Biological functions



GOING INTO THE PHYSIOLOGICAL PATHWAYS OF RESISTANCE



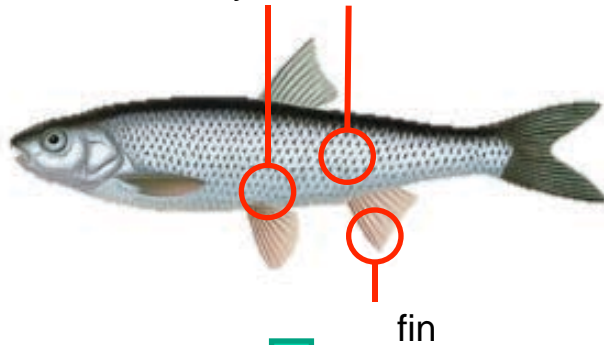
Eglantine
Mathieu-Bégné

4 healthy vs.
4 parasitized



kidney

spleen



fin

RNA-seq



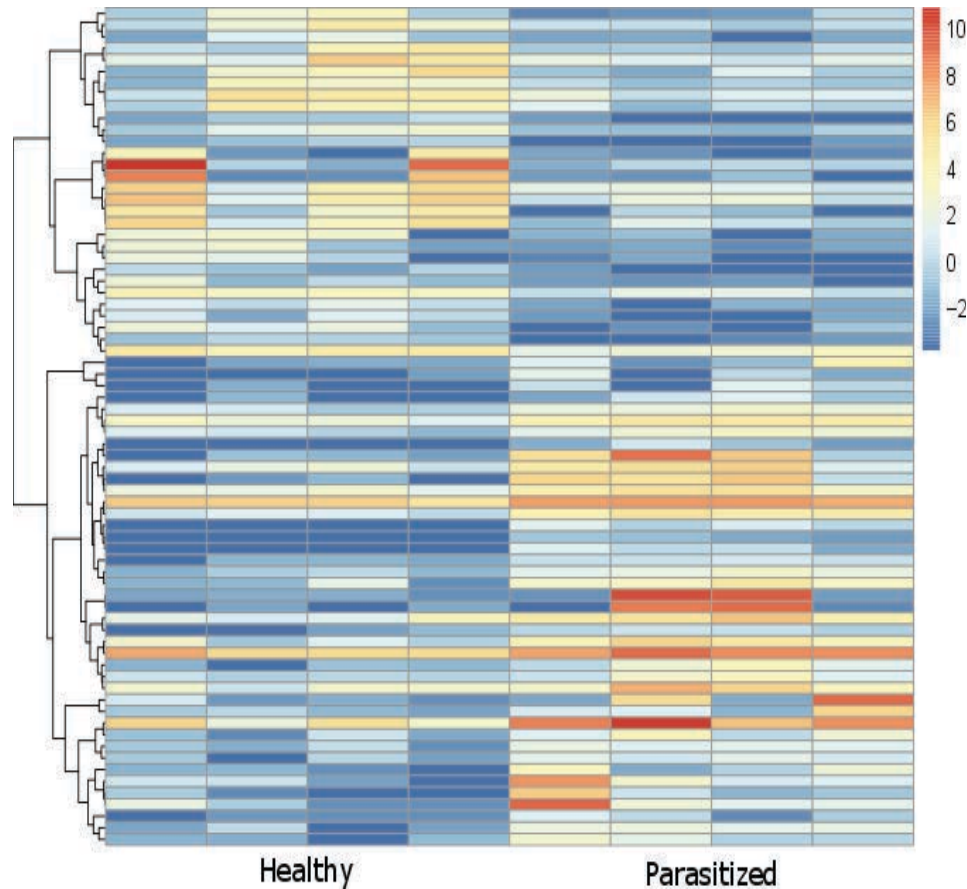
Gene expression profile
Biological functions

Understanding the physiological responses to an ectoparasite:

- **local (fin) response or general response?**
- **If general, is the response tissue-specific?**



Genes expression profile

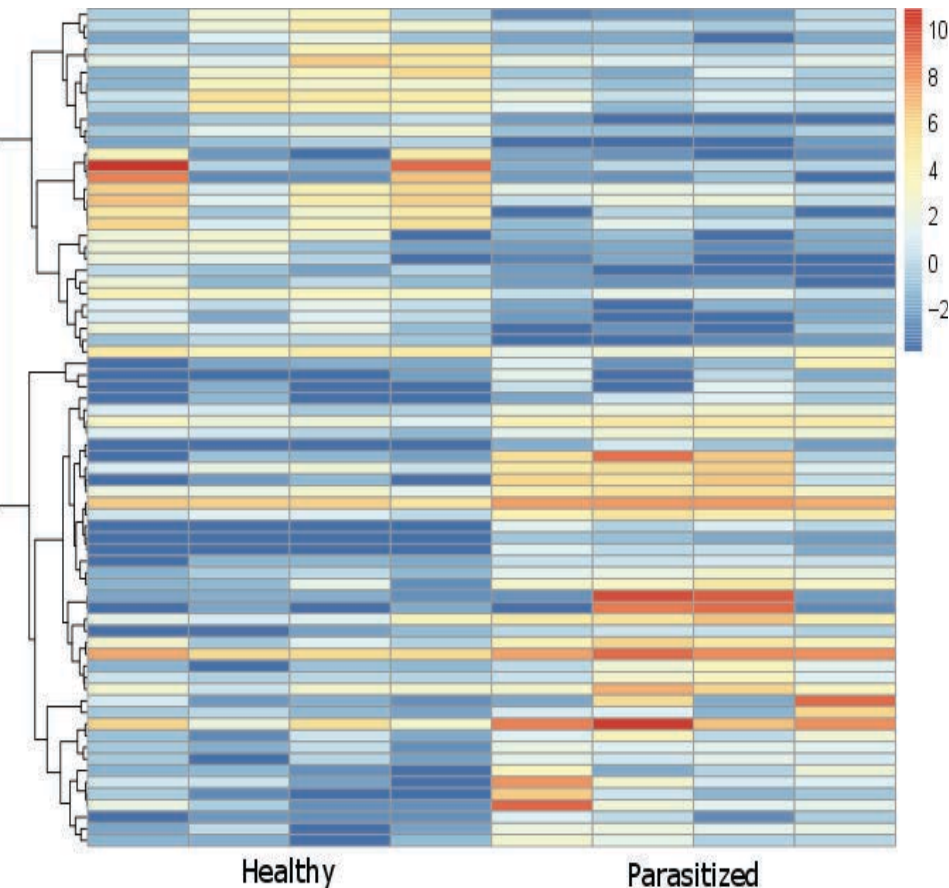


Heat map of genes expression profile:
example of the fin tissue

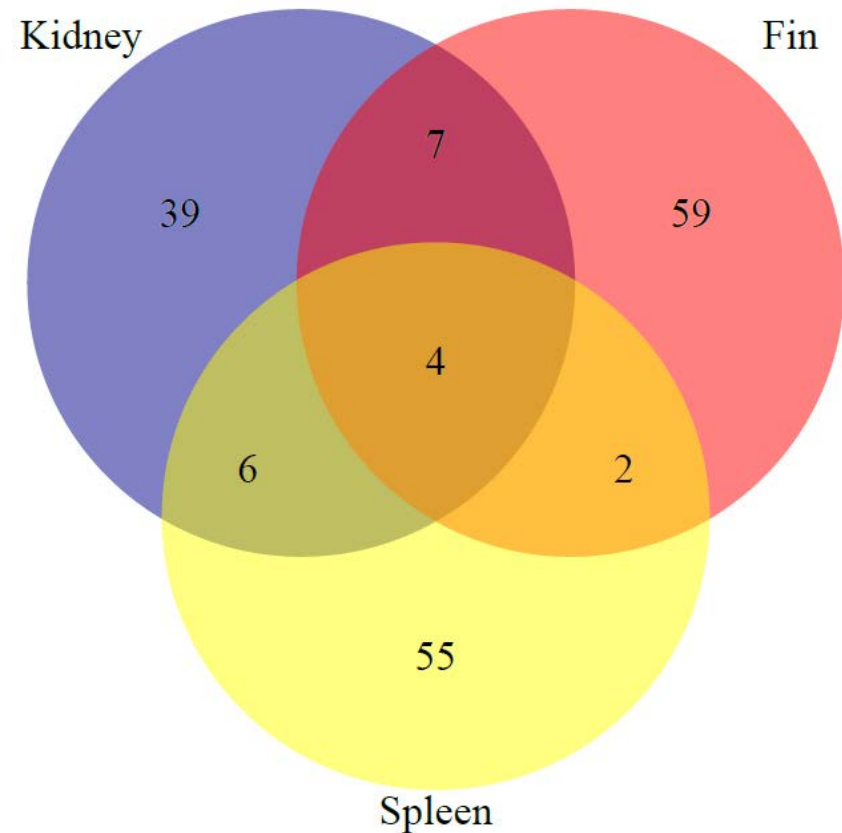
➔ Some genes are differentially expressed depending on the health status



Genes expression profile



Heat map of genes expression profile:
example of the fin tissue

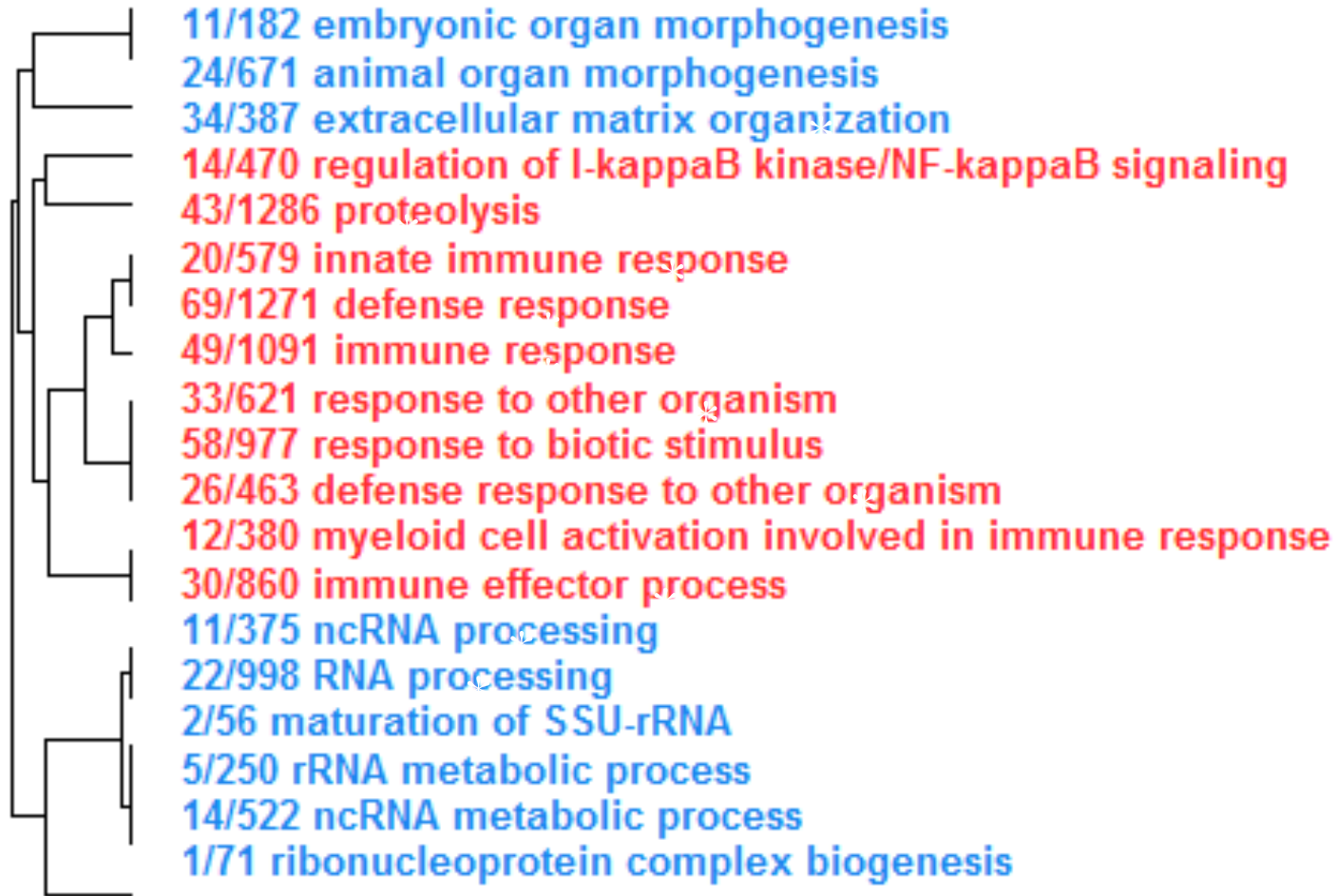


- ➔ Some genes are differentially expressed depending on the health status
- ➔ Host response is tissue specific

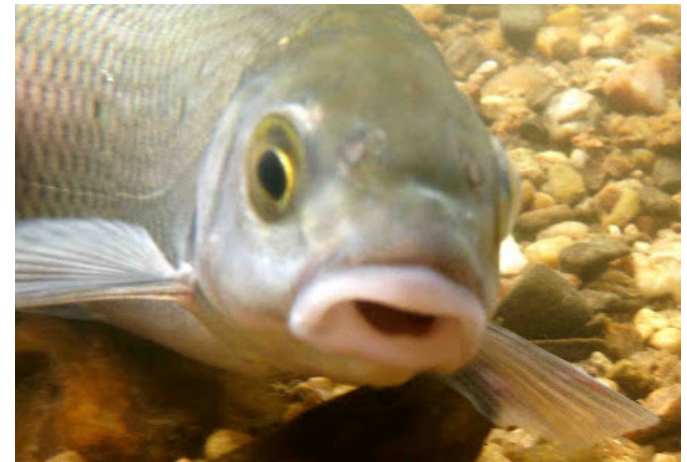


Biological functions involved

Fins



- Generalized immune response in parasitized fishes
- Metabolism/Immunity trade-offs



- ⇒ **A NEWLY INTRODUCED SPECIES (< 200 generations)**
- ⇒ **SIGNIFICANT PATHOGENIC EFFECTS**
- ⇒ **POTENTIAL FOR HOST ADAPTATION**

**PARASITES = NEW SELECTIVE PRESSURE
= REAL-TIME ADAPTATION**



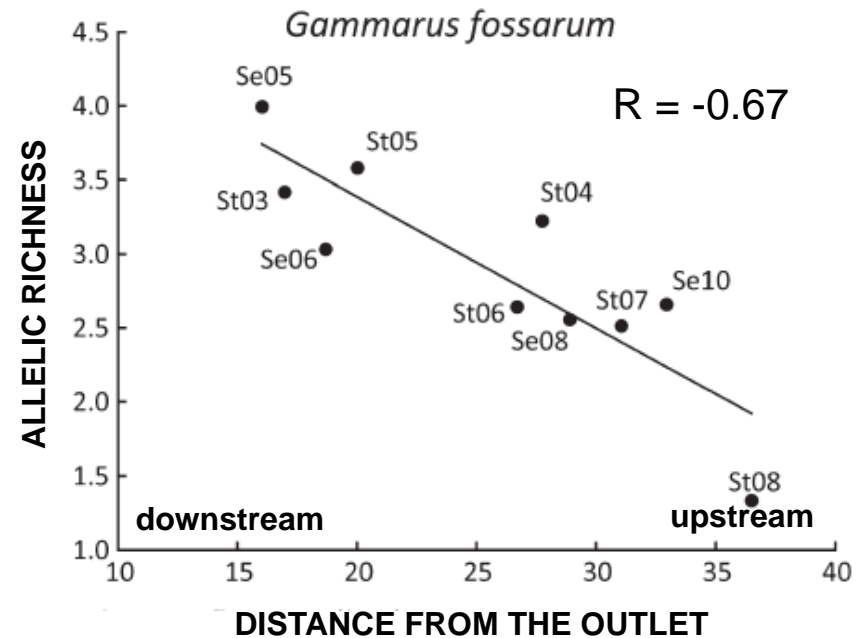
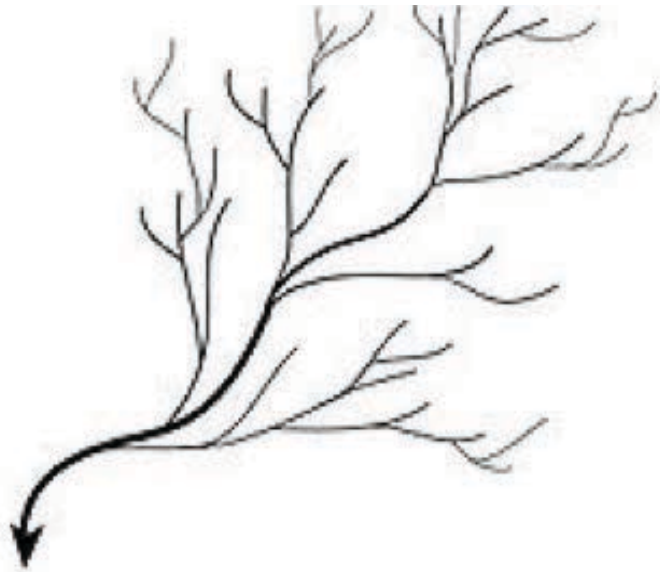
- 1) RIVER NETWORKS AND PATTERNS OF INTRASPECIFIC DIVERSITY**
- 2) ECOLOGICAL CONSEQUENCES OF INTRASPECIFIC DIVERSITY**
- 3) QUANTIFYING THE INFLUENCE OF HABITAT FRAGMENTATION**



Ivan Paz-Vinas

INCREASE IN GENETIC DIVERSITY DOWNSTREAM (IGDD)

Upstream



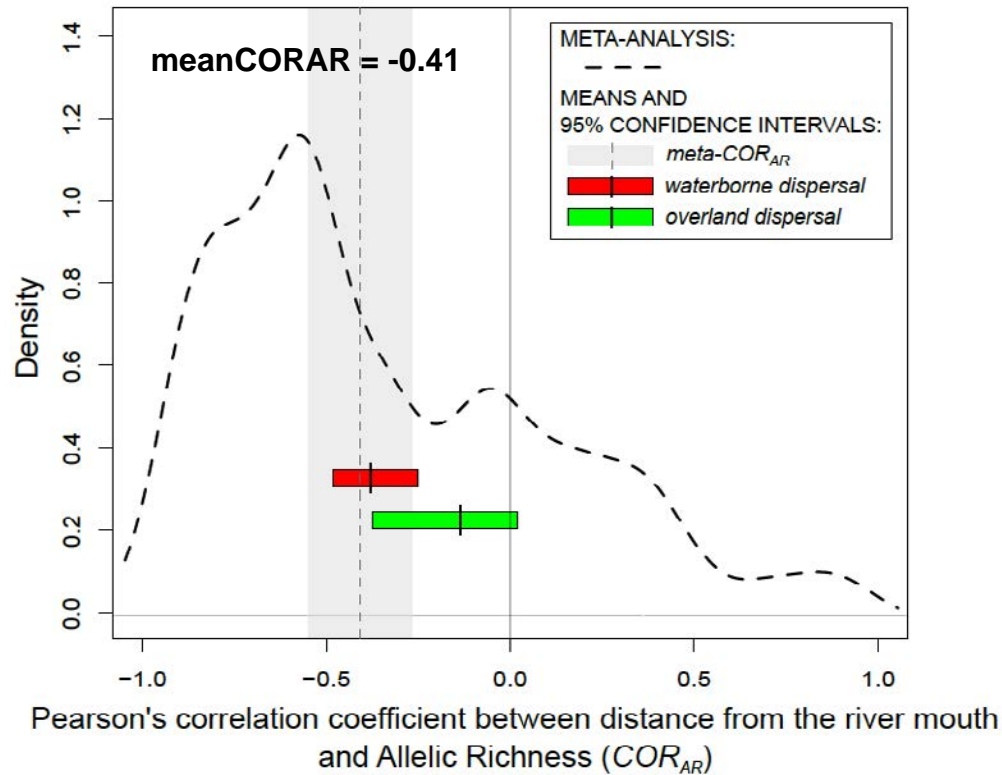
IS THAT A GENERAL PATTERN (RULE) FOUND ACROSS TAXA?



IGDD: A META-ANALYTIC TEST (79 case studies)

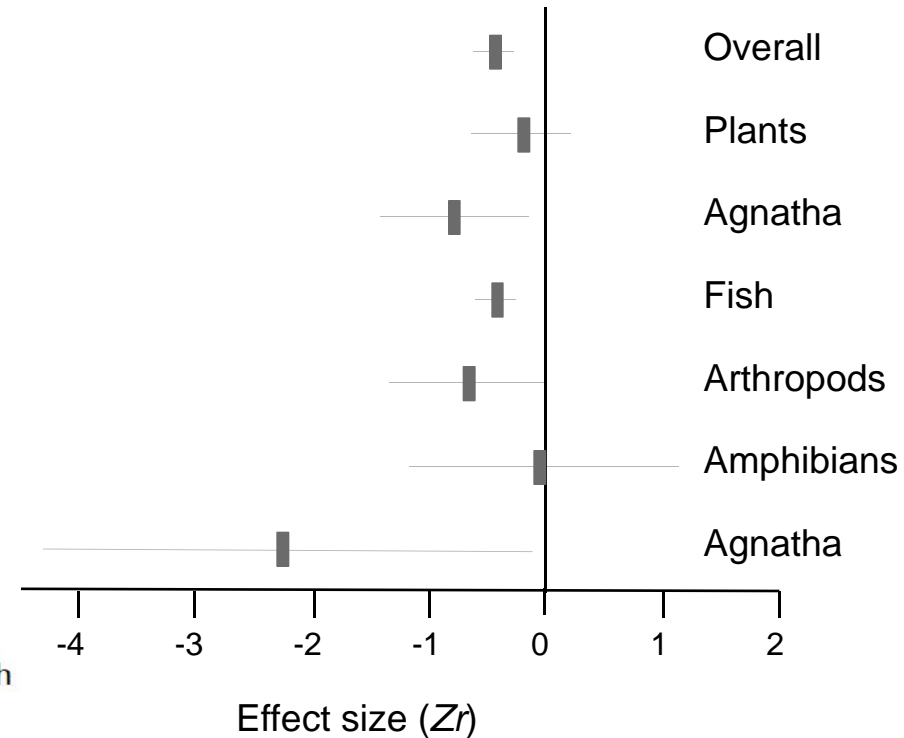
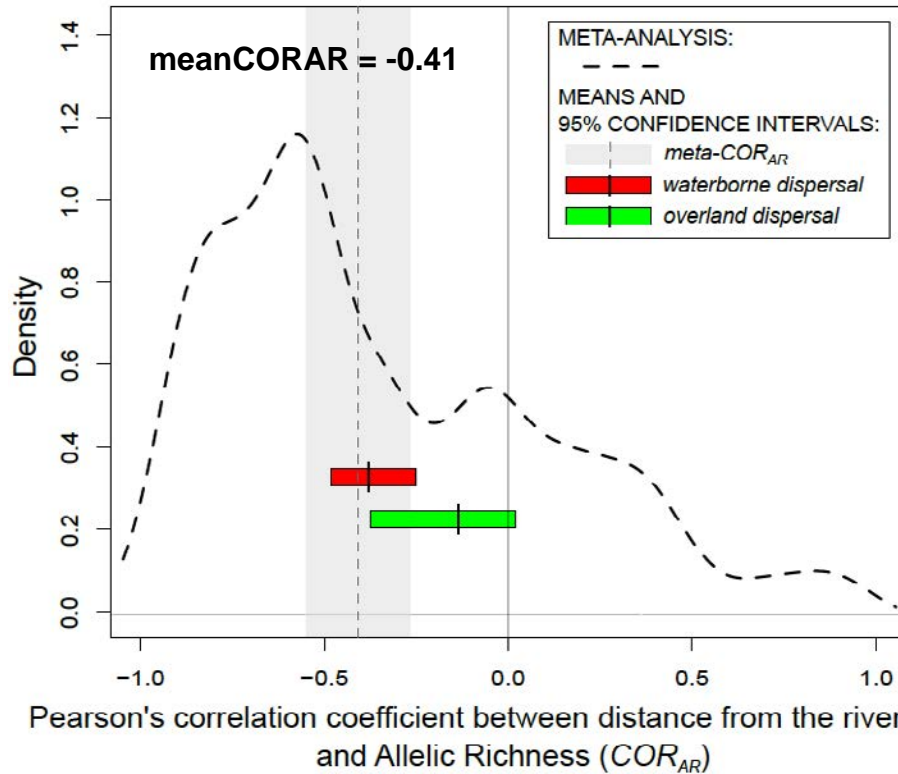


IGDD: A META-ANALYTIC TEST (79 case studies)





IGDD: A META-ANALYTIC TEST (79 case studies)

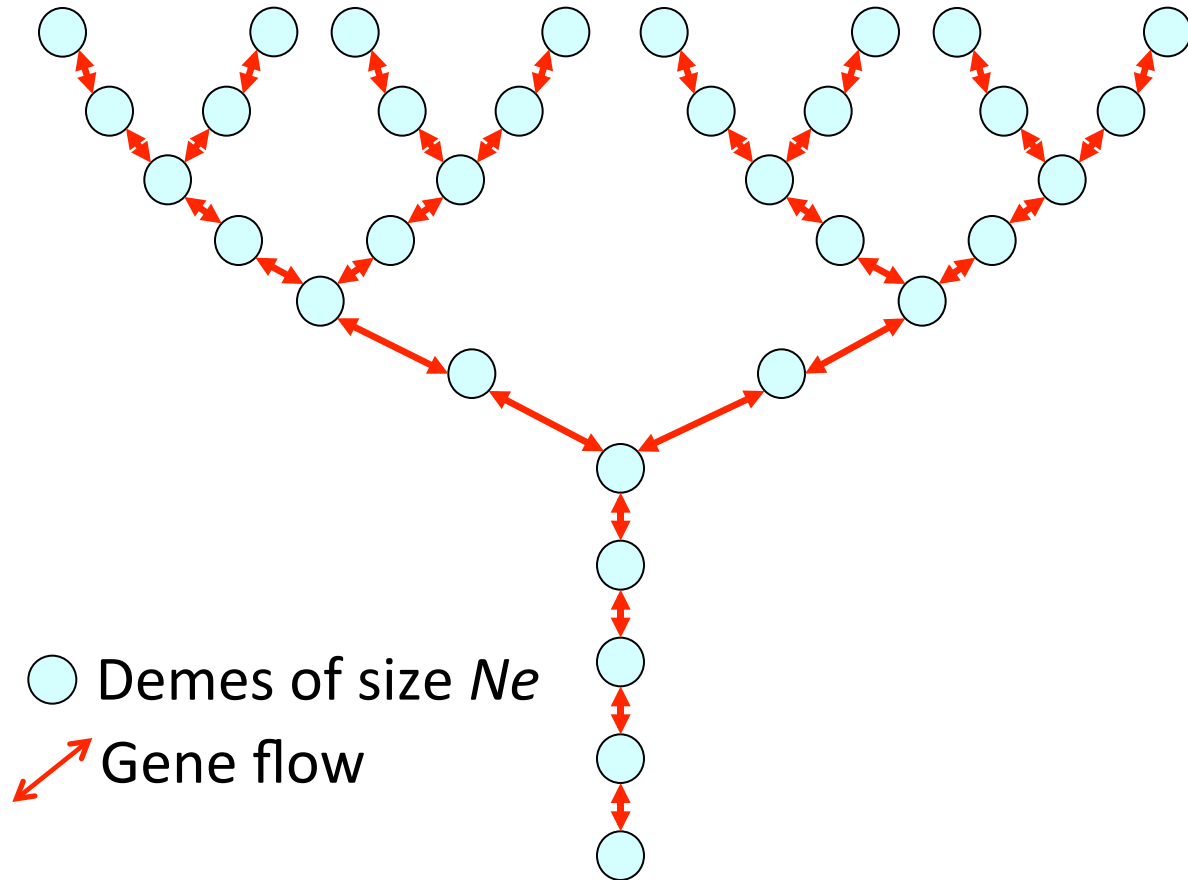


THE PATTERN HOLDS TRUE FOR SEVERAL TAXA

WHAT ARE THE PROCESSES BEHIND THIS GENERAL PATTERN?

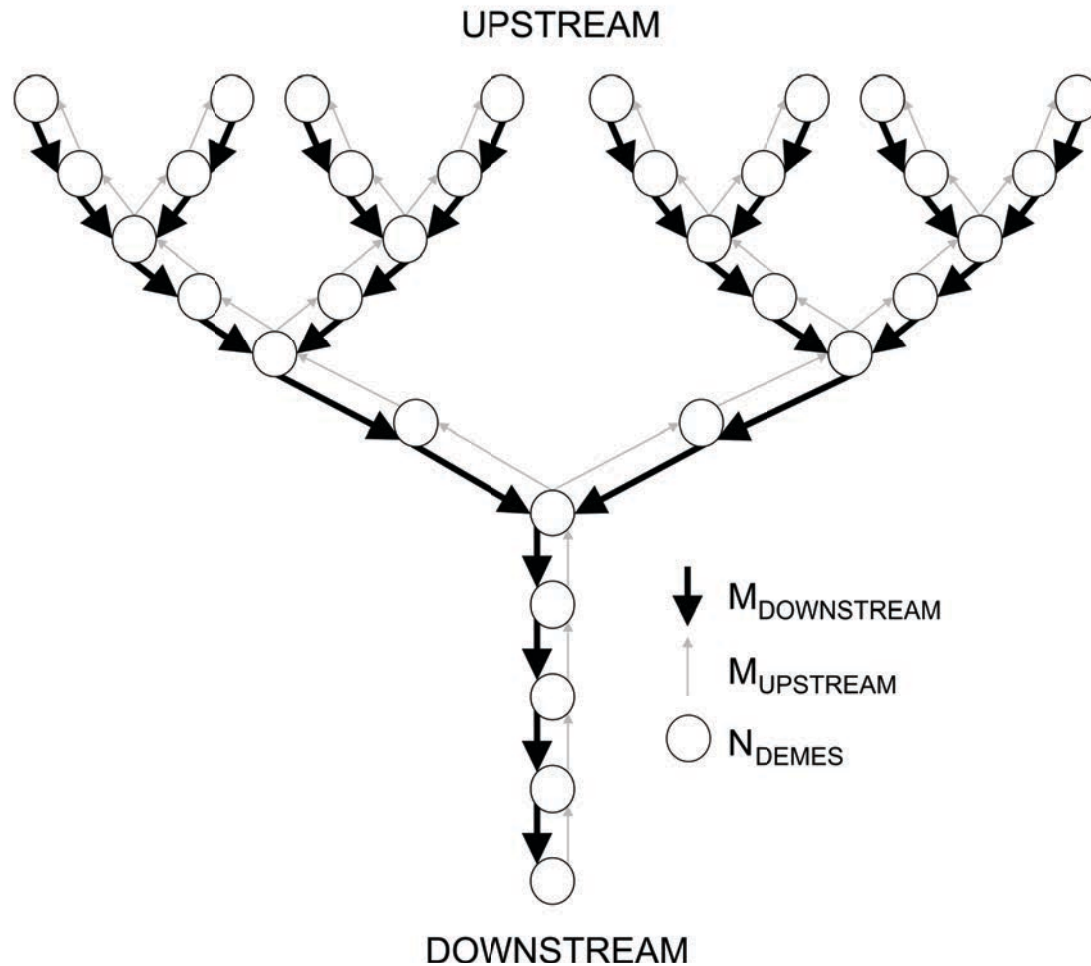


SIMULATIONS OF NEUTRAL GENETIC DIVERSITY IN RIVER NETWORKS





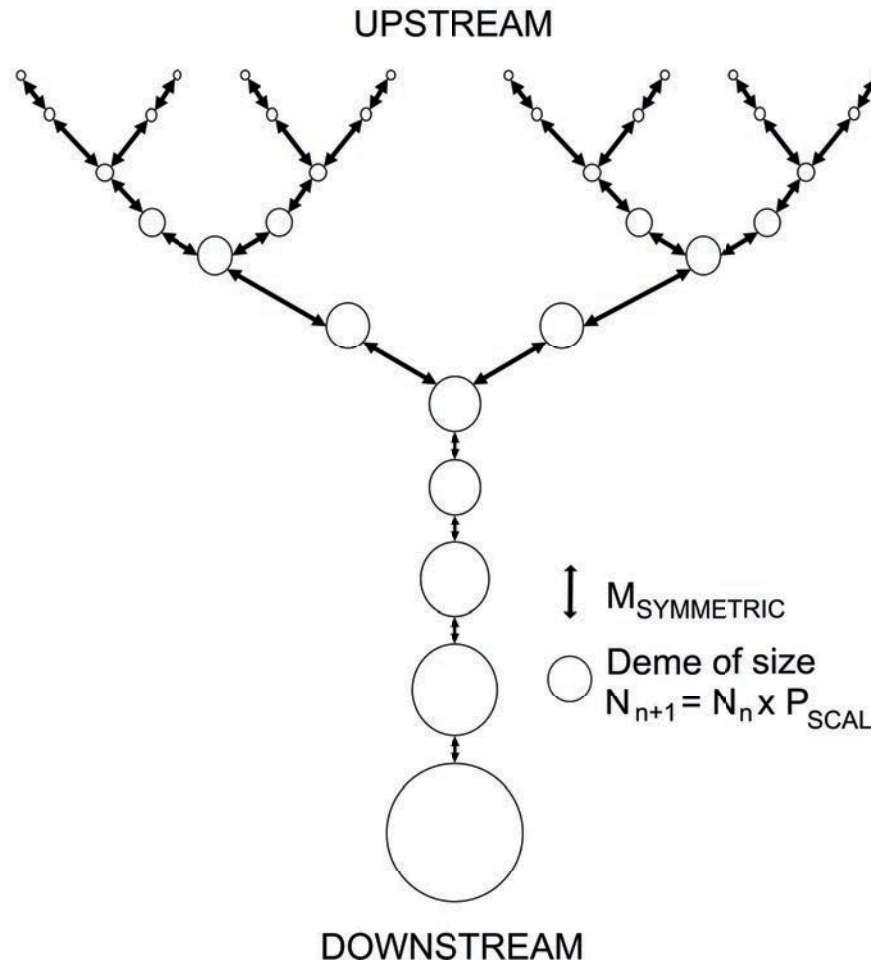
(A) ASYMMETRIC GENE FLOW MODEL



ACCUMULATION OF ALLELES DOWNSTREAM DUE TO ASYMMETRIC GENE FLOW (ASYMM MODEL)



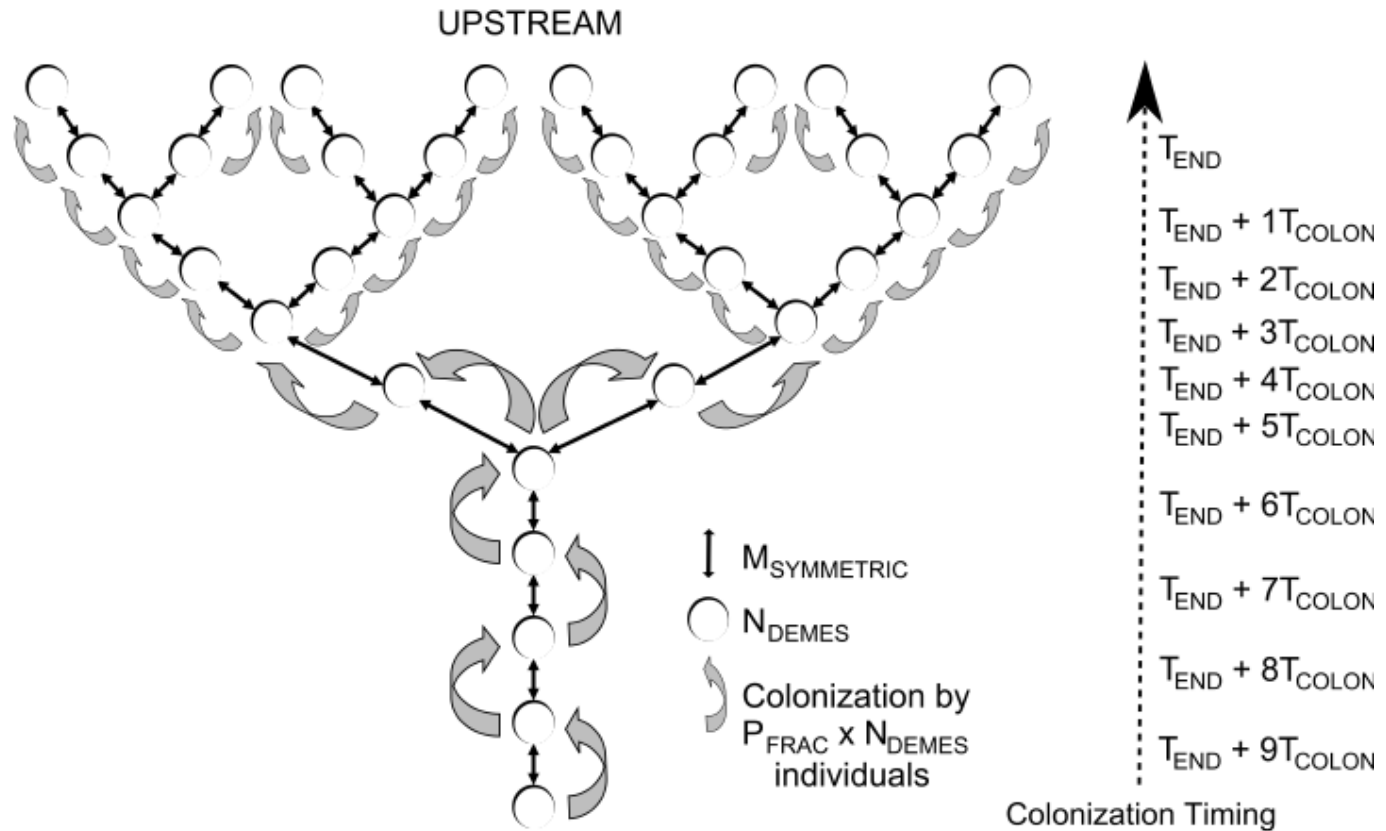
(C) DIFFERENTIAL IN N_e MODEL



HIGHER N_e DOWNSTREAM FAVOUR INCREASE GENETIC DIVERSITY DOWNSTREAM (DIFFNE MODEL)



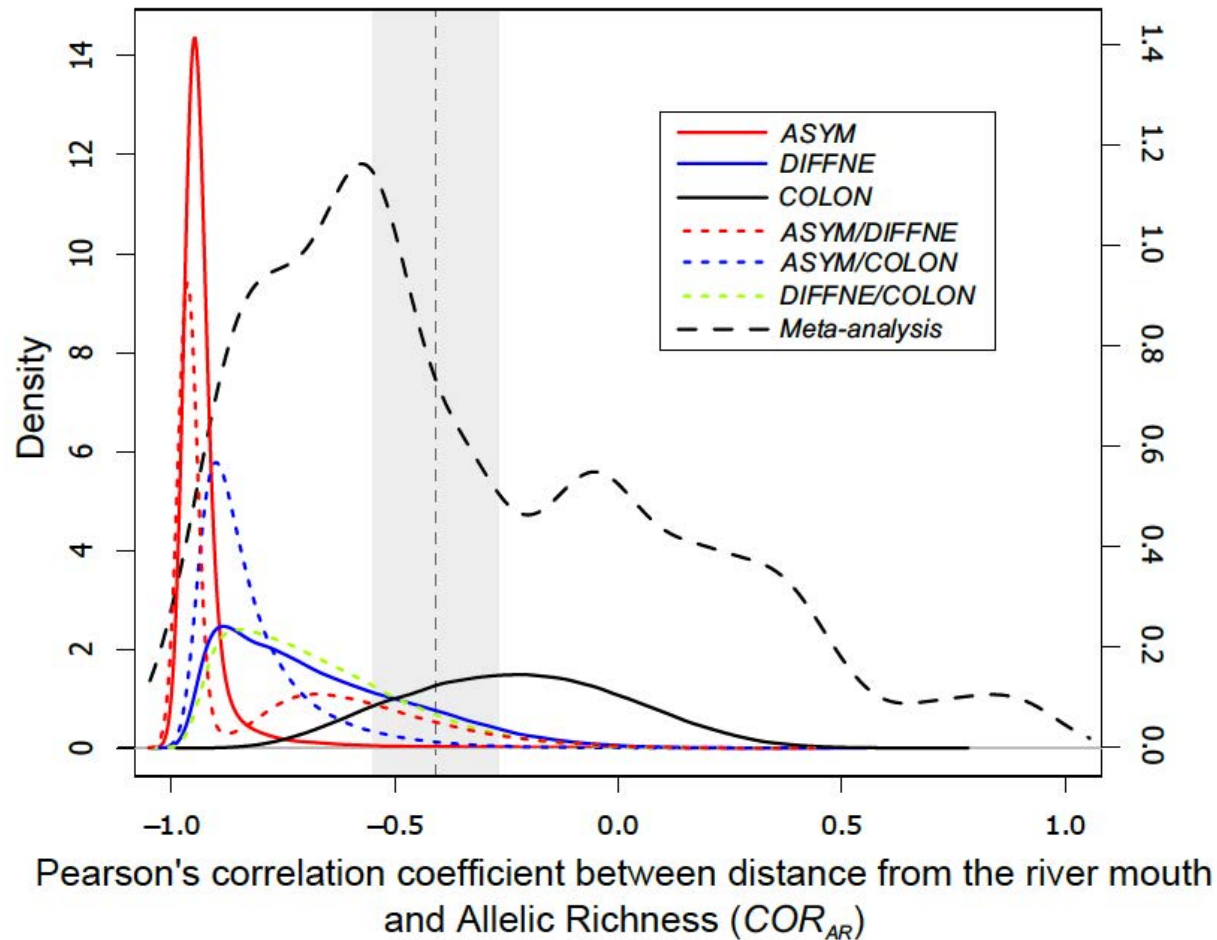
(B) COLONISATION MODEL



COLONISATION PROCESS BEGINNING DOWNSTREAM: LOSS OF ALLELES ALONG THE PROCESS (COLON MODEL)



SIMULATIONS UNDER ALL THESE POSSIBLE MODELS



ALL PROCESSES CAN GENERATE IGDD (IN THEORY), EITHER INDIVIDUALLY OR IN INTERACTION



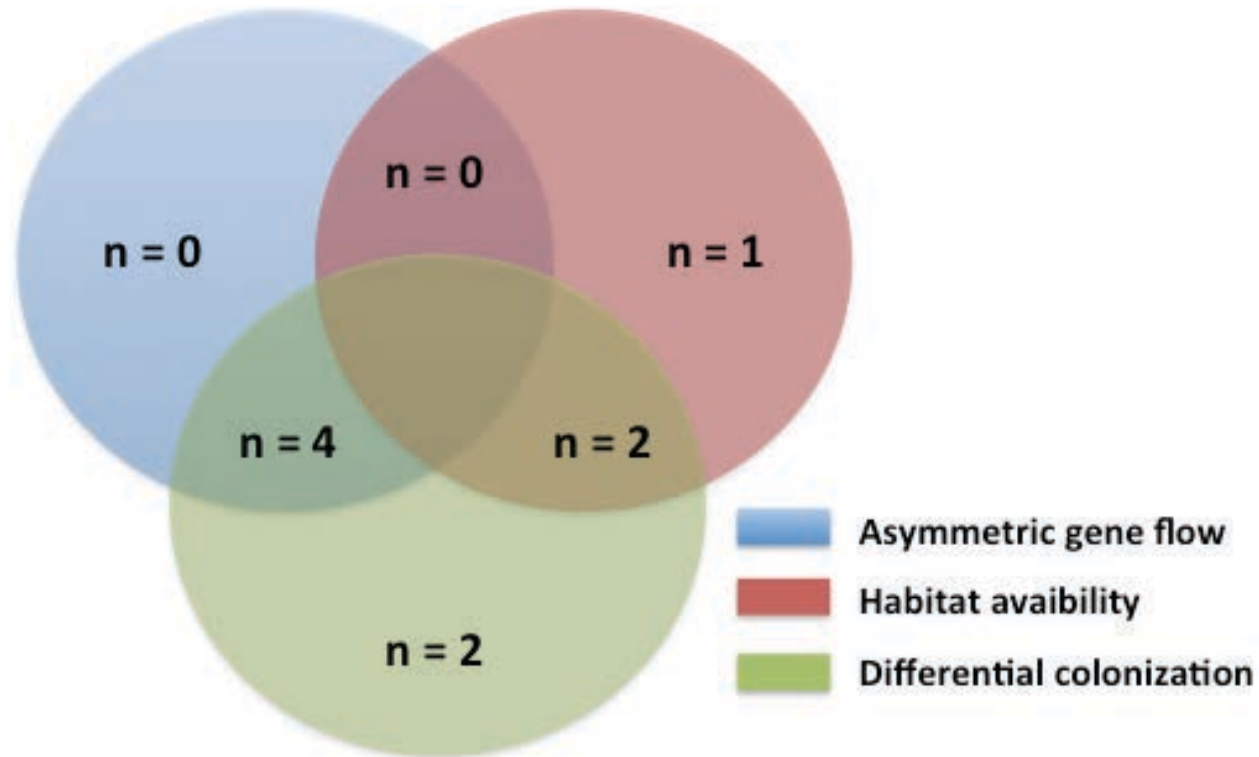
ABC APPROACH TO INFER PROCESSES IN THE WILD:

12 EMPIRICAL CASE STUDIES (fish, invertebrates...)



ABC APPROACH TO INFER PROCESSES IN THE WILD:

12 EMPIRICAL CASE STUDIES (fish, invertebrates...)



**COLONISATION LEGACY AS AN IMPORTANT PROCESS
GENERATING PATTERNS OF GENETIC DIVERSITY IN THE WILD**



INTRASPECIFIC DIVERSITY VARIES SPATIALLY. DOES IT MATTER FOR ECOLOGICAL PROCESSES?

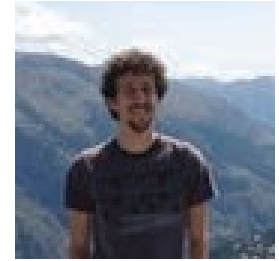


Allan Raffard



INTRASPECIFIC DIVERSITY VARIES SPATIALLY. DOES IT MATTER FOR ECOLOGICAL PROCESSES?

- do diverging populations have similar effects on ecosystem functioning?
- Are these effects of similar magnitude than climatic variation?

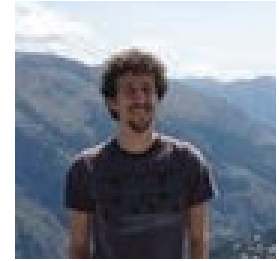


Allan Raffard

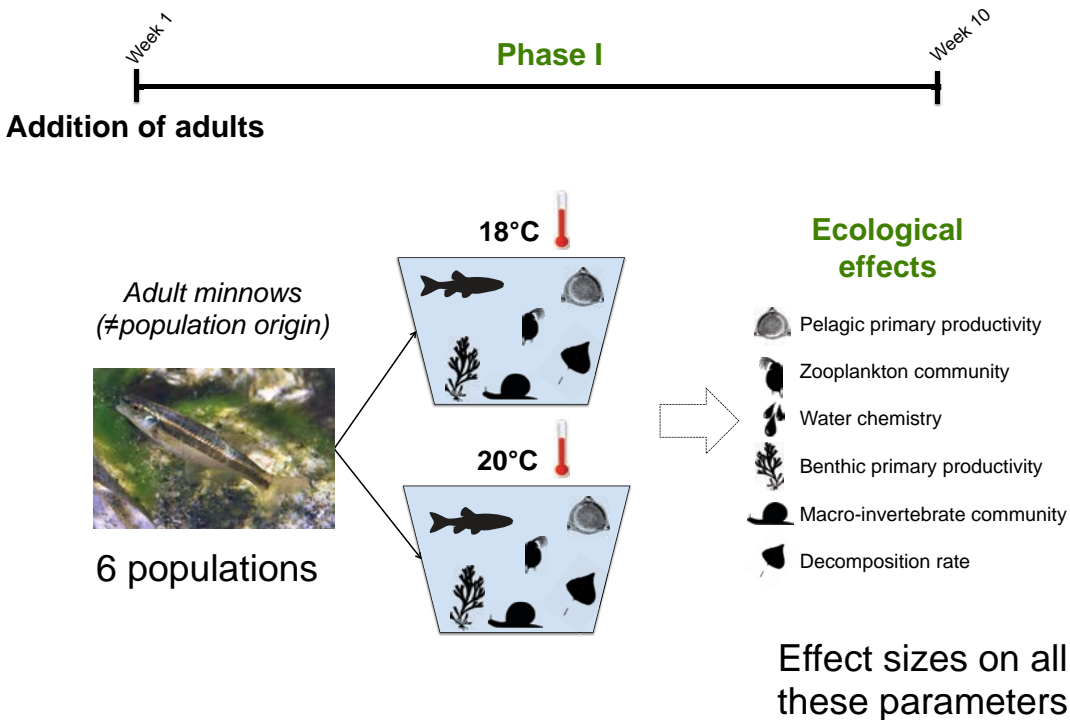


INTRASPECIFIC DIVERSITY VARIES SPATIALLY. DOES IT MATTER FOR ECOLOGICAL PROCESSES?

- do diverging populations have similar effects on ecosystem functioning?
- Are these effects of similar magnitude than climatic variation?



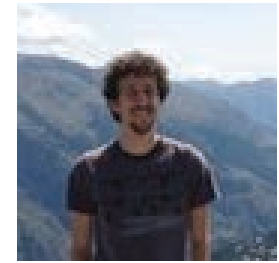
Allan Raffard



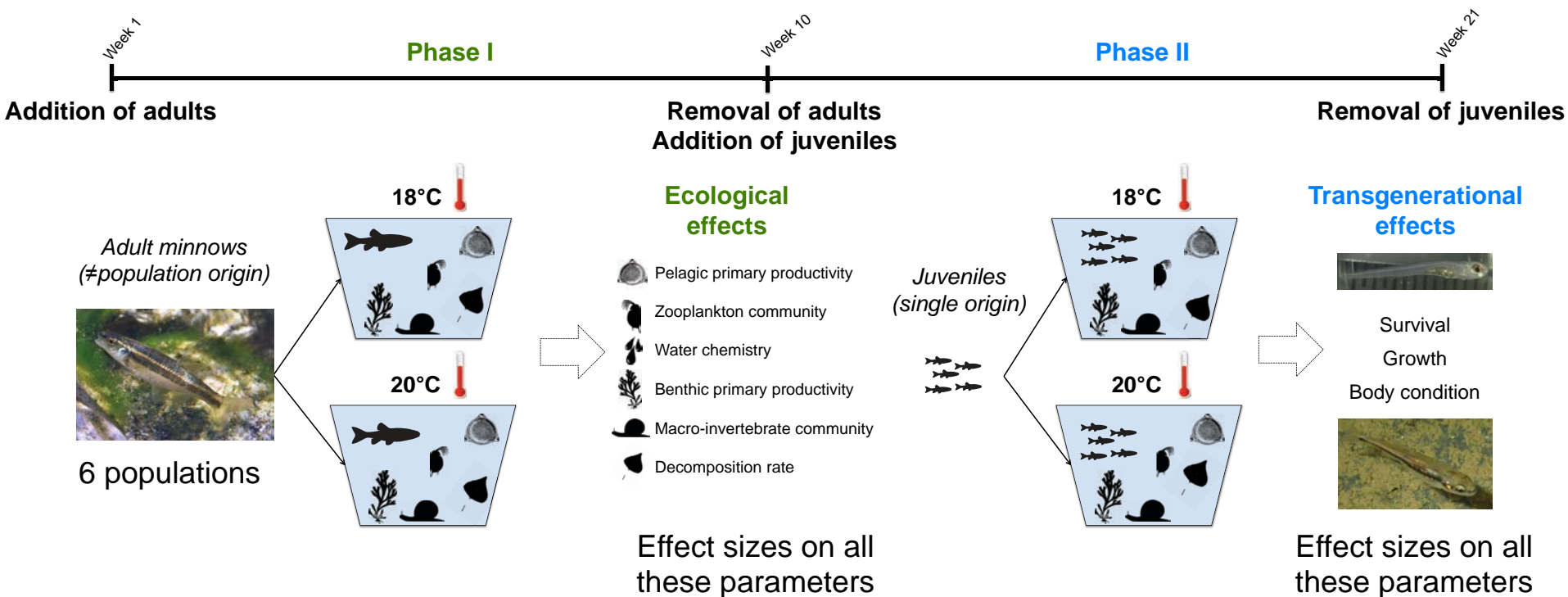


INTRASPECIFIC DIVERSITY VARIES SPATIALLY. DOES IT MATTER FOR ECOLOGICAL PROCESSES?

- do diverging populations have similar effects on ecosystem functioning?
- Are these effects of similar magnitude than climatic variation?

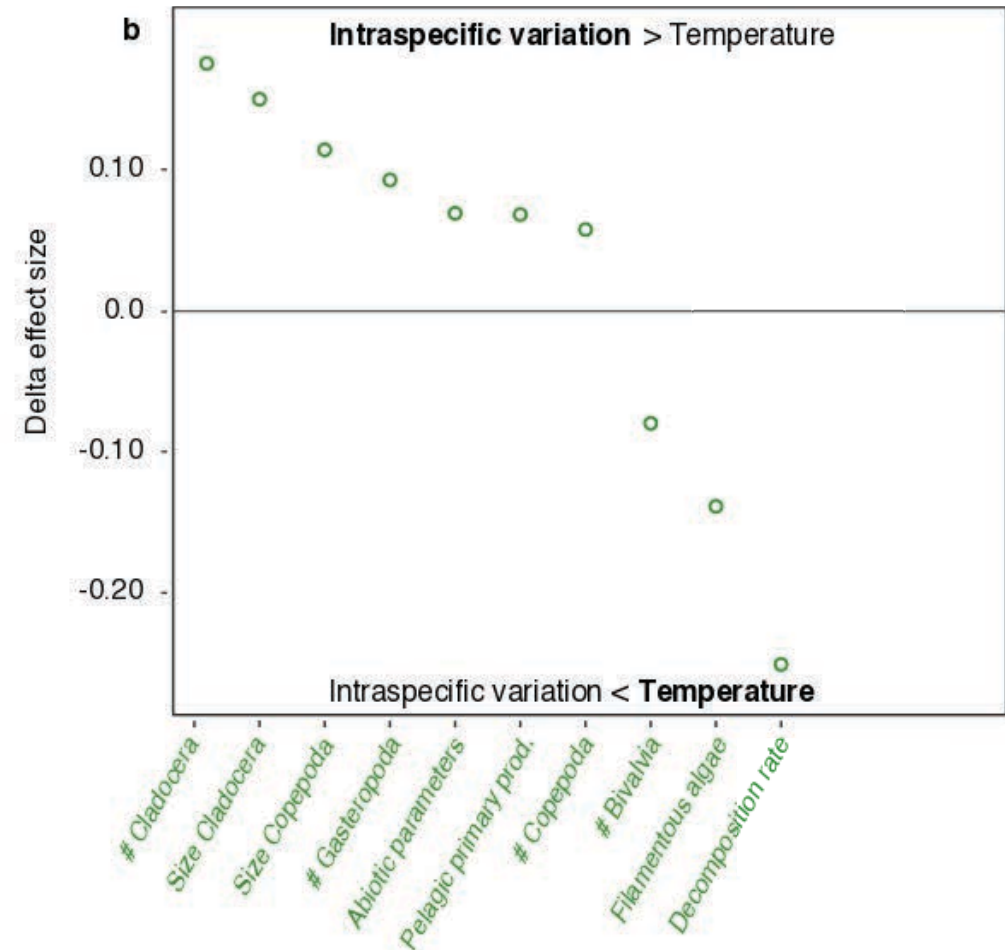
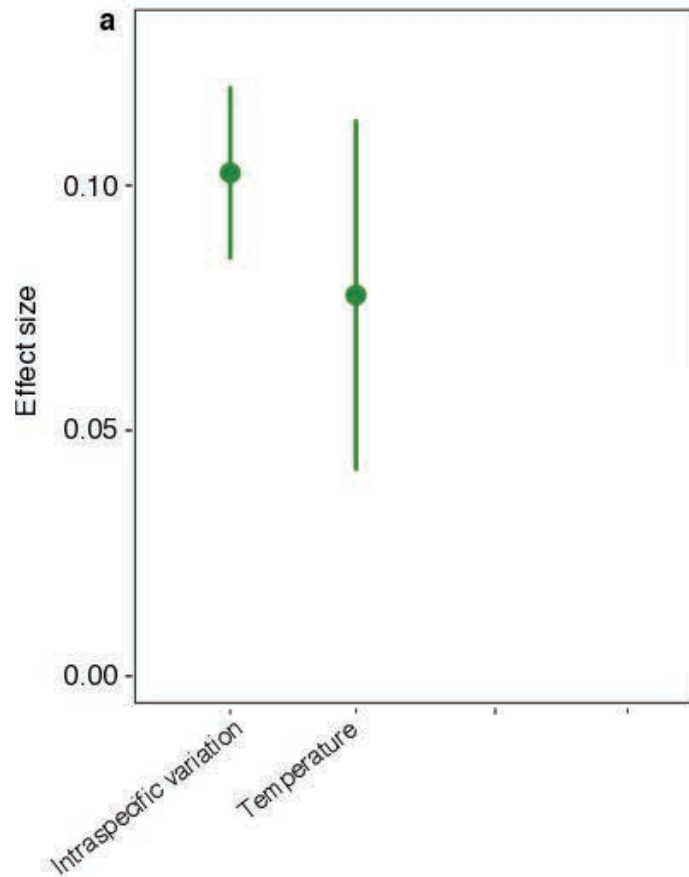


Allan Raffard





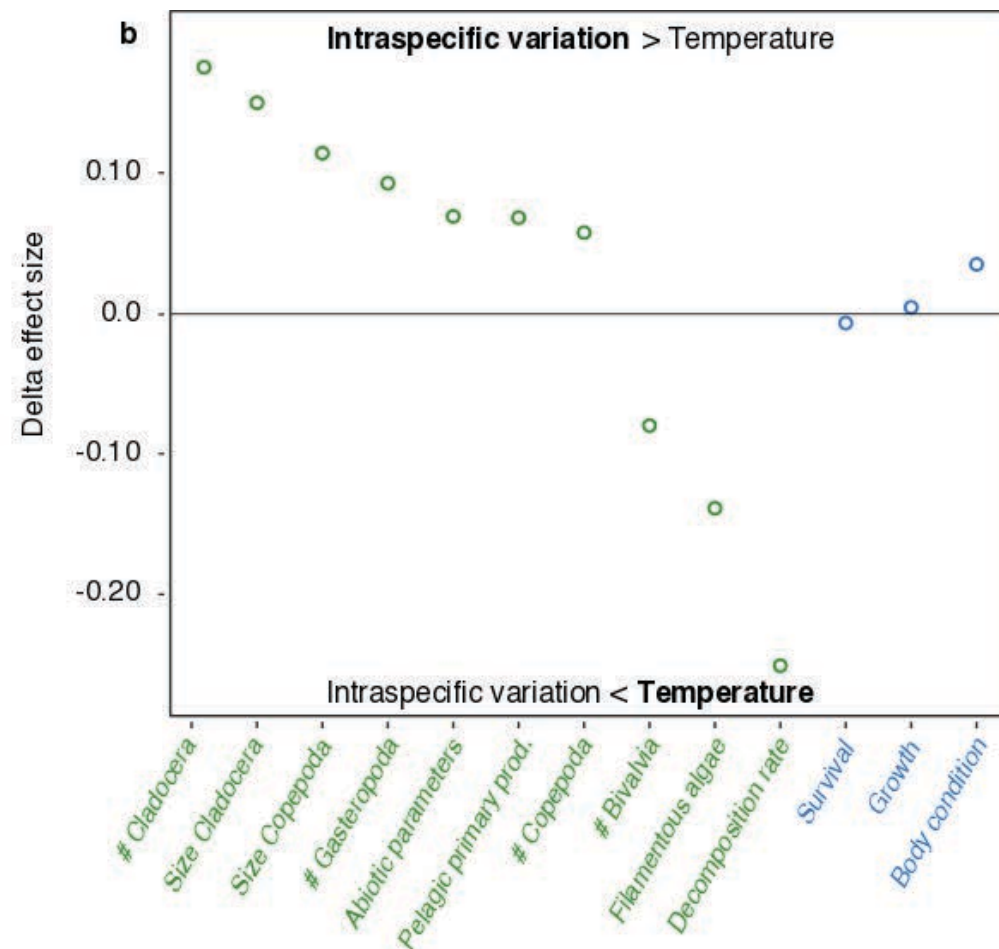
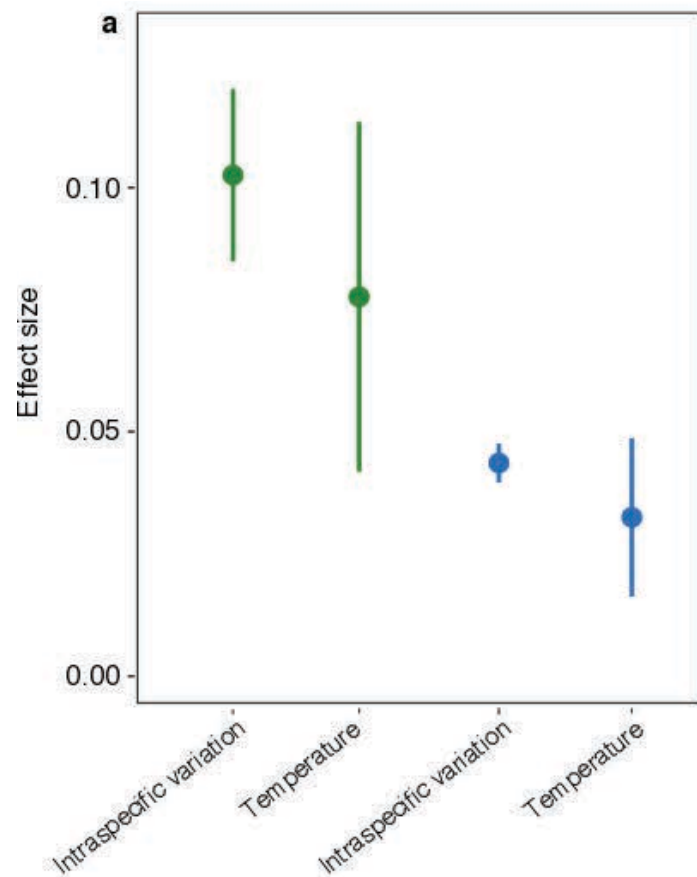
INTRASPECIFIC DIVERSITY STRONGLY IMPACTS ECOLOGY



ID HAS ECOLOGICAL EFFECTS OF SIMILAR STRENGTH THAN WARMING

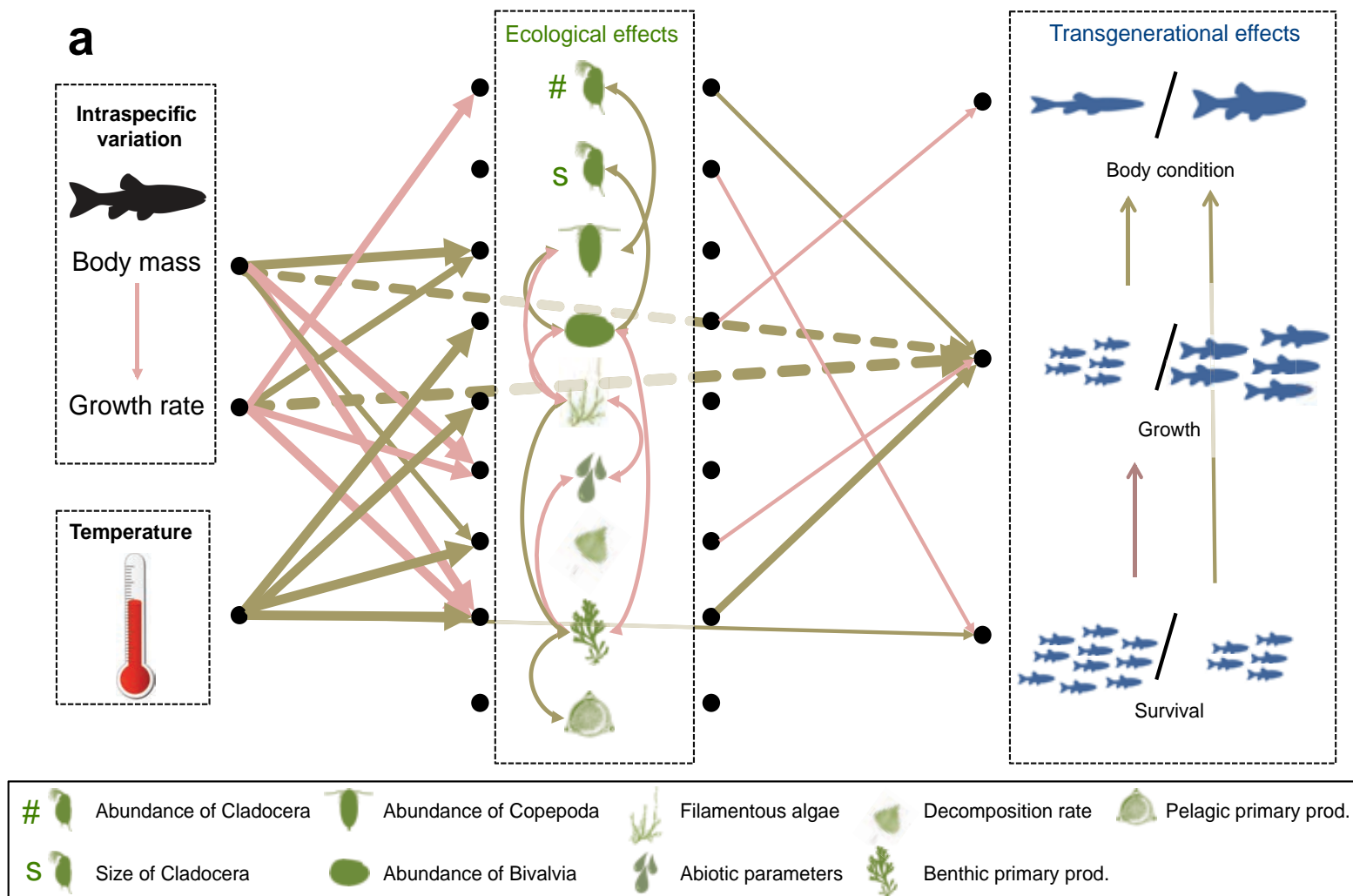


INTRASPECIFIC DIVERSITY STRONGLY IMPACTS ECOLOGY...AND “EVOLUTION”.



ID HAS ECOLOGICAL EFFECTS OF SIMILAR STRENGTH THAN WARMING LEADING TO ECO-EVO FEEDBACKS

(II) ECOLOGICAL CONSEQUENCES OF INTRASPECIFIC DIVERSITY



ID IS NOT DISTRIBUTED HOMOGENEOUSLY, AND THIS CAN HAVE CONSEQUENCES FOR ECOSYSTEM FUNCTIONING (AND EVOLUTION)



MANY OBSTACLES ALONG RIVERS: MAINLY WEIRS

Restoring connectivity by erasing or managing (fishpass) weirs and dams: costly. Hierarchizing actions is needed.



5 m high, 1960



1.2 m high, 1640



MANY OBSTACLES ALONG RIVERS: MAINLY WEIRS

Restoring connectivity by erasing or managing (fishpass) weirs and dams: costly. Hierarchizing actions is needed.



5 m high, 1960



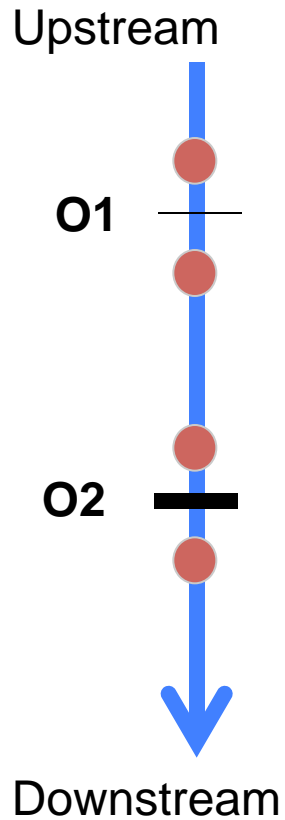
1.2 m high, 1640

Using genetic differentiation (below-to-above design) to identify the most problematic weirs. But!

- How to take into account the age of the obstacle?
- How to take into account the size of populations?



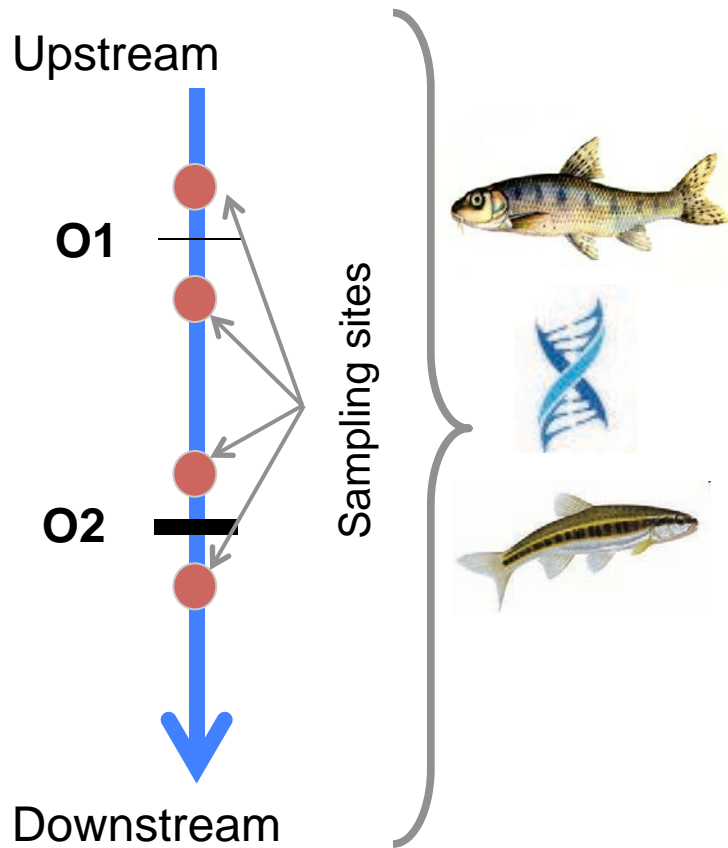
A below-to-above design quantifying genetic differentiation :



(III) ID AS A INDICATOR OF FRAGMENTATION



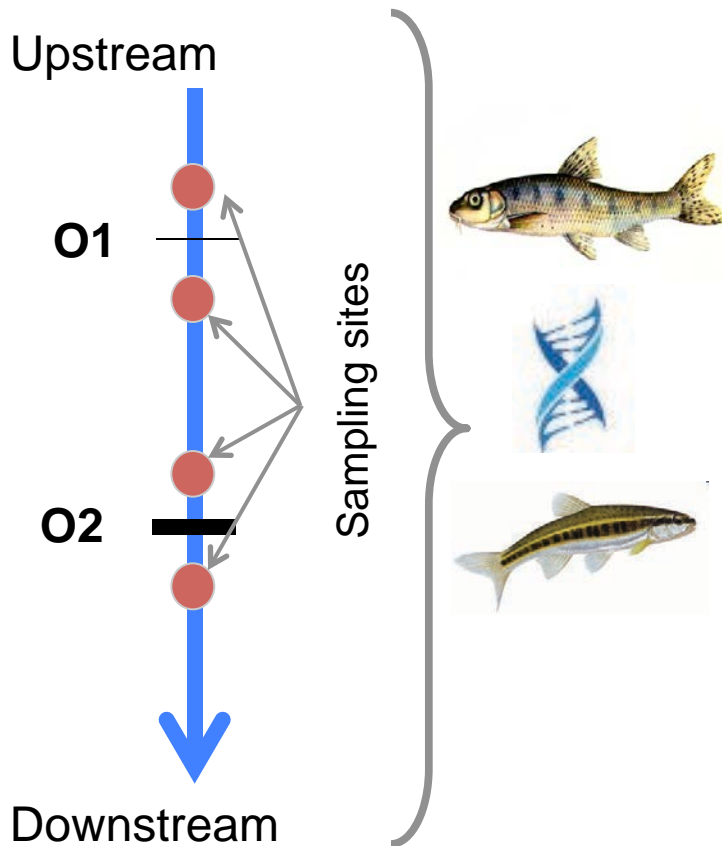
A below-to-above design quantifying genetic differentiation :



$$GD_{O1} < GD_{O2},$$



A below-to-above design quantifying genetic differentiation :



$$GD_{O1} < GD_{O2},$$

if:

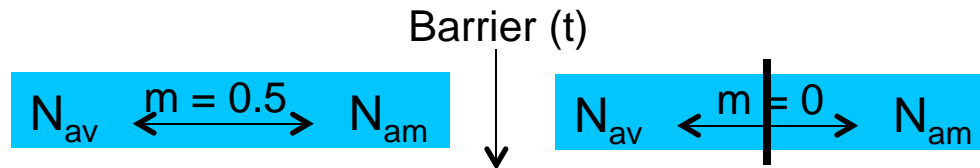
- O1 and O2 of the same age
- Population size in O1 = O2

Designing an index of fragmentation (F) accounting for the age of obstacles and the local population size

(III) ID AS A INDICATOR OF FRAGMENTATION



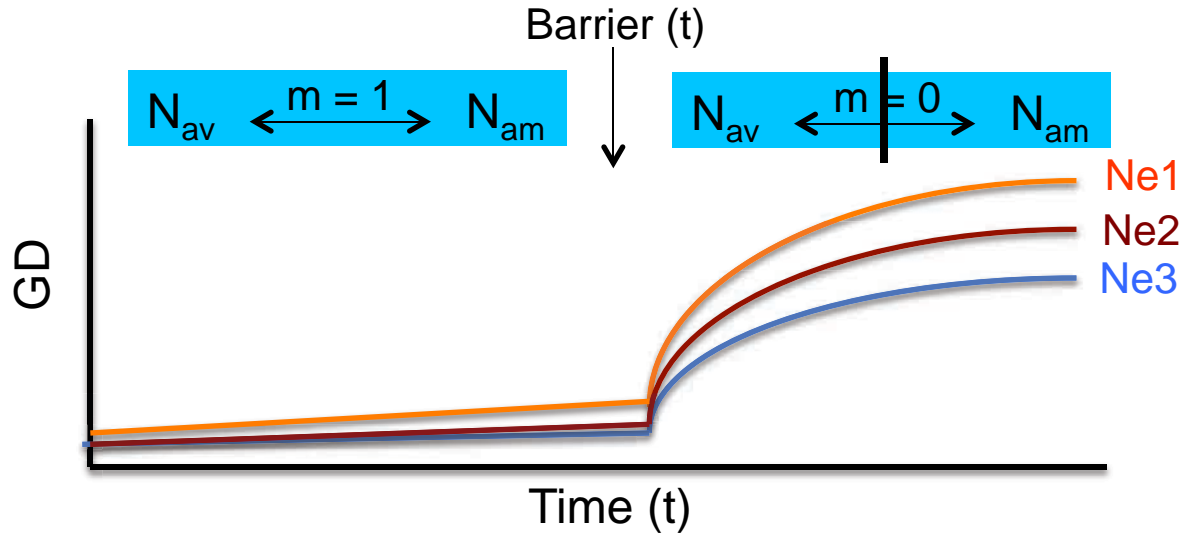
Step 1: Simulations. Calculating expected GD (*eGD*) if the barrier is total ($m=0$, maximum expected effect) according to N_e and Age (t)



(III) ID AS A INDICATOR OF FRAGMENTATION



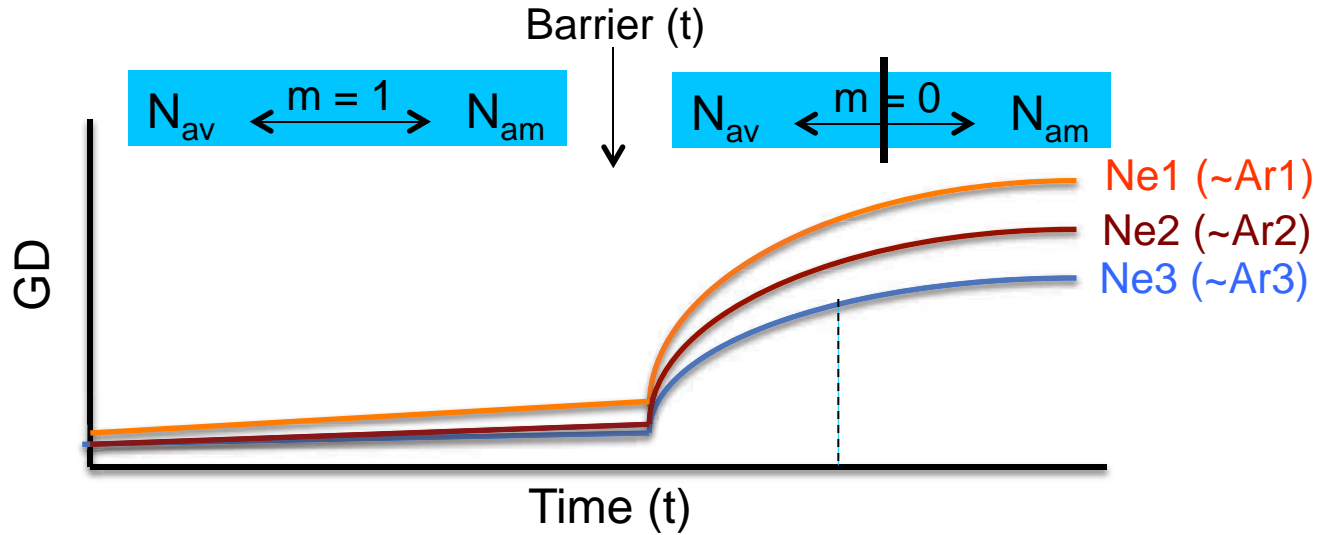
Step 1: Simulations. Calculating expected GD (eGD) if the barrier is total ($m=0$, maximum expected effect) according to N_e and Age (t)



(III) ID AS A INDICATOR OF FRAGMENTATION



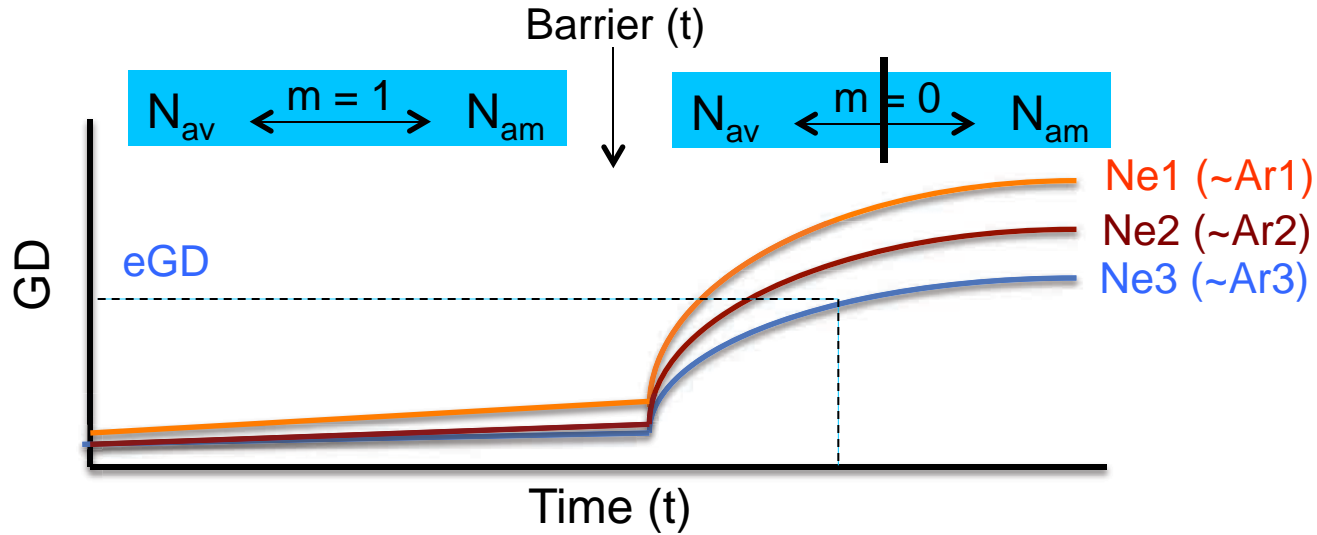
Step 1: Simulations. Calculating expected GD (eGD) if the barrier is total ($m=0$, maximum expected effect) according to N_e and Age (t)



(III) ID AS A INDICATOR OF FRAGMENTATION



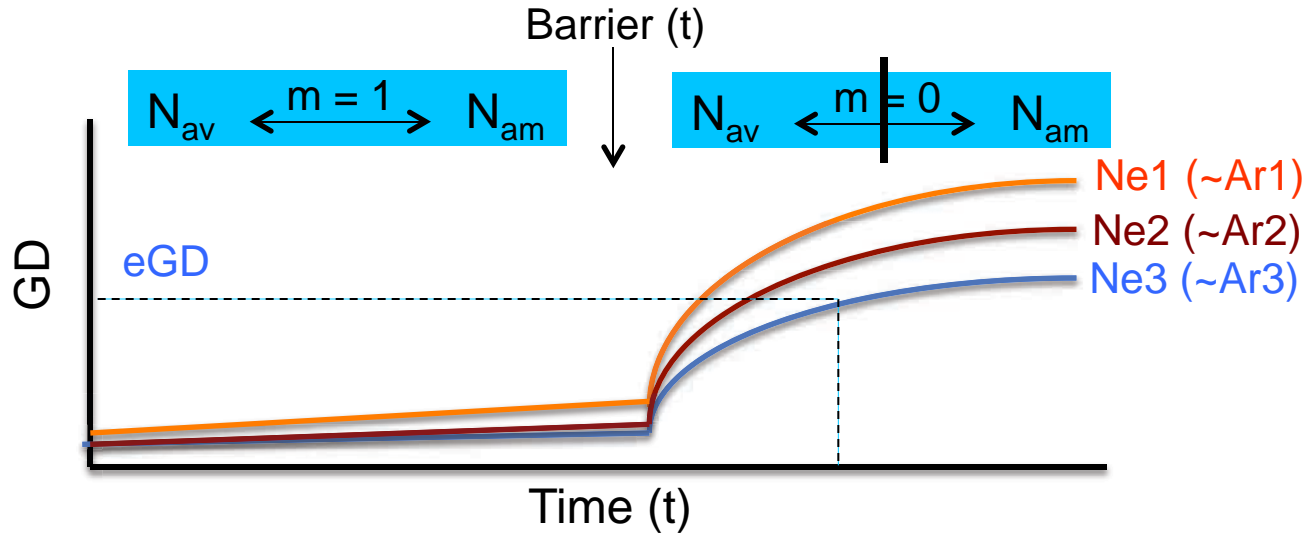
Step 1: Simulations. Calculating expected GD (eGD) if the barrier is total ($m=0$, maximum expected effect) according to N_e and Age (t)



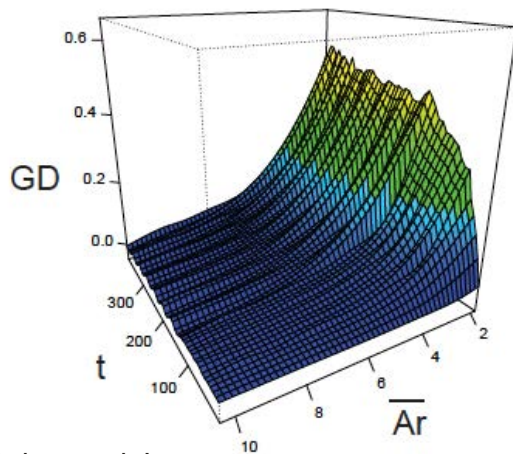
(III) ID AS A INDICATOR OF FRAGMENTATION



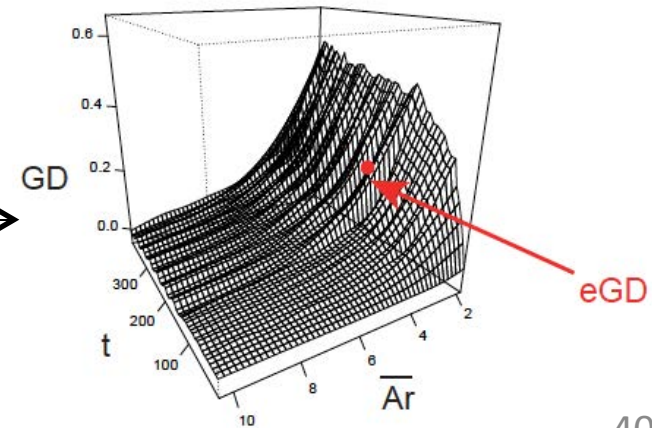
Step 1: Simulations. Calculating expected GD (eGD) if the barrier is total ($m=0$, maximum expected effect) according to Ne and Age (t)



Step 2: Calibrating models (from simulations) to predict eGD from any Ne (Ar) and t



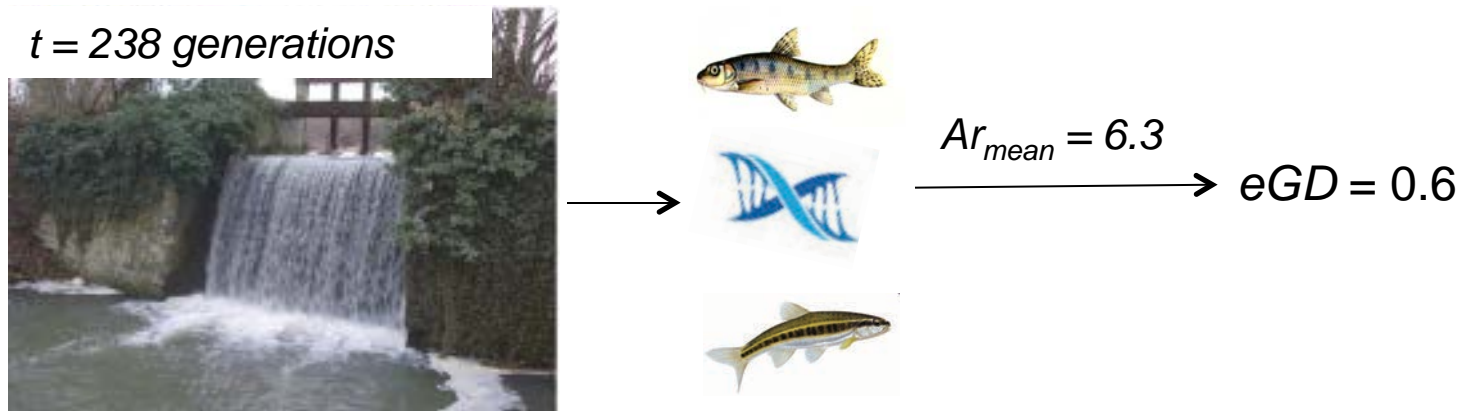
$$eGD \sim Ar_i * t_i \text{ (GAMs)}$$



(III) ID AS A INDICATOR OF FRAGMENTATION



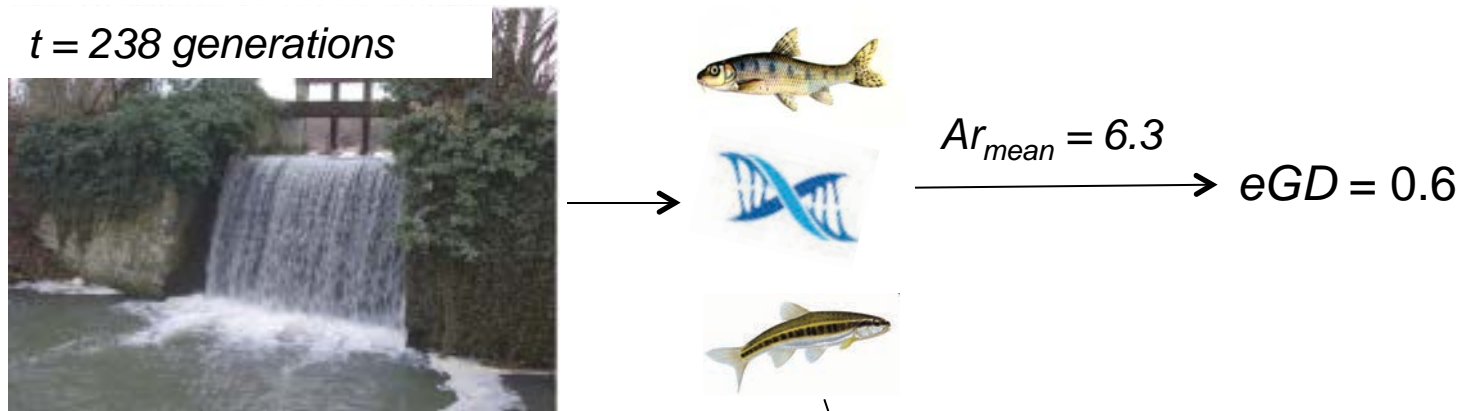
Step 3: using the models to estimate eGD from empirical Ar and t values



(III) ID AS A INDICATOR OF FRAGMENTATION



Step 3: using the models to estimate eGD from empirical Ar and t values



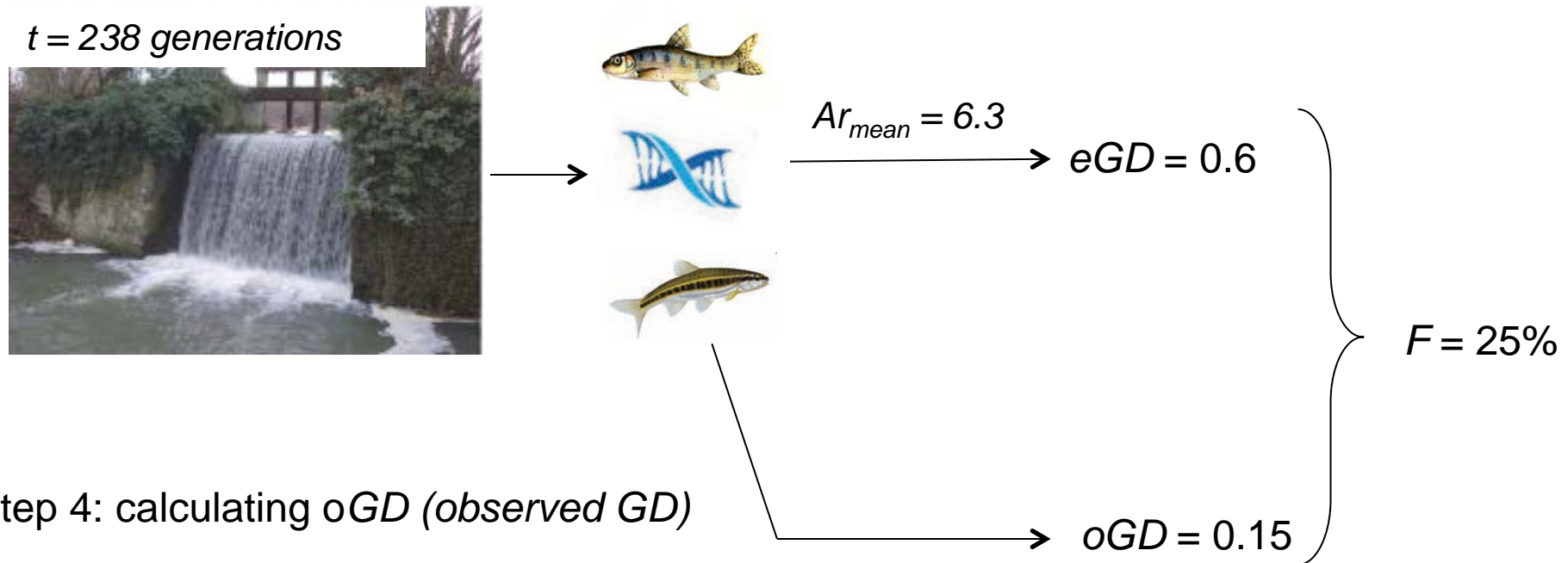
Step 4: calculating oGD (*observed GD*)

$oGD = 0.15$

(III) ID AS A INDICATOR OF FRAGMENTATION



Step 3: using the models to estimate eGD from empirical Ar and t values



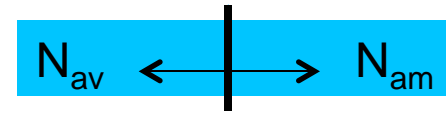
Step 4: calculating oGD (observed GD)

Step 5: calculating the fragmentation index: $F = [oGD/eGD]*100$

Percentage of an expected maximal effect, given N_e and t



Validation using independent simulations.

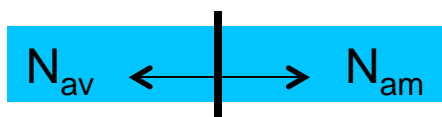


$$m = 0.2 - 0$$

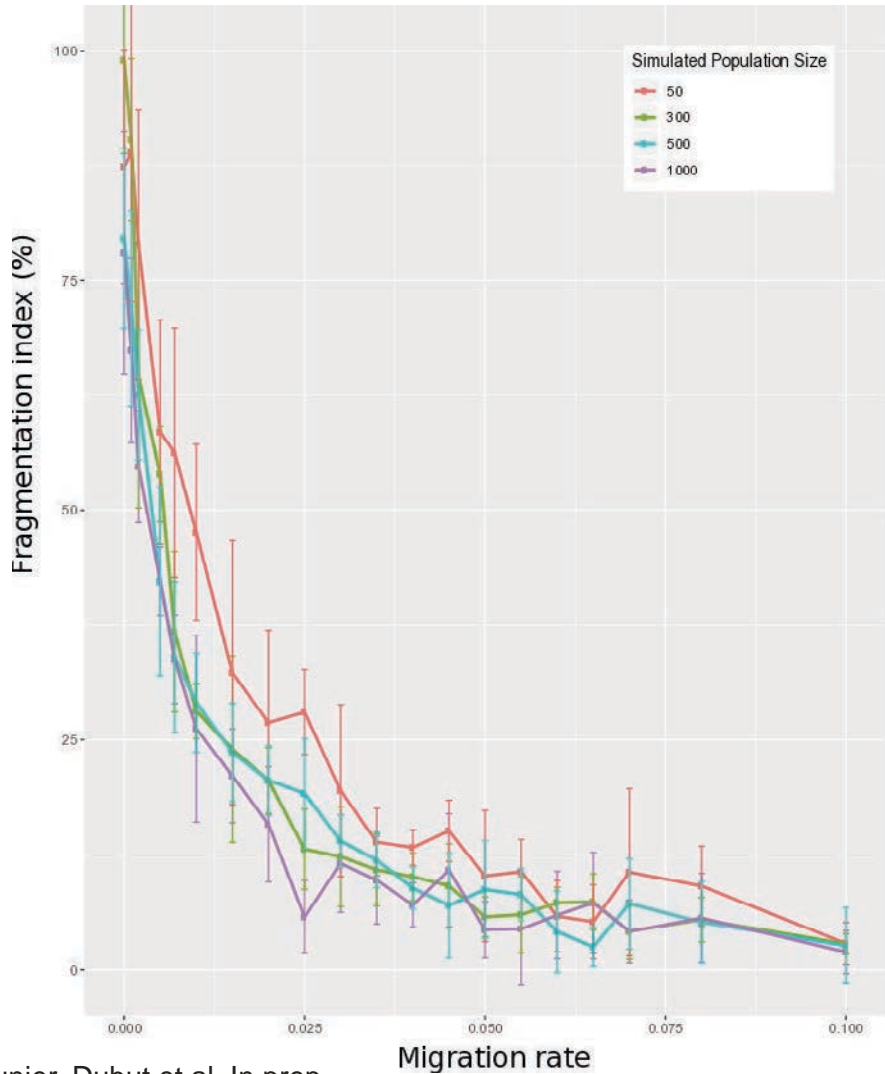
(III) ID AS A INDICATOR OF FRAGMENTATION



Validation using independent simulations.



$$m = 0.2 - 0$$



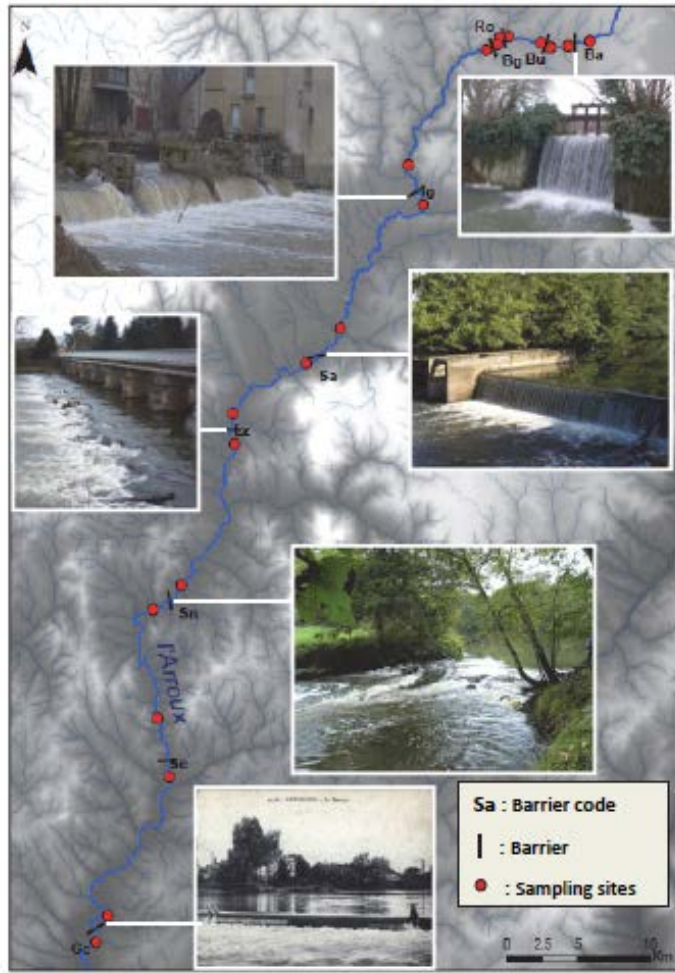
The index is close to 100% when $m = 0$ (total barrier)

The index decreases very quickly and is rapidly closed to 0% (no barrier effect detected when $m > 0.1$)

(III) ID AS A INDICATOR OF FRAGMENTATION



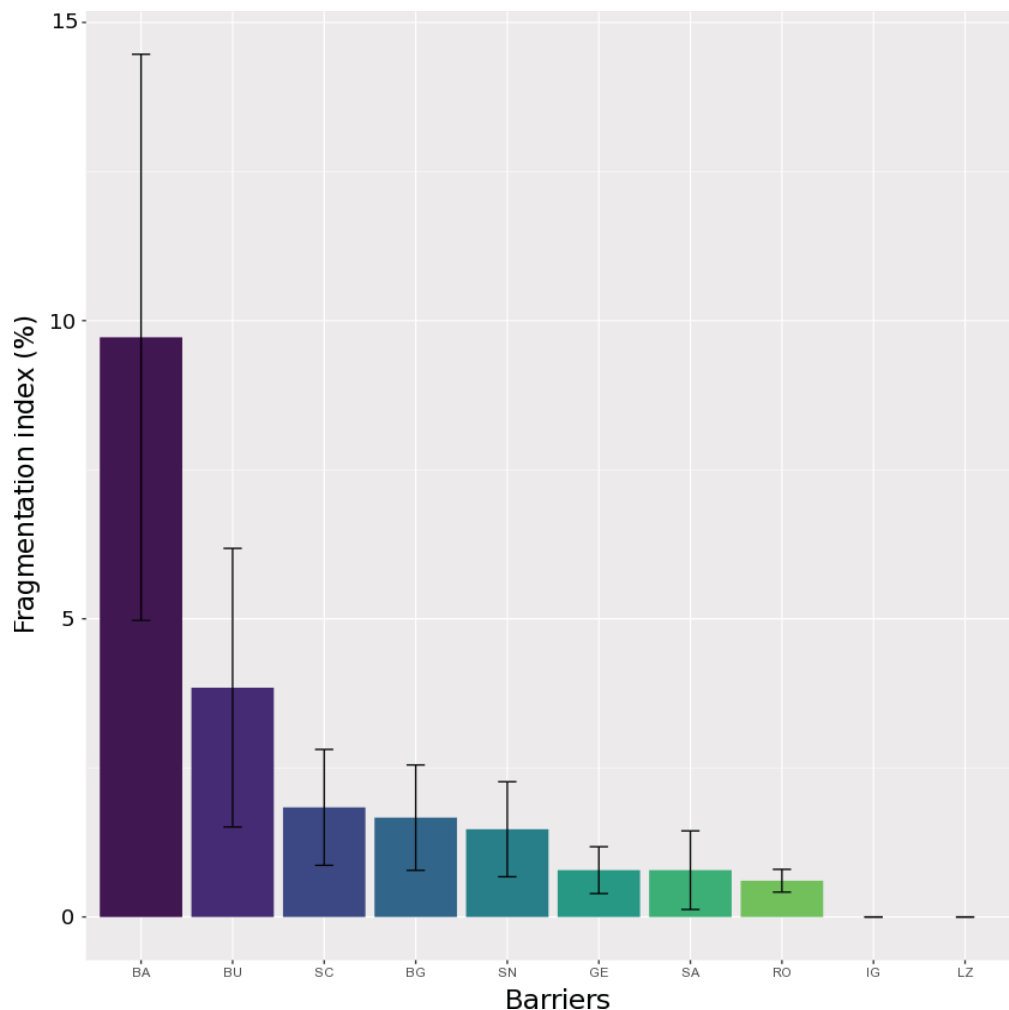
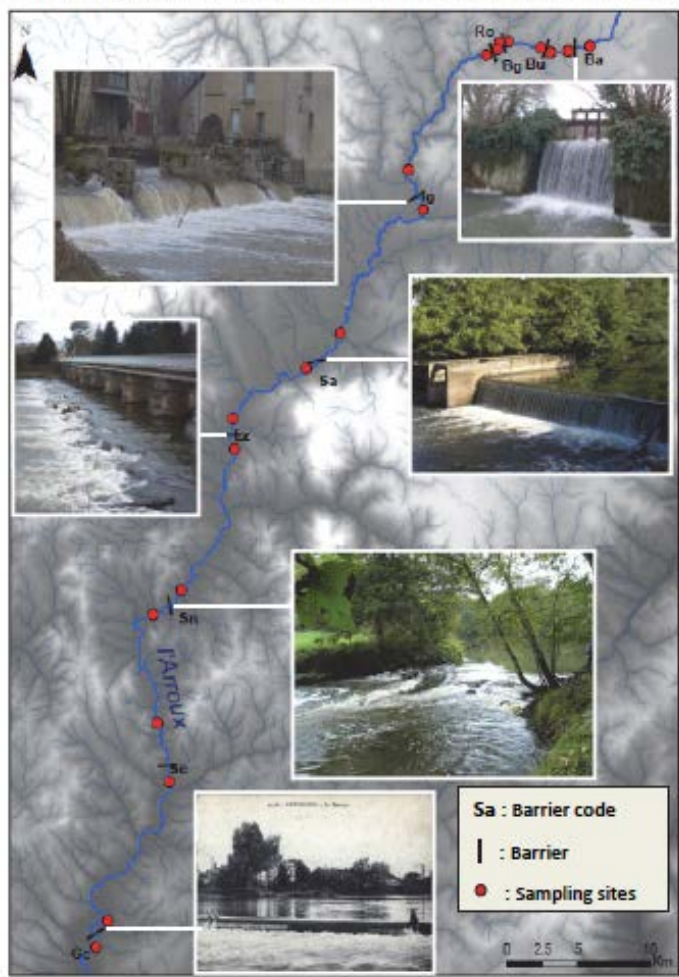
An empirical test on 10 obstacle along a river (Arroux)



(III) ID AS A INDICATOR OF FRAGMENTATION



An empirical test on 10 obstacle along a river (Arroux)



The index has relatively value (weirs are not impassables)

Variability among barriers: hierarchisation



- ⇒ **DIVERSITY VARIES ALONG NETWORKS, PATTERNS ARE REPEATABLE**
- ⇒ **DECIPHERING THESE PROCESSES IS POSSIBLE**
- ⇒ **DIVERSITY IS NOT ECOLOGICALLY TRIVIAL. IT HAS STRONG IMPACTS ON ECOLOGICAL DYNAMICS**
- ⇒ **CONSERVATION OF THESE EVOLUTIONARY AND ECOLOGICAL SIGNIFICANT UNITS IS REQUIRED**
- ⇒ **IT IS ALSO A USEFUL INDICATOR FOR SOME HUMAN PRESSURES**

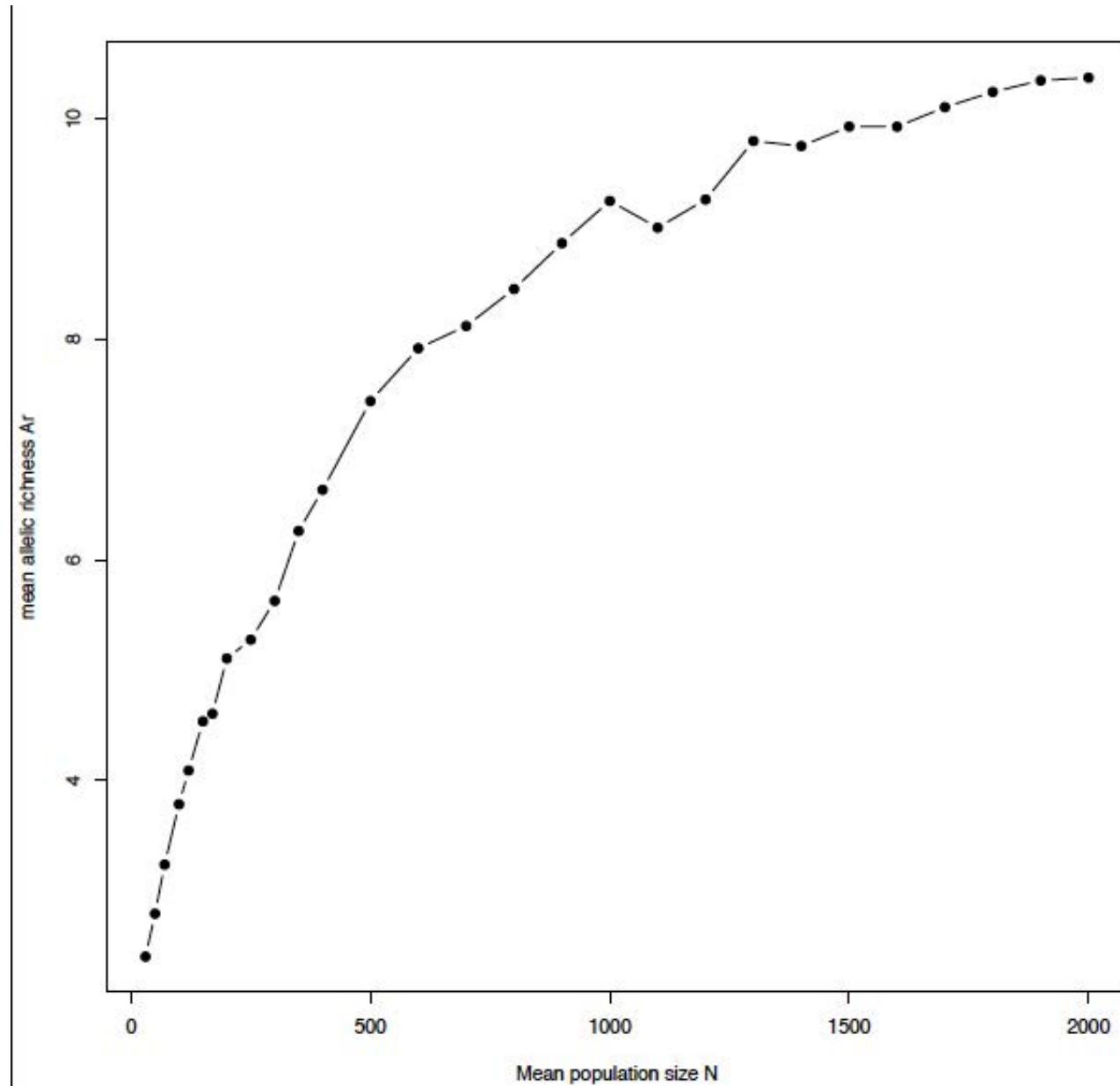
ACKNOWLEDGMENTS



LES FINANCEURS



ET VOUS.





**INTRASPECIFIC DIVERSITY VARIES SPATIALLY AND IT MATTERS FOR ECOLOGICAL PROCESSES.
HOW TO PRIORITIZE CONSERVATION AREAS FOR ID?**

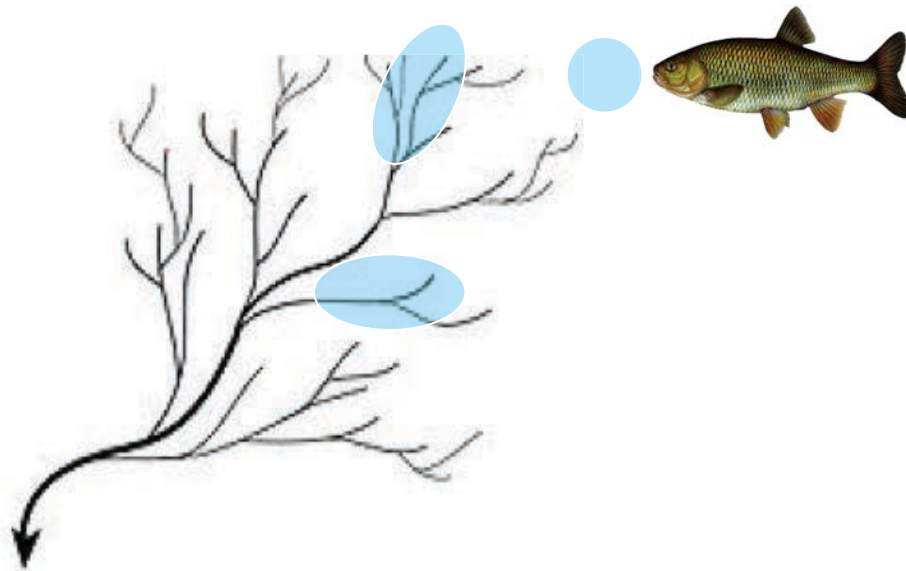




**INTRASPECIFIC DIVERSITY VARIES SPATIALLY AND IT MATTERS FOR ECOLOGICAL PROCESSES.
HOW TO PRIORITIZE CONSERVATION AREAS FOR ID?**



Ivan Paz-Vinas



WHICH PARTS OF THE NETWORK SHOULD WE TARGET TO CONSERVE INTRASPECIFIC DIVERSITY?



**INTRASPECIFIC DIVERSITY VARIES SPATIALLY AND IT MATTERS FOR ECOLOGICAL PROCESSES.
HOW TO PRIORITIZE CONSERVATION AREAS FOR ID?**



Ivan Paz-Vinas



WHICH PARTS OF THE NETWORK SHOULD WE TARGET TO CONSERVE INTRASPECIFIC DIVERSITY?



INTRASPECIFIC DIVERSITY VARIES SPATIALLY AND IT MATTERS FOR ECOLOGICAL PROCESSES. HOW TO PRIORITIZE CONSERVATION AREAS FOR ID?



Ivan Paz-Vinas



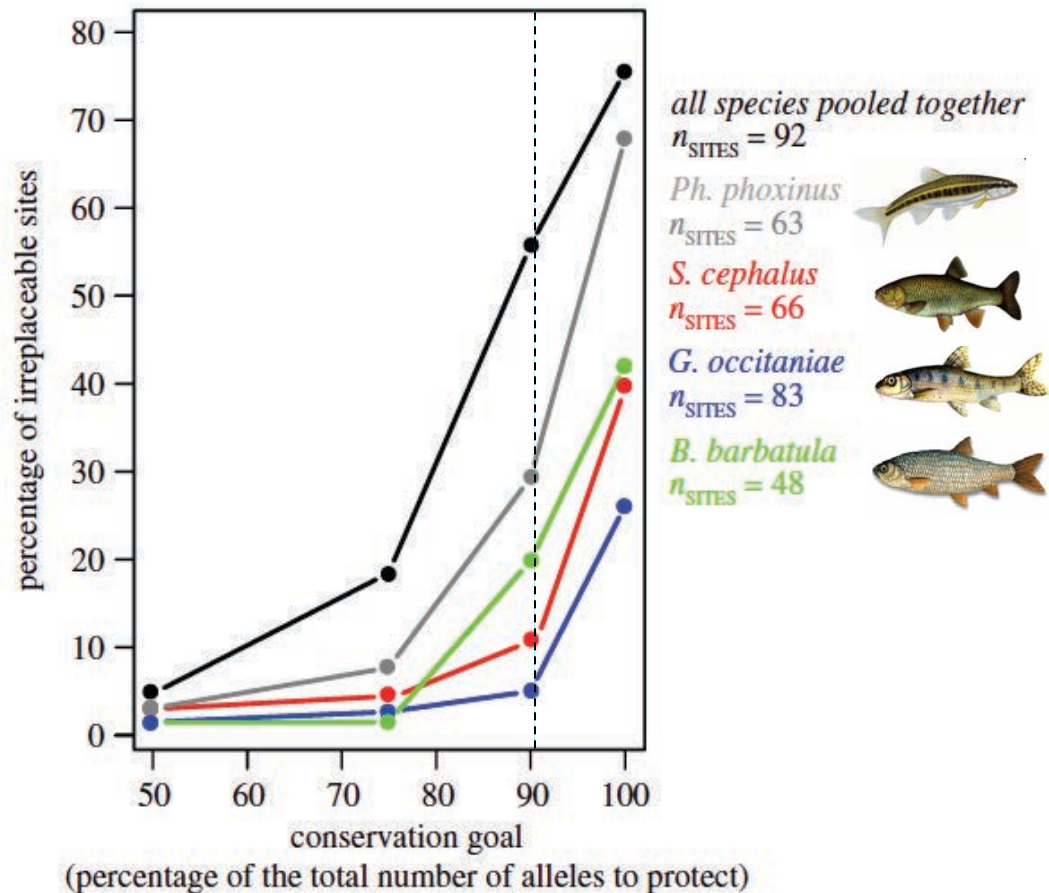
A multispecies conservation planning approach:

- 6 species sampled in 92 sites and typed for microsatellites
- Identifying the sites (*irreplaceable sites*) required to cover a certain amount of genetic diversity (*conservation target*)
- Mapping these sites and testing for congruencies among species

WHICH PARTS OF THE NETWORK SHOULD WE TARGET TO CONSERVE INTRASPECIFIC DIVERSITY?



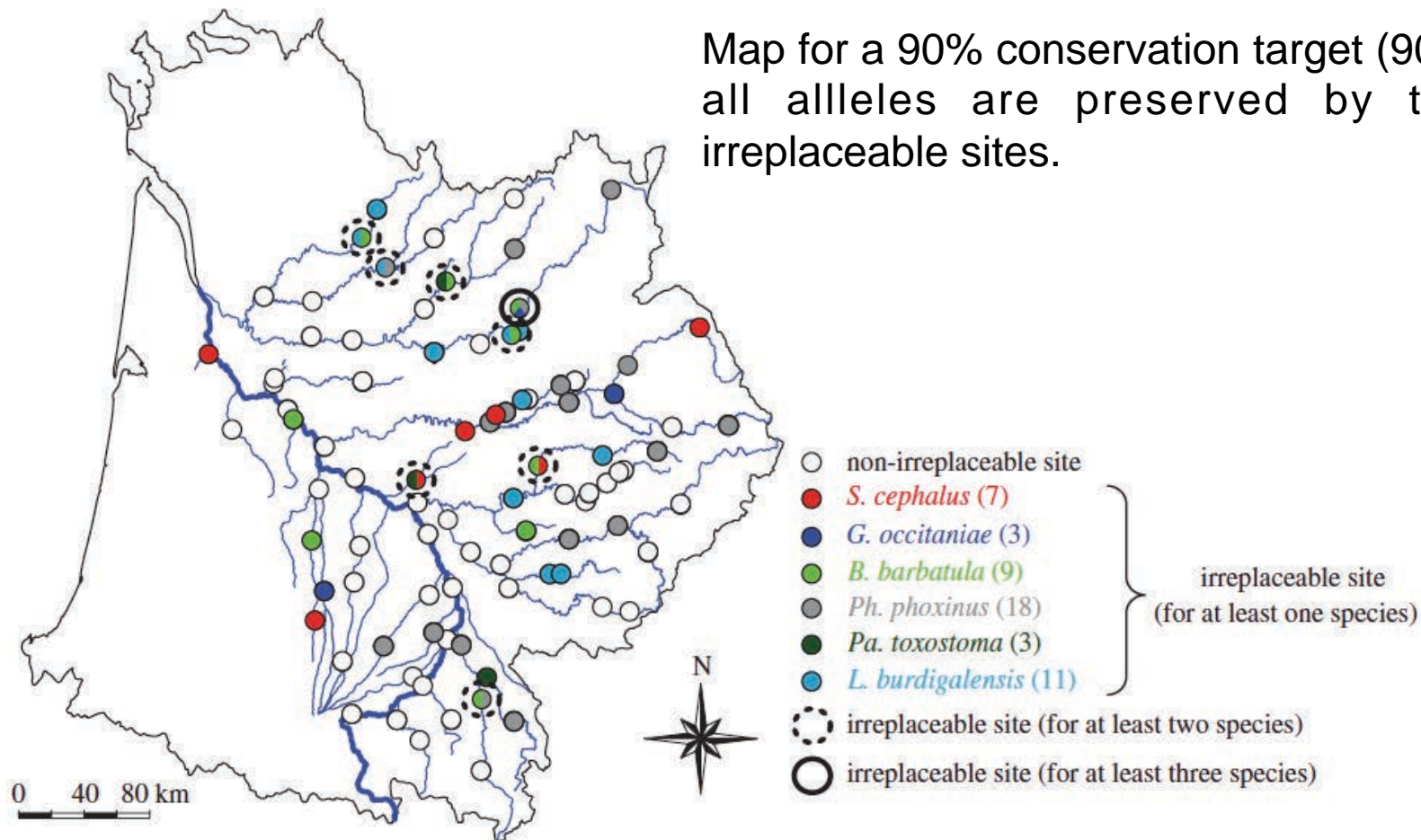
CONSERVATION PLANNING: IDENTIFYING PRIORITY ZONES FOR ID



THE NUMBER OF IRREPLACEABLE SITES VARY ACCORDING TO THE CONSERVATION TARGET AND THE SPECIES



CONSERVATION PLANNING: IDENTIFYING PRIORITY ZONES FOR ID

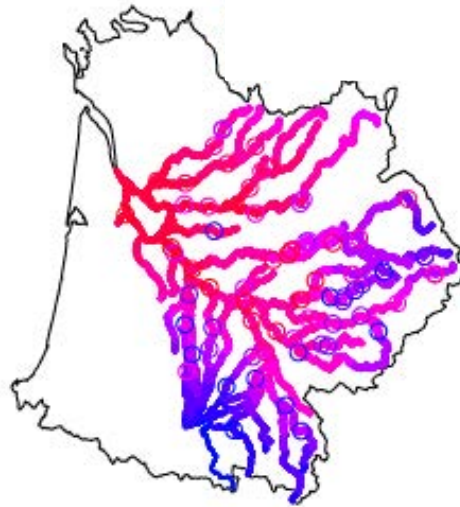


MOST IRREPLACEABLE SITES ARE SPECIES-SPECIFIC. HARD TO CONSERVE ID FOR A POOL OF SPECIES

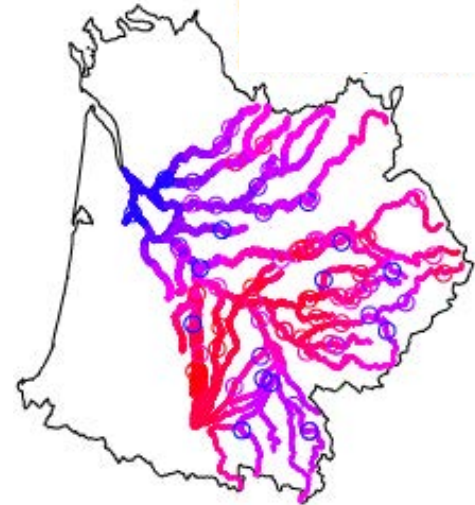


CHUB

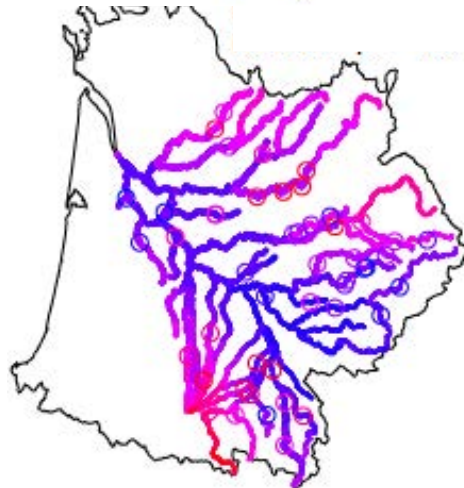
ALLELIC RICHNESS



GENETIC UNIQUENESS (Djost)



LOACH

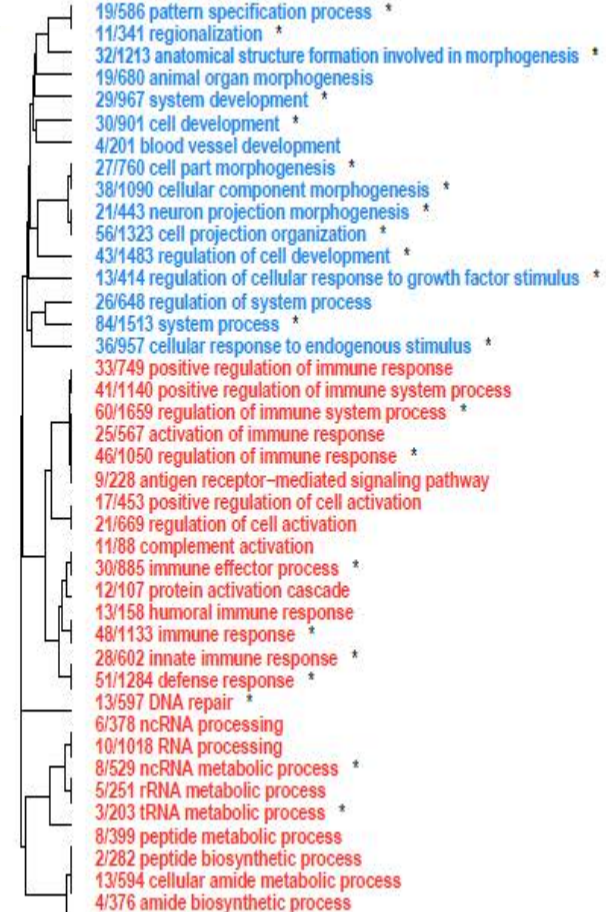
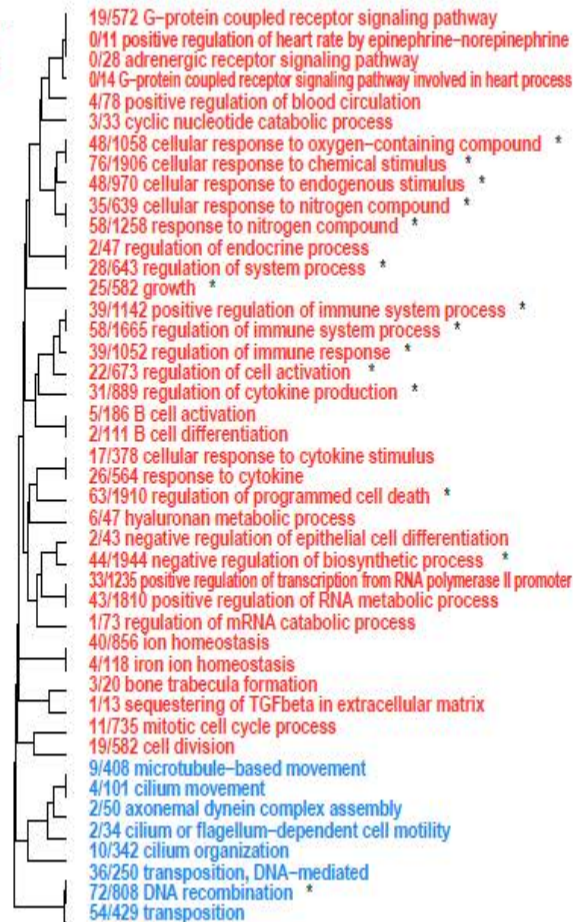
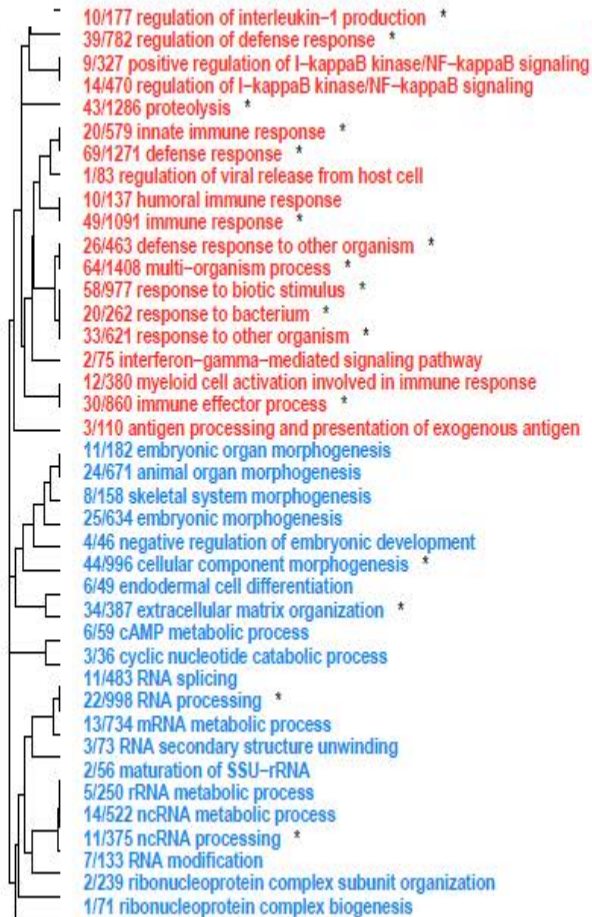


SPATIAL PATTERNS OF GENETIC DIVERSITY ARE NOT CONGRUENT AMONG SPECIES

Molecular level: Gene expression profile & biological functions

involved

DC

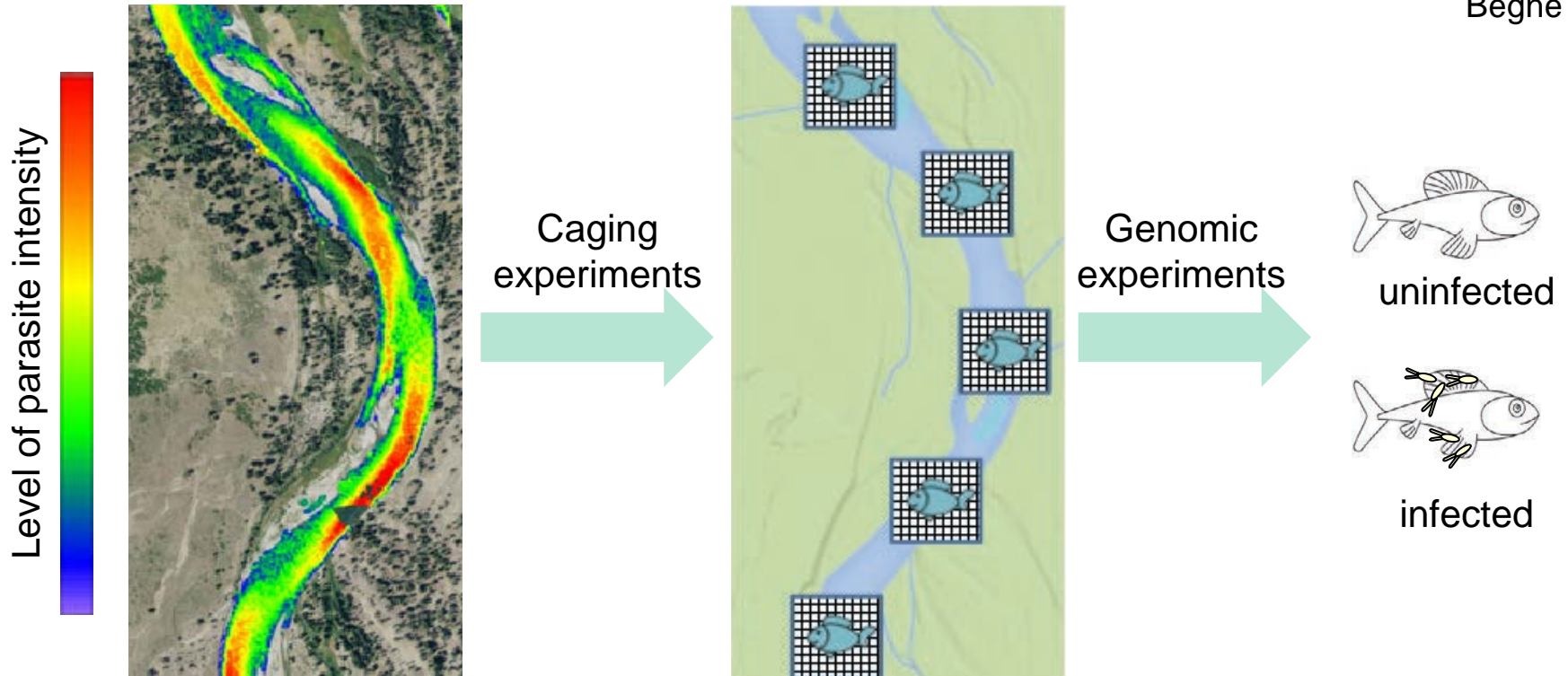




Eglantine
Mathieu-
Bégné

THE ENVIRONMENTAL AND GENOMIC BASES OF RESISTANCE

Meetic: How can an ugly larvae meet a lovely host in a desert?

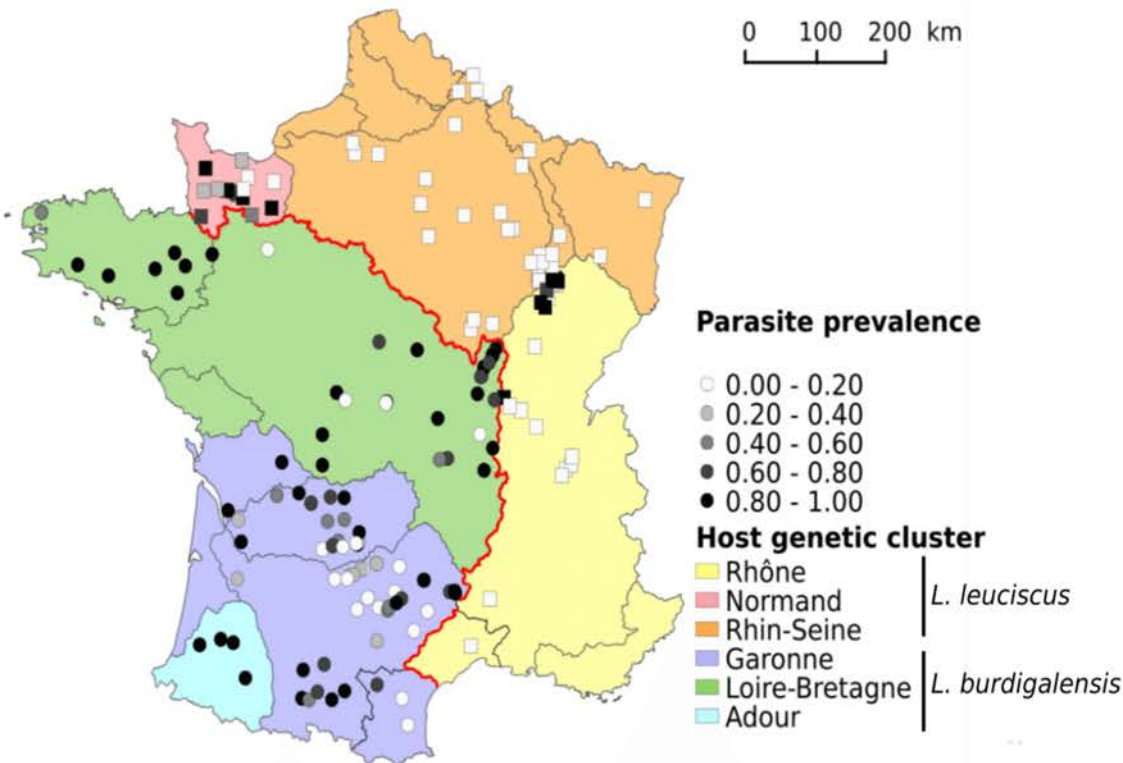


RESOLVING THE “NEEDLE IN A HAYSTACK” PARADOX



Eglantine
Mathieu-
Bégné

THE ENVIRONMENTAL AND GENOMIC BASES OF RESISTANCE



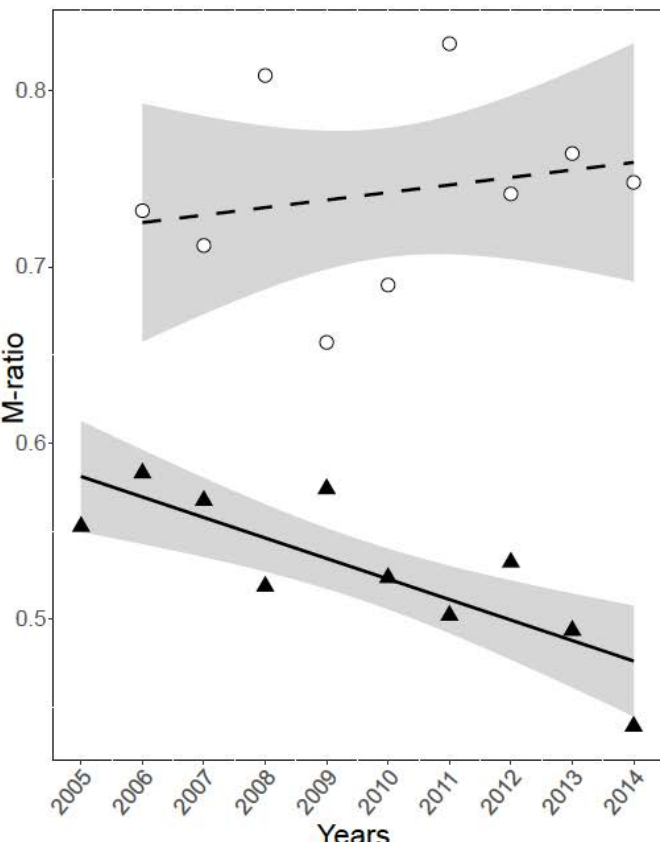
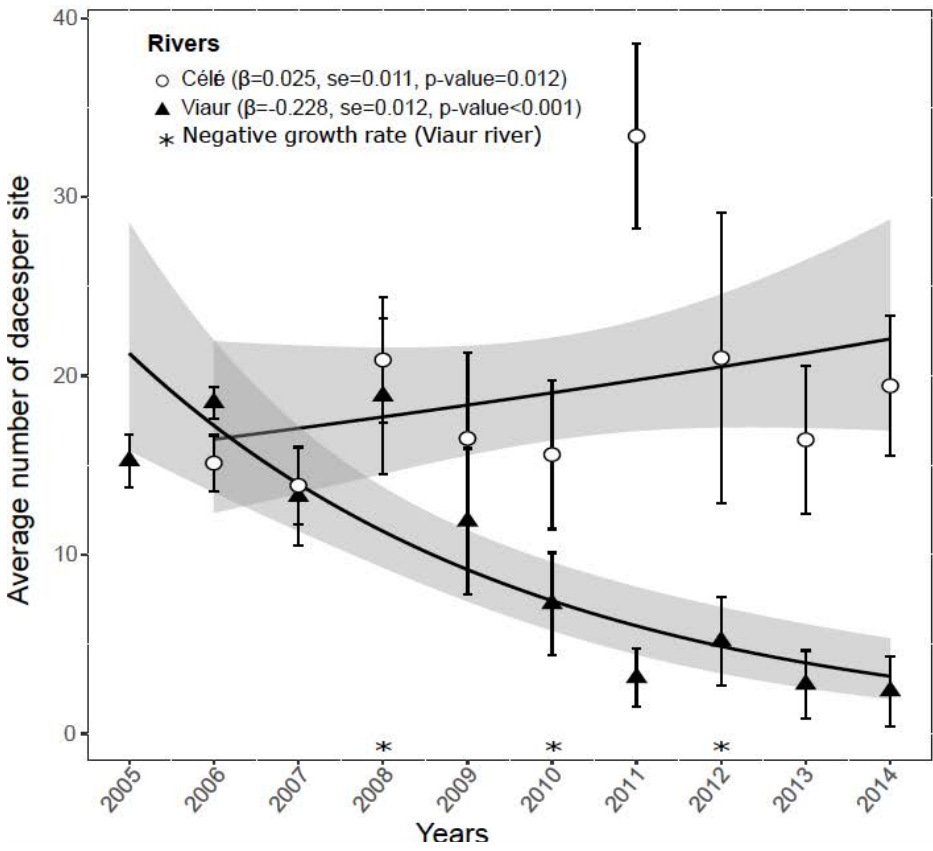
Environmental niche limitations
vs. host resistance in uninfected
areas?
→ Combining niche modeling
and genomic analyses

UNDERSTANDING THE DRIVERS OF RESISTANCE AT LARGE
SPATIAL SCALE



Eglantine Mathieu-Bégné

LONG-TERM CONSEQUENCES

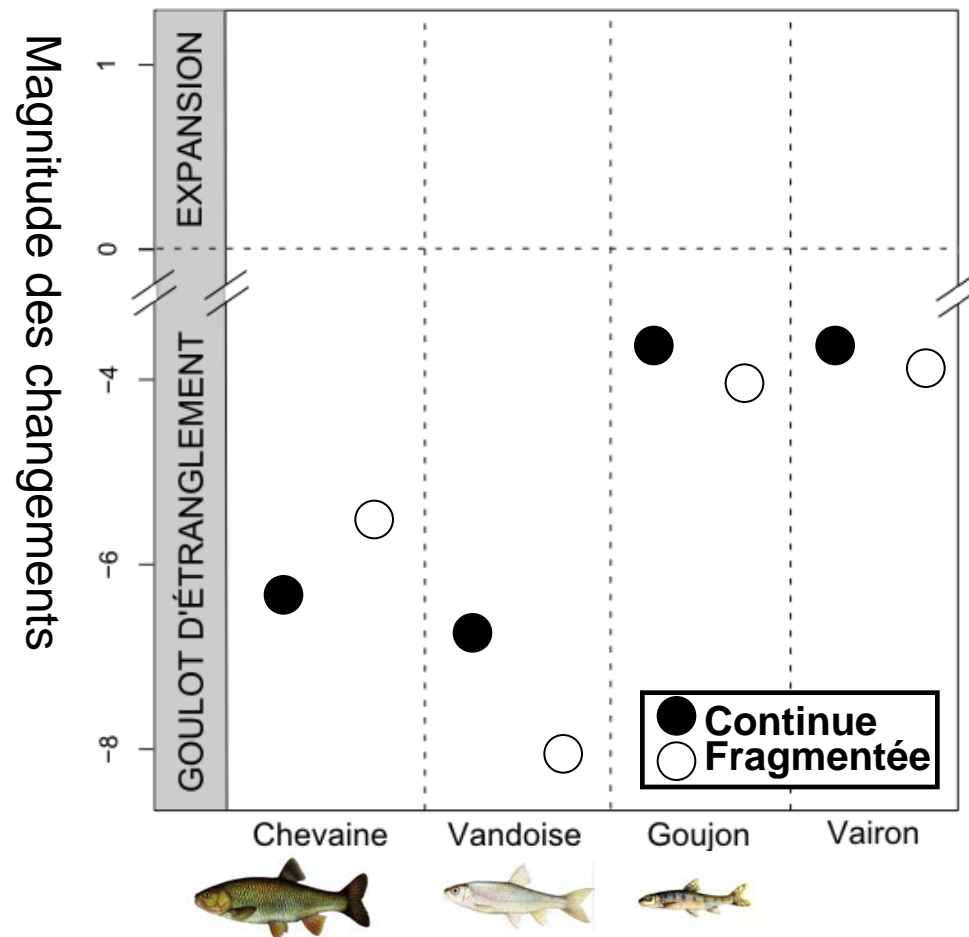


TEMPORAL DYNAMICS OF NEUTRAL AND NON-NEUTRAL GENETIC DIVERSITY IN HOST POPULATIONS

DYNAMIQUE DÉMOGRAPHIQUE DES POPULATIONS :

⇒ Restriction (goulot d'étranglement) attendue en cas de perte d'habitat

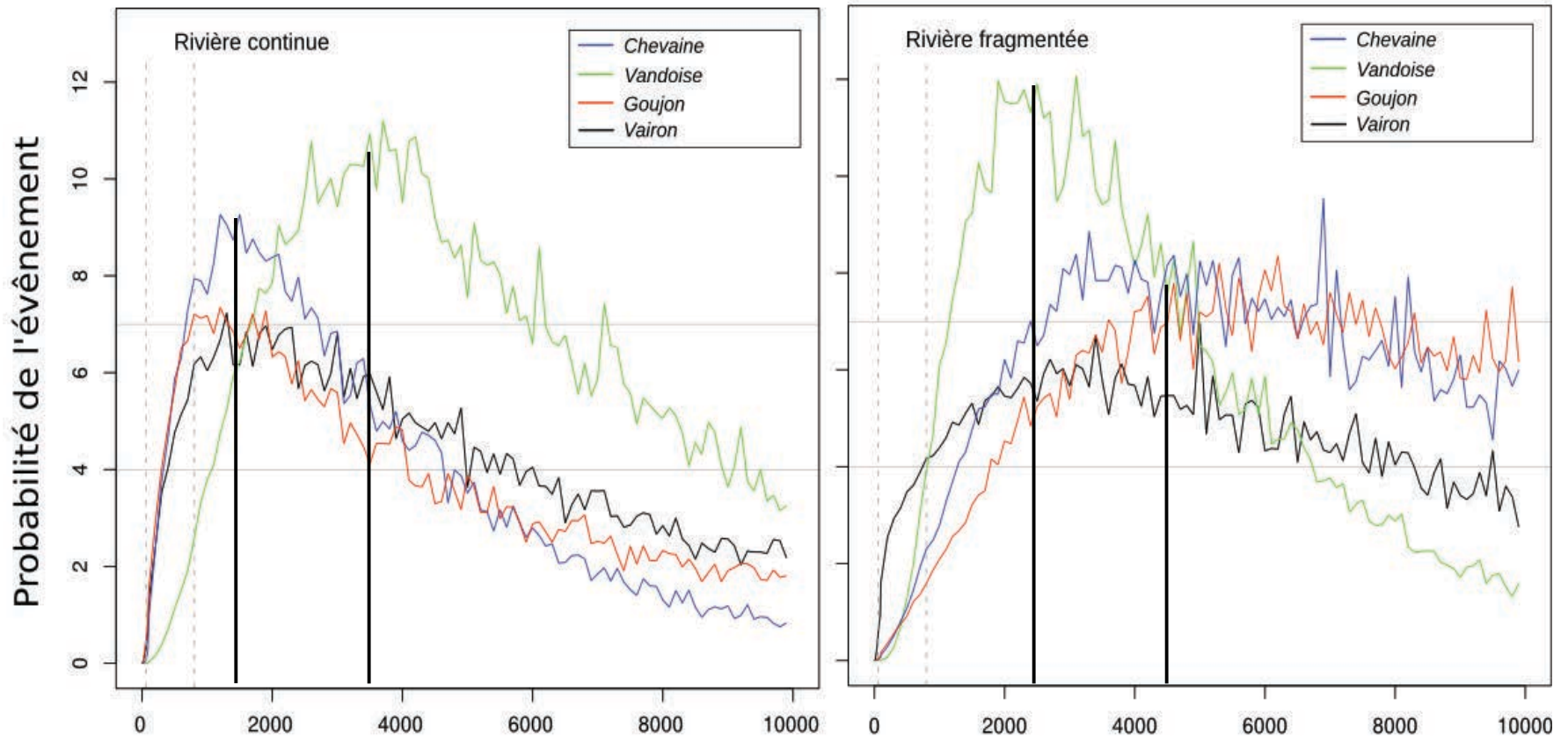
Deux rivières (Viaur et Célé) - Dix stations / rivière



DIFFÉRENCE ENTRE TAILLE DE LA POP ANCESTRALE ET TAILLE DE LA POP ACTUELLE

FORTS GOULOTS D'ÉTRANGLEMENT POUR TOUTES LES ESPÈCES DANS LES DEUX RIVIÈRES

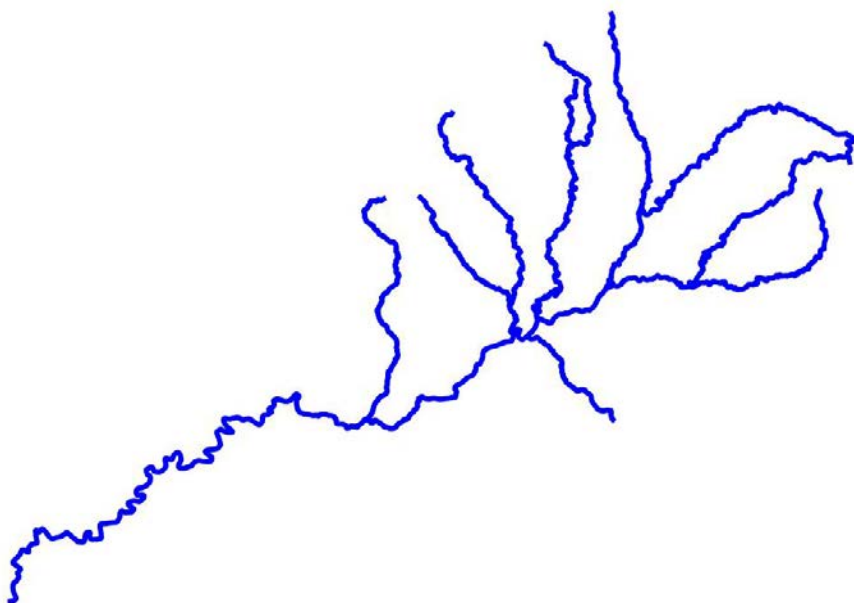
DYNAMIQUE DÉMOGRAPHIQUE DES POPULATIONS :



LA FRAGMENTATION NE SEMBLE PAS ÊTRE À L'ORIGINE DES GOULOTS D'ÉTRANGLEMENT DANS LE VIAUR ET LE CÉLÉ

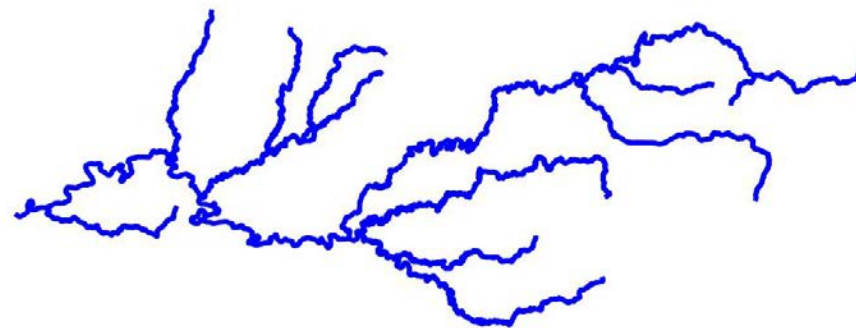
DEUX RÉSEAUX HYDROGRAPHIQUES :

LE CÉLÉ



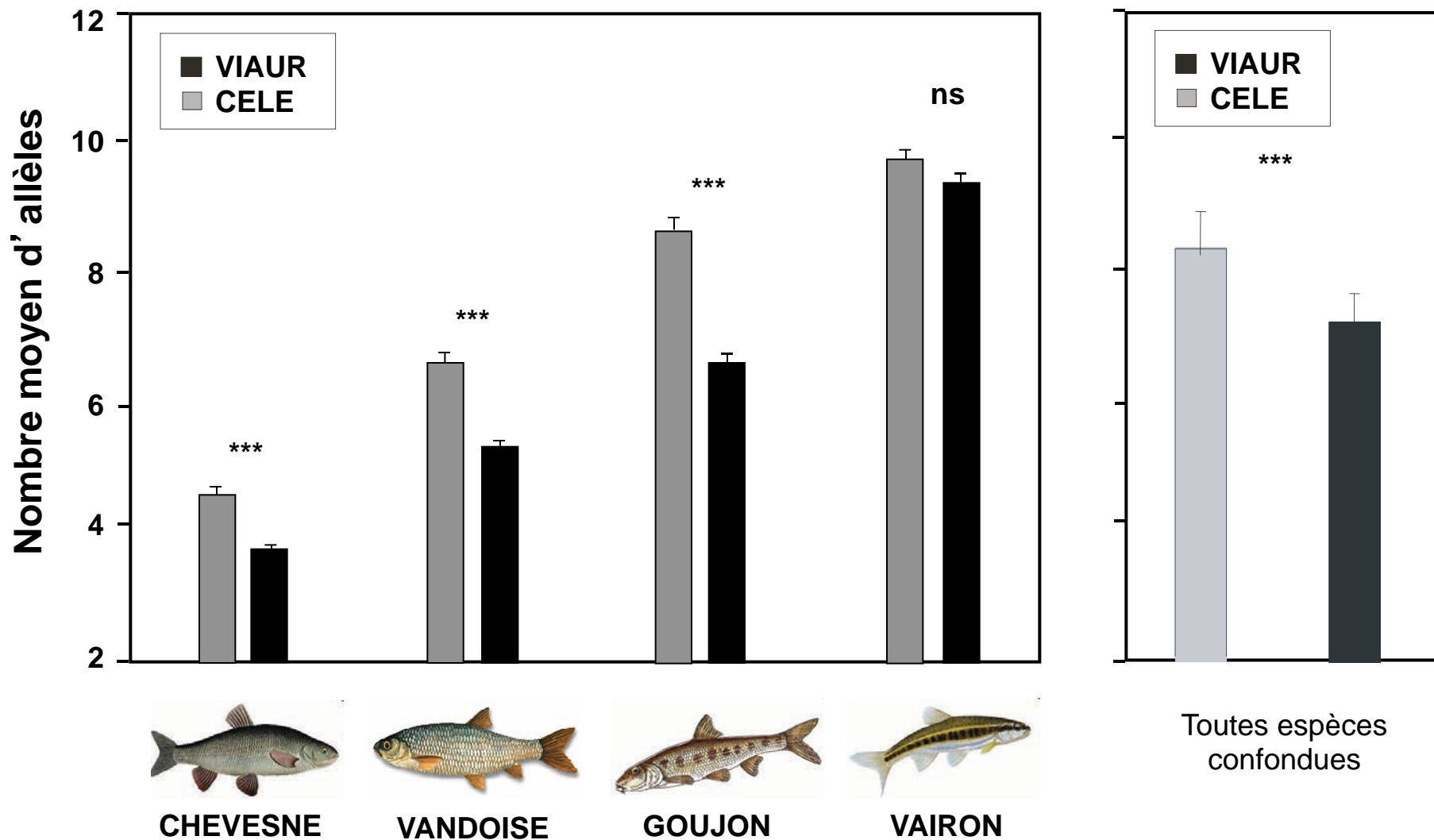
“Faiblement fragmentée”
(~15 chaussées, passes à poissons)

LE VIAUR

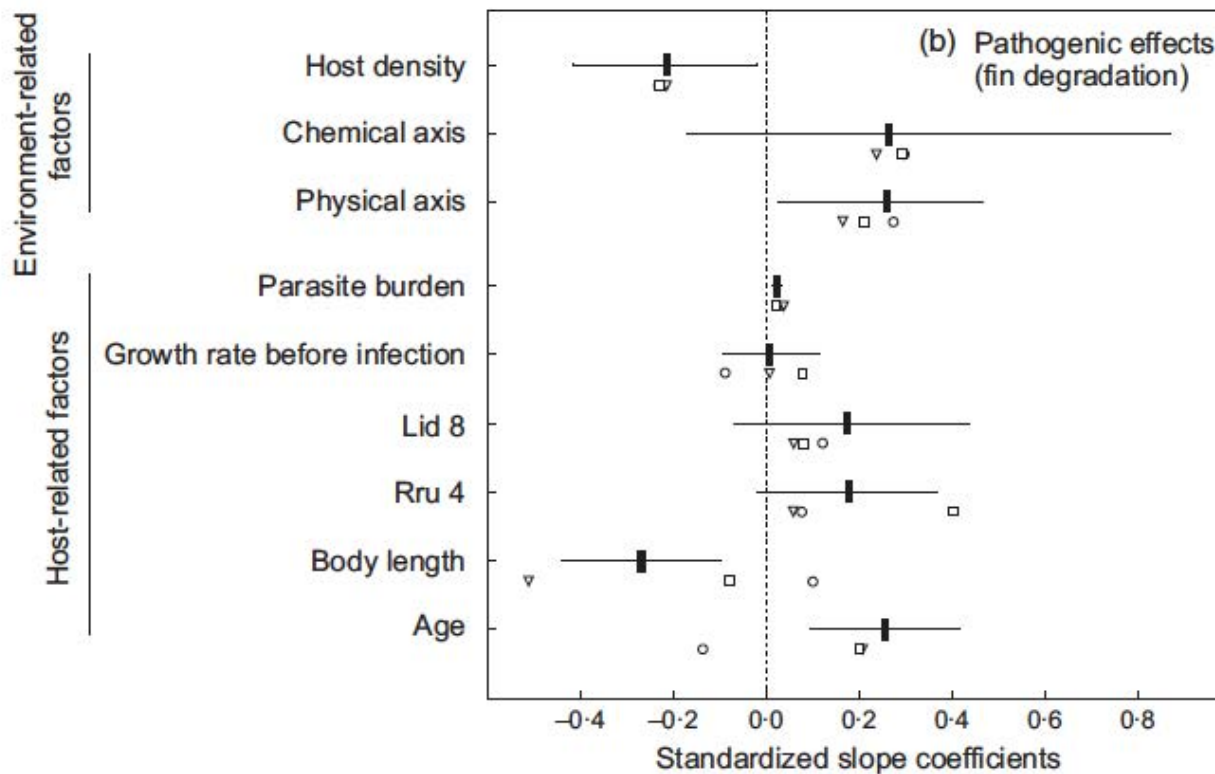
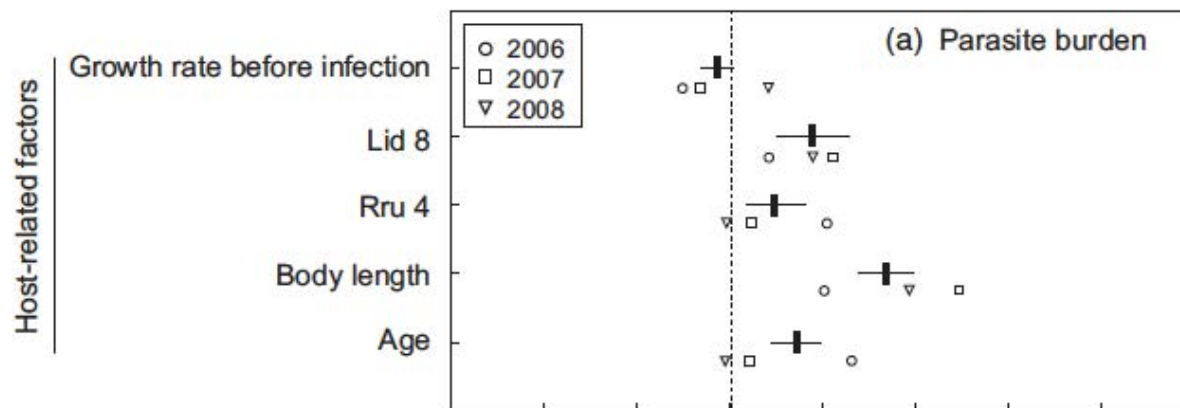


“Fortement fragmentée”
(~50 chaussées, 2 barrages hydro)

“FORTEMENT FRAGMENTÉ” = DIVERSITÉ PLUS FAIBLE

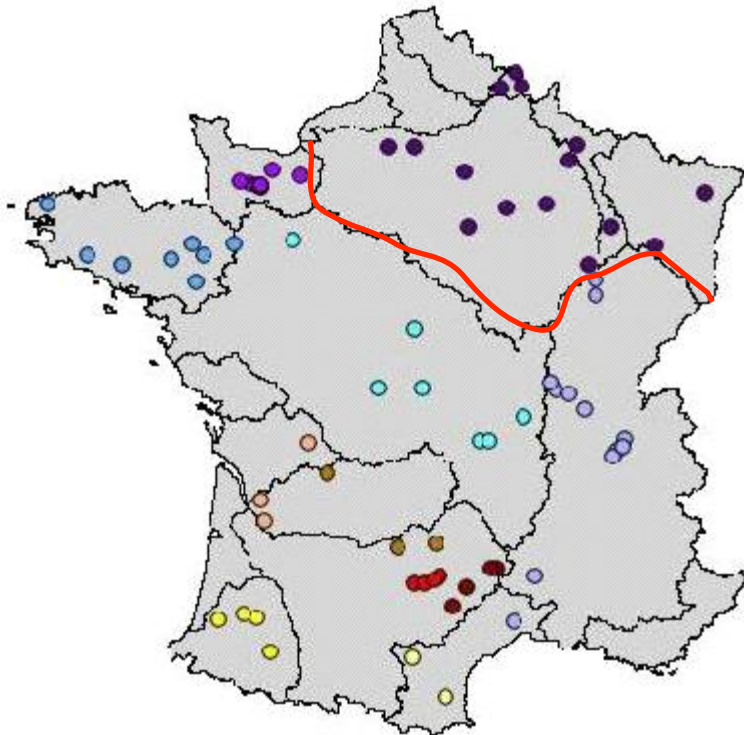


(III) RÉSISTANCE ET TOLÉRANCE

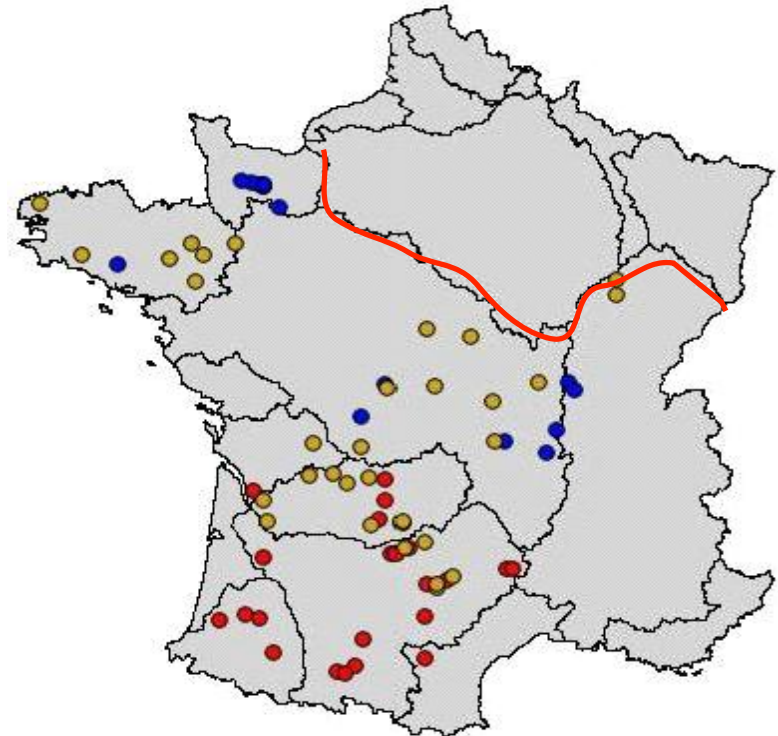


STRUCTURE GÉNÉTIQUE HÔTE ET PARASITE (MICROSATELLITES)

HOST



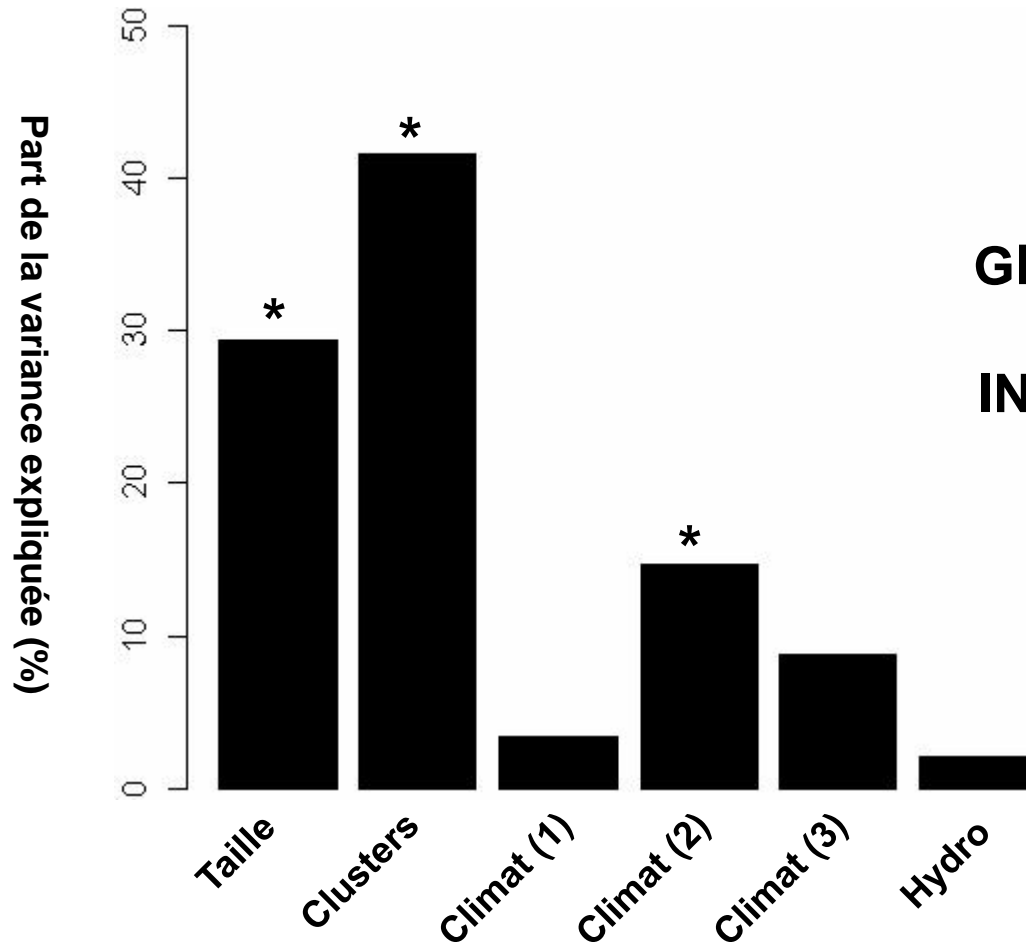
PARASITE



ASSOCIATION GÉNOTYPE – PRÉVALENCE?

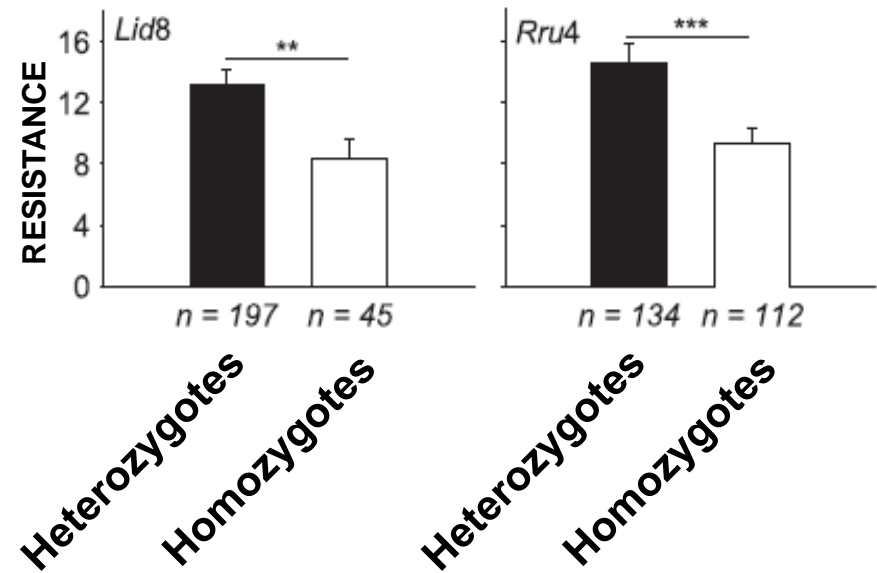
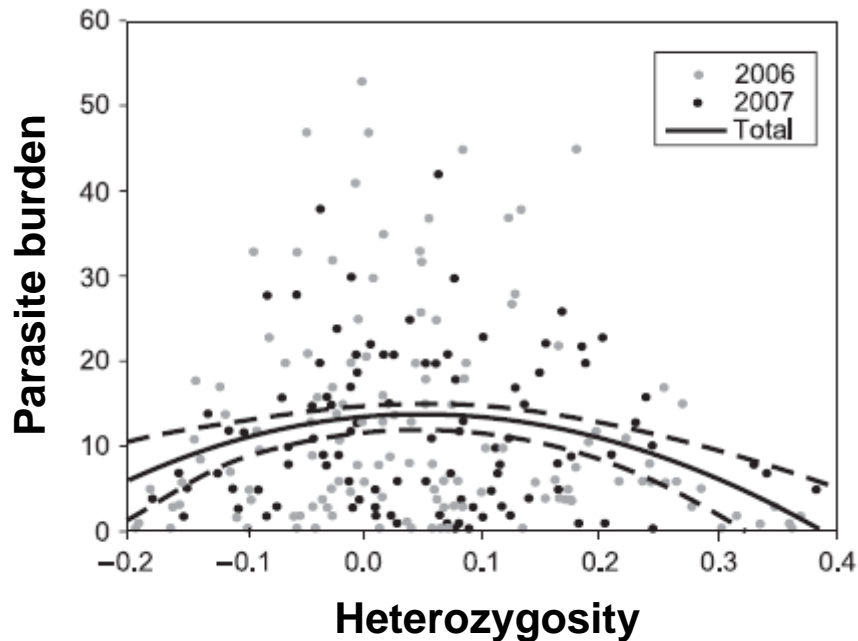
PARTITIONEMENT DE LA VARIANCE

(PRÉVALENCE~TAILLE+CLUSTER GÉNÉTIQUE+HYDROMORPHO+CLIMAT)



**ASSOCIATION
GÉNOTYPE – PRÉVALENCE
SIGNIFICATIVE ET
INDÉPENDANTE D'AUTRES
CO-FACTEURS**

DÉTERMINISMES ENVIRONNEMENTAUX ET GÉNÉTIQUES



RELATIONS SIGNIFICATIVE ENTRE DIVERSITÉ GÉNÉTIQUE (MICROSAT), TOLÉRANCE ET RÉSISTANCE

ET LA RECHERCHE APPLIQUÉE ?

=> CAPACITÉ DE COLONISATION

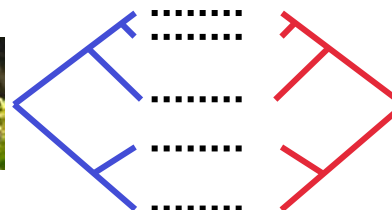


(1) Filtre environnemental

=> Facteurs abiotiques (T°C, courant, oxygénation)

=> Facteurs biotiques (densité de l'hôte)

(2) Capacité de dispersion



(3) Adaptation locale

=> Expérimentation en mésocosmes



(4) Capacité de transfert d'hôtes

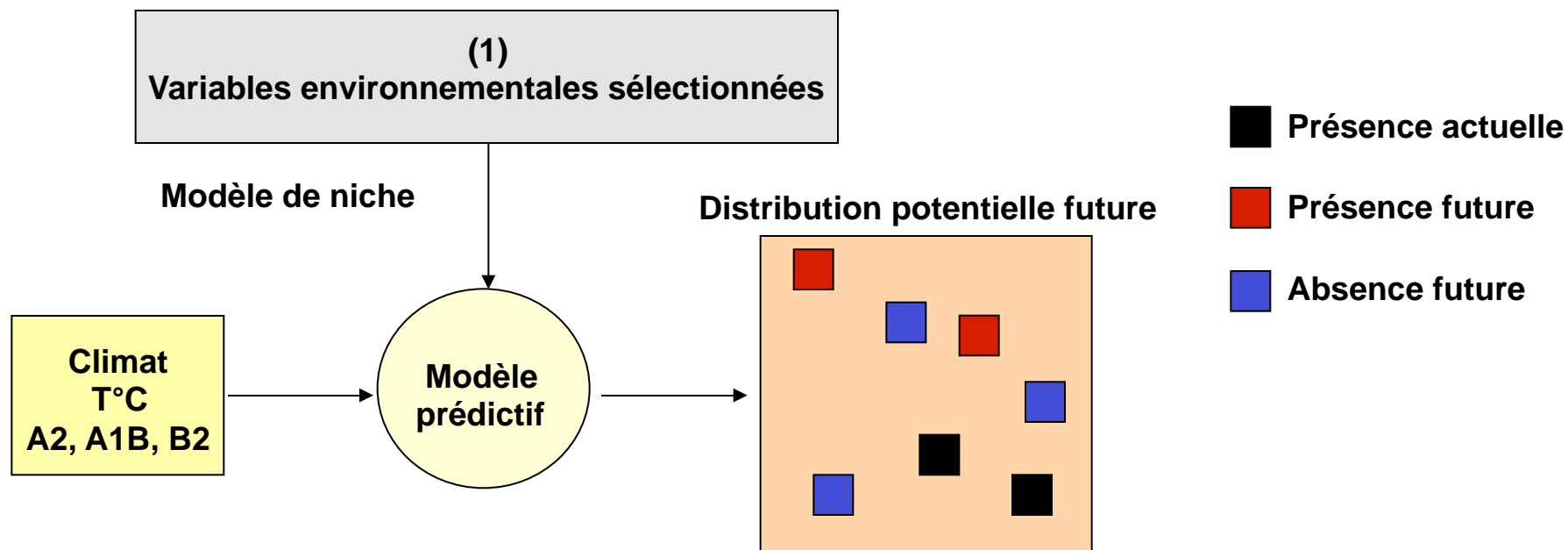
=> Etude descriptive comparative

=> Etude fitness parasite



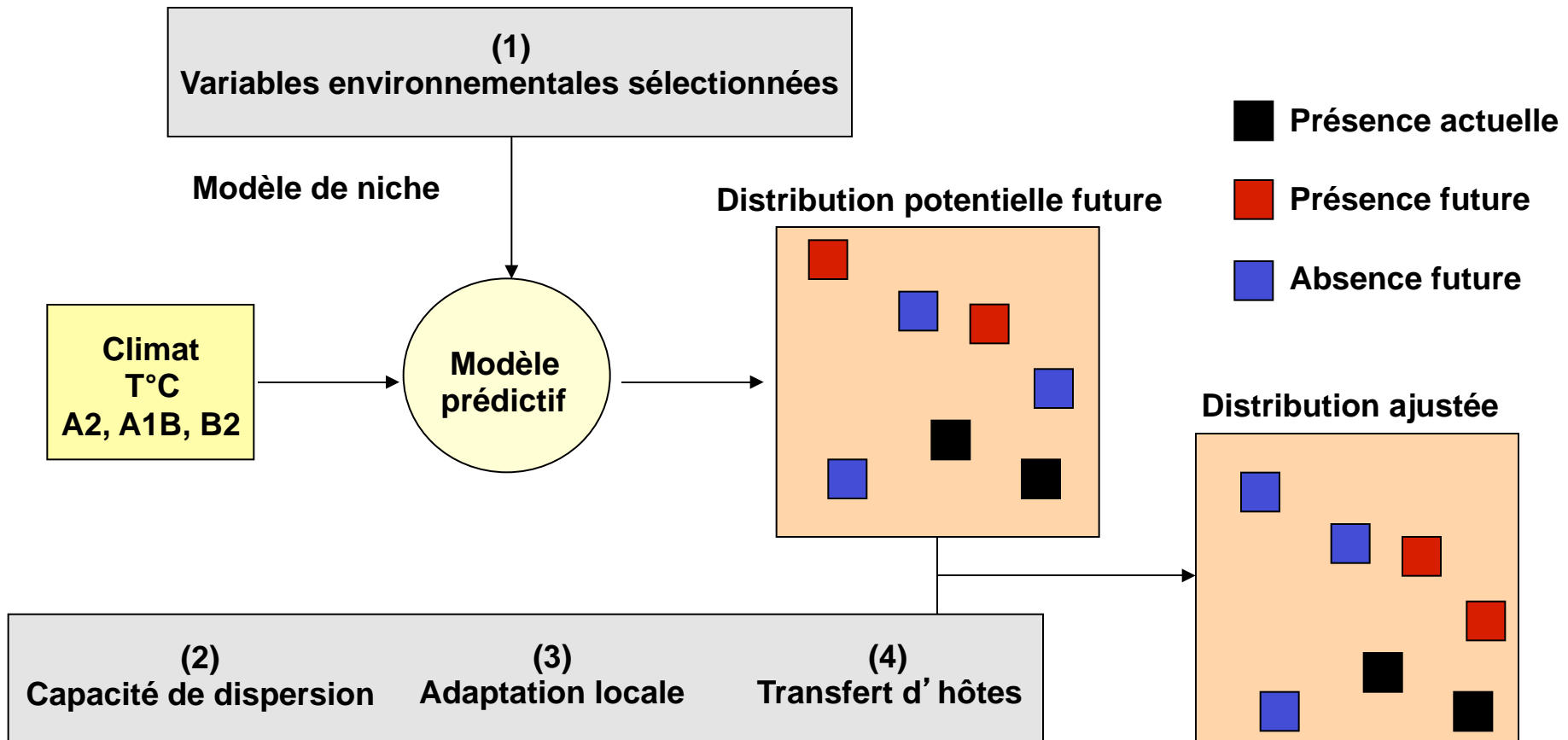
ET LA RECHERCHE APPLIQUÉE ?

=> PRÉDIRE DISTRIBUTION FUTURE



ET LA RECHERCHE APPLIQUÉE ?

⇒ PRÉDIRE DISTRIBUTION FUTURE



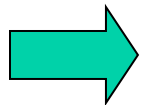
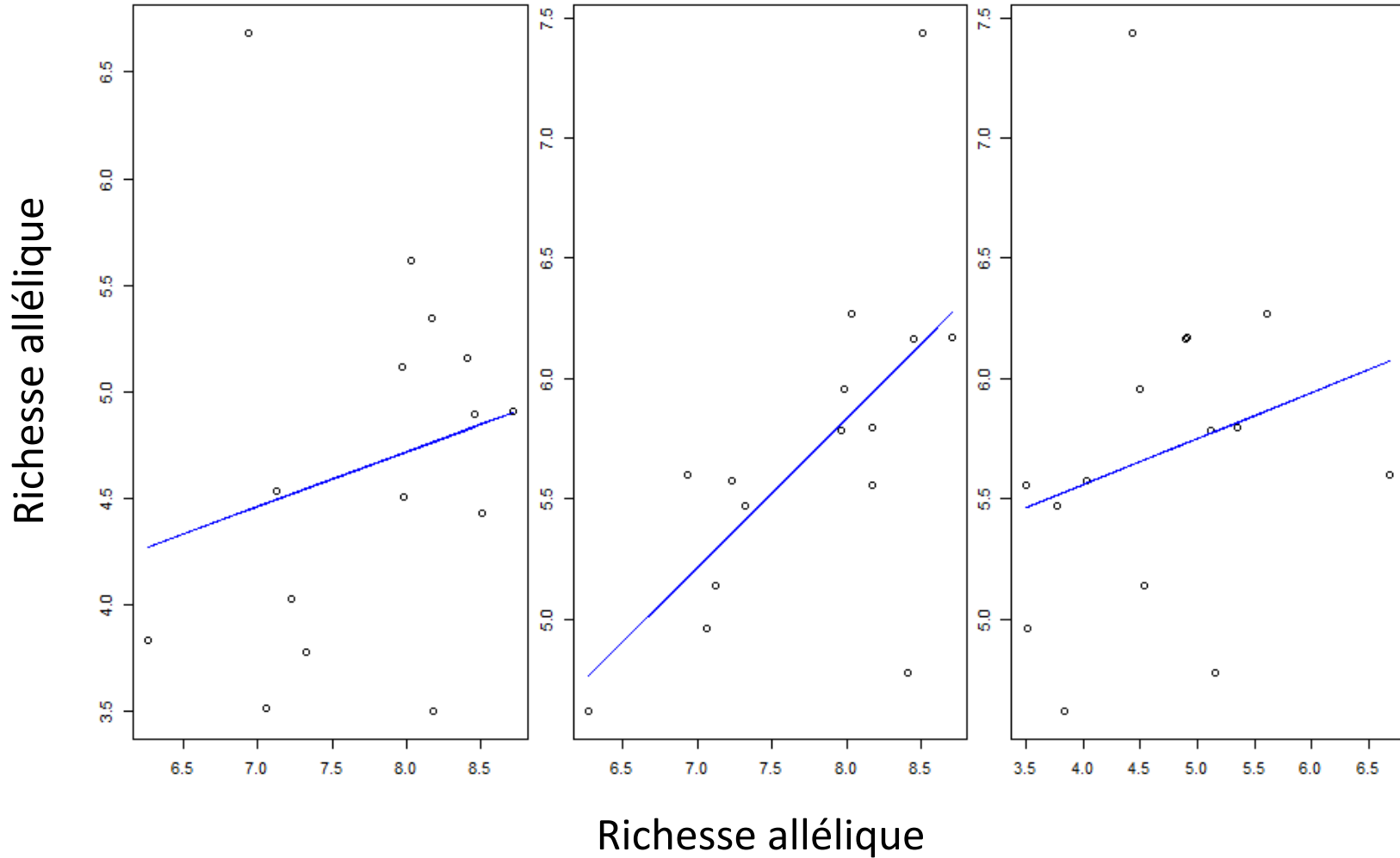
⇒ Carte de distribution future : outil de gestion
⇒ Modèles prédictifs plus réalistes

III - PREMIERS RÉSULTATS

Chevaine vs

Vandoise vs

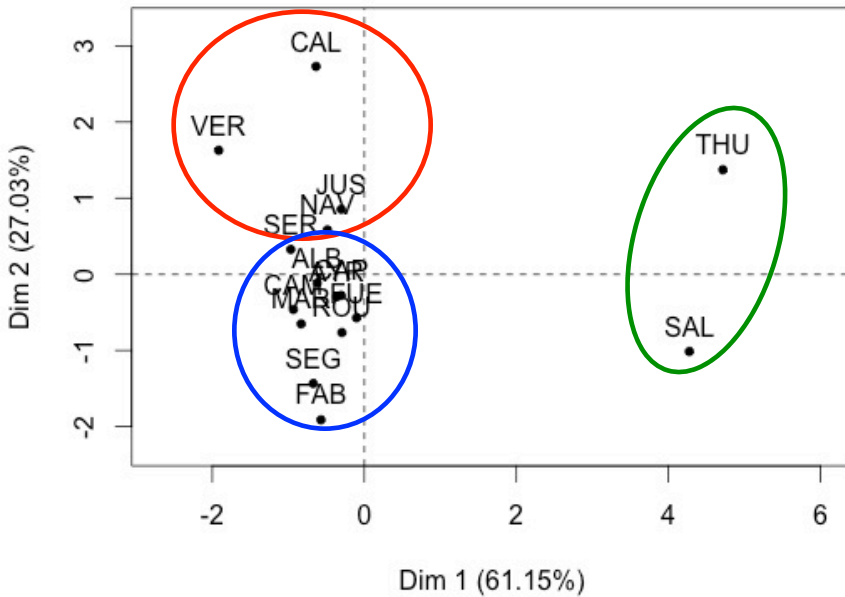
Vandoise vs



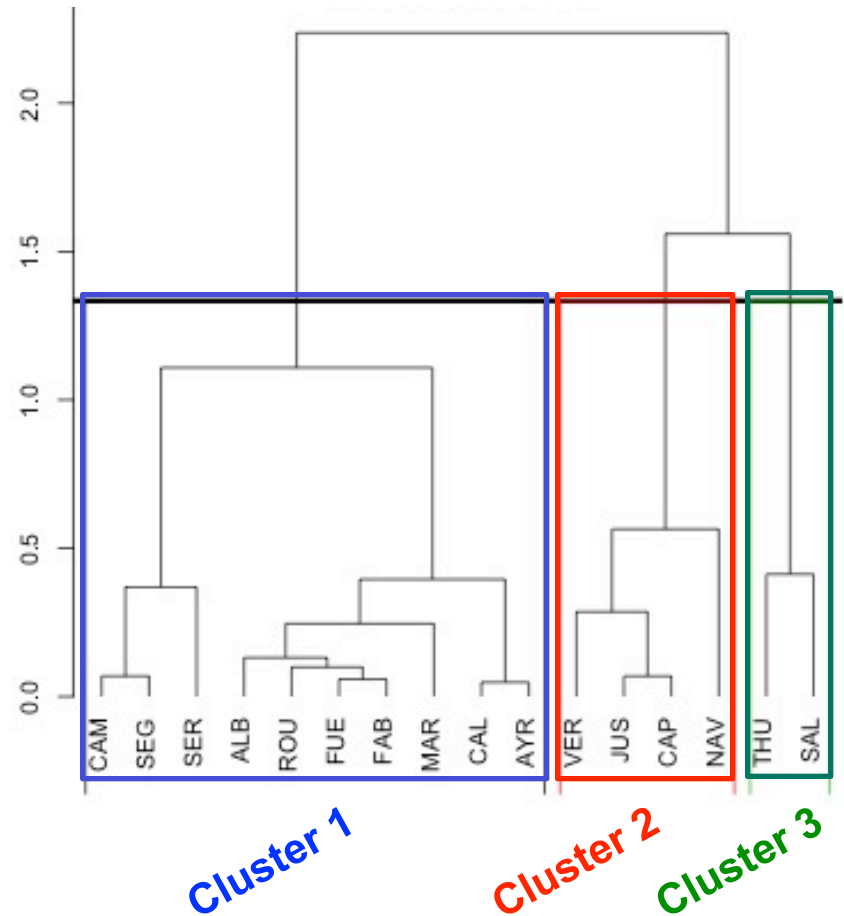
Congruence pour la vandoise et le goujon

VIAUR

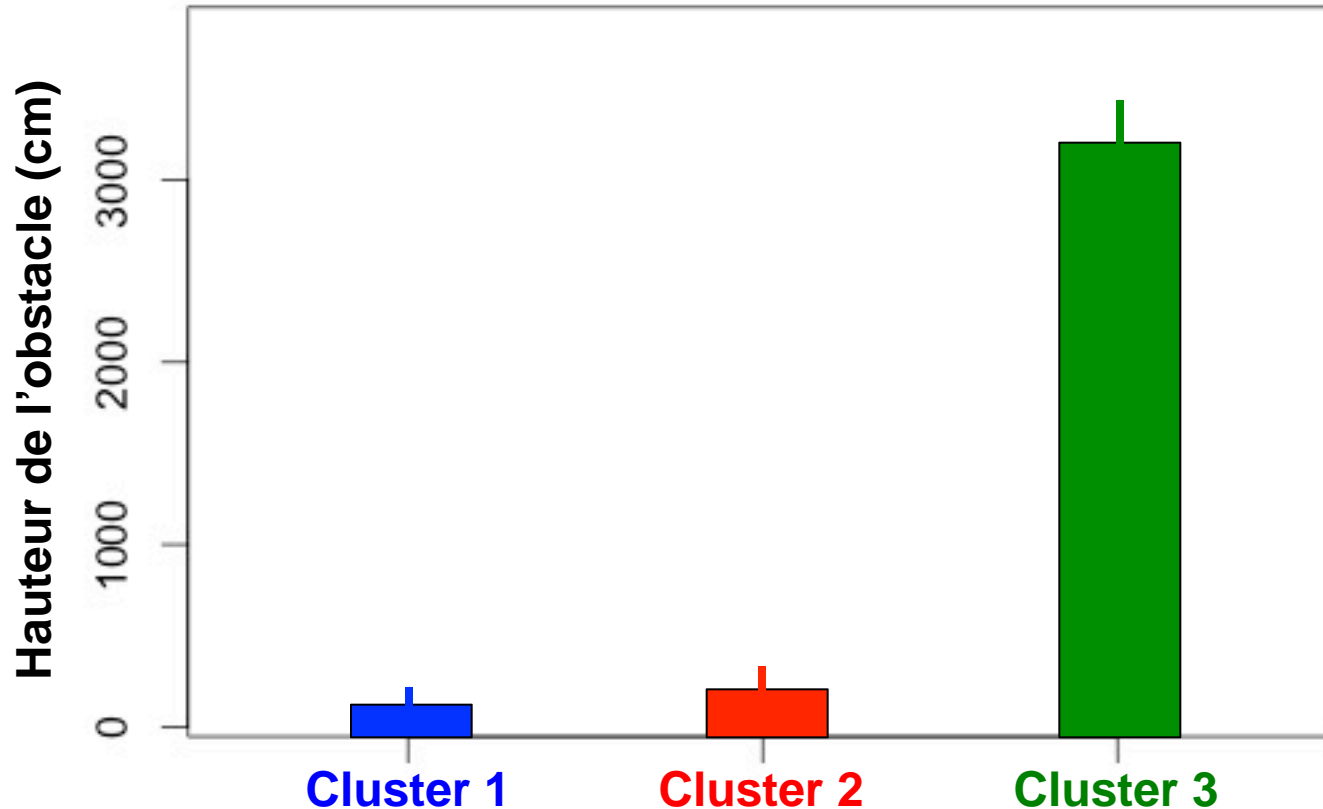
ACP SUR INDICES DE CONNECTIVITÉ



CLUSTERING



VIAUR



⇒ Pas d'effets de distance à la source, pente, tirant d'eau
⇒ Importance de la hauteur de l'obstacle
(barrages hydroélectriques)