

Investigating large scale patterns
to try to understand local processes

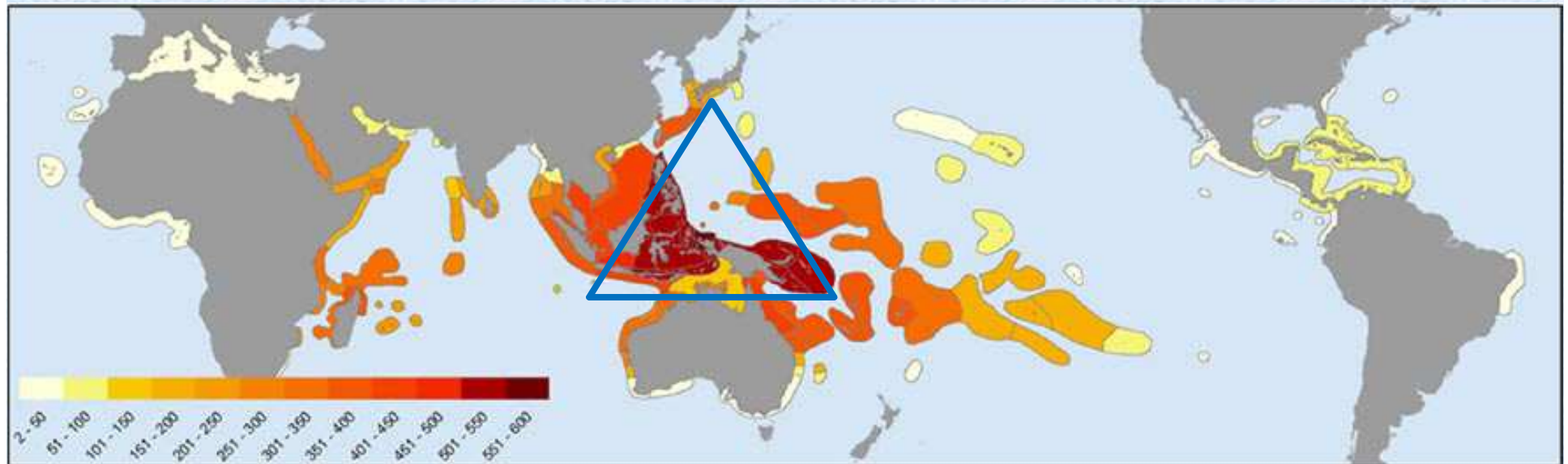
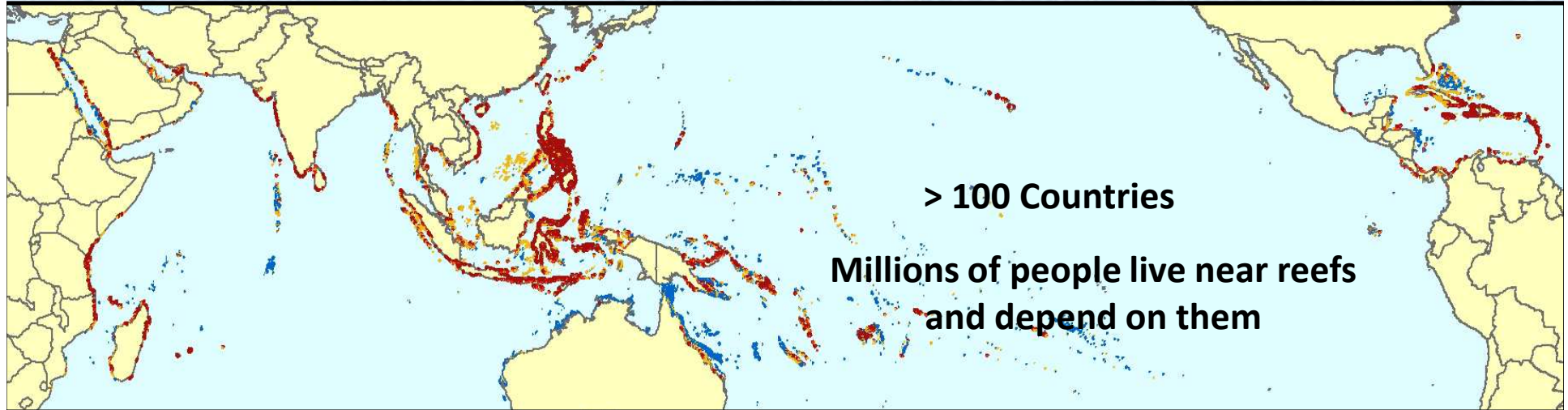
The exemple of Reef Fishes

The GASPAR Group
presented by Michel KULBICKI – IRD- UR 227 – CoReUs – Banyuls/mer - France

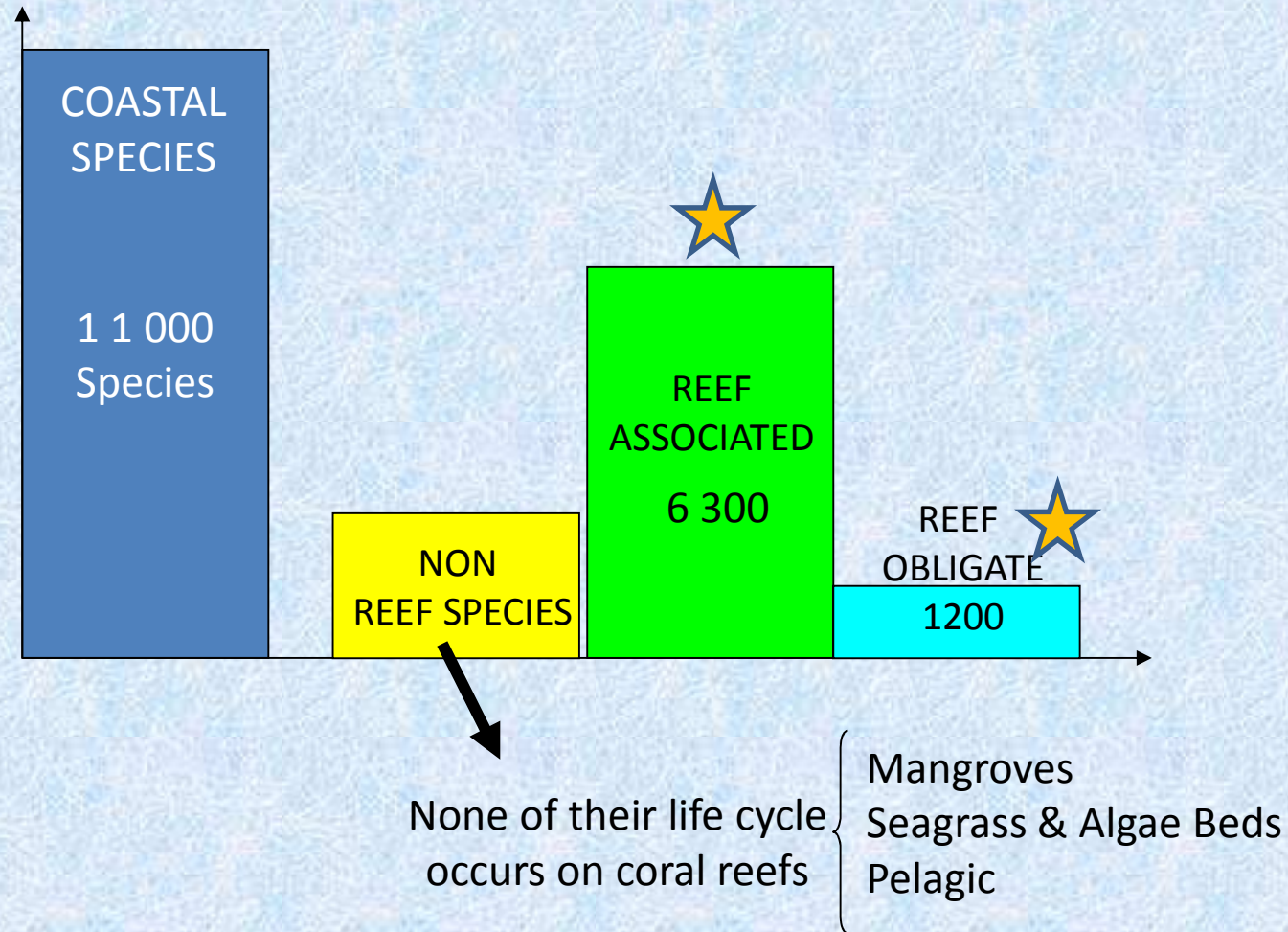
Coral Reefs still exist



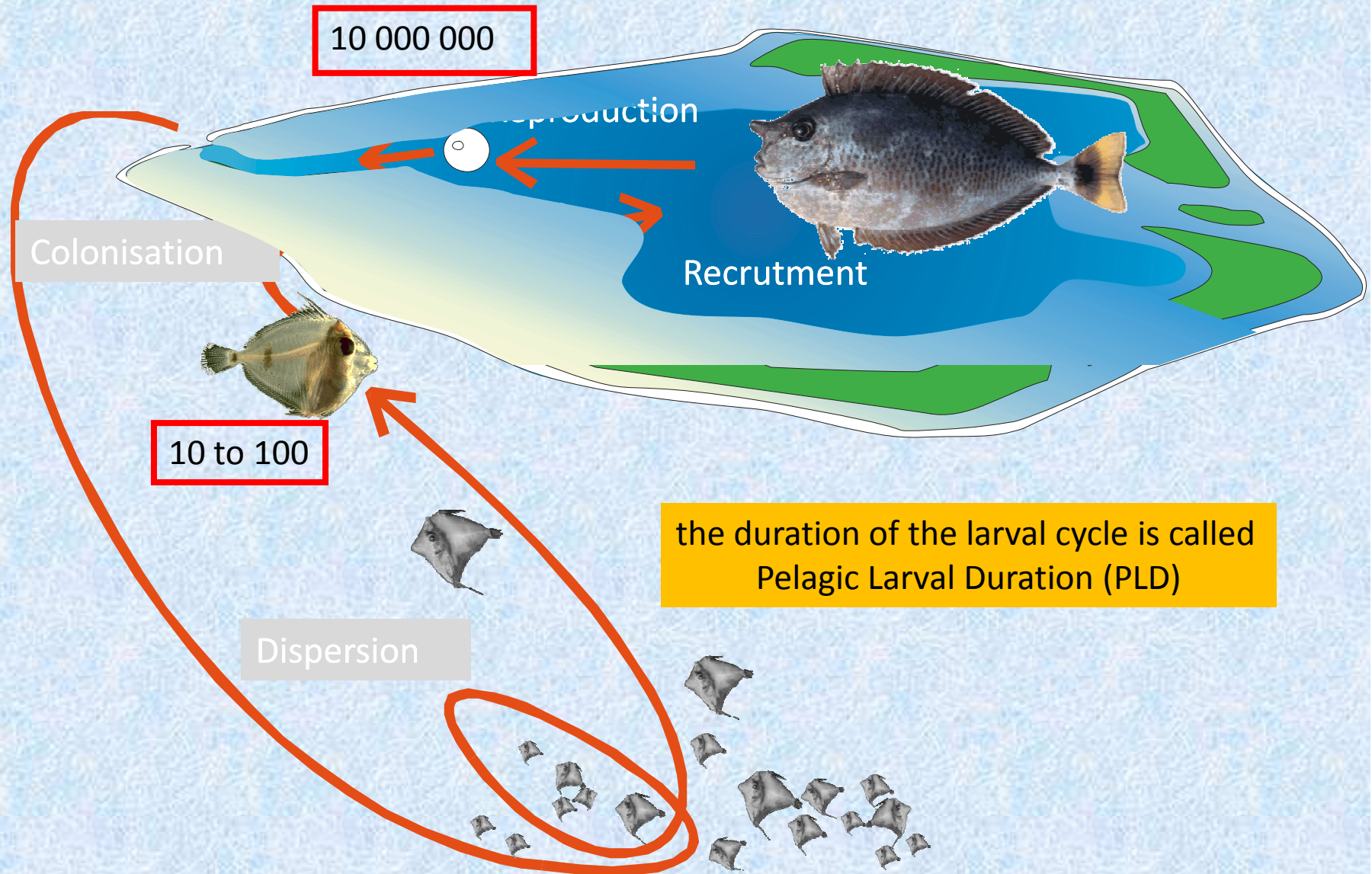
Where are Coral Reefs ?



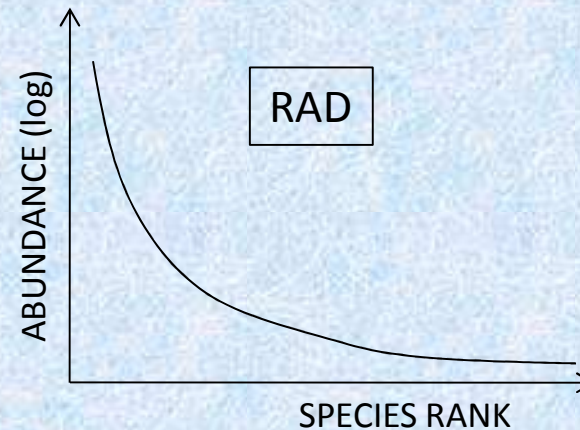
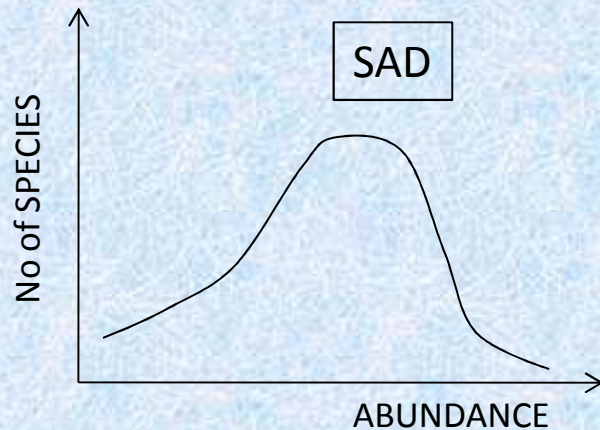
Reef Fishes – How many battalions ?



Life cycle of a reef fish

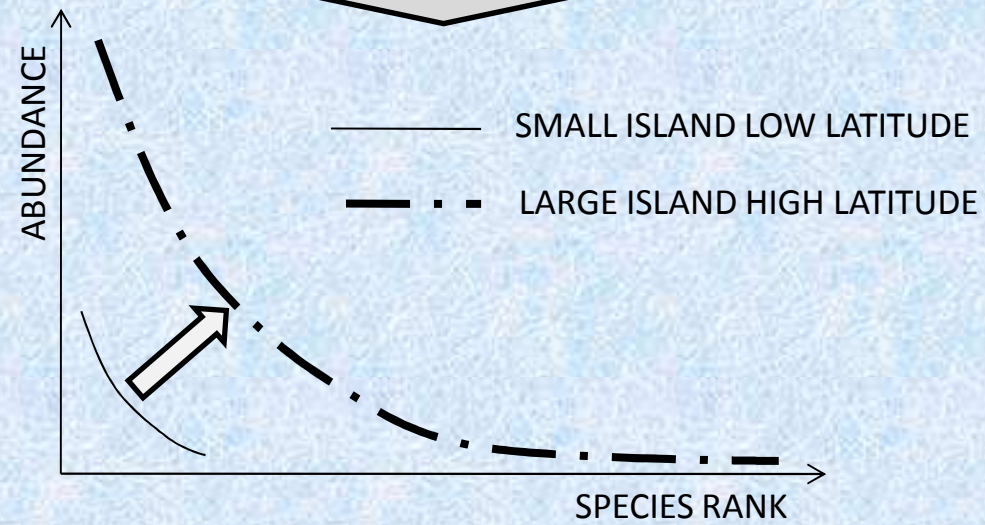
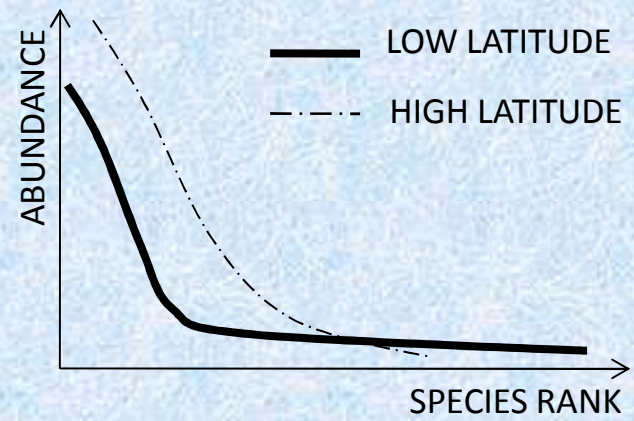
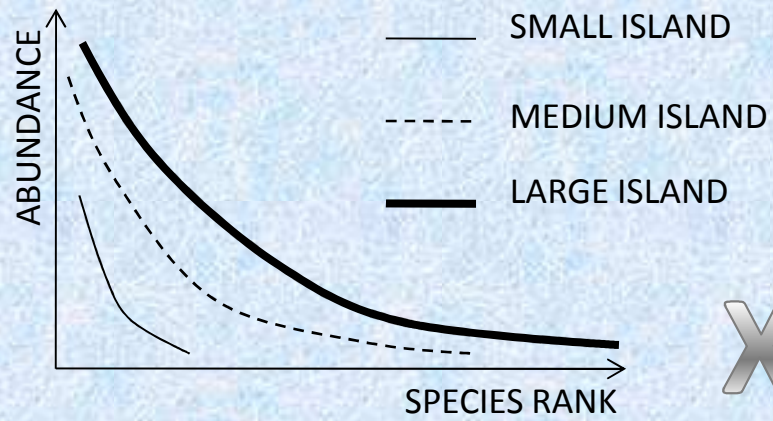


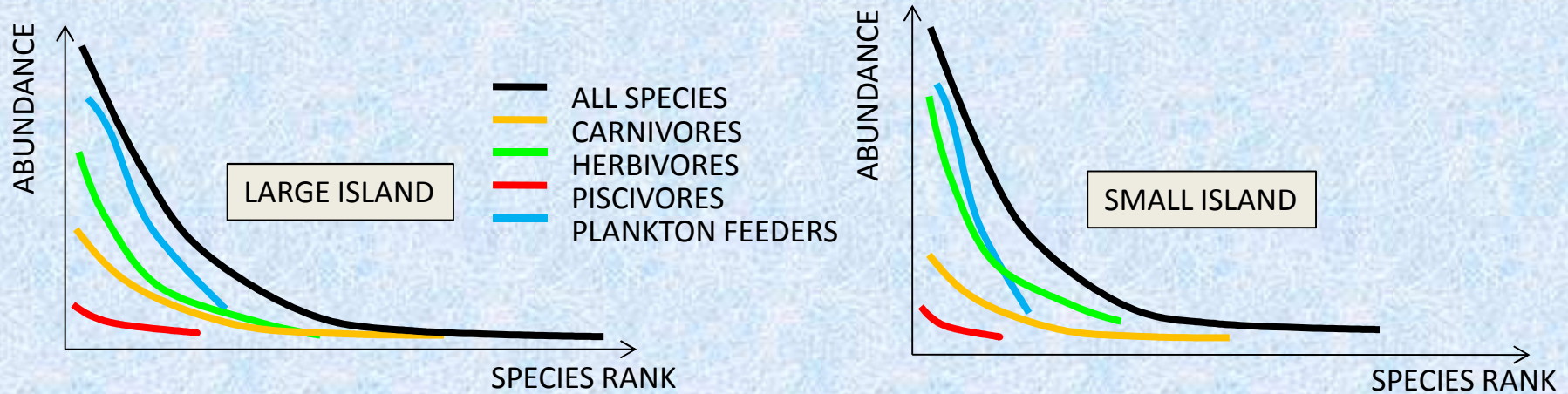
Why were we interested by Reef Fish global diversity ?



Explain local abundance according to both LOCAL & REGIONAL factors

Include intrinsic factors (Life-history traits)





Knowing the Life History Traits of species

Finding out about assembly rules for species & traits

Access reliable abundance data

Access data on the physical, biogeochemical and human factors

3 year
program
GASPAR

Checklists (165) – 6300 teleost species



LHT: Diet, Size, Schooling, Home Range, Level in the water

Other traits for some species (PLD; growth; depth range and so on)

Physical , biogeochemical & human related data

Island and coral reef size

Island isolation

Geomorphological diversity of coral reefs

SST, salinity, chlorophyll and so on

Human density

MPAs : location, surface, status

Field Observations (UVC)

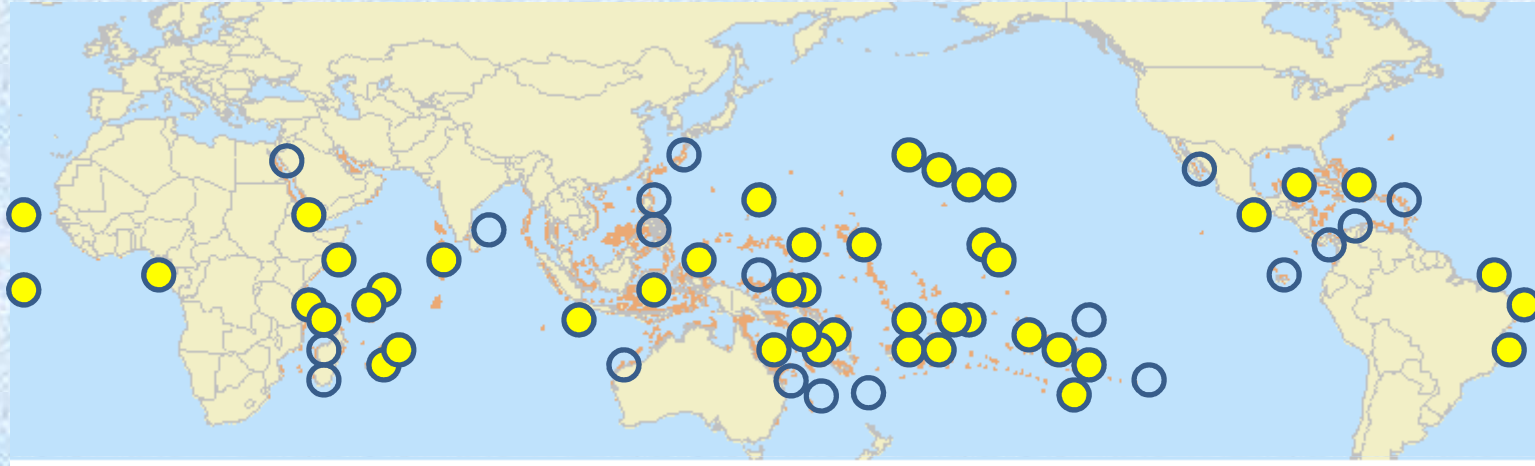
(1.5 million records + ~ 1 million additional)



REGIONS FOR WHICH THE TEAM HAS DATA



OTHER REGIONS POTENTIALLY AVAILABLE



➤ 12 000 transects

Species composition

Abundance

Size distribution

Benthic landscape

Biomass ; ~ Production

Biogeographical

Need for a global framework to organize our data

Existing limits (e.g. Spalding et al., Veron et al.) were not adapted to reef fish

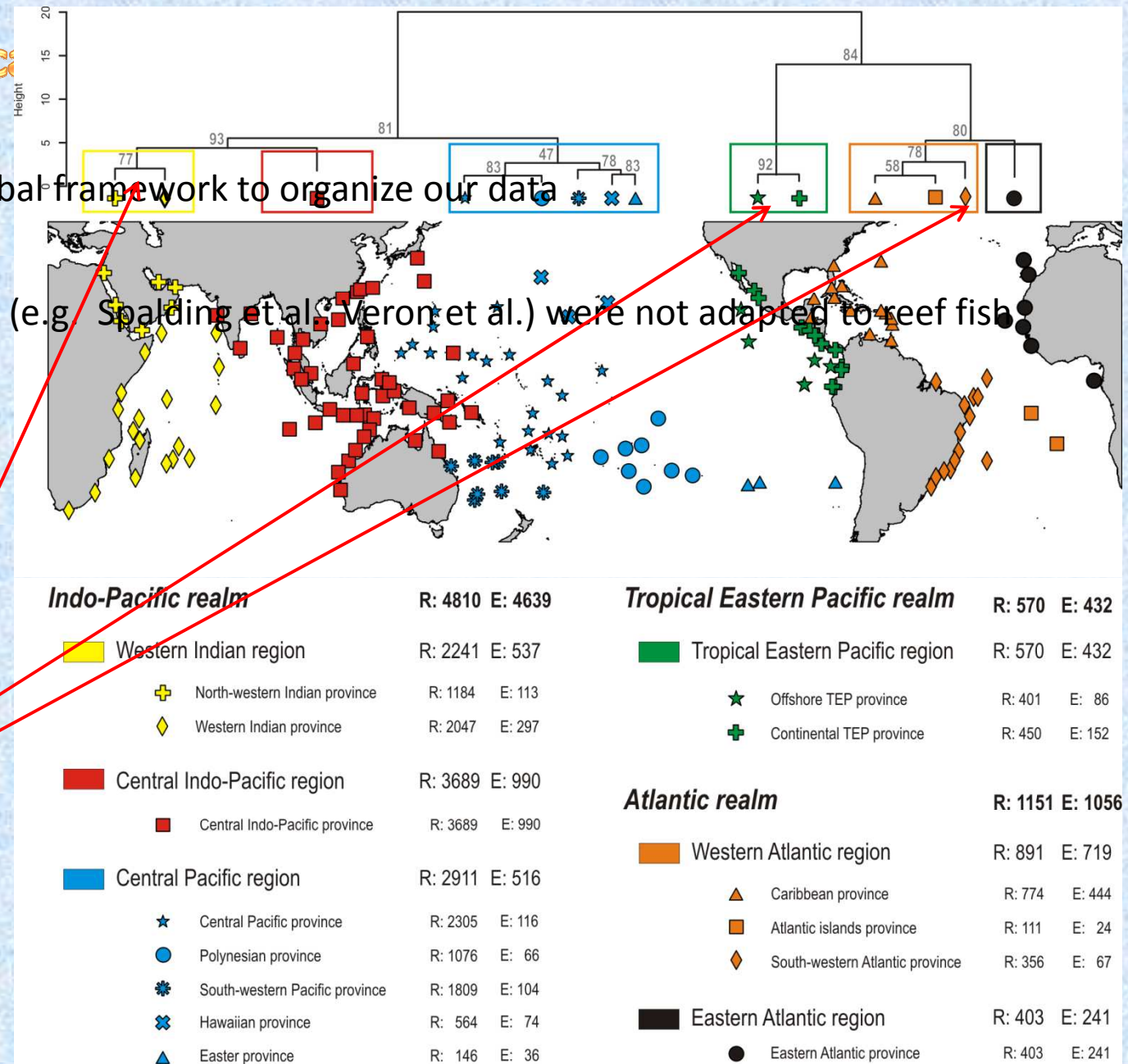
6300 species

Disimilarity

3 Realms

6 regions

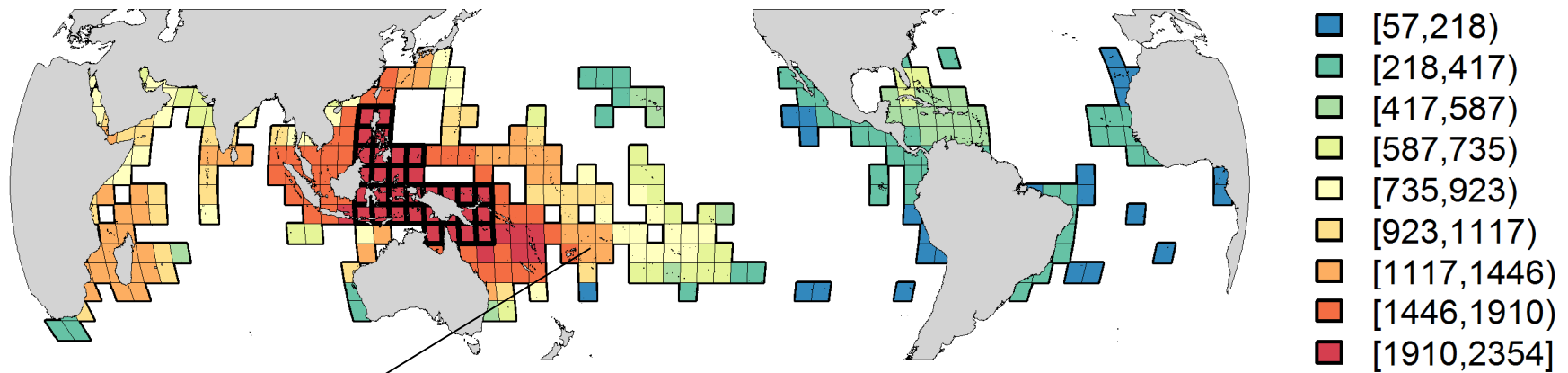
14 provinces



Kulbicki et al. (2013) Global Biogeography of Reef Fishes: A Hierarchical Quantitative Delineation of Regions. Plos One 8(12)

Mouillot et al. . 2013 The challenge of delineating biogeographical regions: nestedness matters. Journal of Biogeography

Total richness

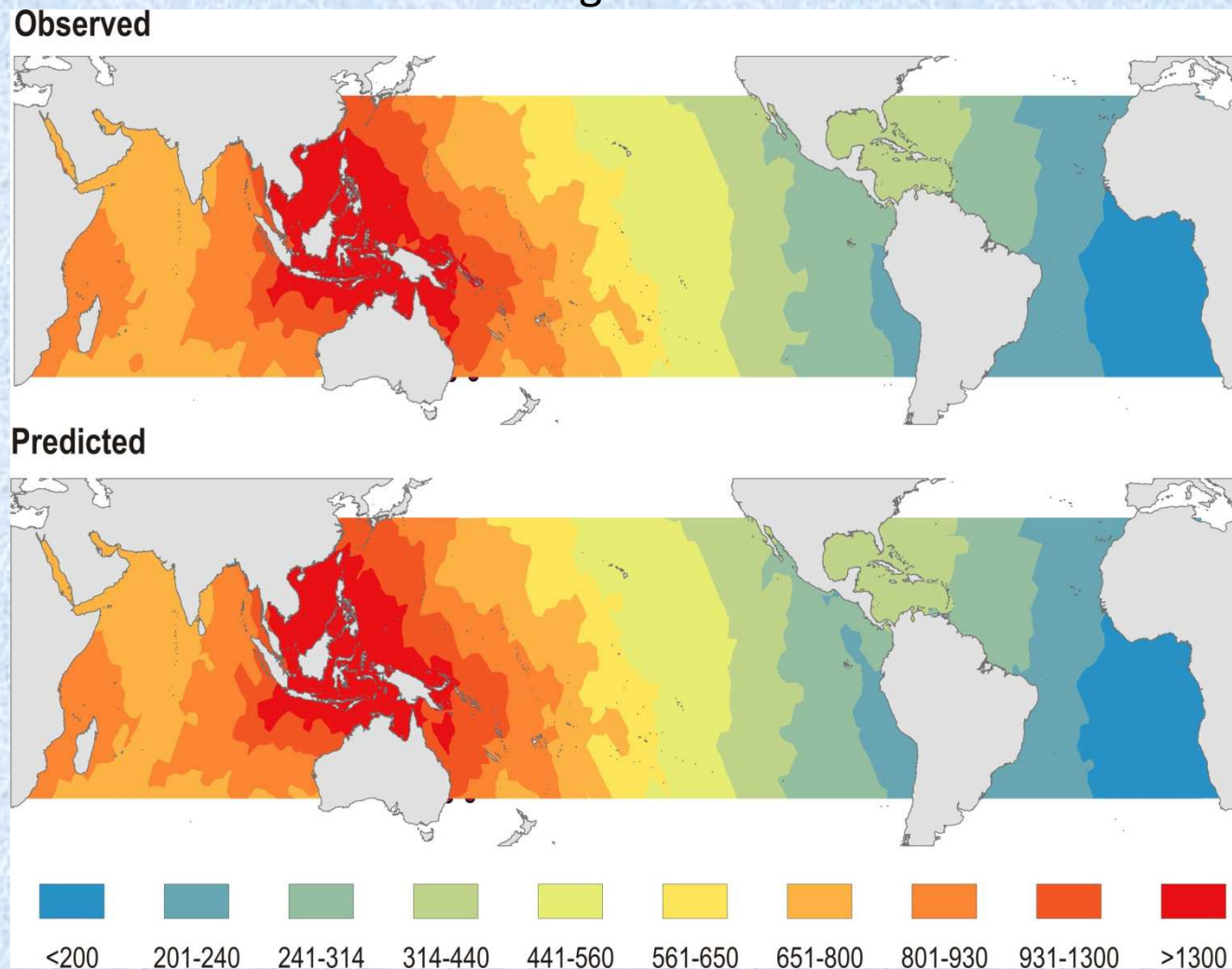


For each cell:

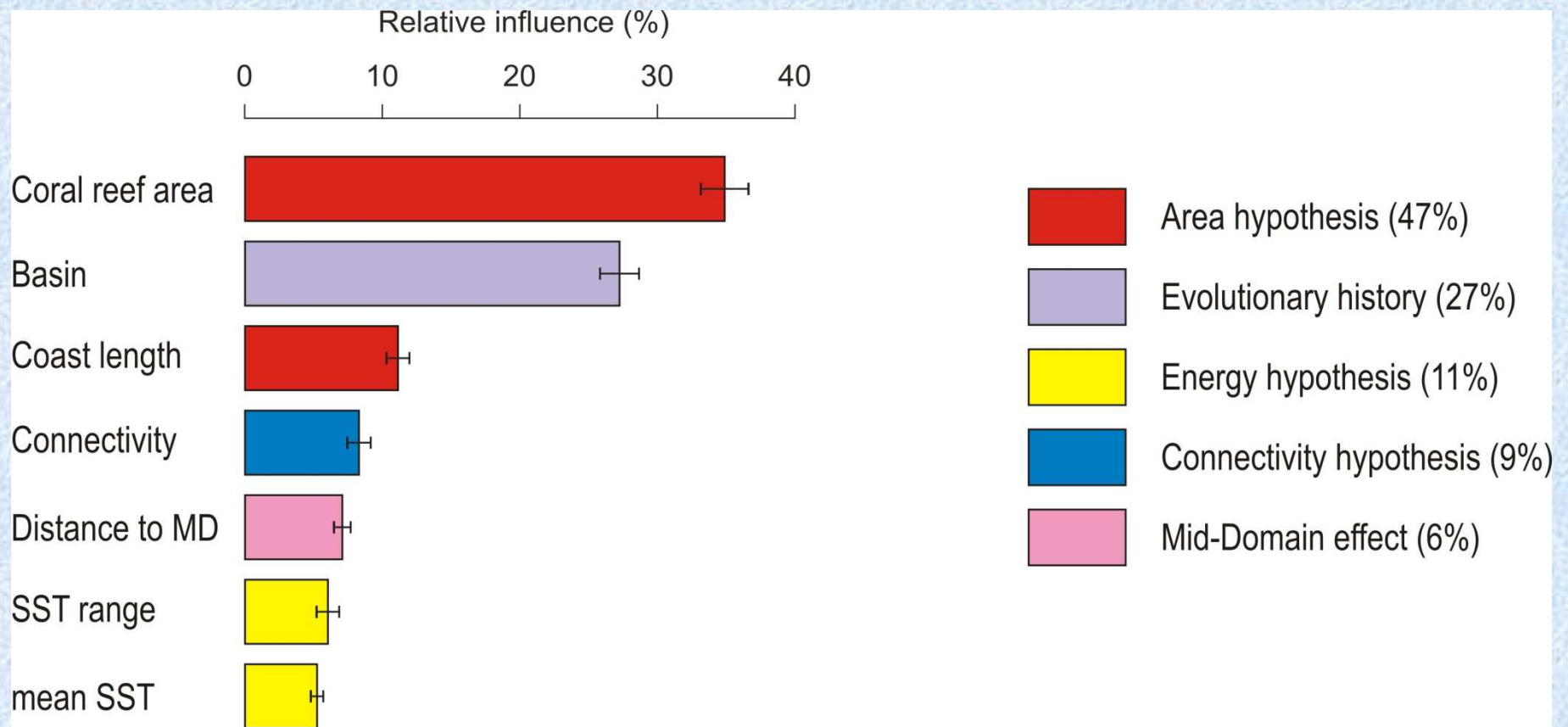
- Species composition
- Life-History Traits distribution
- Physical and biogeochemical factors
- Human associated data

Global patterns and predictors of reef fish diversity

Few and easily accessible variables are enough to accurately predict the richness of reef-fishes at global scale

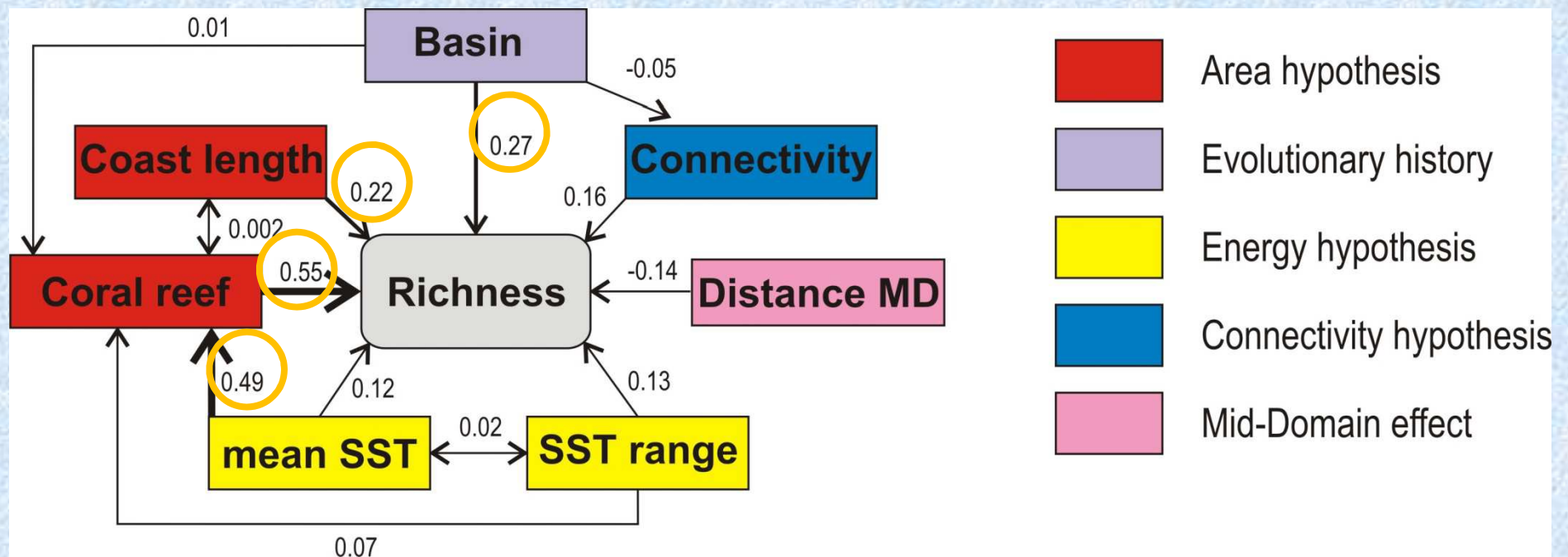


Coral reef area was the most influential variable, followed by evolutionary history, energy, connectivity and the mid-domain effect



More about interactions.... direct and indirect effects

Results of a Structural Equation Model (SEM) which allows to disentangle direct effects from indirect ones by combining several equations within the same model



The role of life-history traits

Several of the previous hypotheses imply an important role for:

1- dispersal capacities

2- colonizing capacities

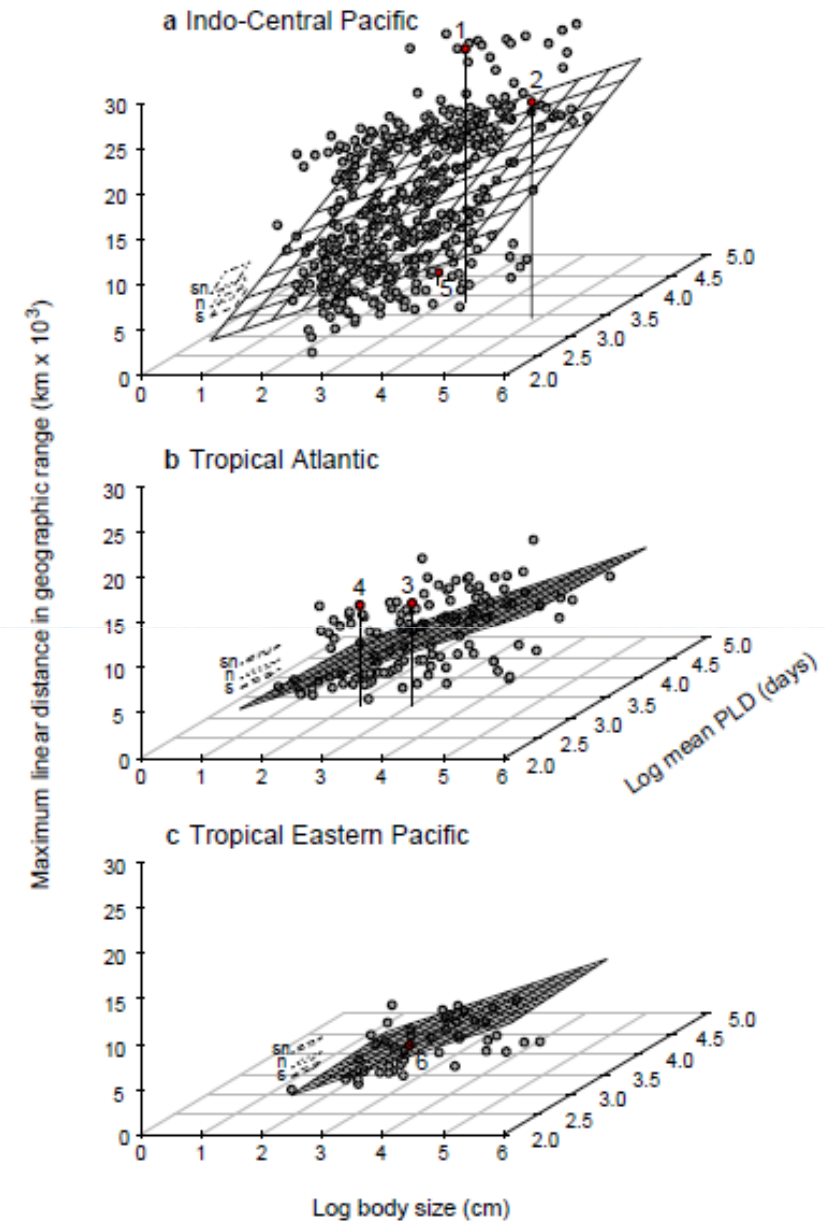
These capacities were known to be linked to

SIZE

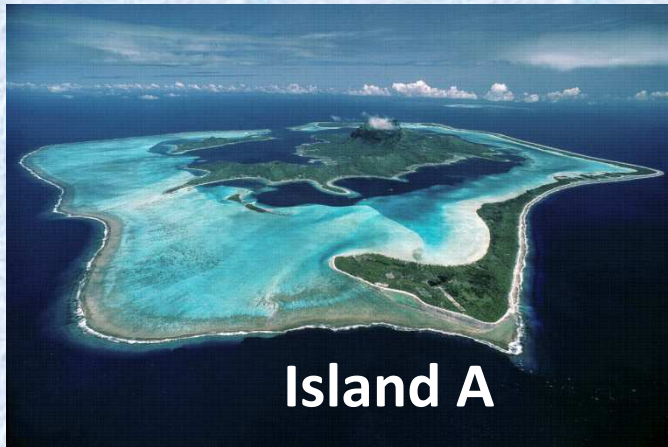
PLD

GEOG RANGE (AOO)

We analyzed the relationships between these parameters taking biogeographical realms into account



Two major strategy types for Dispersion + Colonisation



Strategy 1

Massive but rare egg batches →

Long PLD

Adults: Habitat Generalists ○

Adults : Diet Generalists

Species Large– Solitary – Above bottom

Strategy adapted to colonising
isolated islands

Strategy 2

Small but frequent egg batches →

Short PLD

Adults : Habitat Specialists ●

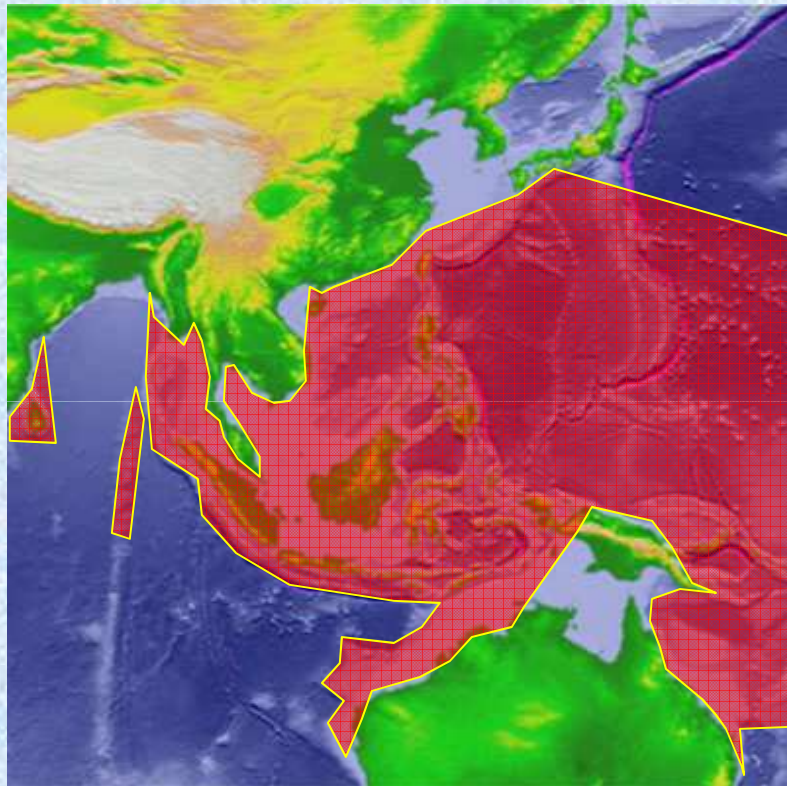
Adults : Diet Specialists

Species Small – Gregarious – Bottom dwelling

Strategy adapted to colonising
in a stepping stone way

The role of evolutionary history

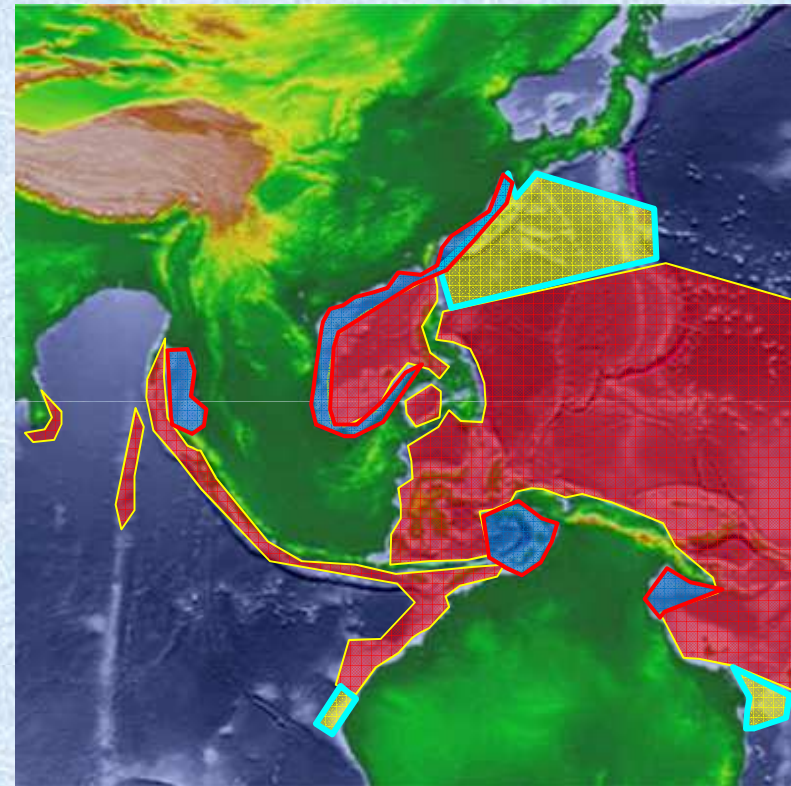
During ice-age periods coral reefs « shrink », sea level goes down



Today



conditions favorable to coral reefs



Lowest Sea Level during the quaternary



too cold



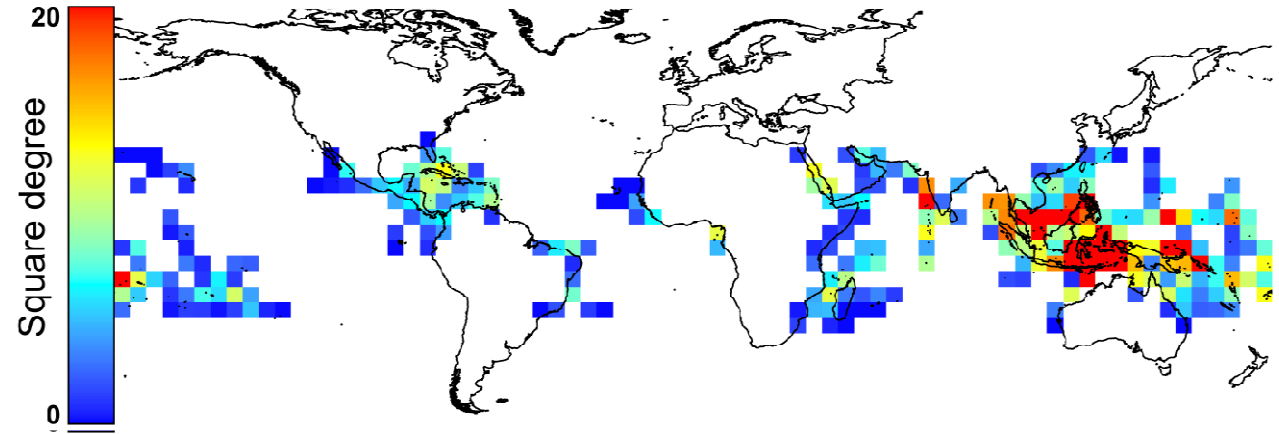
unfavorable salinity

Frequency of refugia
(there were many
changes during the last 3
MY – 1 value each 1000
years)

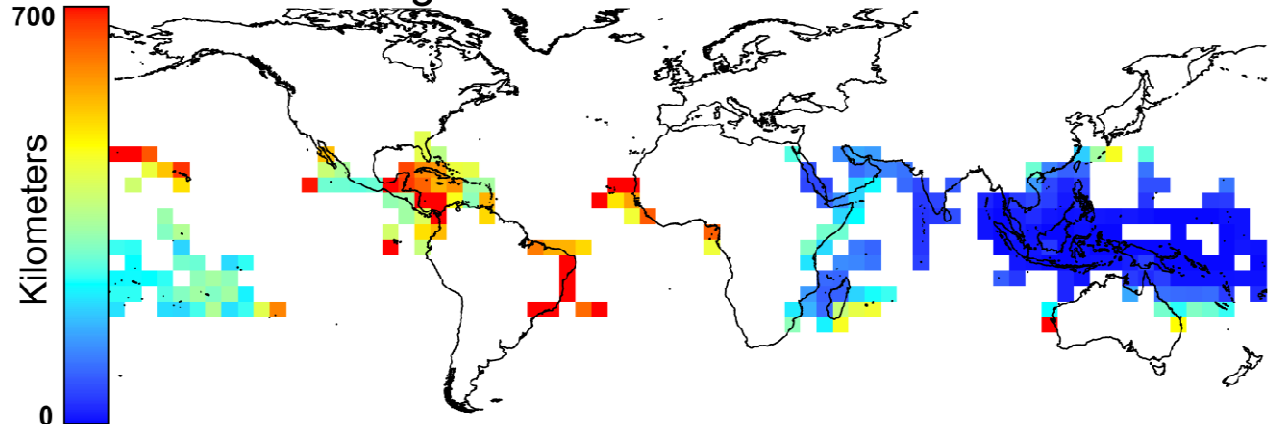
Distance of each cell to
the nearest refugia.
Average over 3 MY

Today's distribution of
reef fish

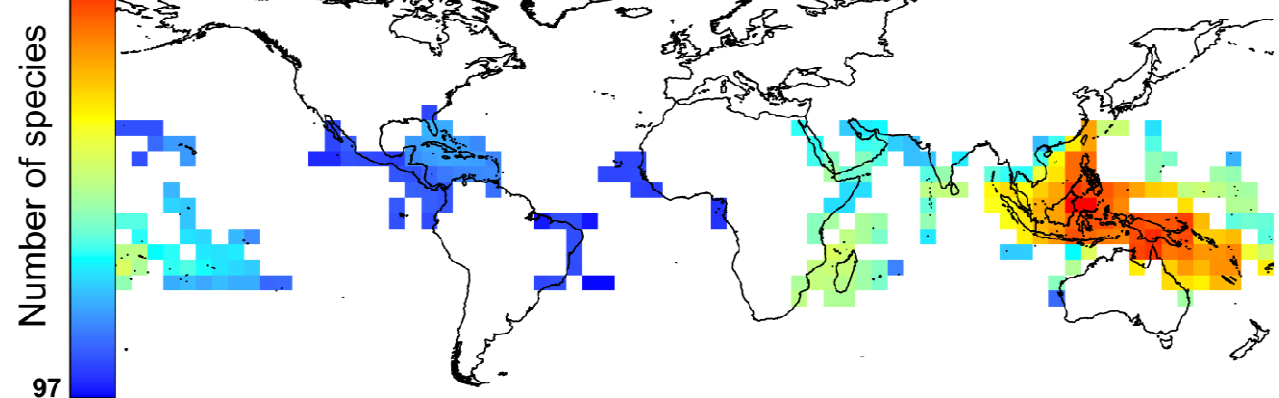
A Past coral reef area



B Isolation from refugia



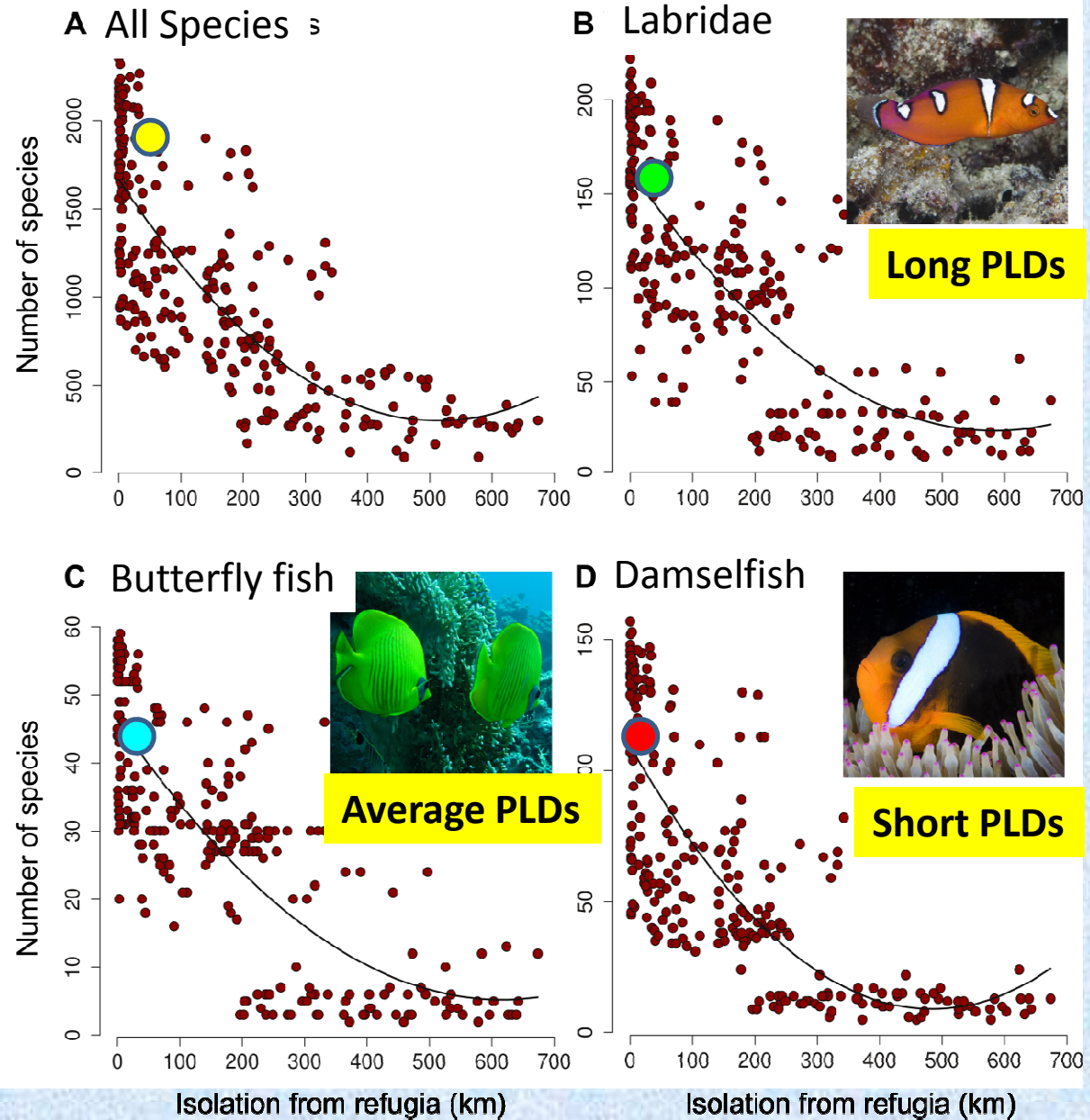
C



Species richness decreases with distance to refugia. This distance explains ~60% of the spatial distribution of today's reef fish



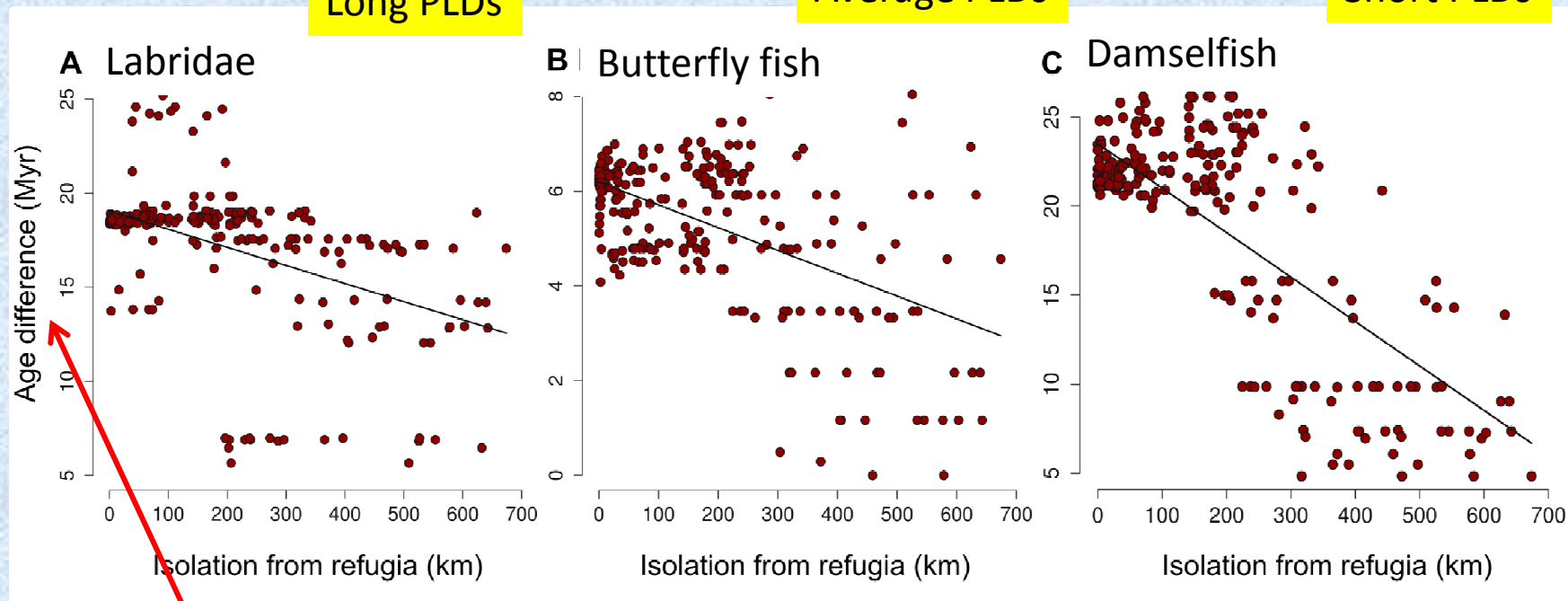
Colonizing capacity is essential to explain the present day diversity distribution



Long PLDs

Average PLDs

Short PLDs



Measure of the dispersion of species ages (difference between the 95th and the 5th percentile for each site) **within each geographical cell**

The further away from refugia and more ages tend to be similar

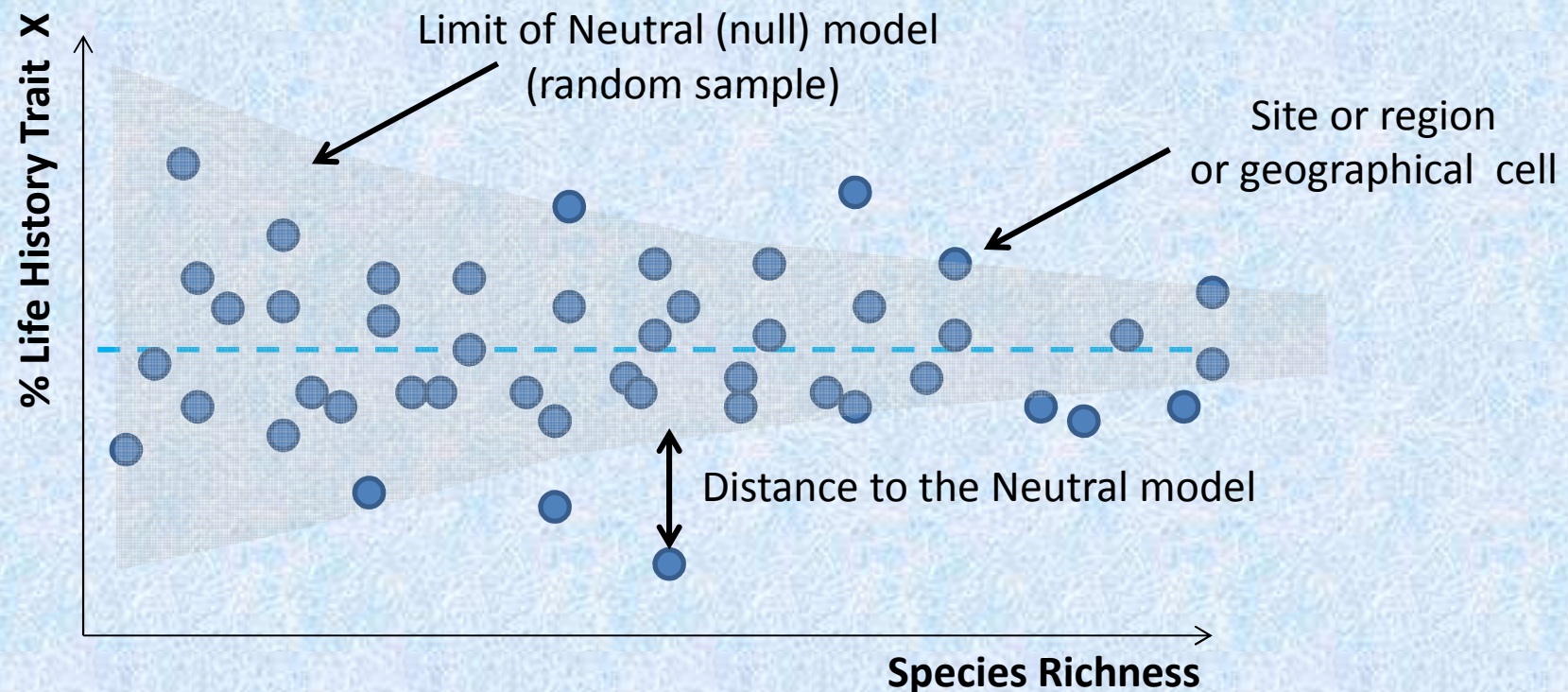
The slope is proportional to average PLD

From Species to Traits

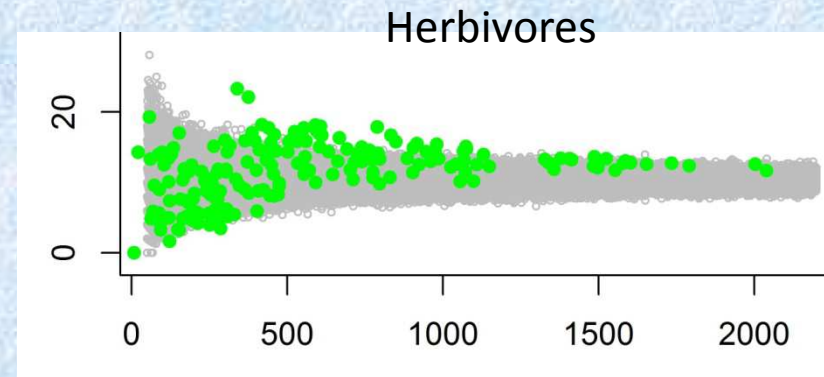
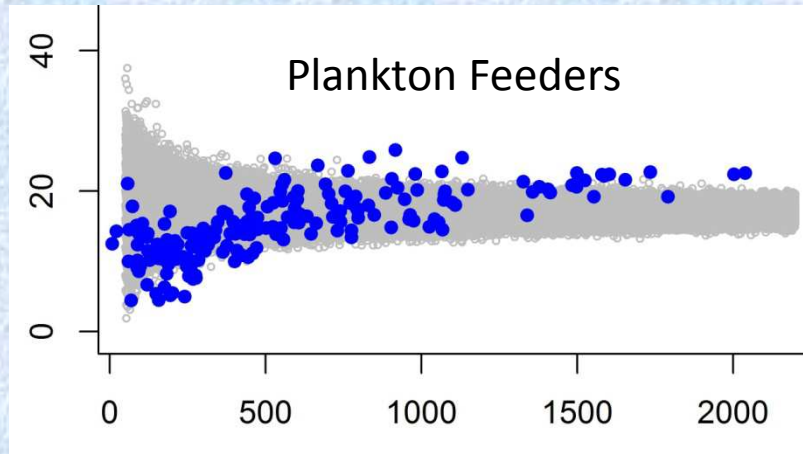
Hypothesis: all reef fish assemblages function along the same principles and therefore should have the same structure on any reef in the world



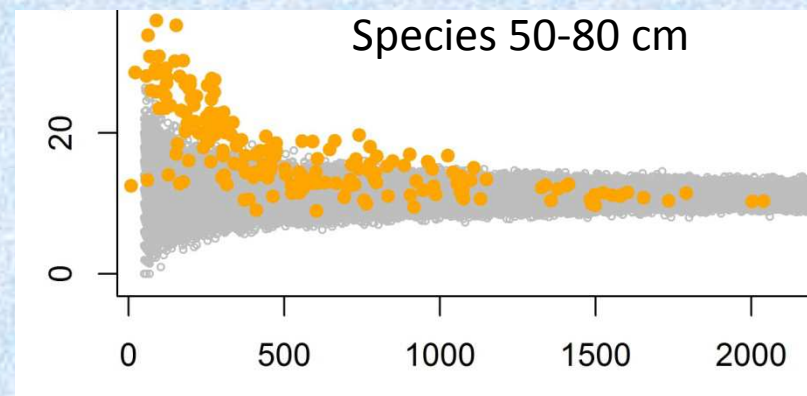
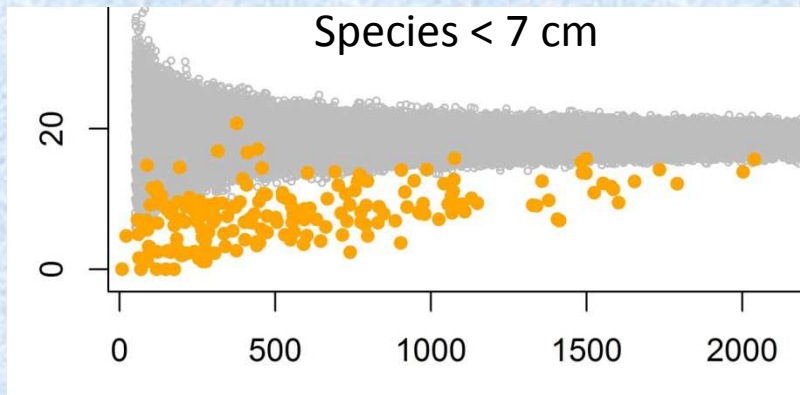
The proportion (% of all local species) of a group – e.g. herbivores – is the same everywhere



Some example from the Indo Pacific

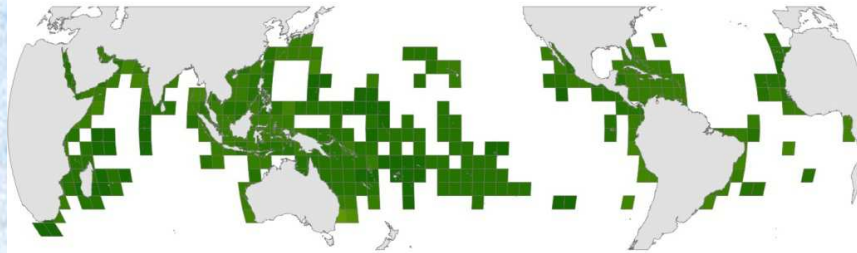


Number of Species / Site



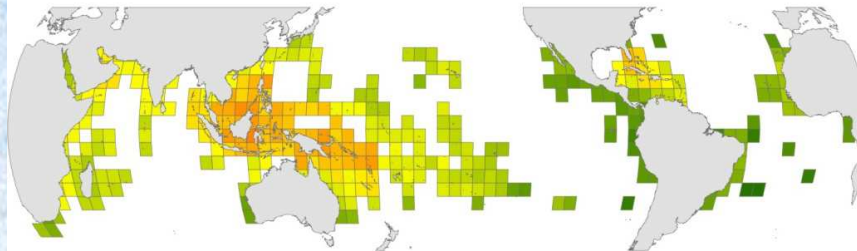
Number of Species / Site

Family based



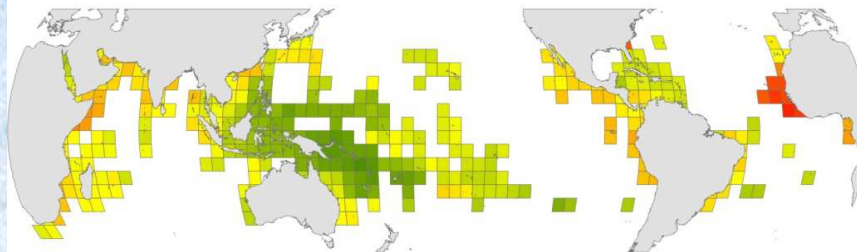
The global deviation (sum of all deviation from Neutral) was constant for families $\rightarrow$ very stable % of each family within a realm

Diet based



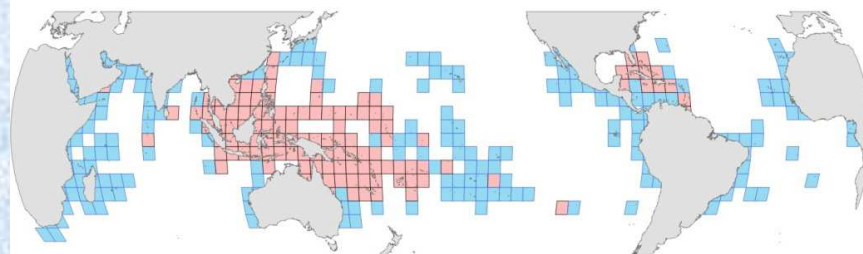
Trophic structure is the furthest from the Neutral model in regions with high diversity

Size based



Size Structure is the furthest from the Neutral model in isolated and peripheral regions

Dominant deviation

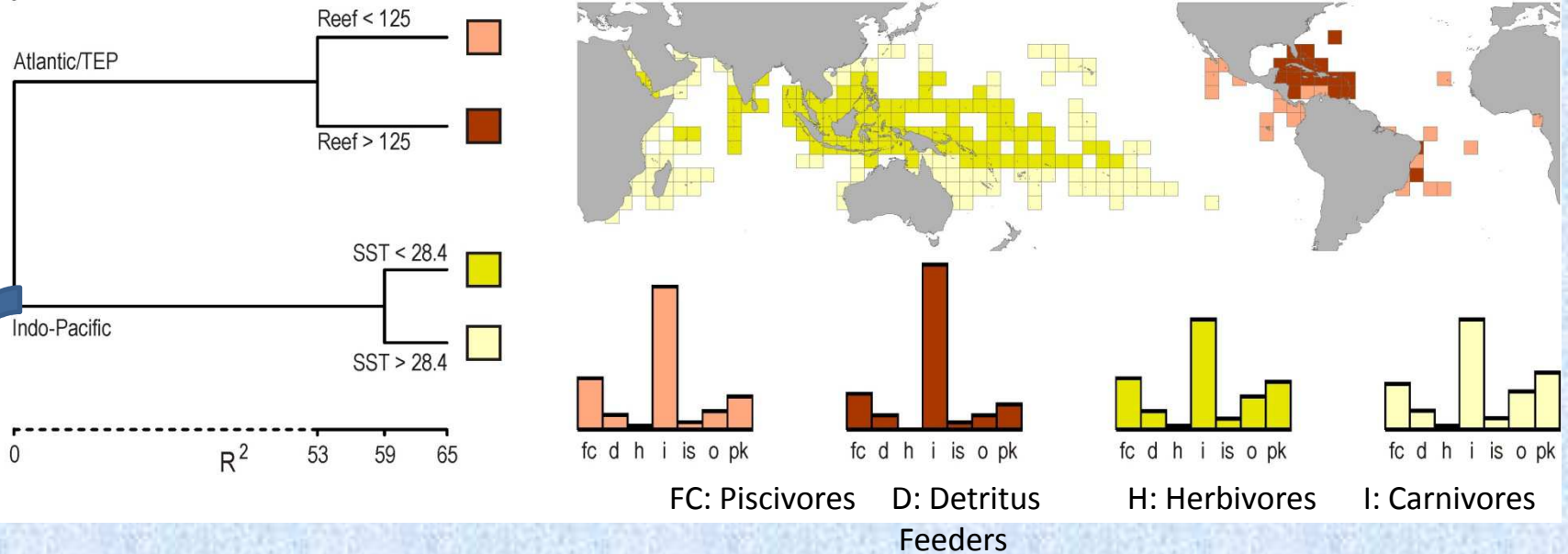


Size-dominated deviation

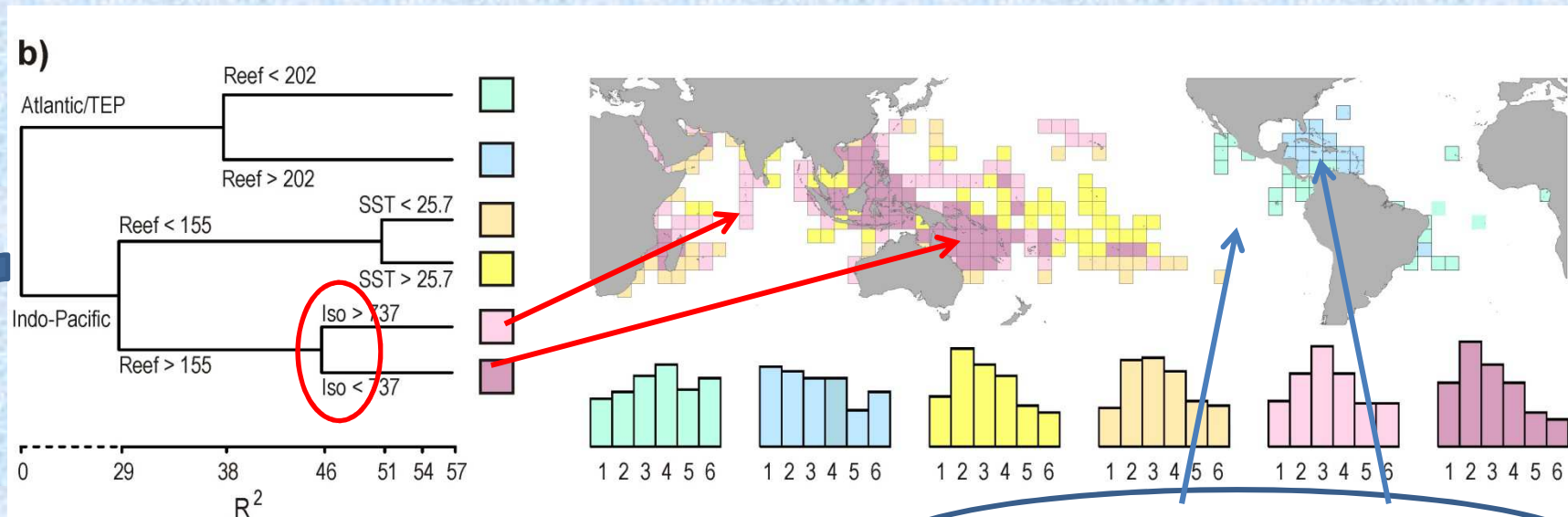
Diet-dominated deviation

Trophic and Size Structures have geographically distinct distributions

a)



No significant effect of island size or isolation, therefore of physical connectivity, on trophic structures



1: < 7cm ; 2: 8-15 cm; 3: 15-30 cm, 4: 30-50 cm; 5: 50-80 cm; 6: > 80 cm

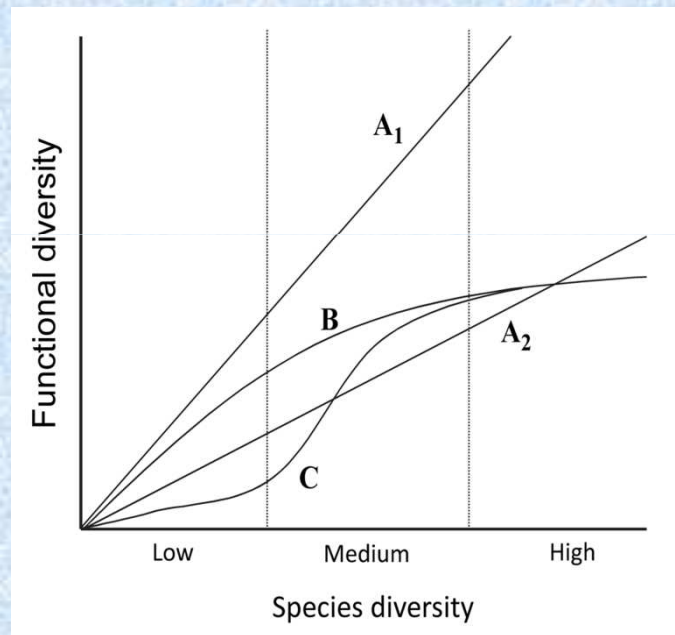
« Isolation » : combinaison of distance and size of the nearest islands, therefore a proxy of physical connectivity

Physical connectivity plays a significant role in the size structure of reef fish assemblages in the Indo-Pacific

From Species to Function

Why are we interested in functions ?

Do functions have the same distribution as species ?



A1: 1 species $\leftrightarrow$ 1 function

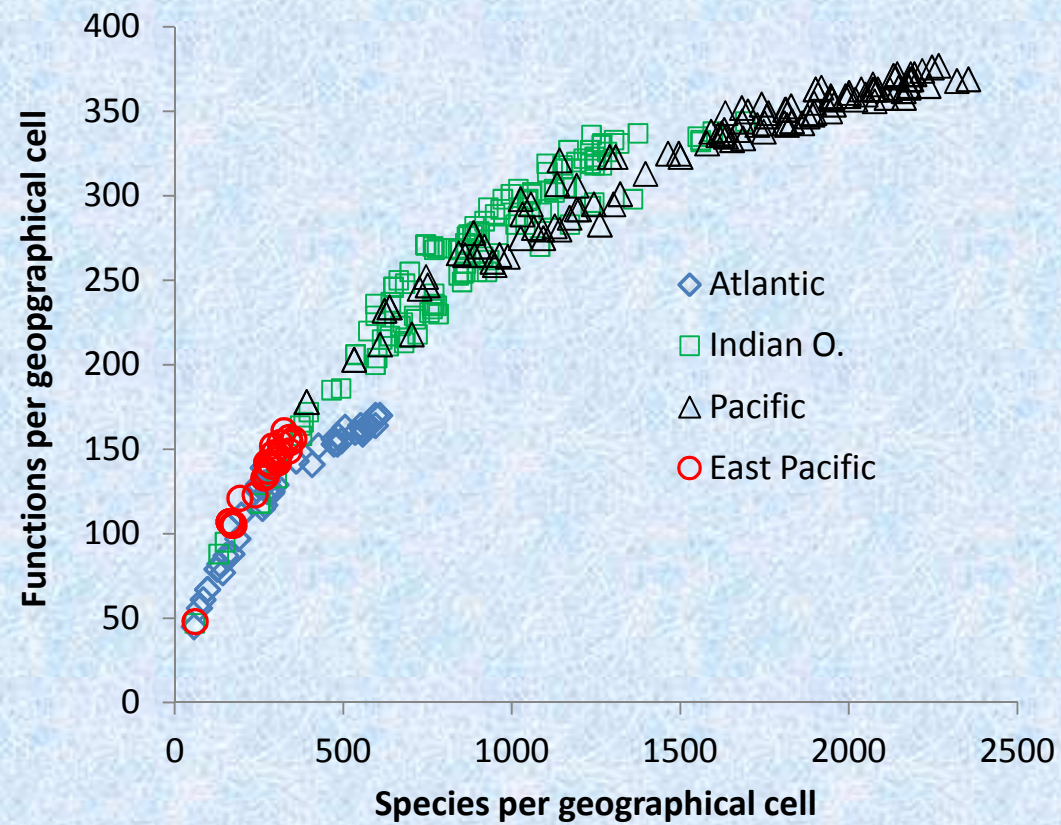
B: increasing redundancy as species richness increases

C: ecological threshold beyond which functions are more diversified

from Michaeli & Halpern 2005

Mouillot et al. 2014 Functional over-redundancy and high functional vulnerability in global fish faunas of tropical reefs. PNAS

Kulbicki et al. in press Patterns and processes linked to body size. C.Mora ed. "Ecology of fishes on coral reefs". Oxford Univ. Press

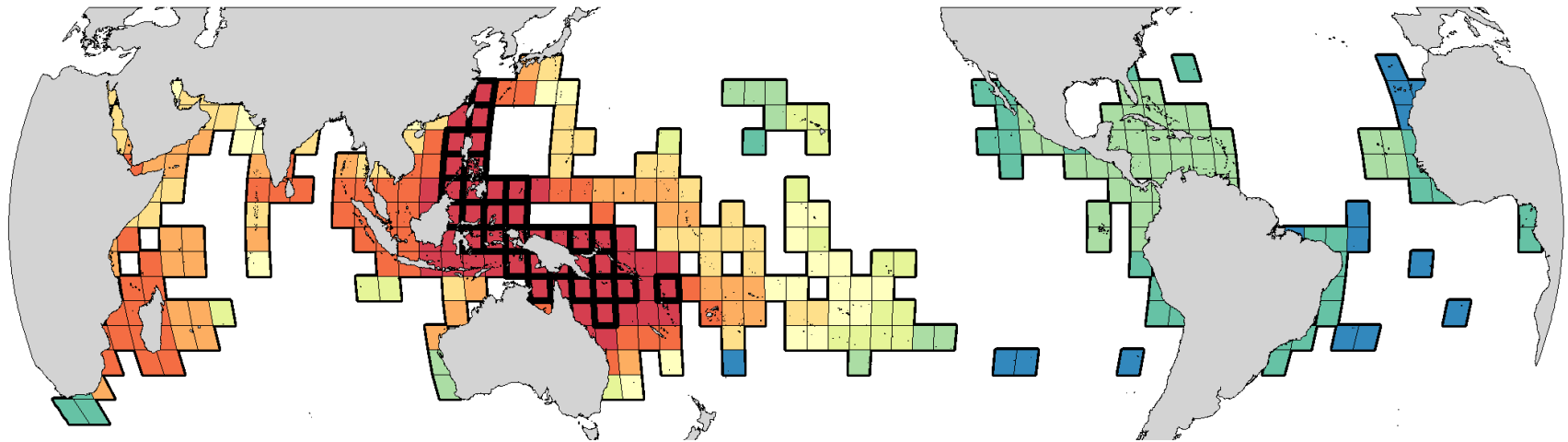


Relation B from previous slide, at large scale, for all realms

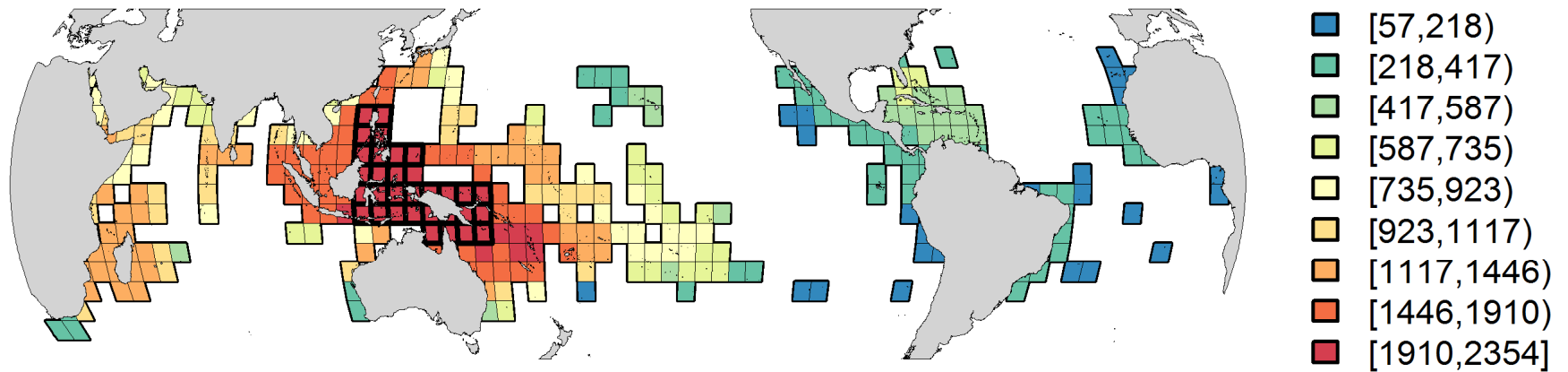
Same is true at a local scale

Guillemot et al. (2011) Functional Redundancy Patterns Reveal Non-Random Assembly Rules in a Species-Rich Marine Assemblage. PLoS ONE 6(10): e26735.

Functional richness



Total richness



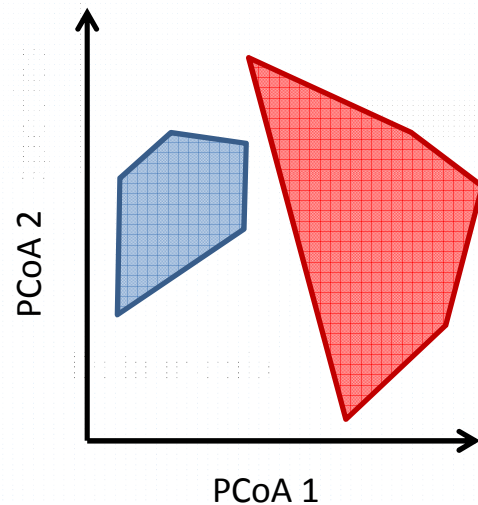
No major mismatch

Functional Volumes

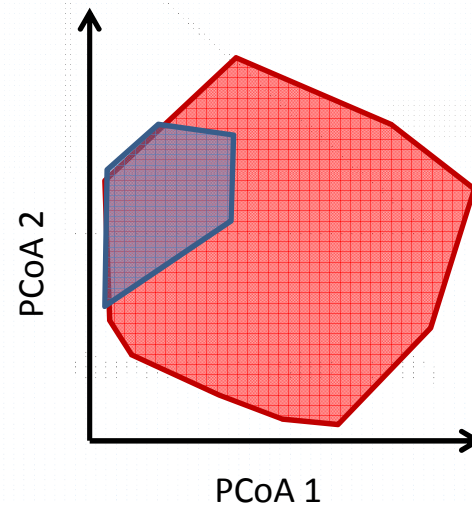
with N Life-History traits → multispace volume
e.g. an assemblage with H, Pis & Carn. has a larger
FV than an assemblage with only H

Hypotheses

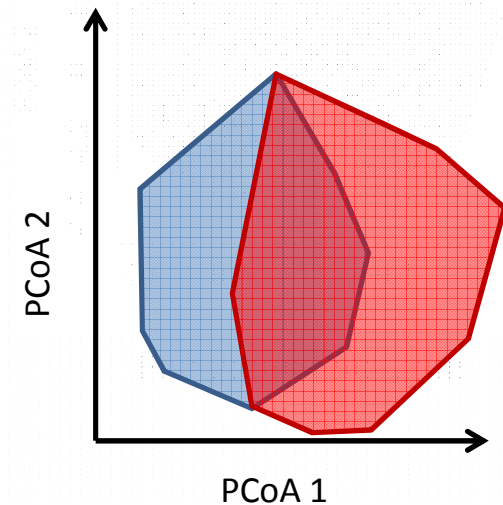
-  Poor region (e.g. Eastern Atlantic)
-  Rich region (e.g. Central Indo-Pacific)



Different Functional richness
Different Functional structure

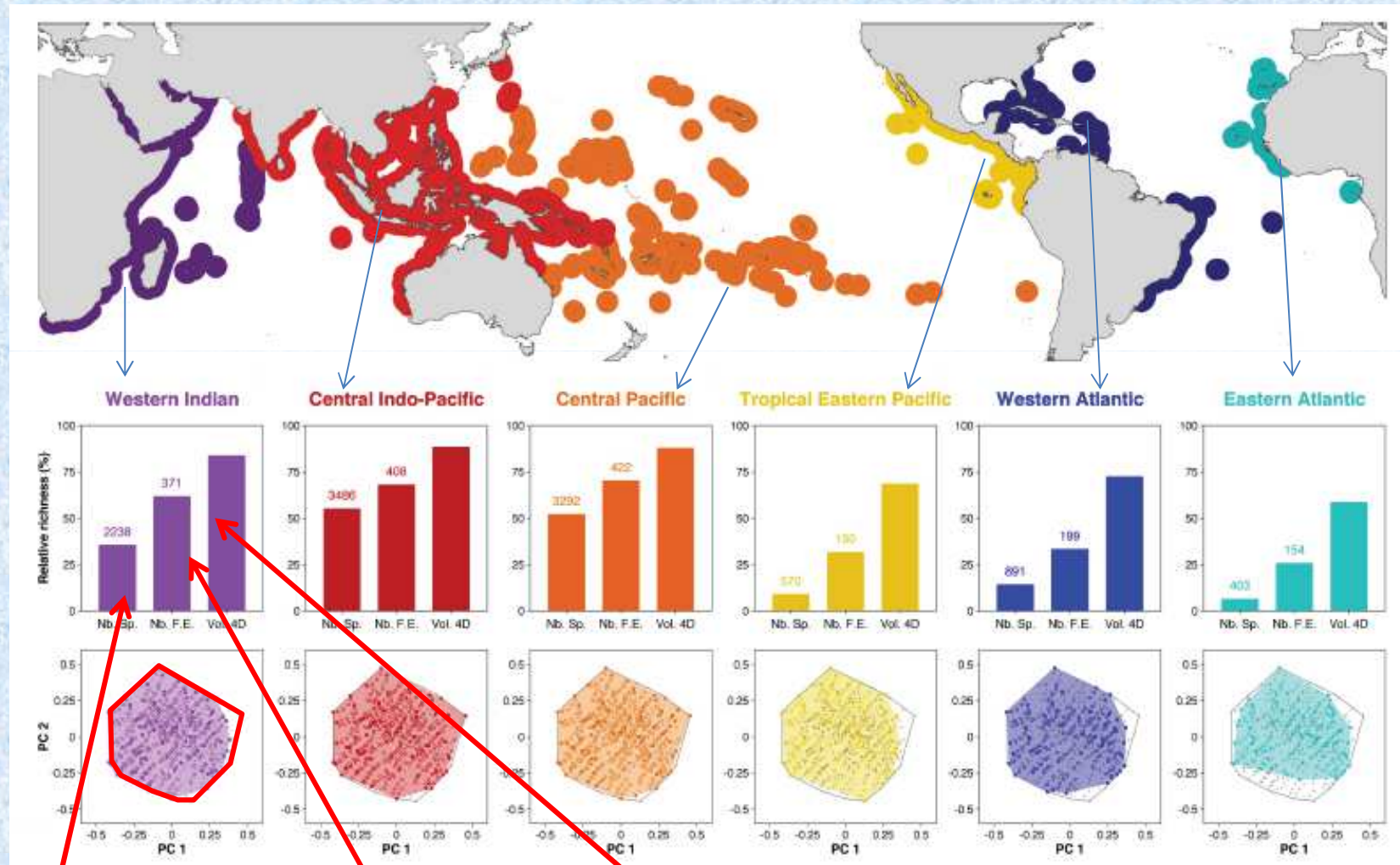


Different Functional richness
Nested Functional structure



Similar Functional richness
Overlapping Functional structure

Despite large differences in species richness, all biogeographical regions share the same functional space



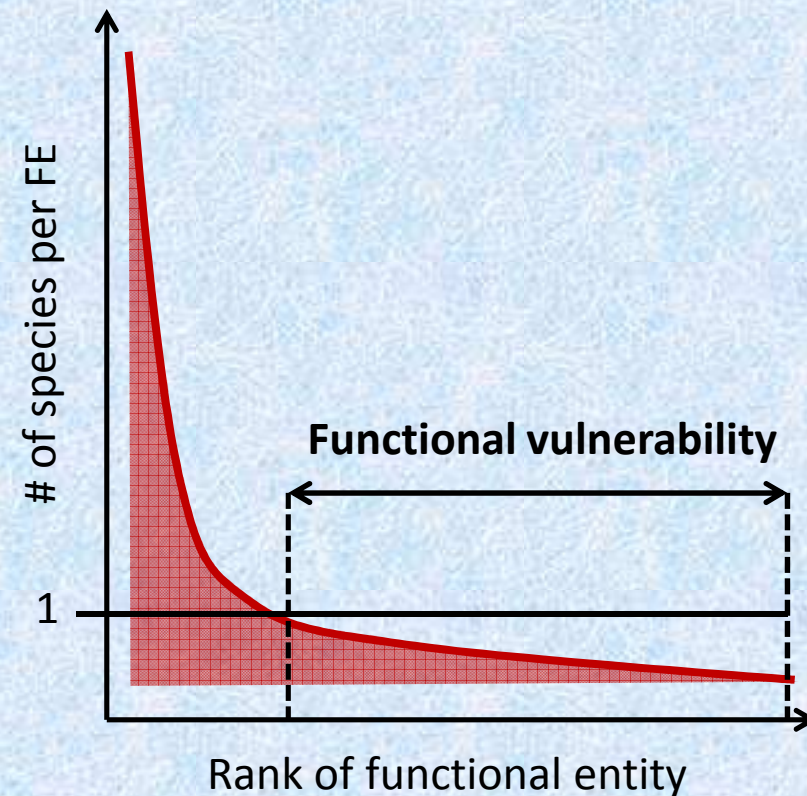
% Nb. Species

% FG

% FV

Vulnerability : Species versus Functions

Species richness packing within functional entities reveals high functional vulnerability of reef fish assemblages



Functional vulnerability: The percentage of functional entities with no redundancy, i.e. composed by only one species

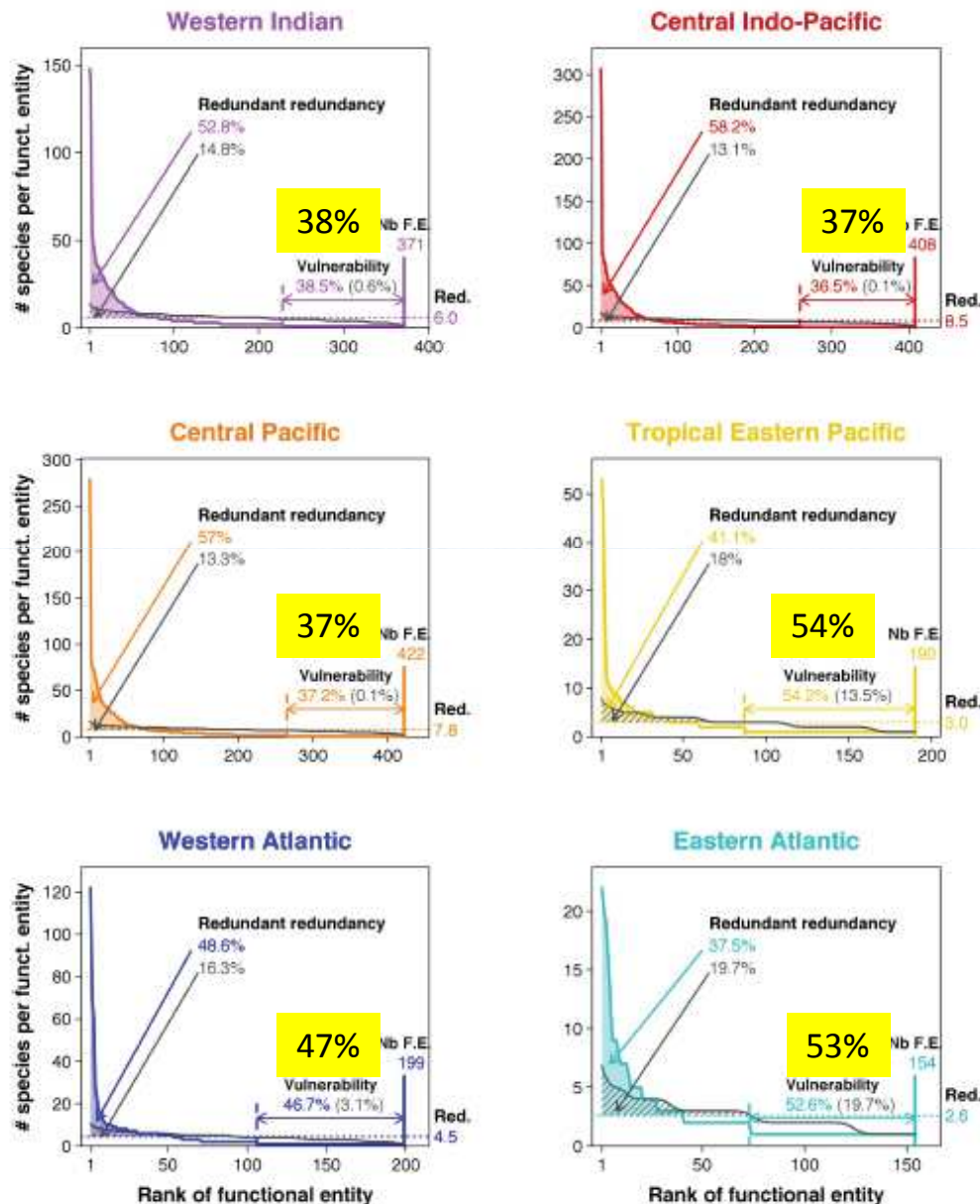
Species richness packing within functional entities reveal high functional vulnerability of reef fish assemblages

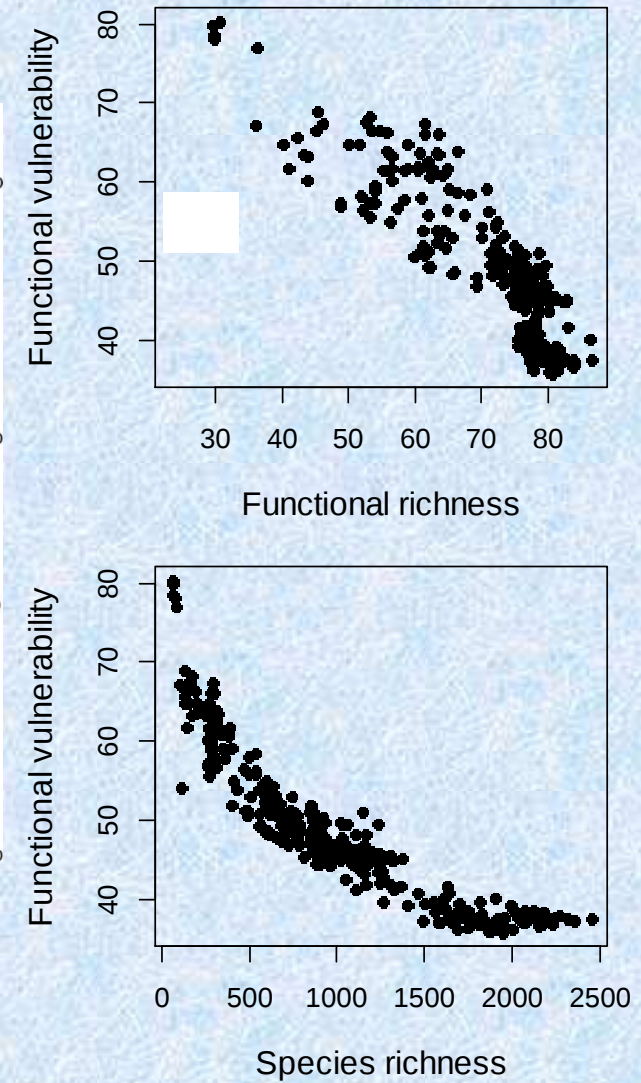
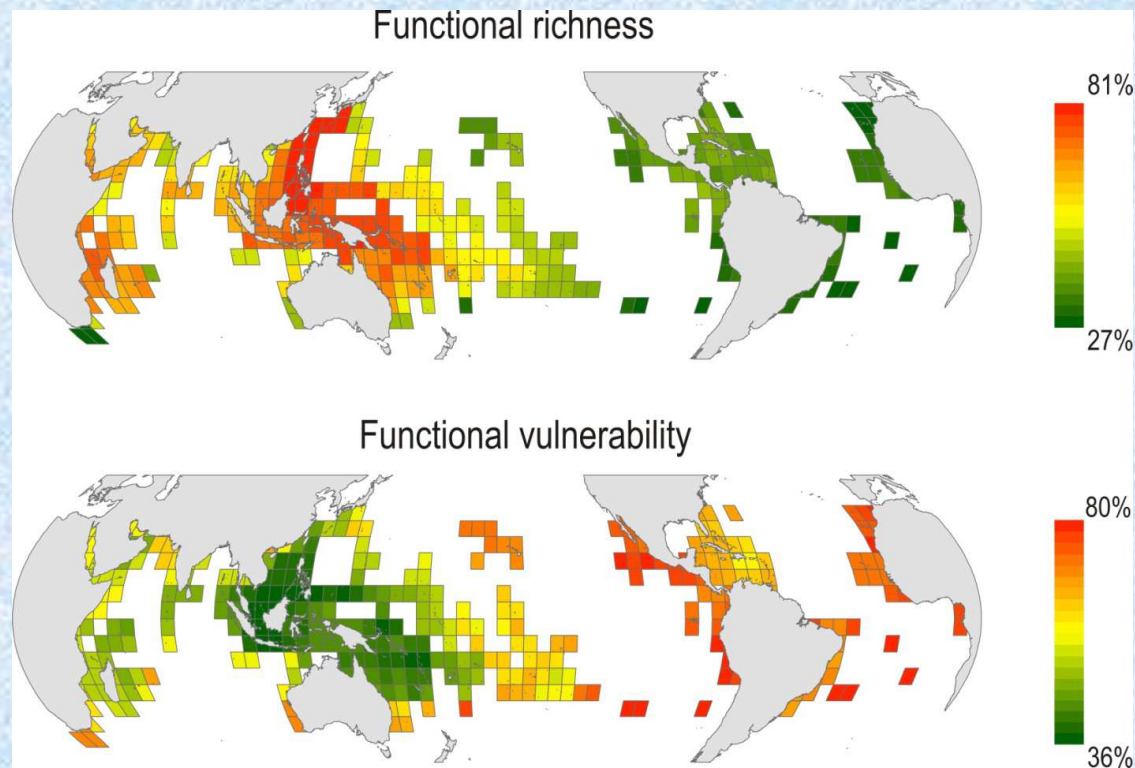
Even the richest regions shows an extremely high functional vulnerability (at least 36%)

The local extinction of those species may have dramatic effect on the functional structure of assemblages

Functional attributes and functional vulnerability, in particular, are important criteria to be taken into account for conservation

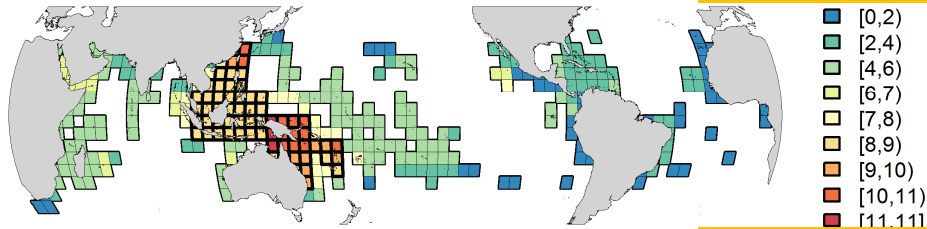
What is the functional vulnerability of local communities?





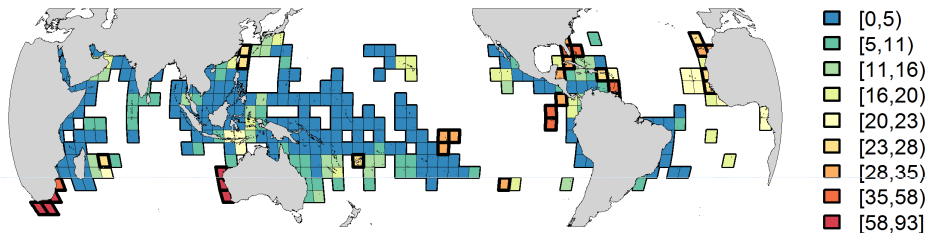
Matching Protection and Vulnerability on a global scale

IUCN richness



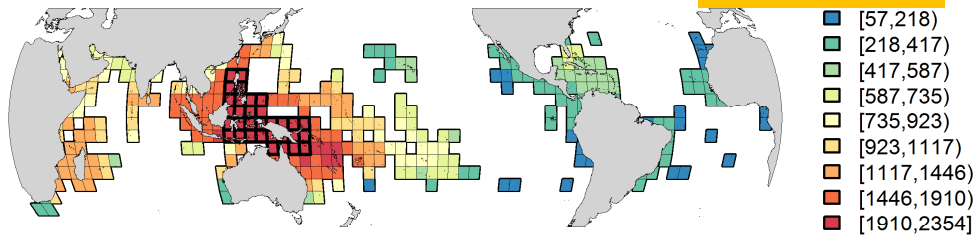
IUCN vulnerable Species

SR richness



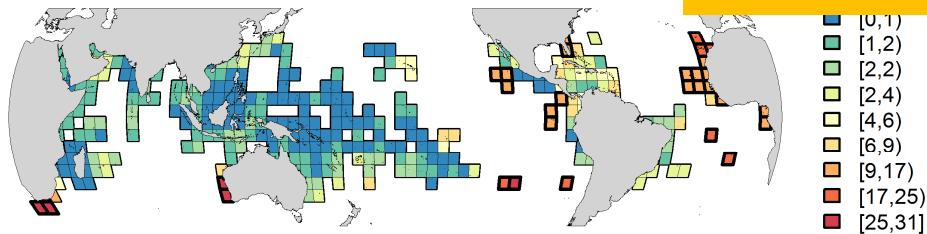
Endemic Species

Total richness



All Species

Taxonomic sensitivity



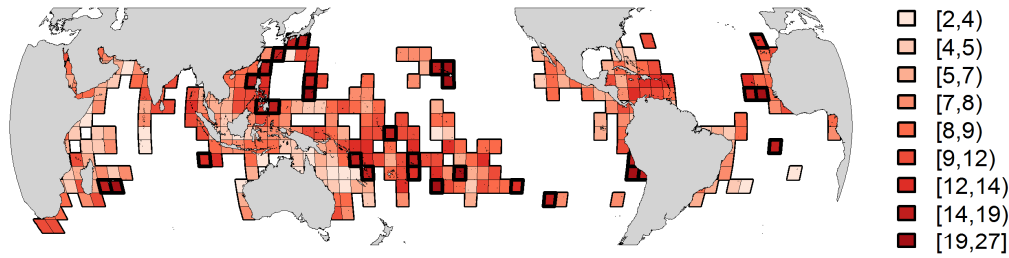
% IUCN & Endemic

VULNERABILITY

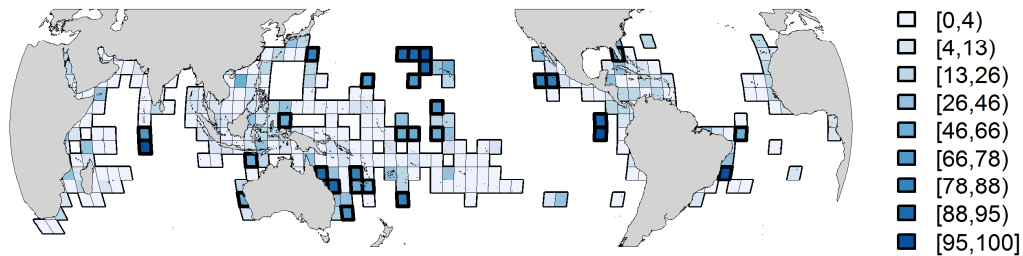
=

f (SENSITIVITY; THREAT)

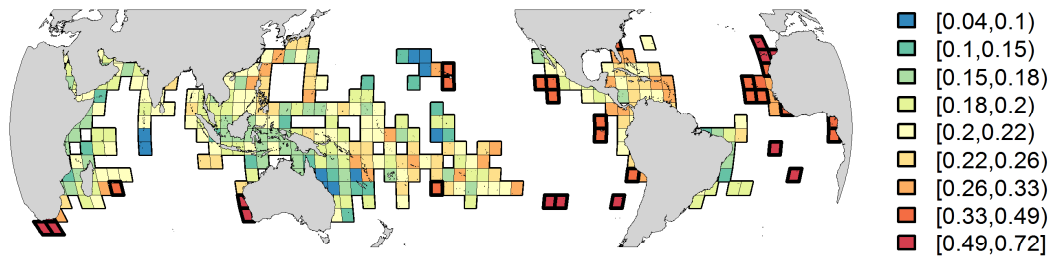
Exposure to human threats



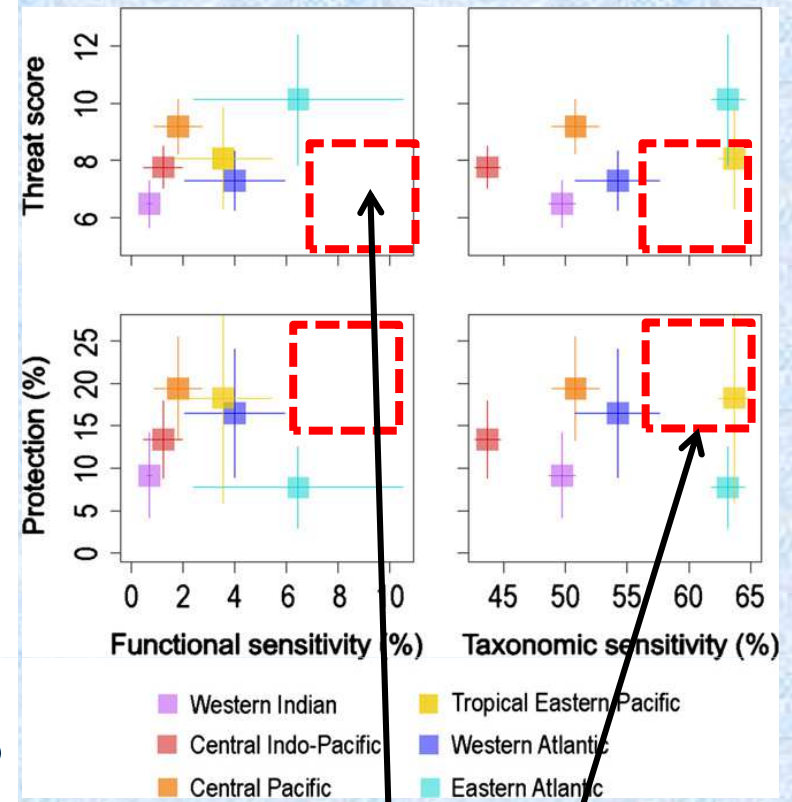
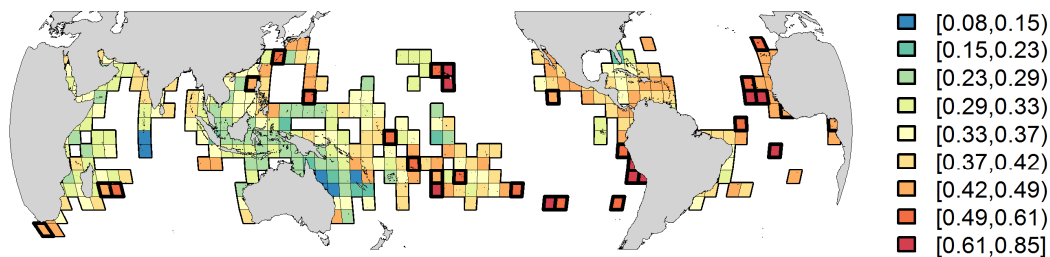
Protected habitat (%)



Taxonomic Vulnerability



Functional Vulnerability



Mismatch between protection
& vulnerability

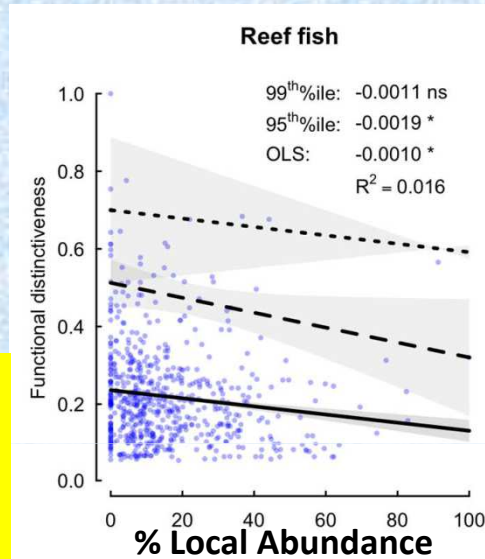
To be effective one
would assume:

Protection max & Threat min.
for the highest sensitivities

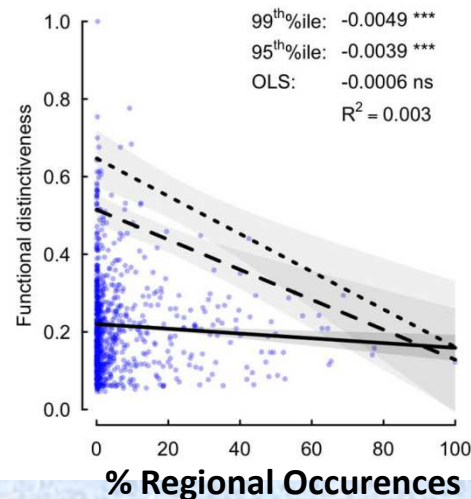
Some other sides of Vulnerability

So far we looked at presence/absence but vulnerability involves also abundance and connectivity

Distinctiveness

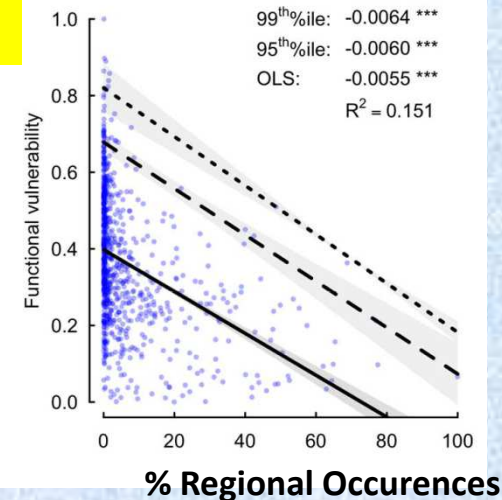
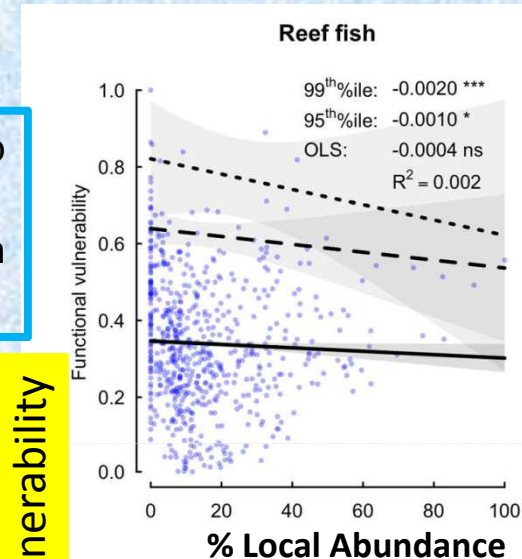


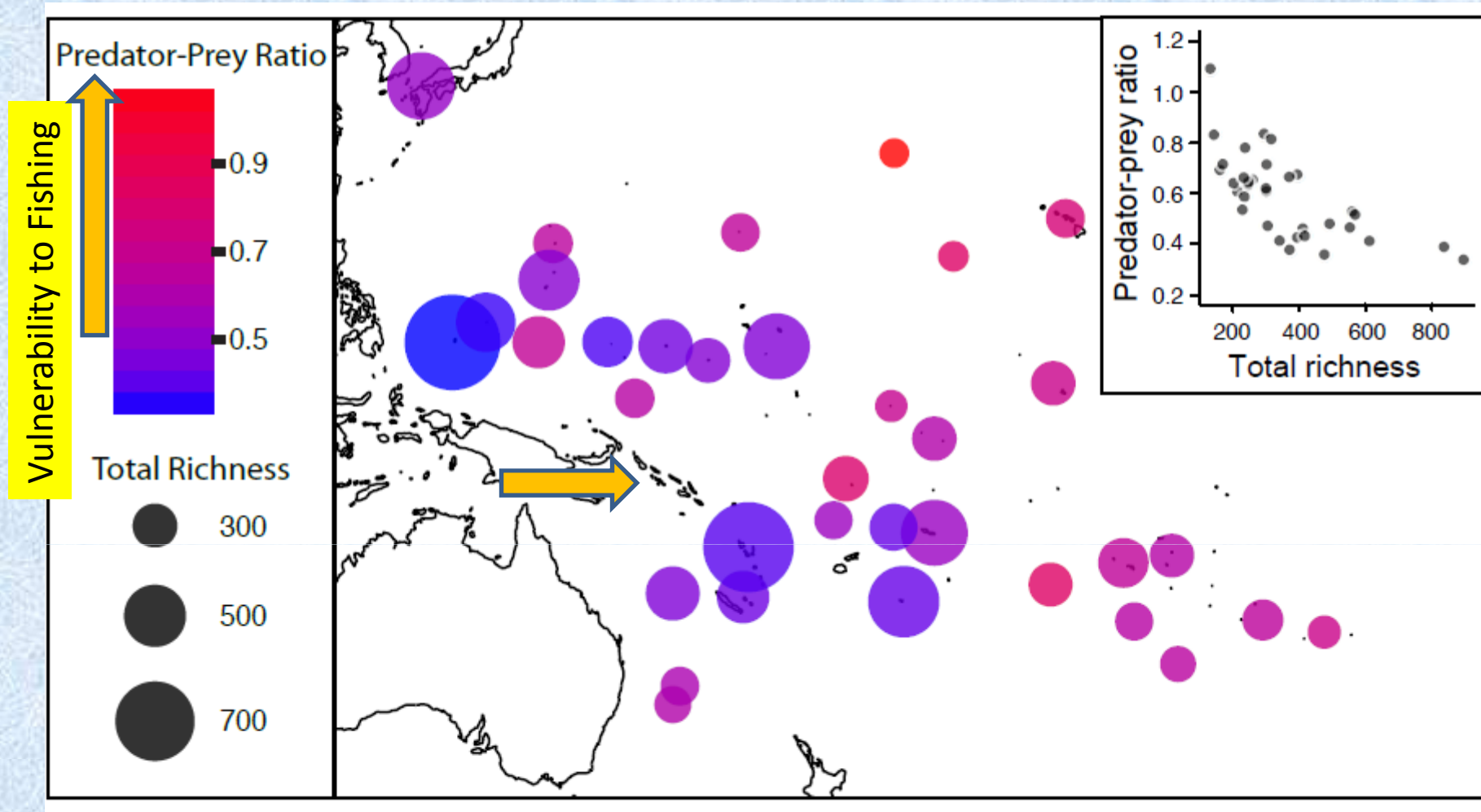
The rarest species tend to have specific functions more often than common species



The rarest species tend to belong to vulnerable functions more often than do common species

Vulnerability

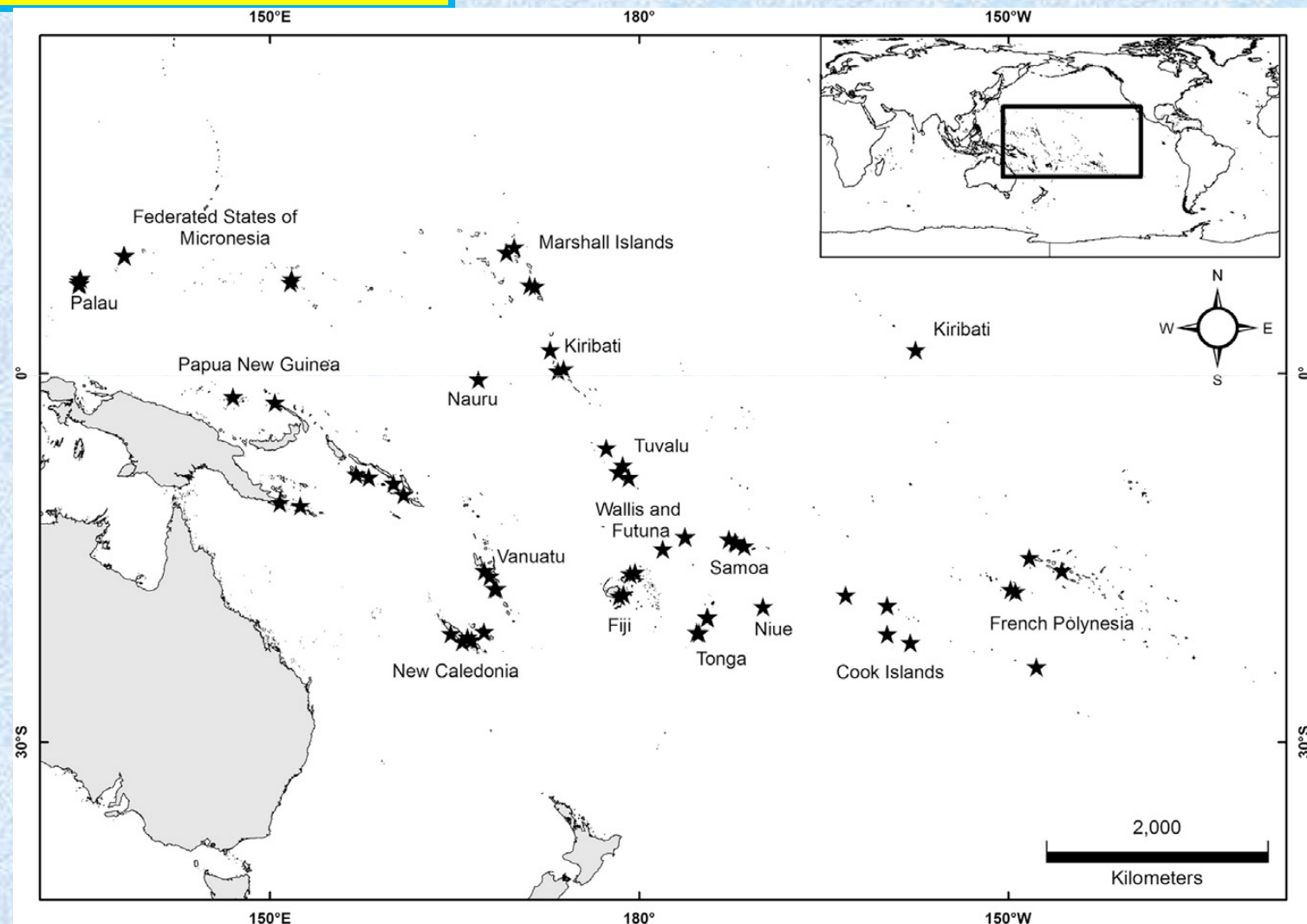




The proportion of predators increases as islands become smaller & isolated
Vulnerability to fishing increases similarly as predators are 1st targets

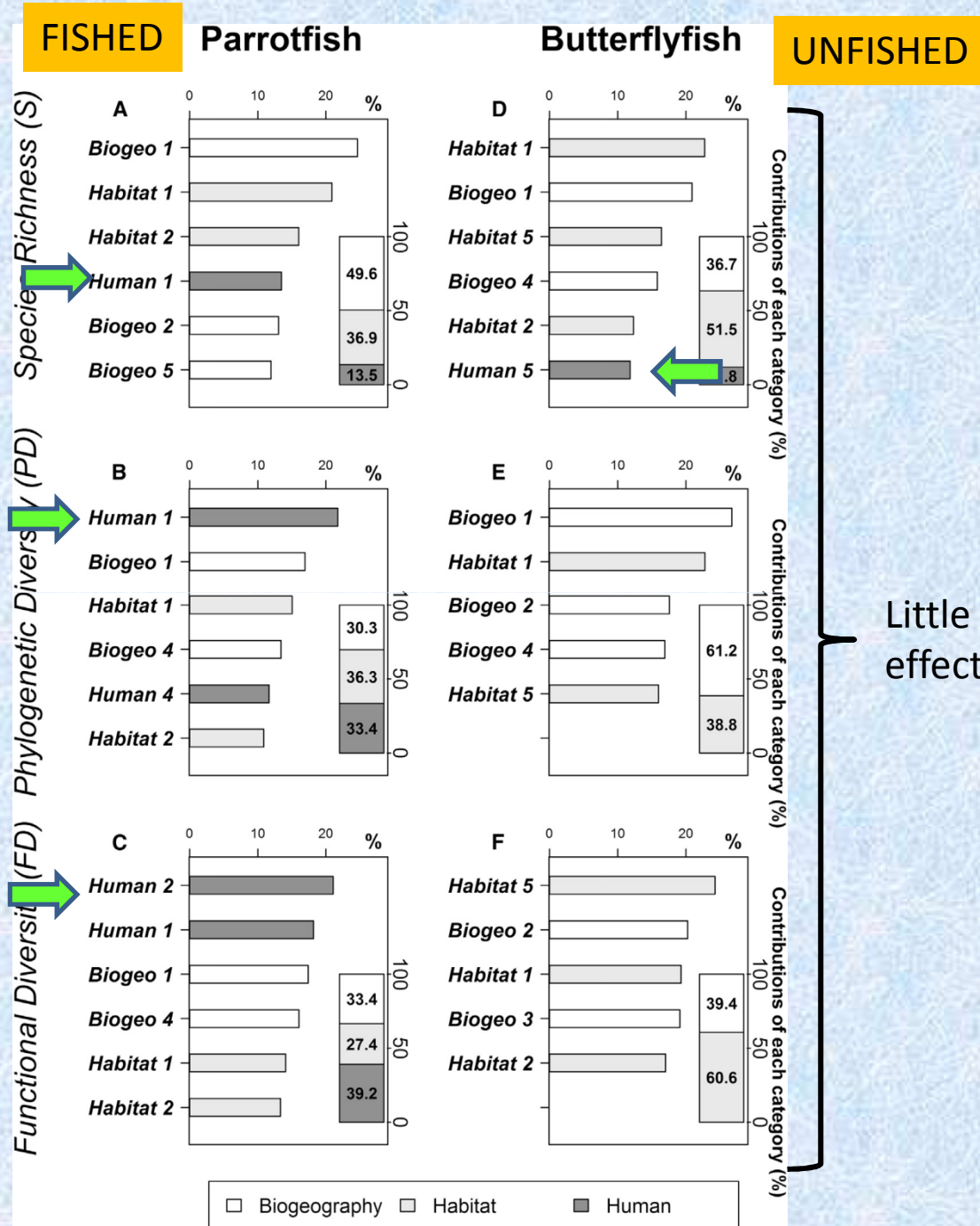
First analyses on the local effects of man

Are functions more robust than species to perturbations ?



Little effect of man on Species Richness

Man important for both Phylogenetic & Functional Diversities

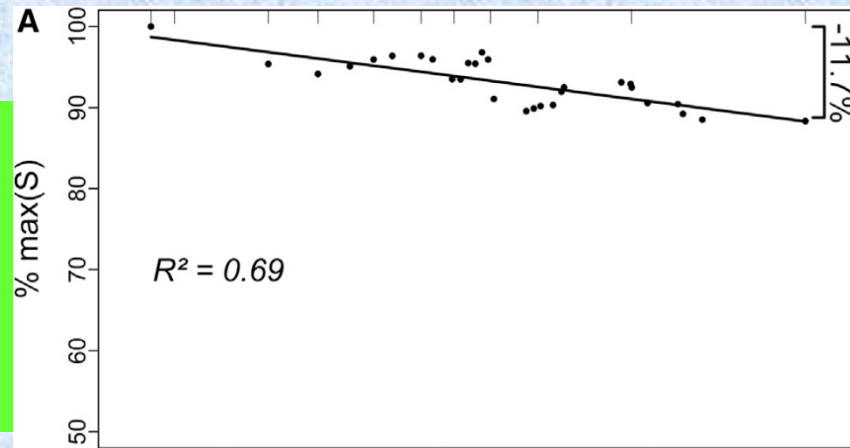


Little or no effect of Man

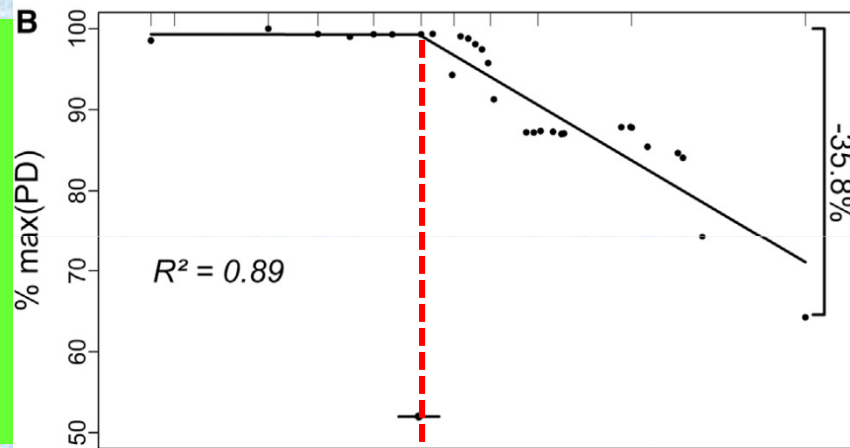
PARROT FISH

Thresholds found at rather low human densities (10-30 persons/km² of Reef)

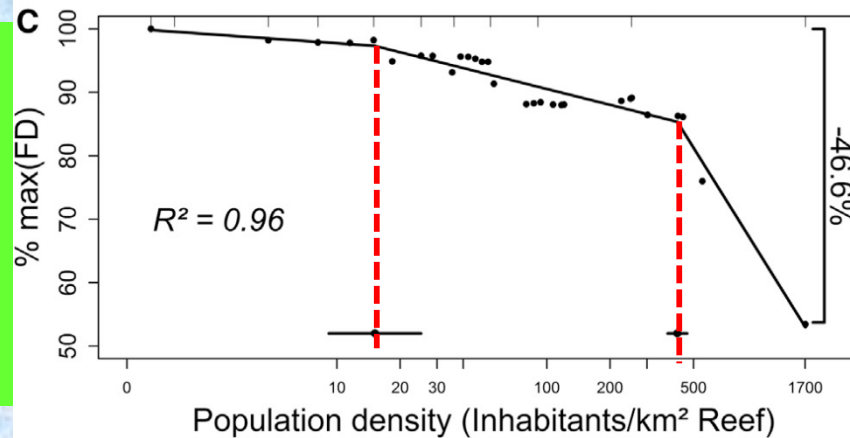
Species Diversity



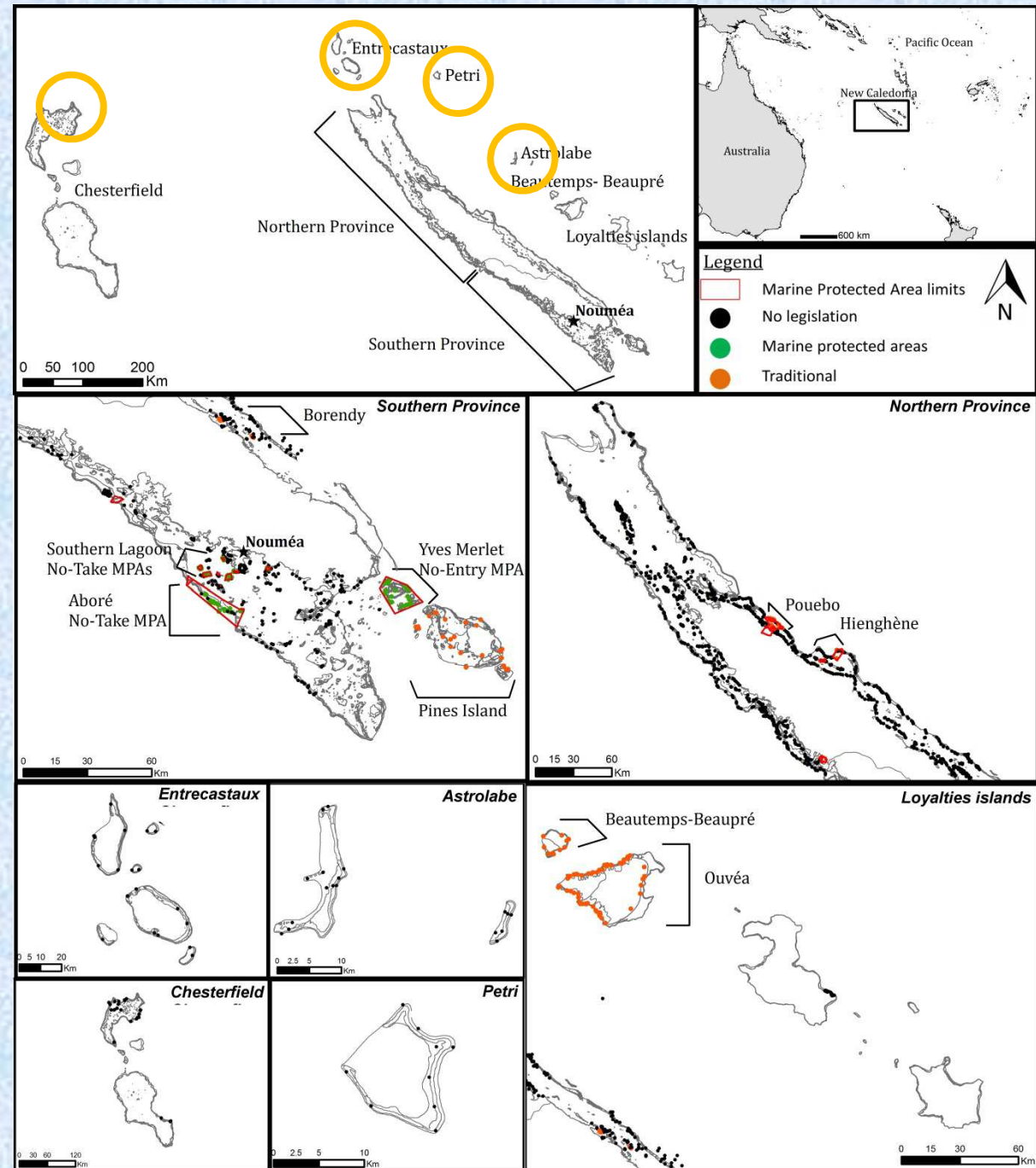
Phylogenetic Diversity

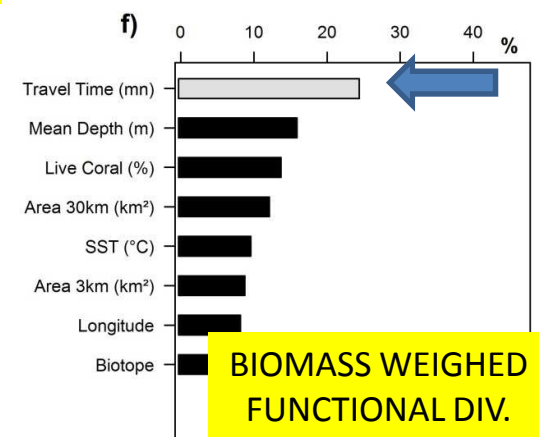
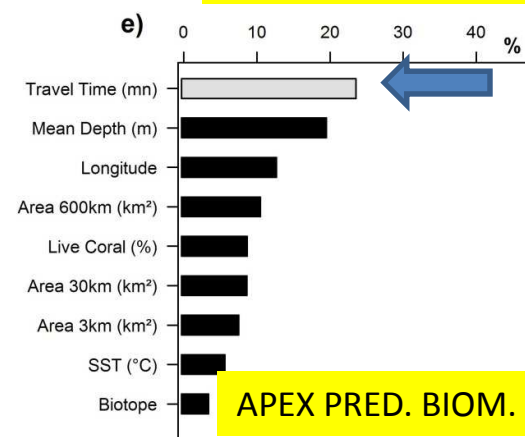
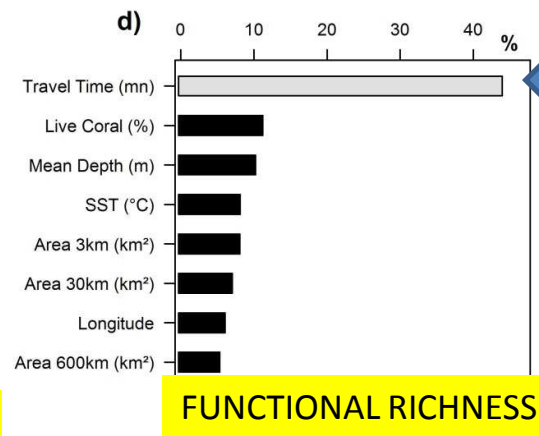
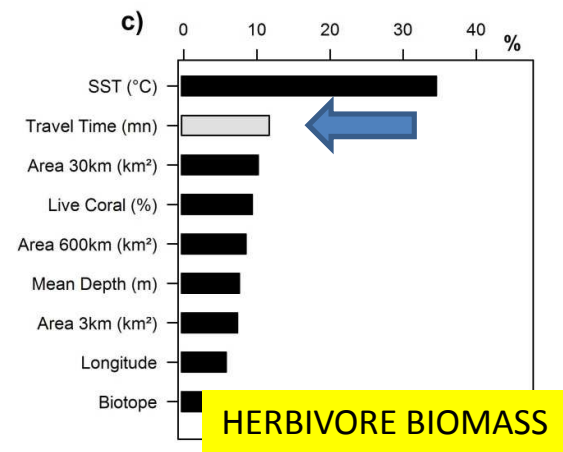
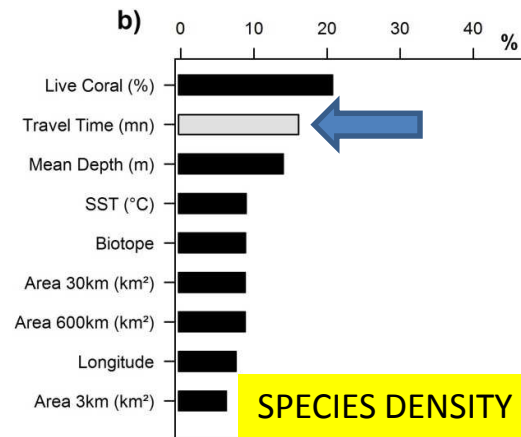
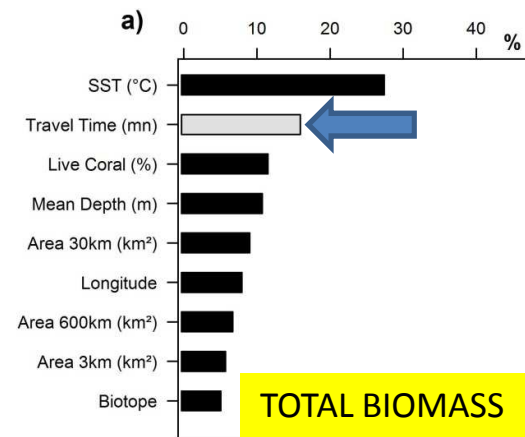


Functional Diversity

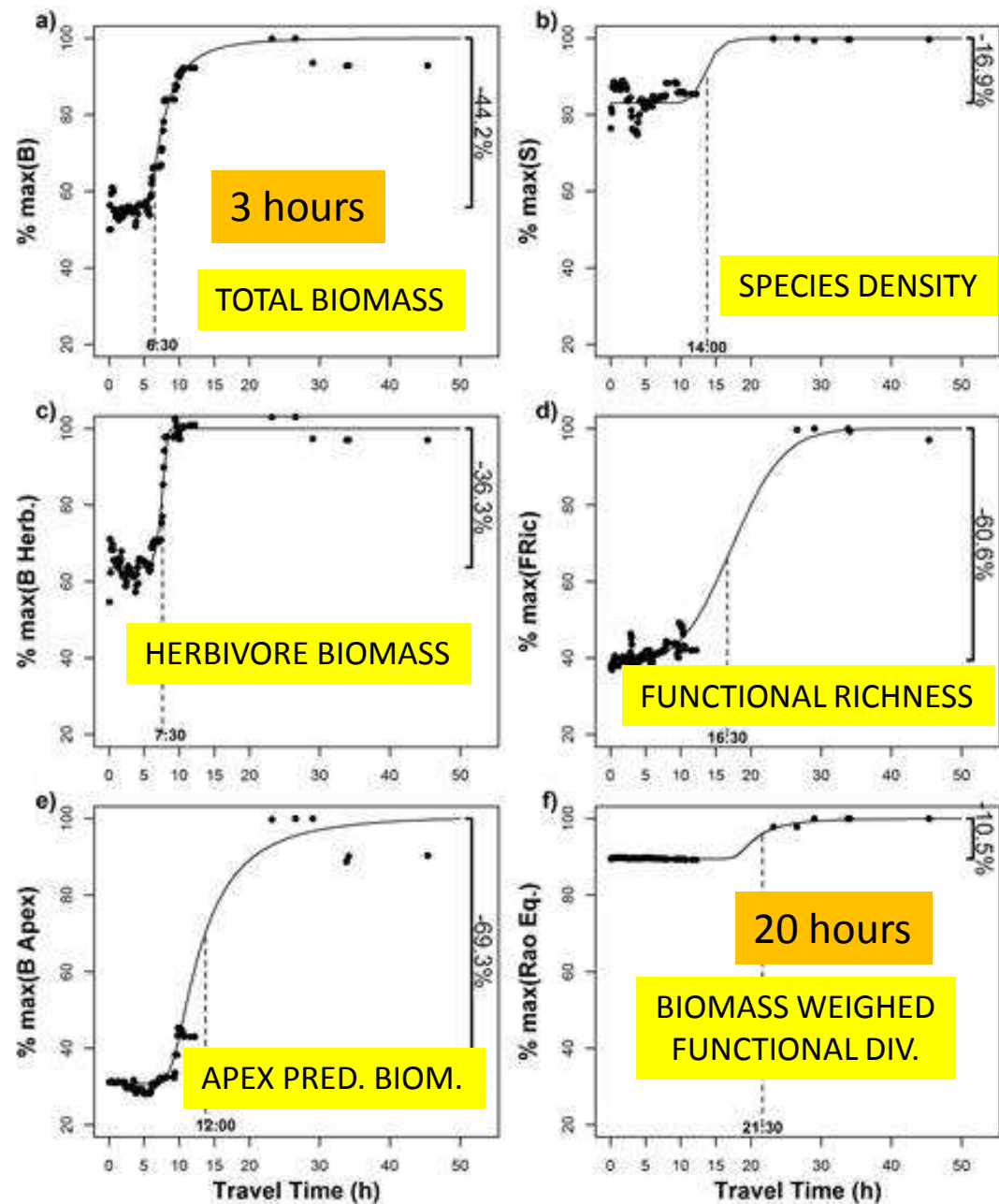


« PRISTINE » ZONES
do they represent
a baseline for MPAs ?





Travel time



Thresholds
for Travel Time



Beyond a limit there
are sudden and very
significant changes in the
characteristics of the
fish assemblages

Performances of various MPAs versus Pristine Areas



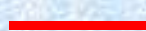
Pristine areas



Exploited areas



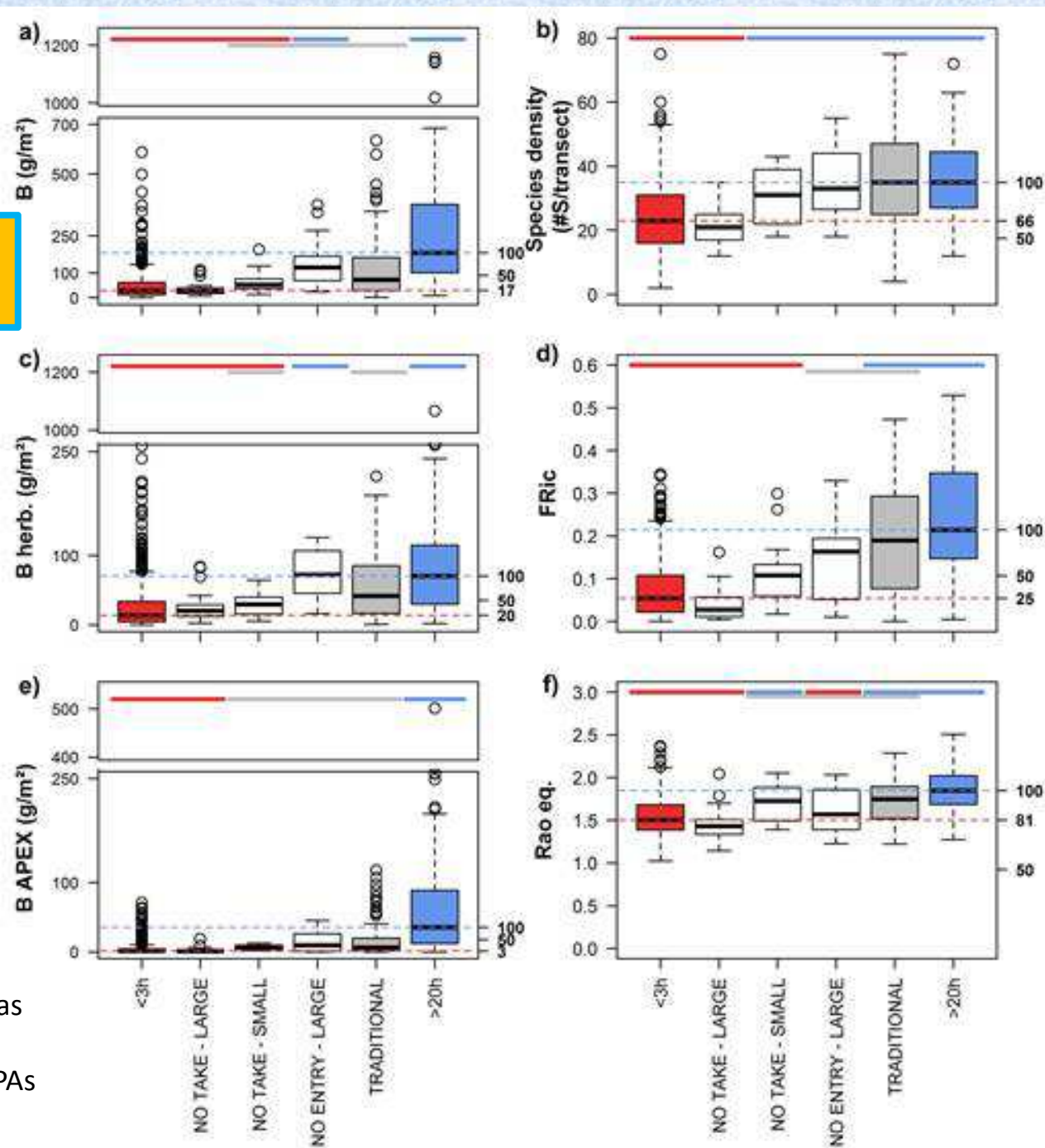
similar to Pristine Areas



similar to exploited Areas

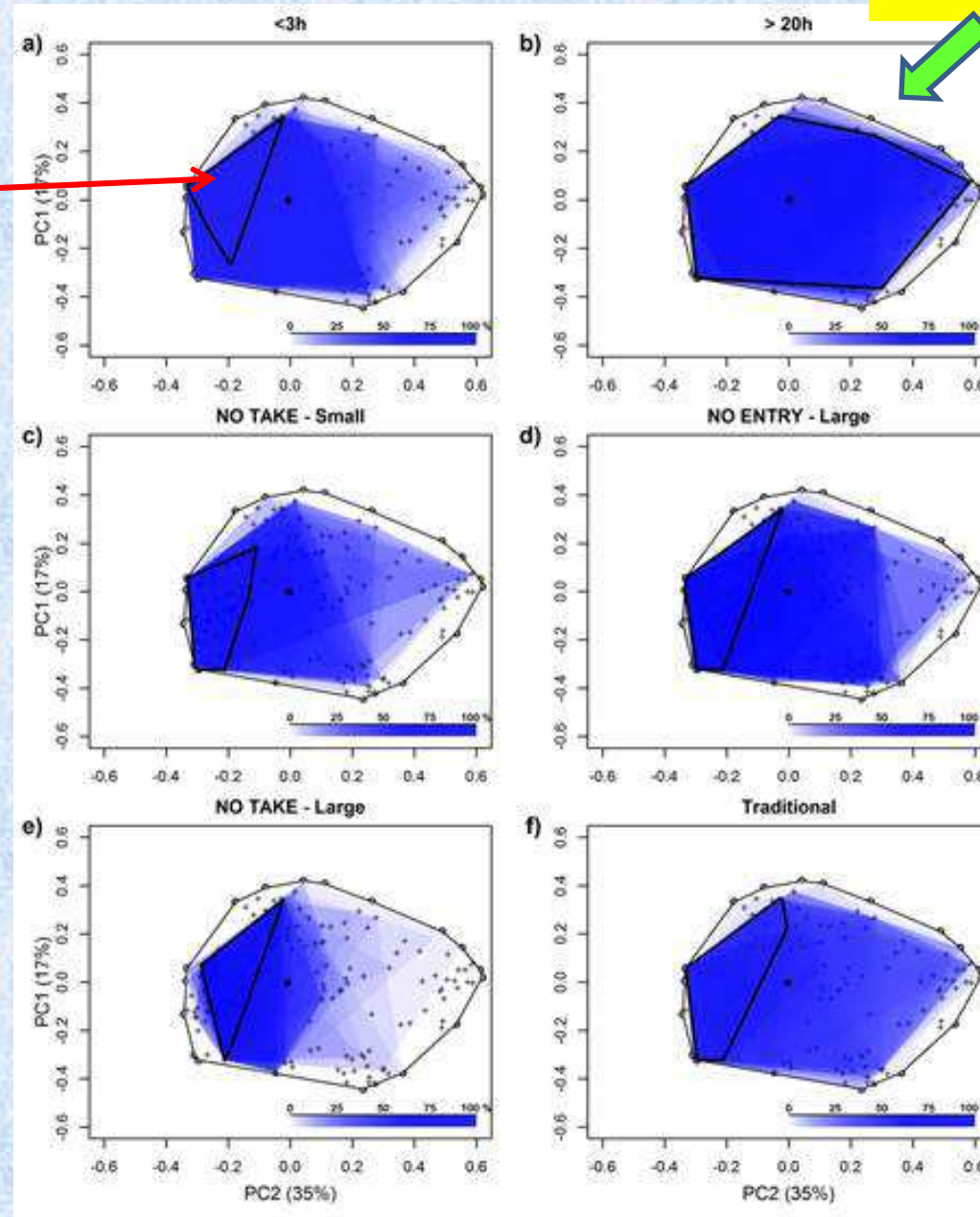


similar to traditional MPAs



Functional volume
filled by > 50% of
the assemblages

There is a huge loss in
functional space from
Pristine Areas to any
management type



What next ?

We have so far mainly focused on diversity (taxonomic & functional)

Need to integrate more Abundance, Biomass and Production

Need to integrate the social & economical aspects to a much larger extent

Need to build robust, pragmatic and transferable predictive models

Reef Fish prove to be an excellent model for general questions in Ecology

Comparative analyses with other vertebrates (birds, mammals, FW fish)

Increasing interest from theoretical ecologists because unique Data Base

Small is at times beautiful



THANK YOU