

# Is sympatric speciation more important in the ocean?

04/22/2021

UROP Online Symposium

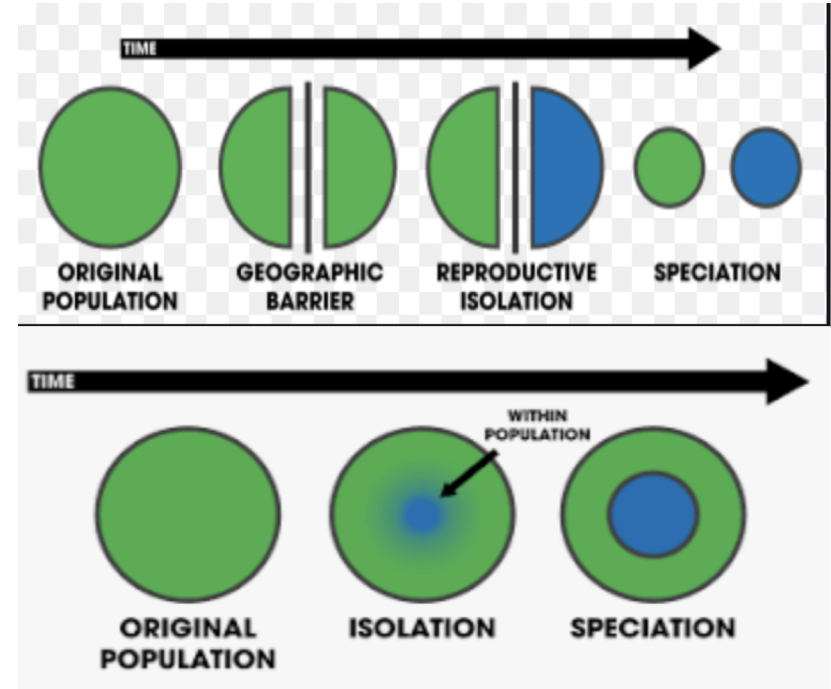
Xiaoyang Song

# Introduction & Background

- Terrestrial environment: existence of hard barriers, geographical isolation, rich resources, high complexity, etc.
- Marine environment: lack of absolute barriers, dispersal ability of larvae of invertebrates, importance of ocean current, etc.
- ***It's reasonable to expect more sympatry in the ocean.***

# Allopatric vs. Sympatric Speciation

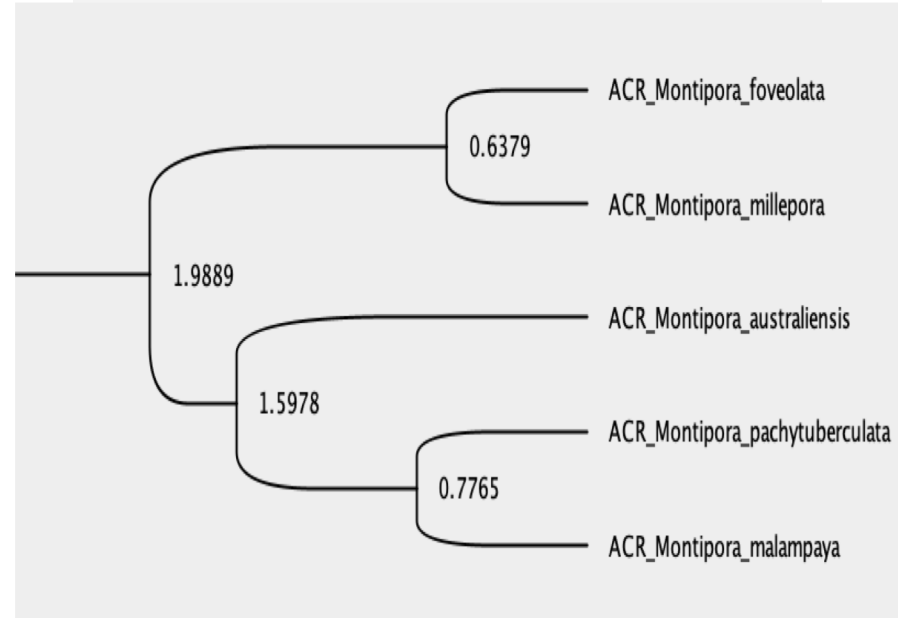
- ***Allopatric Speciation***: speciation that happens when two populations of the same species become isolated from each other due to geographic changes.
- ***Sympatric speciation***: speciation that occurs when two groups of the same species live in the same geographic location. (high range overlap )



Top: allopatric speciation vs. Bottom: sympatric speciation

# Phylogenetic data

- Morphological data (e.g. character matrix)
- Genetic information (e.g. DNA sequences)
- **The Principle of Parsimony:** the simplest explanation that can explain the data is to be preferred (i.e. a hypothesis (tree) of relationships that requires the smallest number of character changes is most likely to be correct.)
- Other concepts: ***node age***, ***species pairs***, etc.

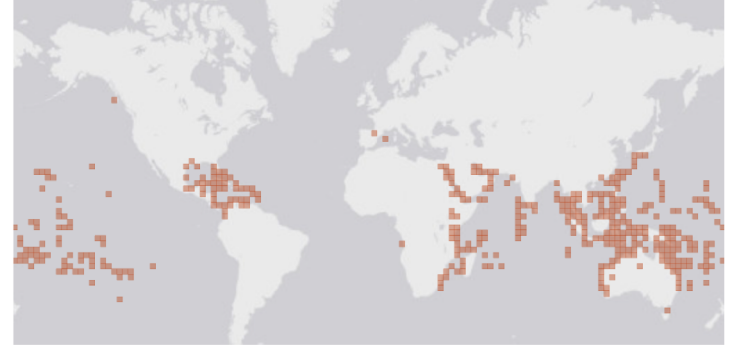


Example of Evolutionary tree for Corals. The numbers above represent the age of the node in the tree..

# Other data?

- **Biogeographic data**
- Help identify range overlap of sister species pairs
- OBIS (Ocean Biodiversity Information System) provides distribution and occurrences data for most known species.
- R library: obistools, georange

DISTRIBUTION



RECORDS



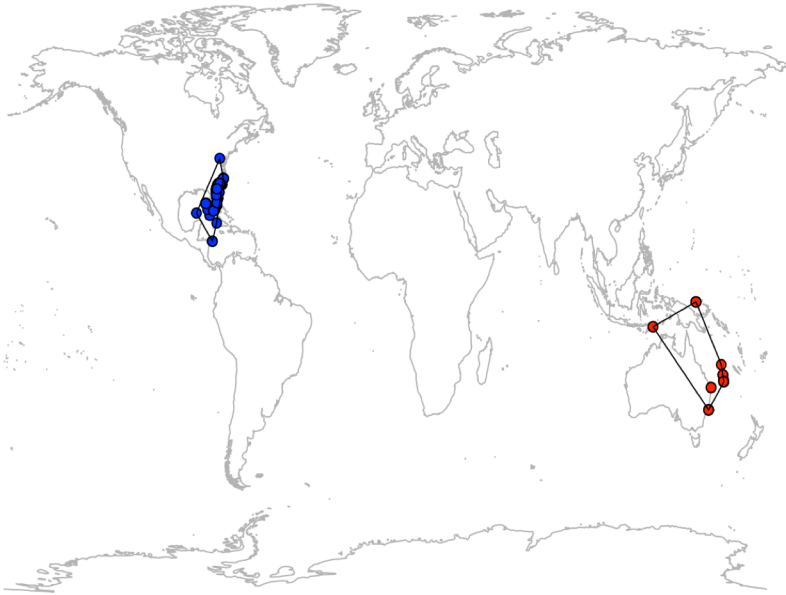
# Methodology

---

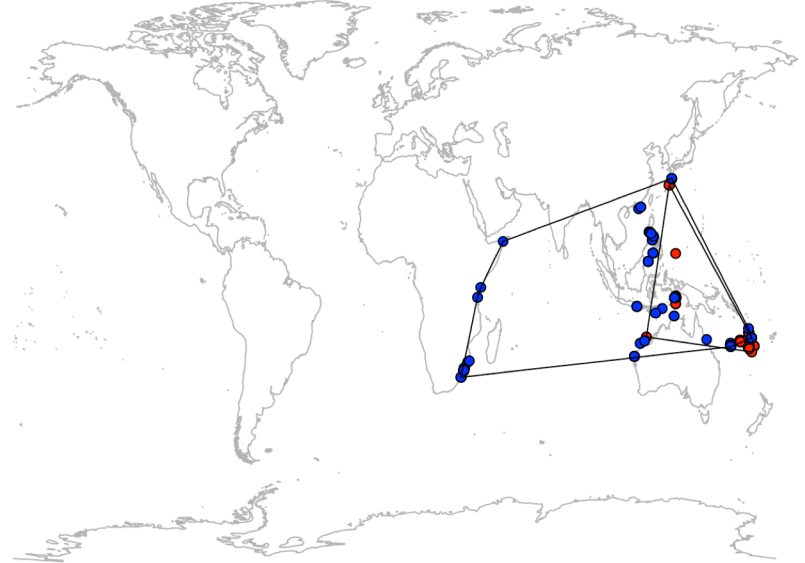
- Build a dataset of marine species with their phylogenetic and range information (including marine mammals and invertebrates).
- Compute the **range overlap** and examine the pattern of speciation
  - a) Identify and discard Outliers
  - b) We compute range overlap by computing the convex hull of data points and the intersected areas of two species.
  - c) Formula: **range overlap** =  $area_i / \min(area1, area2)$ , where **area<sub>i</sub>** is the area of intersection of ranges of two species.
- Examine the relationship between range overlap and node age (i.e. age of the species)

# Example: How we compute range overlap

---



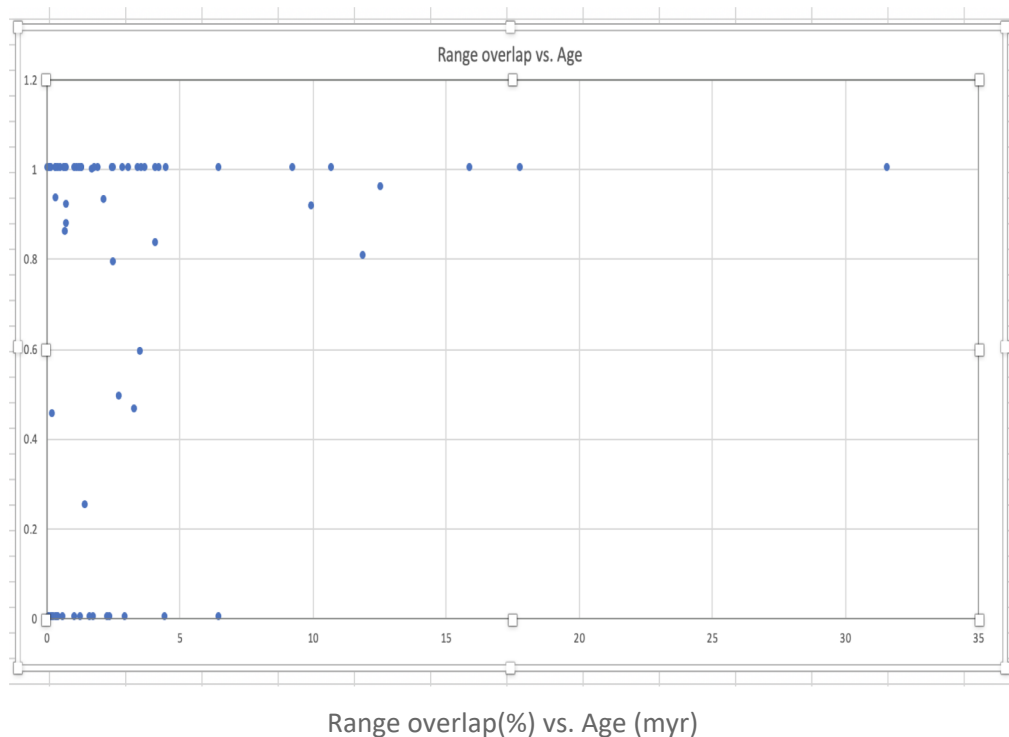
Ex 1: Species pairs with overlap of 0%  
(*Thecopsammia elongata* vs. *Thecopsammia socialis*)



Ex 2. Species pairs with overlap of 0.97  
(*Tropidocyathus labidus* vs. *Tropidocyathus lessonii*)

# Results / Discussion

- > **50%** species pairs in our dataset have a **range overlap** > **0.8**, while for genus with two species, most of them have perfect overlap.
  - Highly sympatry of our dataset
- **Perfect overlap** for really old species
  - Time is long enough for speciation signal to be lost (i.e. meaningless to make inference based on that)
  - Random distribution (influence of ocean current and migrations)





# Results & Graphics

---

**Will add something  
later!**



# Conclusion

---

- Sympatric speciation is the hidden speciation mechanisms for some invertebrates and mammals in the ocean, and it happens for many marine species.
  - Lower speciation rates since sympatric speciation is generally slower than allopatric speciation.
  - However, speciation mechanism for old species remains unknown.
- The increasing importance of sympatry may help explain the fact that ocean has lower biodiversity than the land.
- May add something

**Thanks!**