Genetic Variability in Acropora

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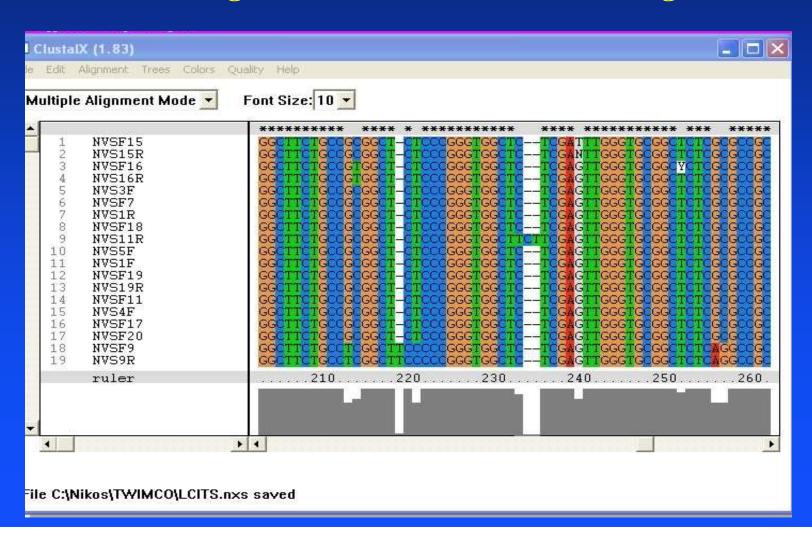
University of Puerto Rico, Mayagüez





■ Fact: genetic variation is the raw material of evolution

Visualizing DNA variation at the ITS-1 region



Observation

Primary mode of reproduction colony fragmentation (low rates of sexual reproduction)

Prediction

Asexual reproduction will result in reduced (local) genetic diversity

Acropora cervicornis

A. palmata



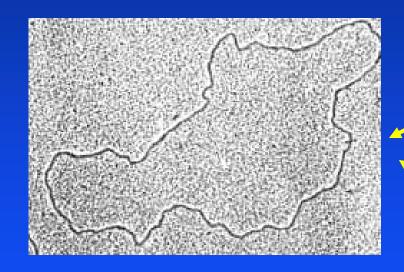


Goal of this proposal

Evaluate the genetic variability of A. cervicornis and A. palmata at different levels of tissue organization

A. palmata

mtDNA





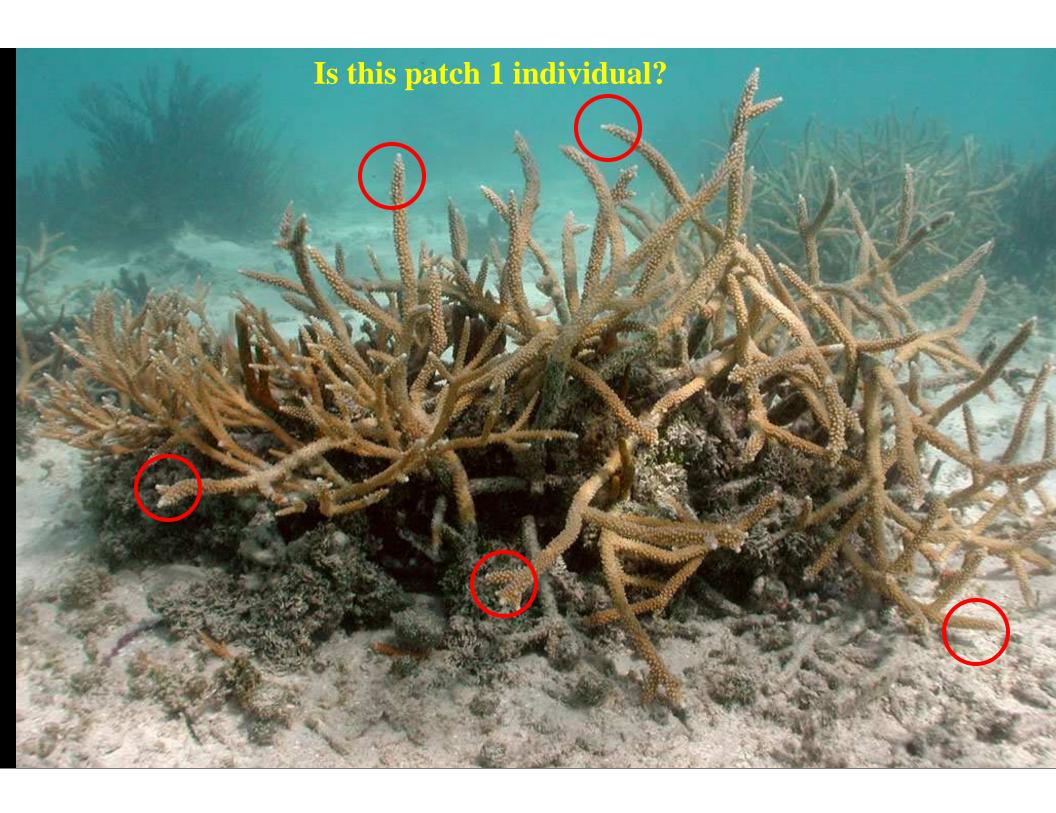
A. cervicornis

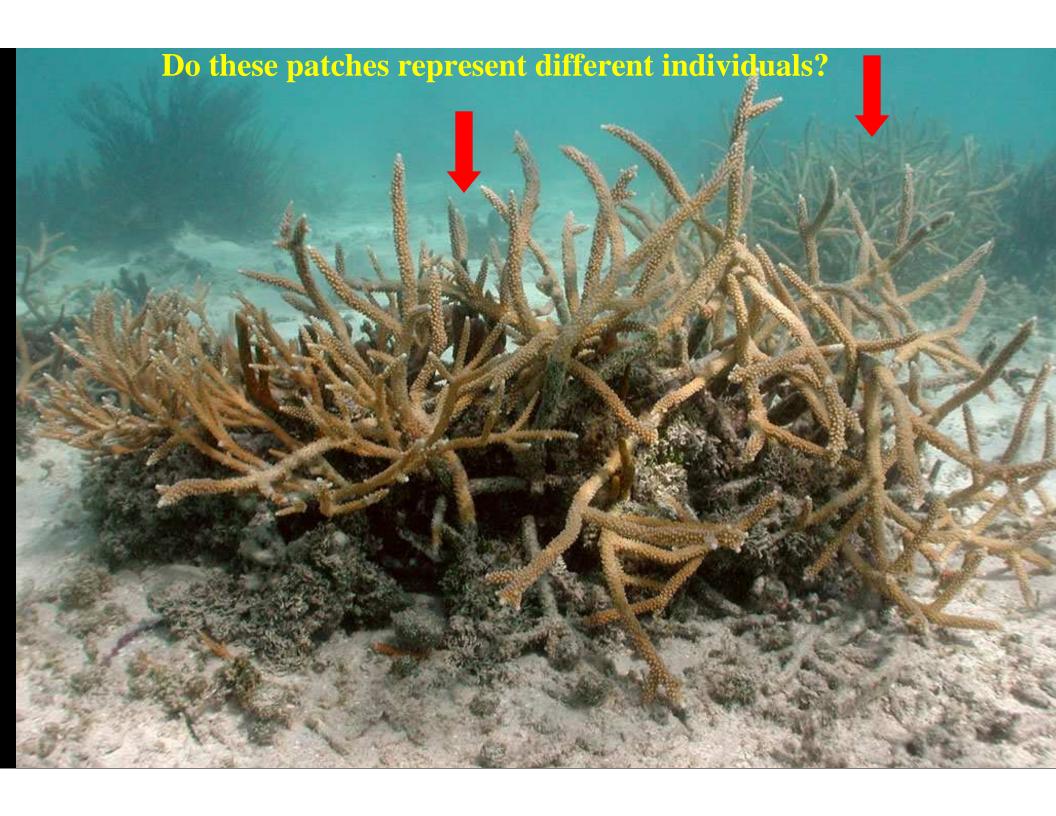


Hierarchical Design

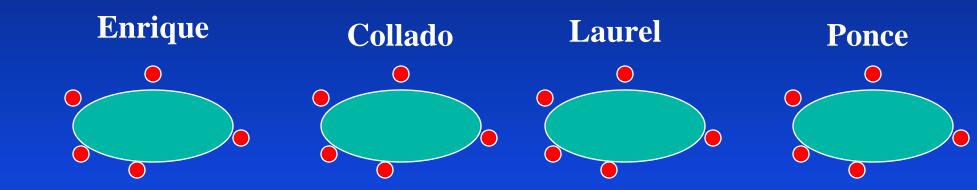
We will evaluate levels of genetic diversity:

- 1) Within discrete patches of *Acropora*
- 2) Among discrete patches of *Acropora* within sampling locations
- 3) Among sampling locations within islands
- 4) Among different islands



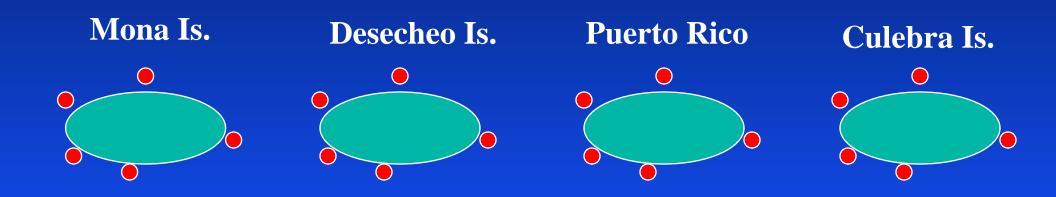


Genetic diversity of *Acropora* between sampling locations



Acropora patch

Genetic diversity of *Acropora* between islands



Acropora patch

Proposed Sampling Locations of *Acropora*



Sampling Locations

Disturbed: Media Luna, Turrumote, Laurel, Enrique, Margarita, Collado, Guanica, Ponce, Rincon.

Non-disturbed: Desecheo Is., Mona Is., Culebra Is.

Materials and Methods

Candidate Genes:

MtDNA: Cytochrome b, putative control region.

Nuclear DNA: ITS-1, and introns from *Pax-C*, calmodulin, and minicollagen

Course of Action

- Develop a reliable DNA extraction technique in Acropora
- Optimize PCR amplification conditions for 4-6 genes per specimen
- **Expand collection of** *Acropora* **to other locations**
- Analyze data

Proposed Schedule

■January 2005-Summer 2005

Training of graduate student

Primer ordering

Collection of fresh coral tissue (local)

DNA extraction

PCR optimization

■ Fall 2005-Summer 2006

Collection of Acropora from all locations

Collection of data

Data Analysis

Manuscript Preparation

People involved with this project

Joselyd Garcia: molecular work, field samples

Sequence divergence in minicollagen, calmodulin, and PaxC (0.6%-2.1%). In ITS-1 is up to 13.2%.

