

# Are you my mother? A test of matrilineal social organization in mass strandings and living groups of rough-toothed dolphins

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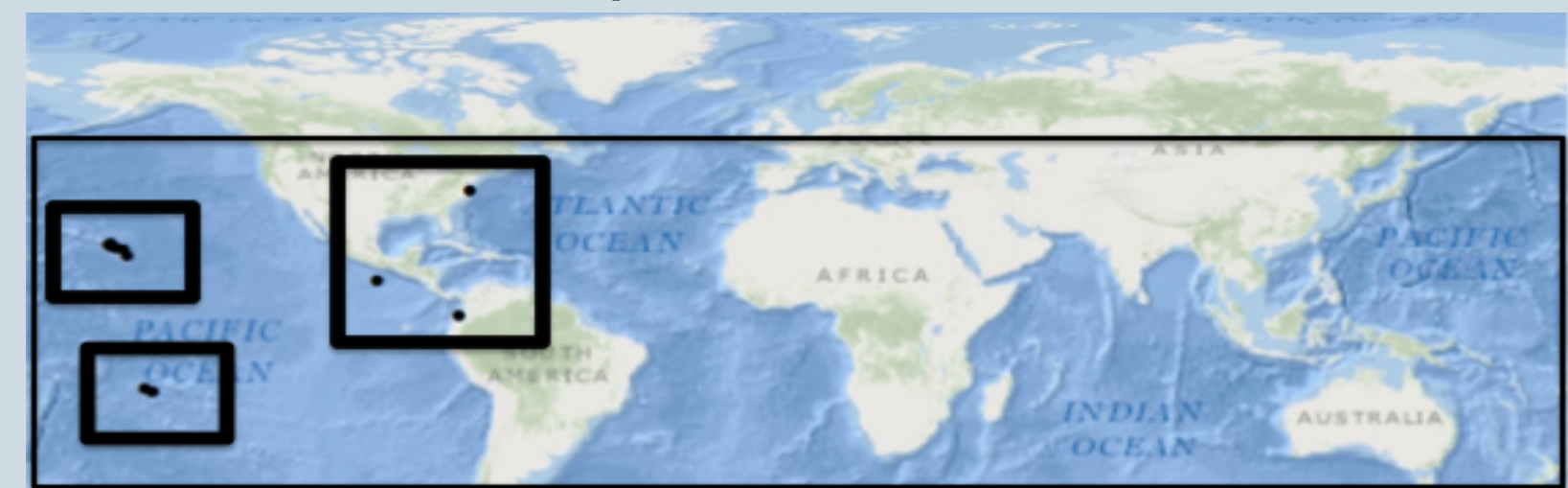


## Introduction



Rough-toothed dolphins  
*Steno bredanensis*

- Form isolated communities containing stable groups around some oceanic islands<sup>1</sup>. These groups generally range from 8-15 individuals, but "super groups" have been observed with as many as 90 individuals<sup>2</sup>.
- Show clear differences in habitat use between island groups, and limited movement among islands<sup>2</sup>.
- Exhibit social characteristics such as cooperative foraging and care-giving behavior similar to species with matrilineal structure like killer whales and pilot whales<sup>3,4</sup>.

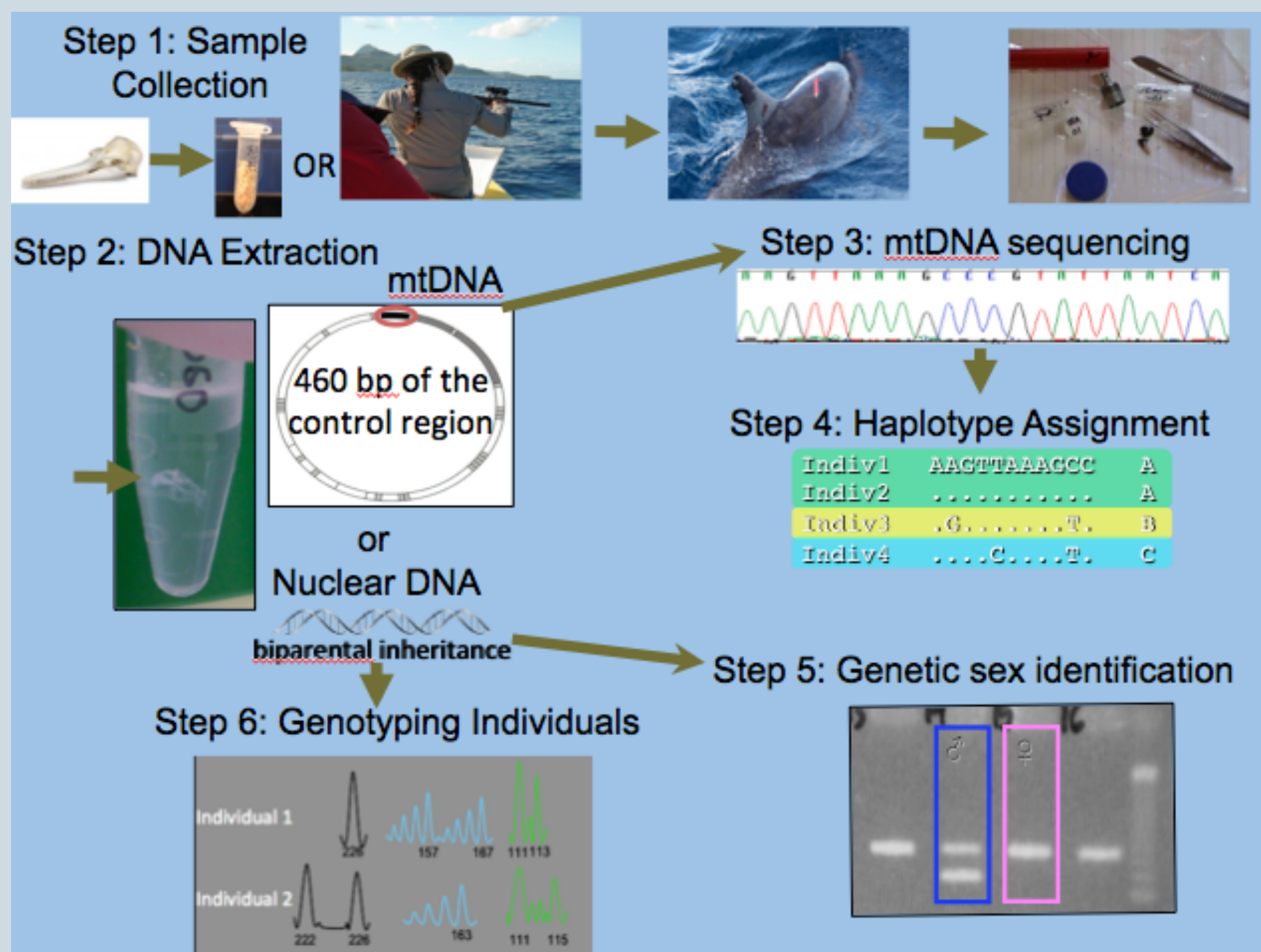


Habitat range and specific study areas

## Methods

### Sample collection spans 1976-2010

- 4 groups (3 mass strandings and 1 bycatch) n=28 teeth samples from adults
- 14 groups; n=106 skin samples from adults



### Markers

- 460bp (biopsies) 350bp (teeth) revealed 24 mtDNA haplotypes.
- 14 microsatellite loci used in genotyping biopsy samples were assessed for null alleles, HWE and linkage disequilibrium, replicates deleted<sup>5</sup>.

### Bi-parental Relatedness (biopsy samples only)

- Relatedness within groups versus between groups was assessed in GenAlEx<sup>6</sup> using the Queller and Goodnight estimator<sup>7</sup>.

### Maternal Relatedness

- An AMOVA was performed in Arlequin to test population structure<sup>8</sup>.
- Mantel test of correlation was used to test if individuals in a group were more likely to share a haplotype than expected by chance<sup>6</sup>.

## Research Questions

- Q1. Is there geographic structure among island communities?  
Q2. Do rough-toothed dolphins form groups with extended matriline?  
Q3. Are groups composed of close kin?

## Biopsy samples results

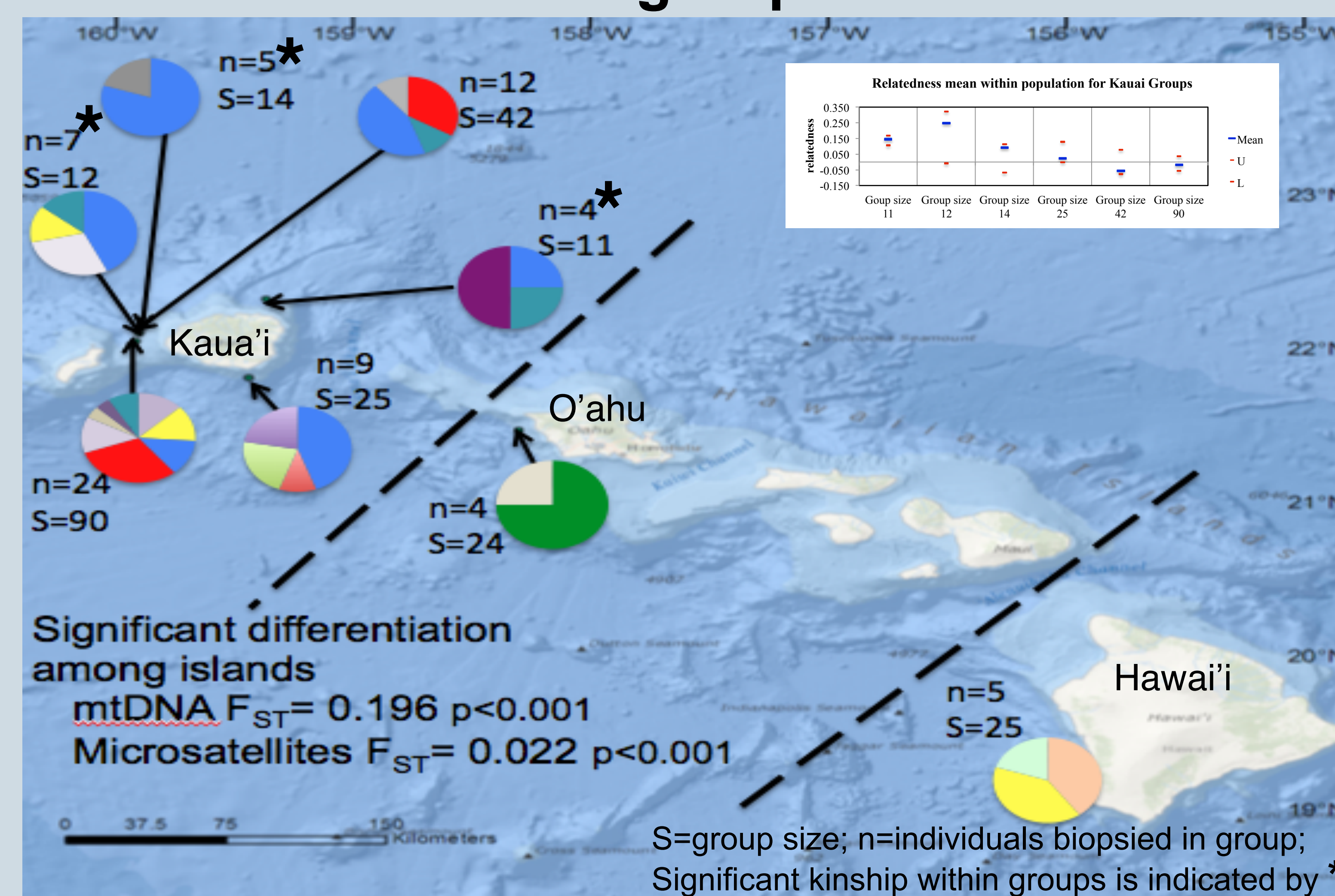
### Hawaiian Islands 8 Groups

8 groups with multiple matriline

### Differentiation within Kauai

mtDNA  $F_{ST}=0.054$   $p=0.08$  Microsatellite  $F_{ST}=0.022$   $p=0.003$

Relatedness within groups  $R = 0.007-0.246$



Significant differentiation among islands

mtDNA  $F_{ST}= 0.196$   $p<0.001$   
Microsatellites  $F_{ST}= 0.022$   $p<0.001$

S=group size; n=individuals biopsied in group; Significant kinship within groups is indicated by \*

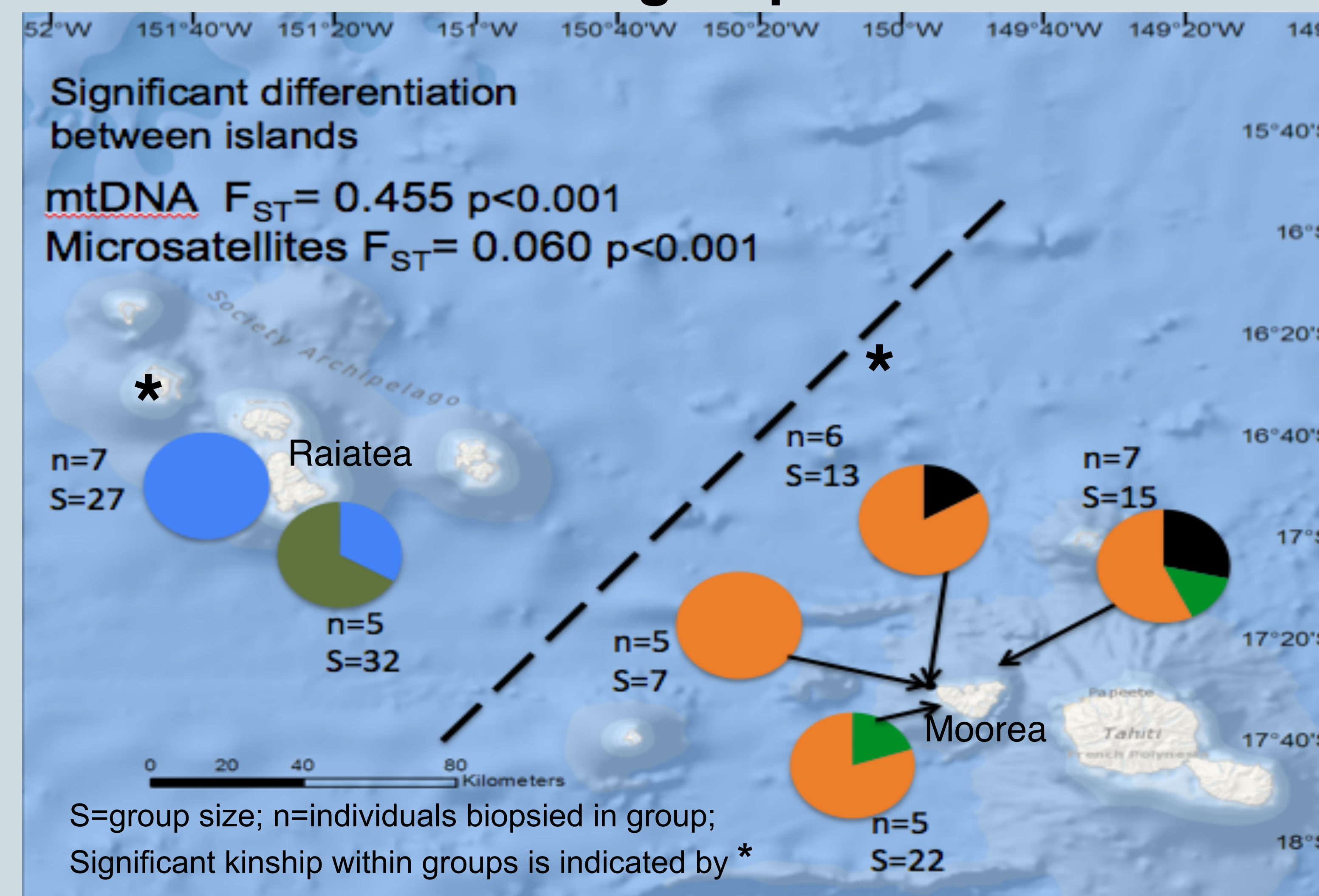
### French Polynesia 6 Groups

4 groups with multiple matriline, 2 groups with single matriline

### Differentiation within Moorea

mtDNA  $F_{ST}=0.002$   $p=0.594$ ; Microsatellites  $F_{ST}=0.051$   $p=0.004$

Relatedness within groups  $R= 0.001-0.124$



Significant differentiation between islands

mtDNA  $F_{ST}= 0.455$   $p<0.001$   
Microsatellites  $F_{ST}= 0.060$   $p<0.001$

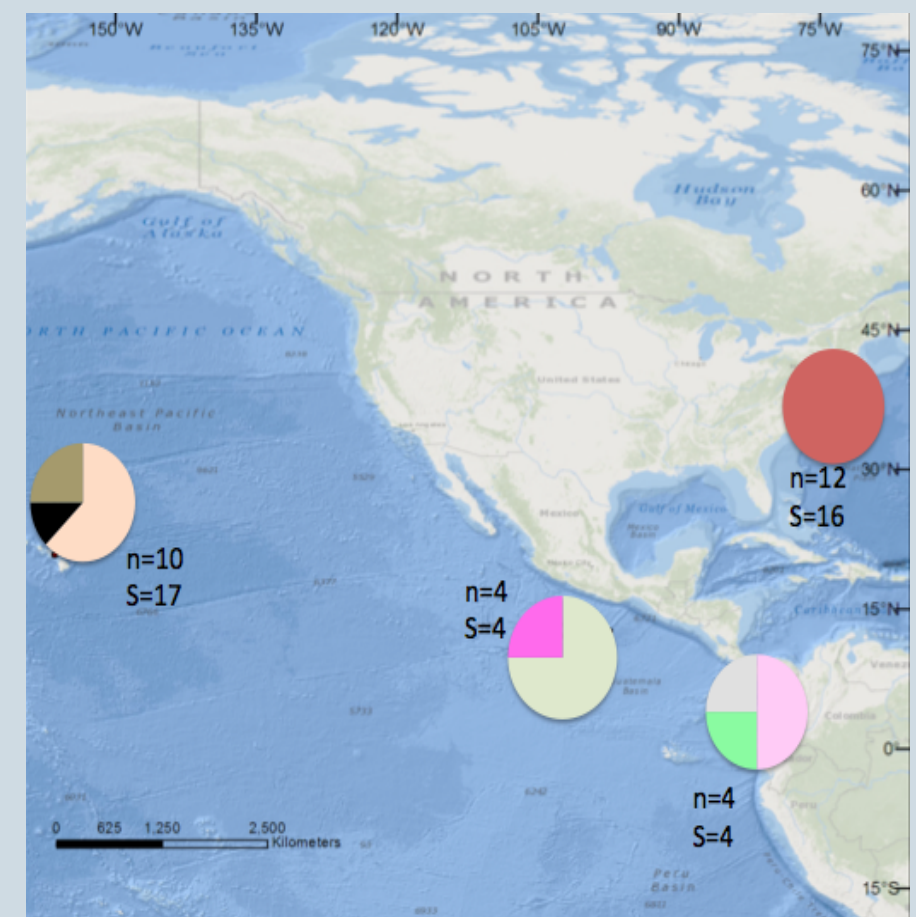
S=group size; n=individuals biopsied in group; Significant kinship within groups is indicated by \*

## Teeth sample results

### Mass Strandings

- 1 group with a single matriline
- 3 groups with multiple matriline

There was no significant difference in maternal structure in live or stranded groups.



## Conclusion



A1. Significant genetic differentiation between island communities but not between all groups within islands indicates local fidelity may drive geographic structure.



A2. Multiple matriline and significant relatedness within some groups suggests genetic structure among groups, but this structure is not strictly matrilineal.



A3. Kinship was significant in 5 of 14 groups. Estimated relatedness was markedly lower in large groups. These groups may be composed of several small groups, similar to the pattern seen in long-finned pilot whales.



Significant relatedness in small groups is consistent with kinship based social behavior such as caregiving or cooperative foraging.

## Literature Cited

1. OREMUS, M. et al. 2012. Pelagic or insular? Genetic differentiation of rough-toothed dolphins in the Society Islands, French Polynesia. *Journal of Experimental Marine Biology and Ecology* 432-433: 37-46.  
2. BAIRD, R. W. et al. 2008. Site fidelity and association patterns in a deep-water dolphin: Rough-toothed dolphins (*Steno bredanensis*) in the Hawaiian Archipelago. *Marine Mammal Science* 24: 535-553.  
3. JEFFERSON, T. A. 2008. Rough-toothed dolphin. In W. F. PERRIN, B. WURSIG and J. G. M. THEWISSEN eds. *Encyclopedia of Marine Mammals*. Elsevier Inc, Burlington, MA.  
4. OREMUS, M. et al. 2013. Genetic Evidence of Multiple Matriline and Spatial Disruption of Kinship Bonds in Mass Strandings of Long-finned Pilot Whales, *Globicephala melas*. *The Journal of Heredity* 104: 301-311.

5. VAN OOSTERHOUT et al. 2006. Estimation and adjustment of microsatellite null alleles in nonequilibrium populations. *Molecular Ecology Notes* 6: 255-256.  
6. PEAKALL, R. and P. E. SMOUSE. 2012. GenAlEx 6.5: Genetic analysis in Excel. Population genetic software for teaching and research - an update. *Bioinformatics*.  
7. QUELLER, D. and K. GOODNIGHT. 1989. Estimating Relatedness Using Genetic Markers. *Evolution* 43: 258-275.  
8. EXCOFFIER, L. and H. E. L. LISCHER. 2010. Arlequin suite ver 3.5: a new series of programs to perform population genetics analyses under Linux and Windows. *Molecular Ecology Resources* 10: 564-567.

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