Employing Genetics To Study Whales

An Informal Introduction

Of Course Not Because...





Lunch break in Samana Bay, the Dominican Republic 1990

Nor?



Camp in Uummannaq District, Northwest Greenland 1991

Whales Are A Resource

- Many populations and species depleted due to past commercial whaling operations.
- Subsistence hunting still ongoing in many parts of the world, as well as scientific whaling, and some unknown degree of illegal whaling.
- Whale watching a source of income in many developed as well as underdeveloped countries.



...But Because...



...And Because

- Very different system compared to "standard" terrestrial or fresh water models. Also different from many marine organisms
 - Active dispersal at all life stages
 - Wide ranges of movement in an environment with few barriers
 - Long life-span and presumably some degree of "culture" among individuals

Basic Aspects

Advantages

- Taxonomically closely related species-complex
- Many truly cosmopolitan species, i.e., many different levels of evolutionary divergence
- Mating and foraging tempo-spatially separated in many species

Basic Aspects

Disadvantages

- Difficult to observe and tag directly
- Highly political due to very different cultural views on exploitation of whales
- "Pop science"

Marine Mammals – Very Different Creatures Than Bugs and Plants
Finite population sizes
Overlapping generations
Small litter size

– For instance:

Humpback whales in the north Atlantic number some 10,000 individuals. Females mature at the age of six and give birth to one calf every second year. The life expectancy is supposedly 30 years, but not known.

Interacting Levels and Processes

Ecosystem Changes in abundance & structure

> *Organismal* Mode and rate of dispersal & gene flow

Molecular

Mode and rate of mutation

Biopsy Collection









....And Into Pickle Juice



Genetic Analyses Conducted



Sequencing mtDNA

Analysis of Maternally and Biparentally Inherited Genetic Markers



The Humpback Whale Megaptera novaeangliae

- Cosmopolitan species
- Weight: ~40 tonnes
- Length ~45' or 15 meters
- Age at maturity: 6-7
- Extensively harvested in all oceans during the 19th and early-mid 20th century
- Protected in the US under the endangered species as well as the marine mammal protection act
- International trade regulated by CITES (for all cetacean species)

East-west Cline in Estimates of Genetic Diversity in the Maternally Inherited Mitochondrial Genome



Genealogy Of Mitochondrial



Area	% of each clade			
	Common	western		
Barents Sea	97	3		
Iceland	86	14		
West Greenland	70	30		
Labrador	74	26		
Newfoundland	68	32		
Gulf of St. Lawrence	62	38		
Gulf of Maine	67	33		

Antarctic alleles

Common North Atlantic alleles western North Atlantic alleles

Western and eastern North Atlantic areas



Divergence Estimates

	<u>Nuc</u>	clear DNA	<u>mtDNA</u>		
Area	F _{ST} (mean)	Range	H _{ST} (mean)	Range	
eastern NAtl.	.0022	-	.009	-	
western NAtl.	.0002	.00010005	.009	.000026	
Barents Sea - wNAtl.	.0038	.00090078	.040	.015085	
Iceland - wNAtl.	.0014	.00030027	.025	.010042	

Nuclear DNA estimates based on six loci

MtDNA divergence among feeding grounds





Calves Stay With Their Mother the 1st Year



Identification Of Mother & Calf Relations



Maternally-directed Site-fidelity to Summer Feeding Grounds

- Individual humpback whales appear to return to the same high-latitude feeding ground every spring throughout its life
- The feeding ground of choice is that to which the calf migrated to with its mother
- Cultural transmission



Two Breeding Populations



Effects of Pleistocene Glaciations On Genetic Diversity





Individual-based Analyses

Insights on an ecological time scale

- Individual identification
 - 6-15 microsatellite loci
 - Estimation of abundance and individual ranges of movement
- Identification of close relatives
 - 20+ microsatellite loci for parent-offspring detection
 - Estimation of abundance
 - Reproductive success -> selection
 - Population structure
 - Estimation of demographic parameters

North Atlantic Humpback Whale

Area	Period	Samples	<i>I</i> [#] (x 10⁻7)	95% CI limits [§] (x 10 ⁻⁷)	Genotypes	Males	Females
Barents Sea	1992-1993	36	8.46	4.9 - 38	35	13	22
Gulf of St. Lawrence	1990-1995	65	1.94	1.26 - 5.52	56	28	28
Gulf of Maine	1990-1995	292	1.38	1.02 - 2.11	256	118	138
Iceland/Jan Mayen	1991-1993	112	1.42	0.88 - 3.28	100	50	50
Newfoundland/Labrador	1991-1995	572	1.34	1.07 - 1.86	464	237	227
West Greenland	1988-1994	189	1.23	0.89 - 2.10	148	75 ^{&}	72 ^{&}
West Indies							
Minus inter-area recaptures		2,491	1.51	1.34 - 1.75	2,368		
Unique genotypes only		2,368	1.51	1.32 - 1.72		1,331 ^{&}	1,033 ^{&}

[#]Probability of identical genotype²⁶ across all loci calculated from all samples (including recaptures). [§]Estimated from 1,000 bootstrap samples. [&]No gender was obtained for a total of four samples

Different Number of Males and Females on the Breeding Grounds

- Abundance estimates of each sex were estimated from with the samples collected on the breeding ground during 1992 and 1993
 - 4,804 males (95% CI: 3,374 7,123)
 - 2,804 females (95% CI: 1,776 4,463)
- Even sex ration in calves and among feeding ground samples
- Difference in male and female abundance probably due to "temporal fidelity" in migration timing among females

Temporal Fidelity



Thank You