Identifying resilient acroporid populations for effective restoration: Mapping genotypic distributions and connectivity in the U.S Virgin Islands

Principal Investigators:

Ron Hill, <u>ron.hill@noaa.gov</u> Hannah Nylander-Asplin, <u>hannah.nylander-asplin@noaa.gov</u> Jennifer Doerr, <u>jennifer.doerr@noaa.gov</u>

> NOAA National Marine Fisheries Service Southeast Fisheries Science Center Galveston Lab 4700 Avenue U Galveston, TX 77551

The distribution of *Acropora* spp. at three sites (Thatch Cay, Lovango Cay, No Name Bay) are well documented based on previous tissue collection and long-term transect analysis (Nylander-Asplin et.al 2021). Rather than mapping by species alone, here we have the unique ability to capture population dynamics with regards to the number of unique genotypes sampled. Mapping the *Acropora spp*. by genotype provides a clearer understanding of the spatial relationships and expanse of distinct genotypes.

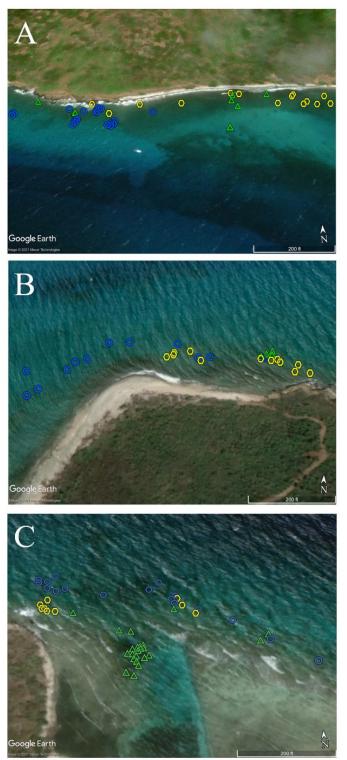


Fig 1. Acropora samples collected around St. Thomas and St. John in 2017. Colonies were sampled at A) Thatch Cay, B) Lovango Cay and C) No Name Bay. Yellow hexagons represent *A. palmata*, blue circles represent *A. cervicornis*, and green triangles represent *A. prolifera*.



Thatch Cay:

N = number of sampled colonies N_g = number of unique genets N_q/N = genotypic richness.

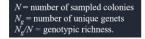
	Thatch Cay		
	N	N_g	Ng/N
A. cervicornis	22	15	0.68
A. palmata	17	9	0.53
A. prolifera	8	6	0.75



Approximate area of enlargement

Fig 2. Genotypic richness of the *Acropora* taxa at Thatch Cay, U.S. Virgin Islands. Circles represent *A. cervicornis*, triangles indicate *A. prolifera* and hexagons detail *A. palmata* distributions. Each unique color represents a unique genotype with similar colors representing more closely related colonies.





	Thatch Cay		
	N	N_g	N_g/N
A. cervicornis	22	15	0.68
A. palmata	17	9	0.53
A. prolifera	8	6	0.75



Approximate area of enlargement

Fig 3. Genotypic richness of the *A. cervicornis* at Thatch Cay, U.S. Virgin Islands. Each color represents a unique genotype where closer colors represent similar, but not identical genotypes. Out of 22 total samples taken, 15 unique genotypes were isolated resulting in a genotypic richness of 0.69 Ng/N.

Thatch Cay: Acropora palmata Genotypic Richness

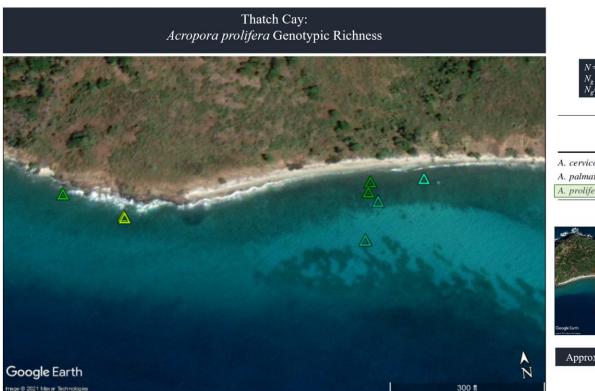


N = number of sampled colonies $N_g =$ number of unique genets $N_g/N =$ genotypic richness.

	Thatch Cay		
	N	N_g	N_g/N
A. cervicornis	22	15	0.68
A. palmata	17	9	0.53
A. prolifera	8	6	0.75



Fig 4. Genotypic richness of the *A. palmata* at Thatch Cay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 17 total samples taken, 9 unique genotypes were isolated resulting in a genotypic richness of 0.53 Ng/N.



 $N_g/N =$ genotypic richness.

number of sampled colonies = number of unique genets

	Thatch Cay		
	N	N_g	N_g/N
A. cervicornis	22	15	0.68
A. palmata	17	9	0.53
A. prolifera	8	6	0.75



Approximate area of enlargement

Fig 5. Genotypic richness of the hybrid *A. prolifera* at Thatch Cay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 8 total samples taken, 6 unique genotypes were isolated resulting in a genotypic richness of 0.75 Ng/N.

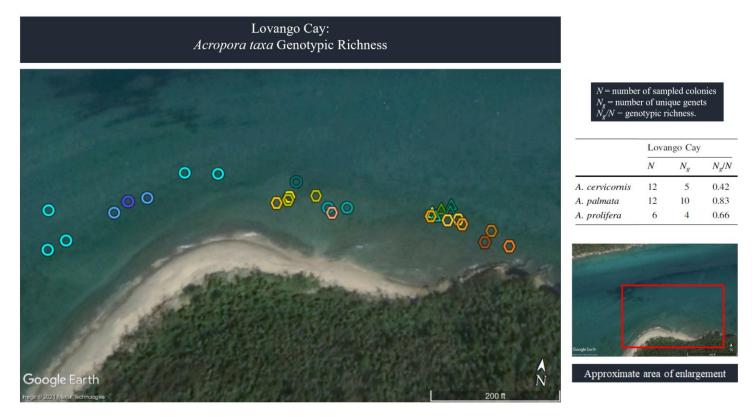
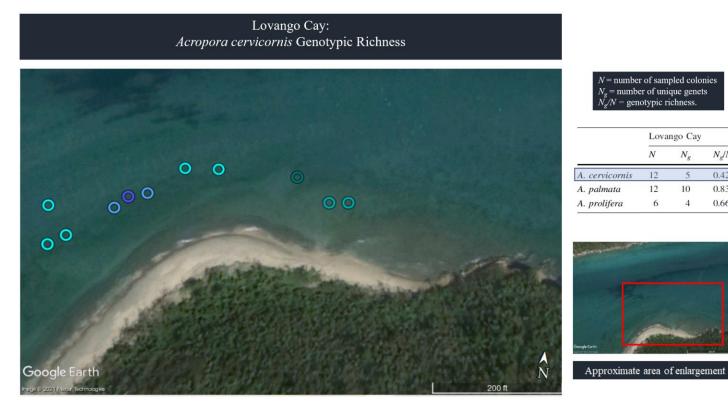


Fig 6. Genotypic richness of the Acropora taxa at Lovango Cay, U.S. Virgin Islands. Circles represent A. cervicornis, triangles indicate A. prolifera and hexagons detail A. palmata distributions. Each unique color represents a unique genotype with similar colors representing more closely related colonies.



NJN

0.42

0.83

0.66

Fig 7. Genotypic richness of the A. cervicornis at Lovango Cay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 12 total samples taken, 5 unique genotypes were isolated resulting in a genotypic richness of 0.42 Ng/N.

Lovango Cay: Acropora palmata Genotypic Richness



Fig 8. Genotypic richness of the *A. palmata* at Lovango Cay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 12 total samples taken, 10 unique genotypes were isolated resulting in a genotypic richness of 0.83 Ng/N.

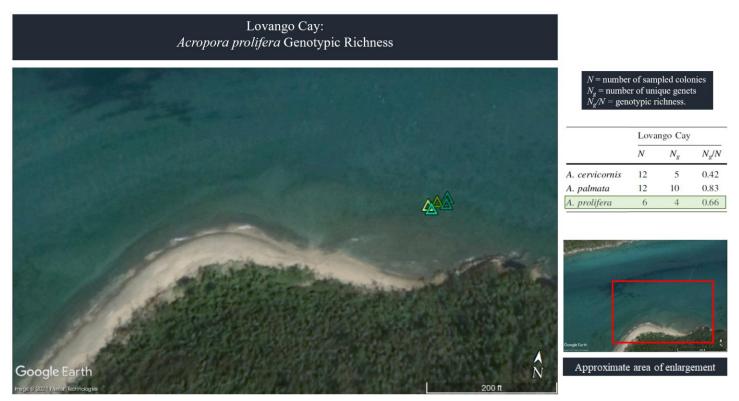
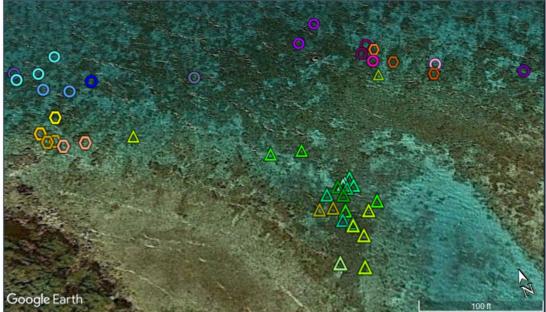


Fig 9. Genotypic richness of the hybrid *A. prolifera* at Lovango Cay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 6 total samples taken, 4 unique genotypes were isolated resulting in a genotypic richness of 0.66 Ng/N.

No Name Bay: Acropora taxa Genotypic Richness



N = number of sampled colonies $N_g =$ number of unique genets $N_g/N =$ genotypic richness.

	No Name Bay		
	N	N_g	Ng/N
A. cervicornis	16	11	0.69
A. palmata	11	8	0.73
A. prolifera	25	25	0.6

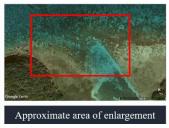


Fig 10. Genotypic richness of the *Acropora* taxa at No Name Bay, U.S. Virgin Islands. Circles represent *A. cervicornis*, triangles indicate *A. prolifera* and hexagons detail *A. palmata* distributions. Each unique color represents a unique genotype with similar colors representing more closely related colonies.



Fig 11. Genotypic richness of the *A. cervicornis* at No Name Bay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 16 total samples taken, 11 unique genotypes were isolated resulting in a genotypic richness of 0.69 Ng/N.

No Name Bay: Acropora palmata Genotypic Richness



Fig 12. Genotypic richness of the *A. palmata* at No Name Bay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 11 total samples taken, 8 unique genotypes were isolated resulting in a genotypic richness of 0.73 Ng/N.

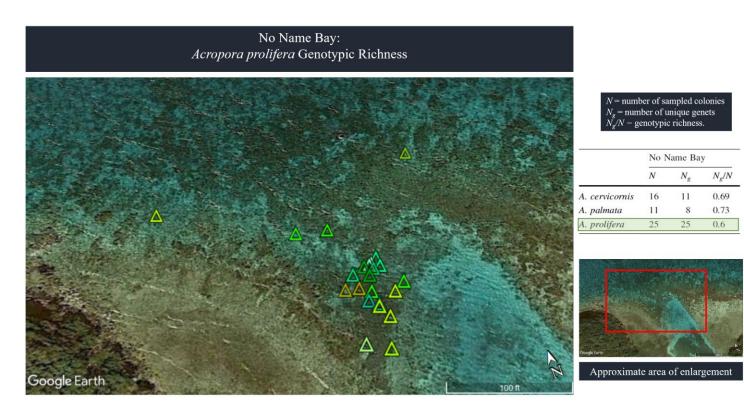


Fig 13. Genotypic richness of the hybrid *A. prolifera* at No Name Bay, U.S. Virgin Islands. Each color represents a unique genotype, where closer colors represent similar, but not identical genotypes. Out of 25 total samples taken, 15 unique genotypes were isolated resulting in a genotypic richness of 0.6 Ng/N.