Nicolas Locatelli

nl532@cornell.edu

EDUCATION

Aug. 2020 - Ph.D. in Biology, Davenport and Baums Labs

Pennsylvania State University, Eberly College of Science

SEPT. 2017 - M.A. in Ecology, Evolution and Conservation Biology

May 2019 Columbia University, Graduate School of Arts and Sciences

Aug. 2012 - B.S. in Molecular Environmental Biology

May 2015 University of California, Berkeley, College of Natural Resources

RESEARCH EXPERIENCE

SEPT. 2024 -	
PRESENT	

Adirondack Fishery Research Program at Cornell University

Postdoctoral Associate

Ithaca, NY | \Developing gene

♦ Developing genetic tools for eDNA surveys of the Adirondack mountains with an emphasis on detecting intraspecific variation in stocked and threatened species.

AUG. 2020 -

Baums and Davenport Labs at Penn State University

Aug. 2024 | Ph.D. in Biology

University Park, PA

♦ Thesis entitled "Genomic insights into coral evolution and adaptation: A comparative study of Caribbean reef-builders"

♦ Focus on genomic resource development and analyses to disentangle the complex relationship of the coral host, its symbiont, and their changing environment.

JUNE 2019 -SEPT. 2020 Ithaca, NY

Cornell University, Department of Natural Resources

Technician

Worked collaboratively with Diana Baetscher and performed metabarcoding analyses and sample acquisition with the goal of understanding which marine stocks are exploited to grow aquacultured species around the world.

Aug. 2018 -May 2019 New York, NY

Melnick Research Group at Columbia University

Graduate Student Researcher

♦ Under guidance from Drs. Don Melnick and Deren Eaton, analyzed whole genome sequence data to understand patterns of introgression amongst wild macaque species.

AUG. 2017 -MAY 2019 New York, NY

The Drew Lab at Columbia University

M.A. in Ecology, Evolution, and Conservation Biology

♦ Thesis project - Population genetic structure, symbiont composition, and clonal prevalence in *Montipora* and *Porites* corals in Kaneohe Bay, Oahu using ddRADseq.

AUG. 2015 -DEC. 2015

Carlson Lab at the University of California, Berkeley

Field Assistant

Branscomb, CA

 Assisted graduate researcher Suzanne Kelson in macroinvertebrate surveys and capturing/recapturing rainbow trout for PIT tagging and growth monitoring

CA

AUG. 2014 -

Richard B. Gump South Pacific Research Station

DEC. 2014 Undergraduate Researcher
Moorea. Research course project s

♦ Research course project surveying two cleaner wrasse species, their fish clients, and surrounding scleractinia in shallow, fringing coral reefs

JAN. 2013 -

Looy Lab at the University of California, Berkeley

JUN. 2013

Undergraduate Research Apprentice

Berkeley, CA | Computer analysis of Cretaceous angiosperm fossils and their damage by arthropods

PUBLICATIONS

SUBMITTED FEB. 2025

Cauvin A, Carne L, Marhaver KL, Vermeij MJA, Locatelli NS, Baums IBB, Paul VJ, and JL Meyer. Pillar coral microbiomes are predominantly composed of endosymbionts. Submitted to Proc. R. Soc. B.

PREPRINT AVAILABLE JAN. 2025

Stankiewicz KH, Valenzuela JJ, Turkarslan S, Wu W, Gomez-Campo K, Locatelli NS, Conn TL, Radice VZ, Parker KE, Alderdice R, Bay LK, Voolstra CR, Barshis DJ, Baums IB, and NS Baliga (2025) Heat-responsive dynamic shifts in alternative splicing of the coral Acropora cervicornis. bioRxiv. https://doi.org/10.1101/2025.01.21.634199.

PUBLISHED FEB. 2025

Locatelli NS and IB Baums (2024) Genomes of the Caribbean reef-building corals Colpophyllia natans, Dendrogyra cylindrus, and Siderastrea siderea. G3: Genes/Genomes/Genetics https://doi.org/10.1093/g3journal/jkaf020

Assemblies available at Zenodo https://doi.org/10.5281/zenodo.13323986

PUBLISHED Nov. 2024

Locatelli NS, Kitchen SA, Stankiewicz KH, Osborne CC, Dellaert Z, Elder H, Kamel B, Koch HR, Fogarty ND, Baums IB (2024). Chromosome-level genome assemblies and genetic maps reveal heterochiasmy and macrosynteny in endangered Atlantic Acropora. BMC Genomics. https://doi.org/10.1186/s12864-024-11025-3.

Assemblies available at Zenodo https://doi.org/10.5281/zenodo.12021087

PUBLISHED JUN. 2024

Howlett, L, Camp, EF, **Locatelli NS**, Baums IB, Strudwick P, Rassmussen S, Suggett DJ (2024). Population and clonal structure of Acropora cf. hyacinthus to inform coral restoration practices on the Great Barrier Reef. Coral Reefs. https://doi.org/10.1007/s00338-024-02520-w

PUBLISHED OCT. 2023

Baetscher DS, **Locatelli NS**, Won ET, Fitzgerald T, McIntvre PB, and NO Therkildsen (2023). Optimizing a metabarcoding marker portfolio for species detection from complex mixtures of globally diverse fishes. Environmental DNA. https://doi.org/10.1002/edn3.479

Published SEPT. 2022

Baums IB, Chamberland VF, **Locatelli NS**, and TL Conn (2022). Maximizing Genetic Diversity in Coral Restoration Projects. In: van Oppen MJH, Aranda Lastra M. (eds) Coral Reef Conservation and Restoration in the Omics Age. Coral Reefs of the World, vol 15. Springer, Cham. https://doi.org/10.1007/978-3-031-07055-63.

PUBLISHED SEPT. 2021

Drew JA, Kahn BM, **Locatelli NS**, Airey ME, and AT Humphries (2021) Examining stakeholder perceptions of oyster ecosystem services using fuzzy cognitive mapping. Conservation Science and Practice 3(11), e531. https://doi.org/10.1111/csp2.531

PUBLISHED Nov. 2020

Locatelli NS, McIntyre PB, Therkildsen NO, and DS Baetscher (2020) GenBank's reliability is uncertain for biodiversity researchers seeking species-level assignment for eDNA. Proceedings of the National Academy of Sciences 117(51) 32211-32212. https://doi.org/10.1073/pnas.2007421117

AVAILABLE DEC. 2019

PREPRINT | Locatelli NS and JA Drew (2019) Population structure and clonal prevalence of scleractinian corals (Montipora capitata and Porites compressa) in Kaneohe Bay. Oahu. bioRxiv. https://doi.org/10.1101/2019.12.11.860585

INVITED TALKS

JUNE 29, 2022

"Tracing biogeography of fishes used in global aquaculture feed with DNA metabarcoding". DS Baetscher, **Locatelli NS**, Fitzgerald T, Lepak RF, Won ET, Therkildsen NO, and PB McIntyre. **EPA Great Lakes Toxicology and Ecology Division Laboratory**

Note: Co-presented with Diana Baetscher of NOAA Alaska Fisheries Science Center. Presented binary classification (scikit-learn) of feed constituents vs. contamination.

Presentations & Posters

APR 1. 2025

"Insights into Adirondack fish communities and adaptation: Lessons from environmental DNA and whole genome sequencing"

Adirondack Research Forum

Podium talk - presented postdoctoral work exploring community shifts using environmental DNA (eDNA), as well as brook trout genomics.

AUG 10.

"Unintentional artificial selection during ex-situ breeding of corals"

2023

Ecological Society of America Annual Meeting

Podium talk - presented work with NOAA and SECORE to understand the impacts of ex-situ culture on the genetic diversity of sexually-derived coral recruits intended for restoration outplanting.

MAY 26,

"Unintentional artificial selection during ex-situ breeding of corals"

2023

Huck Life Sciences Symposium, Penn State

Lightning talk - presented work with NOAA and SECORE to understand the impacts of ex-situ culture on the genetic diversity of sexually-derived coral recruits intended for restoration outplanting.

MAY 9.

"Patterns of Genetic Relatedness in Corals Across Small-Scale Seascapes"

GSAS Master's SynThesis Competition, Columbia University 2019

Speedtalk - Finalist in competition.

MAY 7.

"Phylogeography of two Hawaiian corals, Montipora capitata and Porites compressa"

E3B Master's Thesis Poster Symposium, Columbia University 2019

Poster session - presented results of thesis research.

APRIL 5.

2019

"Genetic structuring of two Hawaiian corals, Montipora capitata and Porites compressa"

The Earth Institute Student Research Showcase, Columbia University

Poster session - presented results of thesis research.

OCTOBER 24.

2018

"Genetic structuring of two Hawaiian corals, Montipora capitata and Porites compressa" **Student Conference on Conservation Science**, American Museum of Natural History Poster session - presented preliminary results of thesis research.

SUMMER 2023 | Dr. John Randall Shuman Troxell Memorial Scholarship in Biology

Penn State University

Nomination-based supplemental funding for outstanding graduate students in the Biology PhD program.

FALL 2021 - | Computation, Bioinformatics, and Statistics NIH Training Grant

SUMMER 2023

Penn State University

♦ NIH T32-funded grant that supports and trains PhD candidates involved in computational, bioinformatics, and statistical research.

SPRING 2021

University Graduate Fellowship

Eberly College of Science, Penn State University

♦ Prestigious fellowship program awarded to outstanding incoming graduate students.

FALL 2020 | Stephen B. Brumbach Distinguished Graduate Fellowship II

Eberly College of Science. Penn State University

Prestigious fellowship program awarded to outstanding incoming graduate students.

FALL 2020 | Braddock Award

Eberly College of Science, Penn State University

♦ Award for exemplary incoming students to the Eberly College of Science.

FALL 2018 | Graduate Student Research Award

Society of Systematic Biologists

Received funding from the Graduate Student Research Award program, a competitive research award for graduate students conducting research in systematics.

SPRING 2018 | The Earth Institute Travel Grant

The Earth Institute, Columbia University

Received funding from the Earth Institute Travel Grant Program, a competitive research award. Funding received for master's thesis on coral population genetics.

SPRING 2018

GSAS Thesis Research Matching Award

Graduate School of Arts and Sciences, Columbia University

> A competitive research award that provides MA students in the Graduate School of Arts and Sciences with matching funds.

SPRING 2018

Ecology, Evolution & Environmental Biology (E3B) Departmental Grant

Columbia University

♦ A competitive research grant providing funding for graduate students in the E3B department. Funding received for master's thesis work on coral population genetics.

FALL 2017

Swiss Benevolent Society General Scholarship

Swiss Benevolent Society of San Francisco

♦ A merit and need-based scholarship aimed at helping Northern Californians of Swiss descent obtain a higher education in any field of study.

FALL 2014 | CNR Dean's List

University of California, Berkeley

♦ Awarded to the top 4% of undergraduates in terms of GPA each semester.

TEACHING EXPERIENCE

SPRING 2023 UNIVERSITY

The Pennsylvania State University

Teaching Assistant

Park, PA ♦ Course: BIOL 230W: Molecules and Cells

> ♦ Taught two, 30-student weekly lab sections focusing on introductions to molecular and microscopy work.

AND 2019 New York, NY

Spring 2018 | Columbia University

Teaching Assistant

♦ Course: First Year Seminar in Ecology, Evolution and Environmental Biology

Graded undergraduate student assignments and organized weekly faculty speakers

COMMUNITY INVOLVEMENT

Peer Reviewer

Reviewer for the following journals:

♦ PCI Genomics

♦ Conservation Genetics

OTHER EMPLOYMENT

FEB. 2016 -

National Geographic Magazine

MAR. 2017 Various Photography Assistant

Assisted photographer in equipment setup and camera trapping

Assisted researchers in mist-netting and bat GPS tagging

AUG. 2012 -JUN. 2015 Berkeley, CA

University of California Botanical Garden

Horticulture and Propagation Assistant

Assisted in the propagation and growth of plants for fundraising sales

Educated visitors about collections and assisted clients with plant care inquiries

SKILLS

FIELD SKILLS

- ♦ Underwater Science
 - ▶ AAUS Scientific Diver
 - ▶ Coring and fragmentation of branching and massive/mounding coral colonies
- ♦ Stony and soft coral husbandry
- Short-term acute temperature stress experiments (Coral Bleaching Automated Stress) System, CBASS)

MOLECULAR | \diamond Preparation of shotgun, Pool-seq, and Hi-C Illumina libraries

BIOLOGY

- ♦ Standard and high molecular weight DNA extractions
- ♦ RNA extraction and library preparation SKILLS
 - ♦ Microsatellite amplification

COMPUTATION SKILLS

- - ▷ Illumina DNA: shotgun WGS, mate-pair, Hi-C, metabarcoding, and ddRAD
 - ▶ Illumina RNA: poly-A selected and total RNAseq
 - ▷ Oxford Nanopore (PromethION)
 - ▶ PacBio HiFi and CLR
 - > Affymetrix Microarray Data
 - ▶ Applied Biosystems 3730XL Microsatellite
 - Experience with the following tools and pipelines:
 - ▶ DNA Mapping: bwa, bowtie2, and minimap2
 - ▶ RNA Mapping: STAR and bowtie2
 - ⊳ SAM/BAM Processing: samtools, sambamba, and samblaster
 - ▷ Variant Calling: Freebayes, bcftools, and GATK4
 - ▶ Gene Expression Analyses: htseq, DESeq2, and limma-voom
 - ▶ Metagenome and Genome Assembly: Flye, hifiasm, Metaplatanus, metaFlye, hifiasm-meta, and MEGAHIT
 - ▶ Genome Annotation: funannotate
 - ⊳ GWAS: GEMMA, PLINK 2, and permGWAS
 - ▶ Polygenic Risk Scores: PRSice-2
 - ♦ Familiarity with bash, python, and R
 - ▶ Experience with supervised machine learning in scikit-learn