JONATHAN B. PURITZ JR.

Department of Biological Sciences
University of Rhode Island
120 Flagg Road, Kingston, RI 02881
Email: jpuritz@uri.edu
Website: MarineEvoEco.com

EDUCATION

2011 Ph.D. in Zoology, University of Hawaii at Manoa

Committee: Rob Toonen, Brian Bowen, Steve Karl, Megan Donahue, Shannon Bennett

2005 B.S. in Biology with Honors, Brown University

Advisors: Jon Witman and David Rand

PROFESSIONAL APPOINTMENTS

| PHOI LOS | SIONAL AFFOINTIVILITY |
|----------|--|
| 2023- | Associate Professor Department of Biological Sciences, University of Rhode Island |
| 2017-23 | Assistant Professor Department of Biological Sciences, University of Rhode Island |
| 2016-17 | Postdoctoral Research Associate Marine Science Center, Northeastern University |
| 2014-16 | Postdoctoral Research Associate, Adjunct Graduate Faculty Harte Research Institute, Texas A&M Corpus Christi |
| 2012-14 | Postdoctoral Research Associate Department of Wildlife and Fisheries Science, Texas A&M University |
| 2012 | Postdoctoral Scholar, Lecturer Hawaii Institute of Marine Biology, University of Hawaii at Manoa |

PUBLICATIONS

Statistics (Google Scholar Accessed 05/16/2024)

2721 Citations

38 Peer reviewed journal articles and book chapters

21 h-index

Pre-prints and manuscripts under review

Torres, W.J., Holstein, D.M., Putnam, H.M., Edmunds, P.J., **Puritz, J.B.**, Toonen, R.J., Hench, J.L. Post-disturbance recovery dynamics of connected coral subpopulations. **Under Review at** *Theoretical Ecology*.

Puritz, J.B., Zhao, H., Guo, X., Hare, M.P., He, Y., LaPeyre, J. Lotterhos, K.E., Lundgren, K. M., Proestou, D., Rawson, P., Fernandez Roboledo, J. A., Wedop, B., Witkop, E., Gomez-Chiarri, M. Nucleotide and structural polymorphisms of the eastern oyster genome paint a mosaic of divergence, selection, and human impacts. *BioRxiv*. DOI: https://doi.org/10.1101/2022.08.29.505629

Hart, M. W. Guerra, V., Byrne, M., **Puritz, J.B.** Genomic data improve coalescent inference across a range of demographic parameters and life-histories. **Under Review at** *Evolutionary Applications*. Preprint DOI: 10.22541/au.159990335.58331776

Published and In Press

- Puritz, J.B., Guo, X., Hare, M.P., He, Y., Hillier, L., Lotterhos, K.E., Liu, M., Minx, P., Proestou, D., Rice, E., Tolison, C., Warren, W., Zhao, H., Gomez-Chiarri, M. A second unveiling: haplotig masking of the eastern oyster genome improves population level inference. *Molecular Ecology Resources*. DOI:10.1111/1755-0998.13801
- Bogan, S.N., Johns, J., Griffiths, J.S., Davenport, D., Smith, S.J., Schaal, S.M., Downey-Wall, A., Lou, R.N., Lotterhos, K., Guidry, M.E., Rivera, H.E., McGirr, J.A., **Puritz J.B.**, Roberts, S.B., Silliman, K. A dynamic web resource for robust and reproducible genomics in non-model species: marineomics.io. *Methods in Ecology and Evolution*. DOI: 10.1111/2041-210X.14219
- Klanten, O.S., Gall, M., Barbosa, S.S., Hart, M.W., Keever, C.C., **Puritz, J.B.**, Haratino, J., Toonen, R.J., Selvakumaraswamy, P., Grosberg, R.K., Byrne, M. Population connectivity across east Australia's Bioregions and larval duration of the range extending sea star *Meridiastra calcar.* **Aquatic Conservation: Marine and Freshwater Ecosystems.** DOI: 10.1002/aqc.3973
- Guo, X., **Puritz, J. B.**, Wang, Z., Proestou, D., Allen, Jr., S., Small, J., Verbyla, K., Zhao, H., Haggard, J., Chriss, N., Zeng, D., Lundgren, K., Allam, B., Bushek, D., Gomez-Chiarri, M., Hare, M., Hollenbeck, C., LaPeyre, J., Liu, M., Lotterhos, K.E., Plough, L., Rawson, P., Saillant, E., Rikard, S., Varney, R., Wikfors, G., Wilbur, A. Development and evaluation of high-density SNP arrays for the eastern oyster *Crassostrea virginica*. *Marine Biotechnology*. DOI: 10.1007/s10126-022-10191-3
- Schiebelhut, L.M., Giakoumis, M., Castilho, R., Duffin, P.J., **Puritz, J.B.**, Wares, J.P., Wessel, G.M. and Dawson, M.N., 2022. Minor Genetic Consequences of a Major Mass Mortality: Short-Term Effects in Pisaster ochraceus. *The Biological Bulletin* DOI: 10.1086/722284
- Willis, S.C., Hollenbeck, C.M., **Puritz, J.B.** and Portnoy, D.S. Genetic recruitment patterns are patchy and spatiotemporally unpredictable in a deep-water snapper (Lutjanus vivanus) sampled in fished and protected areas of western Puerto Rico. *Conservation Genetics*. DOI: 10.1007/s10592-021-01426-2
- Portnoy, D.S., Fields, A.T., **Puritz, J.B.**, Hollenbeck, C.M., Patterson, W.F. Genomic Analysis of Red Snapper, Lutjanus campechanus, Population Structure in the U.S. Atlantic and Gulf of Mexico. *ICES Journal of Marine Science*. DOI: 10.1093/icesjms/fsab239
- Trigg, S.A., Venkataraman, Y.R., Gavery, M., Roberts, S.B., Bhattacharya, D., Downey-Wall, A., Eirin Lopez, J.M., Johnson, K.M., Lotterhos, K.E., **Puritz, J.B.**, Putnam, H.M. Invertebrate methylomes provide insight into mechanisms of environmental tolerance and reveal methodological biases. *Molecular Ecology Resources.* DOI: 10.1111/1755-0998.13542
- Modak, T. H., Literman, R., **Puritz, J.B.**, Johnson, K. J., Roberts, E. M., Proestou, D., Guo, X., Gomez-Chiarri, M., Schwartz, R. S. Extensive genome-wide copy number variation in the eastern oyster (*Crassostrea virginica*). *Phil. Trans. R. Soc. B.* DOI: 10.1098/rstb.2020.0164
- Catalano, K. A., Dedrick, A. G., Stuart, M. R., **Puritz, J. B.,** Montes Jr., H. R., Pinsky, M. L. Quantifying dispersal variability among nearshore marine populations. *Molecular Ecology.* DOI: 10.1111/mec.15732
- 2020 Hart, M.W. and **Puritz, J.B**. Correction to 'Extraordinarily rapid life-history divergence between Cryptasterina sea star species'. *Proceedings of the Royal Society B*, *287*(1930), p.20201325. DOI: 10.1098/rspb.2020.1325
- O'Leary, S.J., **Puritz, J. B.**, Willis, S.C., Hollenbeck, C.M. and Portnoy, D.S., 2018. These aren't the loci you're looking for: Principles of effective SNP filtering for molecular ecologists. *Molecular Ecology*. DOI: 10.1111/mec.14792

 A top downloaded paper of 2019

- Schiebelhut, L.M., **Puritz, J. B.**, and Dawson, M.N. Decimation by sea star wasting disease and rapid genetic change in a keystone species, Pisaster ochraceus. **Proceedings of the National Academy of Sciences**. DOI: 10.1073/pnas.1800285115
- Puritz, J. B., and Lotterhos, K. E. Expressed Exome Capture Sequencing (EecSeq): a method for cost-effective exome sequencing for all organisms with or without genomic resources.

 *Molecular Ecology Resources**. DOI: 10.1111/1755-0998.12905
- Puritz, J. B., Keever, C. C., Addison, J. A., Barbosa, S. S., Byrne, M., Hart, M. W., Grosberg, R. K., Toonen, R. J. Life history predicts past and present population connectivity in two sympatric sea stars. *Ecology and Evolution*. DOI: 10.1002/ece3.2938
- Willis, S. C., Hollenbeck, C. M., **Puritz, J.B.**, Gold, J. R., Portnoy, D. S. Haplotyping RAD loci as an efficient method to characterize and filter paralogs and manage linkage disequilibrium. *Molecular Ecology Resources*. DOI: 10.1111/1755-0998.12647
- Puritz, J. B., Portnoy, D. S., Gold, J. R. Fine-scale partitioning of genomic variation among recruits in an exploited fishery: causes and consequences. *Scientific Reports*. DOI: 10.1038/srep36095
- Selkoe, K.A., D'Aloia, C. C., Crandall, E. D., Iacchei, M. I., Liggins, L., **Puritz, J. B.**, von der Heyden, S., Toonen, R. J. A decade of seascape genetics: contributions to basic and applied marine connectivity. *Marine Ecology Progress Series*. DOI:10.3354/meps11792

 Feature Article
- Portnoy, D. S., **Puritz, J. B.**, Hollenbeck, C. M., Gelsleichter, J., Chapman, D., Gold, J. R. Selection and sex-biased dispersal in a coastal shark: the influence of philopatry on adaptive variation. *Molecular Ecology*. DOI: 10.1111/mec.13441
- Gold, J. R., Willis, S. C., Renshaw, M. A., Buentello, A., Walker, Jr., H. J., **Puritz, J. B.,**Hollenbeck, C. M., Voelker G. Phylogenetic relationships of tropical eastern Pacific snappers
 (Lutjanidae) inferred from mtDNA sequences. *Systematics and Biodiversity*.
 DOI:10.1080/14772000.2015.1078857
- Puritz, J.B., Matz, M. V., Toonen, R. J., Weber, J. N., Bolnick, D. I., Bird, C. E. Comment: Demystifying the RAD fad. *Molecular Ecology*. DOI: 10.1111/mec.12965

 2nd Most Accessed Paper for 2014-2015
- 2014 **Puritz, J. B.,** Hollenbeck, C. M., Gold, J. R. *dDocent*: a RADseq, variant-calling pipeline designed for population genomics of non-model organisms. *PeerJ*. DOI: 10.7717/peerj.431
- Puritz, J. B., Renshaw, M. A., Abrego, D., Vega, R. R., Gold, J. R. Reproductive variance of brood dams and sires used in restoration enhancement of spotted seatrout (*Cynoscion nebulosus*) in Texas bays and estuaries. *North American Journal of Aquaculture* DOI: 10.1080/15222055.2014.920751
- Toonen, R. J., **Puritz, J. B.,** Forsman, Z.H., Whitney, J. L., Fernandez-Silva, I., Andrews, K. A., Bird, C. E. ezRAD: a simplified method for genomic genotyping in non-model Organisms. *PeerJ.* DOI:10.7717/peerj.203
- 2013 Keever, C., **Puritz, J. B.,** Addison, J. A., Byrne, M., Grosberg, R. K., Toonen R. J., Hart, M.W. Shallow gene pools in the high intertidal: Extreme loss of genetic diversity in viviparous sea stars (Parvulastra). *Biology Letters.* DOI:10.1098/rsbl.2013.0551
- Andrews, K. R., Perrin, W. F., Oremus, M. Karczmarski, L., Bowen, B. W., **Puritz, J. B.**, Toonen, R. J. The evolving male: spinner dolphin (*Stenella longirostris*) ecotypes are divergent at Y chromosome but not mtDNA or autosomal markers. *Molecular Ecology*. 10.1111/mec.12193
- Barbosa, S., Klaten, S., **Puritz, J. B.,** Toonen, R. J., Byrne, M. Very fine scale population genetic structure of sympatric asterinid sea stars with benthic and pelagic larvae: influence of

mating system and dispersal potential. **Biological Journal of the Linnean Society**. DOI: 10.1111/bij.12006

- 2013 Puritz, J. B. and Toonen, R. J. Next-generation sequencing for high-throughput molecular ecology: a step-by-step protocol for targeted multilocus genotyping by pyrosequencing. Chapter in Methods in Molecular Biology: Microsatellites. Volume Editor: Kantartzi, S. K. Pages 89-99
- Puritz, J. B., Keever, C. C., Addison, J. A., Byrne, M. Hart, M. W. Grosberg, R. K., Toonen, R. 2012 J. Extraordinarily rapid life history divergence between Cryptasterina sea star species. Proc. R. Soc. B. DOI: 10.1098/rspb.2012.1343
- Puritz, J. B., Addison, J. A., Toonen, R. J. Next-Generation Phylogeography: the application of 2012 targeted next-generation sequencing of non-model organisms. PLoS One DOI: 10.1371/journal.pone.0034241
- 2011 Puritz, J. B. and Toonen, R. J. Coastal Pollution Limits Pelagic Larval Dispersal. *Nature* Communications. DOI: 10.1038/ncomms1238
- 2011 Toonen, R. J., Andrews, K. R., Baums, I. B., Bird, C. E., Concepcion, G. T., Daly-Engel, T. S., J. Eble, A. Faucci, M. Gaither, M. lacchei, J. Puritz, J. Schultz, D. Skillings, M. Timmers & B. Bowen. Defining boundaries for ecosystem-based management: A multispecies case study of marine connectivity across the Hawaiian Archipelago. Journal of Marine Biology DOI: 10.1155/2011/460173
- 2009 Keever, C. C., Sunday, J., Puritz, J. B., Addison, J. A., Toonen, R. J., Grosberg, R. K., Hart, M. W. Discordant distributions of populations and genetic variation in a sea star with high dispersal potential. *Evolution*. DOI: 10.1111/j.1558-5646.2009.00801.x
- 2008 Haney, R. A., Dionne, M., Puritz, J. B., Rand, D. M. The comparative phylogeography of east coast estuarine fishes in formerly glaciated sites: Persistence versus Recolonization in Cyprinodon variegatus ovinus and Fundulus heteroclitus macrolepidotus. Journal of Heredity DOI: 10.1093/jhered/esn107
- Walker, A. N., Bush, P., Puritz, J. B., Wilson, T., Chang, E. S., Miller, T., Holloway, K., 2005 Horst, M. N. Bioaccumulation and metabolic effects of the endocrine disruptor methoprene in the lobster, Homarus americanus. Integrative and Comparative Biology. DOI: 10.1093/icb/45.1.118

GRANTS AWARDED

Total funding = \$2,618,780; Current funding= \$2,328,938

Current Funding

1) 2024 Rhode Island Sea Grant \$247,475

The interaction of environment, genotype, and disease on juvenile oyster survival (PI Puritz; Total to Puritz: \$247,475; additional \$123,734 in match Project Total: \$371,209)

2) 2021 **USDA- Agriculture Research Service**

\$250.000

Population genomic evaluation of wild and cultured Eastern oyster populations from the Northeast region (Pls Puritz and Proestou; Total to Puritz: 250,000; additional \$50,000 in match Project Total: \$300,000)

2021 National Science Foundation 3)

\$680,898

How do multiple coastal stressors structure the genomic diversity of marine populations?

(PI Puritz; Total to Puritz: \$680,898; Project Total: \$680,898)

| 4) | 2020 | National Science Foundation EAGER: Development of a tool to rapidly and cost-effectively sequence the exome of any organism (PI Puritz; Total to Puritz: \$298,978; Project Total: \$298,978) | |
|--------|--------|--|-----------|
| 5) | 2019 | US Fish and Wildlife Service Assessing Horseshoe Crab Population Structure within Southern New England (Pls: McManus, Ameral, and Puritz; Total to Puritz \$80,435; additional match: \$43,302 (URI) & \$13,992 DEM; Project Total: \$163,715) | \$80,435 |
| 6) | 2019 | Atlantic Coast Marine Fisheries Council From Sequence to Consequence: Genomic Selection to Expand And improve Selective Breeding for The Eastern Oyster (Eastern Oyster Breeding Consortium; URI Pls: Puritz and Gomez-Chiarri; Total to URI: \$690,000; Project Total: \$4,363,092) | \$690,000 |
| Past I | Fundin | g | |
| 2019 | | Rhode Island Sea Grant How do multiple coastal stressors limit oyster recruitment? (PI Puritz; Total to Puritz: \$249,192; additional \$124,617 in match Project Total: \$373,809) | \$249,192 |
| 2016 | | Harte Research Institute Staff Innovations Grant (Declined) Collaborative proposal to investigate evolutionary impacts of barotrauma in red snapper | \$5,000 |
| 2015 | | Texas Research Development Fund Program Impacts of sewage effluent on genomic diversity and connectivity of marine intertidal communities | \$20,000 |
| 2011 | | NSF East Asian Pacific Science Institute Fellowship Fellowship for two months of research at the University of Sydney | \$8,000 |
| 2011 | | University of Hawaii Graduate Student Organization Travel Grant for 2011 Annual Meeting of the Western Society of Naturalists | \$750 |
| 2010 | | University of Hawaii Graduate Student Organization Travel Grant for 2010 Benthic Ecology Meeting | \$900 |
| 2009 | | University of Hawaii Graduate Student Organization Research grant for sample collection trip to California | \$750 |
| 2008 | | The Charles H. and Margaret B. Edmondson Research Fund Research on the population connectivity of <i>Linckia multifora</i> | \$1,500 |
| 2007 | | Ecology, Evolution, and Conservation Biology Travel Grant Travel Grant for 2007 Annual Meeting of the Western Society of Naturalists | \$750 |
| 2007 | | University of Hawaii Arts and Sciences Advisory Council Award Research expedition to the Chagos Archipelago | \$3,000 |

TEACHING EXPERIENCE

| Classes | |
|---------|--|
| 2024 | Instructor, University of Rhode Island BIO 594: Conservation and Population Genomics: theory and practice |
| 2024 | Instructor, University of Rhode Island BES 582: Biological and Environmental Sciences Colloquium |
| 2023 | Instructor, University of Rhode Island BES 581: Biological and Environmental Sciences Colloquium |
| 2023 | Instructor, University of Rhode Island BIO 354: Invertebrate Zoology |
| 2023 | Instructor, University of Rhode Island BIO 594: Conservation and Population Genomics: theory and practice |
| 2022 | Instructor, University of Rhode Island BIO 354: Invertebrate Zoology |
| 2022 | Instructor, University of Rhode Island BIO 594: Using genomic techniques to examine the evolution of populations |
| 2021 | Instructor, University of Rhode Island BIO 354: Invertebrate Zoology |
| 2021 | Instructor, University of Rhode Island BIO 425G: The origins and fate of marine biodiversity: a larval perspective |
| 2020 | Instructor, University of Rhode Island BIO 354: Invertebrate Zoology |
| 2019 | Instructor, University of Rhode Island BIO 425G: The origins and fate of marine biodiversity: a larval perspective |
| 2019 | Instructor, University of Rhode Island BIO 130: Topic in Marine Biology |
| 2019 | Instructor, University of Rhode Island BIO 354: Invertebrate Zoology |
| 2019 | Instructor, University of Rhode Island BIO 594: Using genomic techniques to examine the evolution of populations |
| 2018 | Instructor, University of Rhode Island BIO 354: Invertebrate Zoology Includes rebuilding curriculum for both lecture and lab |
| 2018 | Instructor, University of Rhode Island BIO 594: Using genomic techniques to examine the evolution of populations New Graduate Class offered for the first time |
| 2017 | Co-Instructor, University of Rhode Island BIO 354: Invertebrate Zoology |
| 2017 | Guest Lecturer, University of Rhode Island BIO 360: Marine Biology (1 class, Fall semester) |
| 2012 | Lecturer, University of Hawaii at Manoa Instructor of record for Biology 404: Advanced Topics in Marine Biology |

Capstone class for seniors majoring in Marine Biology

Workshops

2015-18 Invited Faculty, Winter School, ETH Zürich, Switzerland

Bioinformatics for Adaptation Genomics: Adaptation genomics in the realm of Next-Generation Sequencing data analysis

- Funded by Adaptation to a Changing Environment initiative, ETH Zürich, Switzerland
- Responsible for whole day workshop on "Extracting SNP data from NGS sequencing"

2015 Invited Faculty, Nha Trang University, Vietnam

Training workshop for Next Generation Sequencing

- Funded by the project "Building a Mekong River genetic biodiversity research network" in the PEER program funded by USAID in the process of project implementation, NTU partnered with Old Dominion, and Texas A & M University Corpus Christi.
- Responsible for multiple day workshop on RADseq Bioinformatics

2014 Invited Faculty, De LaSalle University, Manila, Philippines

Pacific Advanced Science Institute (PacASI) sponsored workshop: "Introduction to Genome Data Analysis: Assembly, Annotation, and Application."

- PacASI is a partnership between the Center for Natural Science and Ecological Research at De La Salle University and the National Science Foundation Partnerships for Enhanced Engagement in Research.
- Responsible for several lectures and hands on workshops designed for international students' first engagement with genome scale data.

2013 Invited Faculty, Hawaii Institute of Marine Biology

29th Annual Edwin W. Pauley Summer Program: "Advancing tools for biodiversity studies: Genomics and bioinformatics of cnidarians with a focus on corals"

 Responsible for developing and overseeing student research projects and for a workshop on RAD Sequencing

INVITED PRESENTATIONS

diversity

| 2022 | Coastal pollution, next-generation sequencing, and the evolution of marine populations Department of Biology- Miami University |
|------|---|
| 2018 | Coastal pollution, next-generation sequencing, and the evolution of marine populations Department of Marine Sciences- University of Connecticut Avery Point |
| 2018 | Coastal pollution, next-generation sequencing, and the evolution of marine populations Graduate School of Oceanography- University of Rhode Island |
| 2018 | Coastal pollution, next-generation sequencing, and the evolution of marine populations Ecology and Evolutionary Biology Department- Brown University |
| 2017 | Expressed Exome Capture Sequencing: a method for cost-effective exome sequencing for all organisms Cellular and Molecular Biology Department- University of Rhode Island |
| 2017 | Coastal pollution, next-generation sequencing, and the evolution of marine populations Biology Department- Woods Hole Oceanographic Institution |
| 2016 | Harnessing the power of next-generation sequencing to examine the evolution of marine populations Department of Biological Sciences- University of Rhode Island |
| 2015 | Using next-generation sequencing to examine patterns of coastal genomic |

| | Department of Biology Seminar- University of Louisiana at Lafayette |
|--------------|--|
| 2015 | Using next-generation sequencing to examine patterns of coastal genomic |
| 2013 | diversity |
| | Marine Science Center Seminar- Northeastern University |
| 2015 | The Seascape Genetics of Coastal Pollution Waterscape Genetics- New Perspectives on Connectivity in Fluid Environments Symposium at the International Association of Landscape Ecology World Congress |
| 2013 | Impacts of Coastal Pollution and Life-History on Marine Population Connectivity Biological Sciences Seminar- Old Dominion University |
| 2012 | The Impacts of Coastal Pollution on Marine Population Connectivity Harte Research Institute Seminar Series- Texas A&M Corpus Christi |
| 2012 | The Impacts of Coastal Pollution on Marine Population Connectivity Wildlife and Fisheries Sciences Brown Bag Seminar- Texas A&M University |
| 2009 | "The Frontiers of Conservation Genetics: From Genes to Genomes" Hanauma Bay Education Program Lecture Series- Hanauma Bay Nature Preserve |
| CONFER | ENCE PRESENTATIONS (LAST 7 YEARS) |
| ^undergradua | ate mentee, # post-baccalaureate mentee, * graduate student mentee |
| 2024 | Acute and long-term responses of juvenile oysters to dynamic multiple coastal stressors |
| | Guidry*, M.E., Randall*, C., Puritz, J.B. National Shellfishery Association Meeting. Charlotte, NC. |
| 2024 | Assessing the evolutionary response of the eastern oyster to exposure to coastal acidification and sewage effluent: a CASE study |
| | Puritz, J. B., Zyck*, A., Harvey, J. A., Lotterhos K.E. National Shellfishery Association Meeting. Charlotte, NC. |
| 2023 | The effects of probe and DNA insert length on the <i>de novo</i> assembly and capture efficiency of expressed exome capture sequencing (EecSeq) Green*, J. M., and Puritz J. B. National Shellfishery Association Meeting. Baltimore, MD. |
| 2023 | Investigating the effects of coastal stressors on the genomic variation of oyster |
| | populations in Narragansett Bay Zyck*, A., Stevick, R., Gomez-Chiarri, M., Puritz J. B. National Shellfishery Association Meeting. Baltimore, MD. |
| 2023 | Nucleotide and structural polymorphisms of the eastern oyster genome paint a |
| | mosaic of divergence, selection, and human impacts Puritz, J.B., Zhao, H., Guo, X., Hare, M.P., He, Y., LaPeyre, J. Lotterhos, K.E., Lundgren, K. M., Proestou, D., Rawson, P., Fernandez Roboledo, J. A., Wedop, B., Witkop, E., Gomez-Chiarri, M. National Shellfishery Association Meeting. Baltimore, MD. |
| 2022 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., Harvey, J. A., Lotterhos K.E. Benthic Ecology Meeting. Portsmouth, NH. |
| 2021 | Unlocking the Exome: exploring de novo assembly for capture sequencing Green*, J. M., and Puritz J. B. National Shellfishery Association Meeting. |
| 2021 | From Sequence to Consequence: genomic selection to expand and improve selective breeding for the eastern oyster Guo, X., Allen Jr., S., Proestou, D., Allam, B., Gomez-Chiarri, M., Hare, M., Liu, M., Lotterhos, K.E., Kube, P., Plough, L., Puritz, J.B., et al. National Shellfishery Association Meeting. Virtual. |

| 2021 | Investigating the effects of coastal stressors on the genomic variation of oyster populations in Narragansett Bay Zyck*, A., Stevick, R., Gallagher^#, A., Padro^, N., Gomez-Chiarri, M., Puritz J. B. National Shellfishery Association Meeting. Virtual. |
|------|---|
| 2021 | An assembled genome reference for the eastern oyster: a resource for discovery and innovation Puritz, J.B., Zhao, H., Weedop, B., Modak, T., Roberts, E., Allen Jr., S., Hare, M., Lotterhos, K.E., Rawson, P., Schwartz, R., Proestou, D., Guo, X., Warren, W., Gomez-Chairri, M. National Shellfishery Association Meeting. Virtual. |
| 2021 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., Harvey, J. A., and Lotterhos K. E. National Shellfishery Association Meeting. Virtual |
| 2020 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., Harvey, J. A., and Lotterhos K. E. Western Society of Naturalists. Virtual. |
| 2020 | Unlocking the Exome: exploring de novo assembly for capture sequencing Green*, J. M., and Puritz J. B. Western Society of Naturalists. Virtual. |
| 2020 | Investigating the effects of coastal stressors on the genomic variation of oyster populations in Narragansett Bay Zyck*, A., Stevick, R., Gallagher^#, A., Padro^, N., Gomez-Chiarri, M., Puritz J. B. Western Society of Naturalists. Virtual. |
| 2020 | Unlocking the Exome: exploring de novo assembly for capture sequencing Green*, J. M., and Puritz J. B. National Shellfishery Association Meeting. Baltimore, MD. Cancelled due to Covid-19 |
| 2020 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., Harvey, J. A., and Lotterhos K. E. National Shellfishery Association Meeting. Baltimore, MD. Cancelled due to Covid-19 |
| 2019 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., Harvey, J. A., and Lotterhos K. E. Evolution. Providence, RI. |
| 2019 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., and Lotterhos K. E. Aquaculture 2019. New Orleans, LA. |
| 2018 | Expressed Exome Capture Sequencing: a method for cost-effective exome sequencing for all organisms Puritz, J. B., and Lotterhos K. E. Global Invertebrate Genome Alliance. Curacao. |
| 2018 | Expressed Exome Capture Sequencing (EecSeq): a method for cost-effective exome sequencing of non-model organisms Puritz, J. B., and Lotterhos K.E. National Shellfishery Association Meeting. Seattle, WA. |
| 2017 | Expressed Exome Capture Sequencing: a method for cost-effective exome sequencing for all organisms Puritz, J. B., and Lotterhos K.E. Western Society of Naturalists. Pasadena, CA. |
| 2017 | Expressed Exome Capture Sequencing: a method for cost-effective exome sequencing for all organisms Puritz, J. B., and Lotterhos K.E. Evolution. Portland, OR. |

2017 Expressed Exome Capture Sequencing (EecSeq): a method for cost-effective exome sequencing of non-model organisms

Puritz, J. B., and Lotterhos K.E. National Shellfishery Association Meeting. Knoxville, TN.

CONFERENCE POSTERS (LAST 7 YEARS)

^undergraduate mentee. # post-baccalaureate mentee. * graduate student mentee 2024 What are the impacts of coastal stressors on *Crassostrea virginica* on growth during early life stages? Westbrook[^], C., Zyck^{*}, A., Puritz, J.B. National Shellfishery Association Meeting. Charlotte, NC. 2024 The phenotypic effects of diel-cycling of dissolved oxygen and pH across populations of eastern ovster larvae Randall*, C., Wildes^, E., Guidry*, M.E., Puritz, J.B. National Shellfishery Association Meeting. Charlotte, NC. Understanding how coastal stressors affect selectively bred eastern oysters 2024 (Crassostrea virginica) Wildes', E., Bucci', J., Randall#, C., Guidry*, M.E., Puritz, J.B. National Shellfishery Association Meeting. Charlotte, NC. Exploring genetic diversity and population differentiation of horseshoe 2024 crabs in southern New England through ddRAD sequencing Santiago[^], H., Green^{*}, J.M., Barret^{*}, G., Ameral^{*}, N., Puritz, J.B. SACNAS National Diversity in STEM Conference. Portland, OR. 2023 The effects of diel-cycling acidification and hypoxia across multiple developmental stages of the eastern oyster (crassostrea virginica) Zyck*, A., Puritz J. B. National Shellfishery Association Meeting. Baltimore, MD. 2023 Diel-cycling hypoxia and Acidification increases susceptibility to Roseovarious Oyster Disease in *Crassostrea virginica* Kulesh[^], K., Zyck^{*}, A., Puritz J. B., Gomez-Chiarri, M. National Shellfishery Association Meeting.

Baltimore, MD.

Multiple coastal stressors induce differential RNA expression in larval oysters
Guidry* M.E. Harvey J. A. Lotterhos K.E. Schedl# M. Puritz J. B. National Shellfishery

Guidry*, M.E., Harvey, J. A., Lotterhos, K. E., Schedl*, M., Puritz, J. B. National Shellfishery Association Meeting. Baltimore, MD.

2022 Comparing extracted DNA/RNA concentration from oyster gill & mantle tissues utilized in EecSeq for exome surveying
Santiago^, H., Green*, J.M., Puritz, J.B. SACNAS National Diversity in STEM Conference.
Puerto Rico.

2021 Unlocking the Exome: exploring de novo assembly for capture sequencing
Green*, J. M., and Puritz J. B. AGA Presidential Symposium- Conservation Genomics: Current
Applications and Future Directions. Snowbird, UT.

The effect of environmental parameters on *Crassostrea virginica* shell sizes. Satkowski[^], S., Zyck^{*}, A., Puritz, J.B. National Shellfishery Association Meeting. Virtual.

Investigating the effects of coastal stressors on the genomic variation of oyster populations in Narragansett Bay

Zyck*, A., Stevick, R., Gallagher^*, A., Padro^, N., Gomez-Chiarri, M., Puritz J. B. National Shellfishery Association Meeting. Virtual.

2020 Investigating the effects of coastal stressors on the connectivity of oyster populations in Narragansett Bay
Zyck*, A., Gallagher*, A., Padro*, N., Puritz J. B. National Shellfishery Association Meeting.

Baltimore, MD.

Cancelled due to Covid-19

| | Cancelled due to Covid-19 |
|------|--|
| 2020 | CASE-ing out the transcriptomics of multiple coastal anthropogenic stressors on eastern oyster larvae Schedl*, M., Harvey, J. A., Lotterhos, K. E., Puritz, J. B. National Shellfishery Association Meeting. Baltimore, MD. Cancelled due to Covid-19 |
| 2020 | Understanding the effects of multiple stressors on oyster larvae Tarrant*, M., Zyck*, A., Schedl*, M., Puritz J. B. National Shellfishery Association Meeting. Baltimore, MD. Cancelled due to Covid-19 |
| 2019 | Assessing the evolutionary response of eastern oyster larvae to exposure to coastal acidification and sewage effluent: a CASE study Puritz, J. B., Harvey, J. A., Lotterhos K.E. Gordon Research Conference: Ecological and Evolutionary Genomics. |
| 2019 | Expressed Exome Capture Sequencing: a method for cost-effective exome sequencing for all organisms Puritz, J. B., and Lotterhos K.E. Evolution. Providence, RI. |
| 2019 | CASE-ing out the transcriptomics of multiple coastal stressors Schedl*, M., Harvey, J. A., Lotterhos, K. E., Puritz, J. B. Evolution. Providence, RI. |
| 2019 | Understanding the impacts of sewage effluent on the genomic diversity and population connectivity of the fiddler crab (Uca rapax) Zyck*, A. H., Dimens, P., Willis, S., Portnoy, D., Puritz J. B. Evolution. Providence, RI. |
| 2019 | An Analysis of Population Structure, Genetic Variation and Outlier SNPs in the Eastern Oyster, Crassostrea virginica Weedop, K. B., Freeman, K., Roberts, E., Proestou, D., Puritz, J. B., Gomez-Chiarri, M., and Lotterhos K. E. Evolution. Providence, RI. |
| 2019 | Unlocking the exome: exploring de novo assembly options for expressed exome capture sequencing Green*, J. M., and Puritz J. B. Evolution. Providence, RI. |
| 2018 | Unlocking the Exome: exploring de novo assembly options for capture sequencing Green*, J. M., and Puritz J. B. Western Society of Naturalists. |
| | |

MENTORING Undergraduate

| Ondorgradato | | | | |
|--------------|--|--|--|--|
| 2024- | Juliana Bucci, Cierra Westbrook, Chris Delano, India Greene | | | |
| 2023-2024 | Hector Santiago, Lauren Albanese, Cierra Westbrook, Erin Wildes, Juliana Bucci | | | |
| 2022-2023 | Hector Santiago, Lydia Cross, Ben Poepsel, Hailee Carlson, Halle Peterlin | | | |
| 2021-2022 | Anna Sorgie, Madeline Kistler, Joseph Maiorano, Finn Harty | | | |
| 2020-2021 | Nina Padro, Anna Sorgie, Seraphina Satkowski, Allison Gallagher, Nadia Moss | | | |
| 2019-2020 | Melati Tarrant, Allison Gallagher, Nina Padro | | | |
| 2018-2019 | Emma Ferrante, Kevin Dyer, Marygrace Trousdell | | | |
| 2017-2018 | Elliot Vosburgh, Kate Leiden | | | |
| | | | | |

Primary Graduate Advisor

| 2024- | Olivia Nieves | PhD. | Biological and Environmental Sciences |
|---------|-------------------|------|---------------------------------------|
| 2023- | Cassandra Cerasia | PhD. | Biological and Environmental Sciences |
| 2021-24 | Gabriel Barrett | MS. | Biological and Environmental Sciences |
| 2020- | Megan Guidry | PhD. | Biological and Environmental Sciences |

| 2019-22 | Natalie Ameral | MS. | Biological and Environmental Sciences |
|---------|----------------|------|---------------------------------------|
| 2019- | Jacob Green | PhD. | Biological and Environmental Sciences |
| 2018- | Amaelia Zyck | PhD. | Biological and Environmental Sciences |

Graduate Committees

| 2024- | Karis Kang, PhD. Biological and Environmental Sciences |
|---------|---|
| 2022- | Leann Biancani, PhD. Biological and Environmental Sciences |
| 2022- | Jill Ashey, PhD. Biological and Environmental Sciences |
| 2022- | Danielle Becker, PhD. Biological and Environmental Sciences |
| 2022- | Michelle Hauer, PhD. Graduate School of Oceanography |
| 2022-24 | Eren Ada, PhD. Biological and Environmental Sciences |
| 2020-21 | Tyler Devos, M.S. Biological and Environmental Sciences |
| 2019-24 | Benjamin Ha, PhD. Ecology and Evolutionary Biology (UCLA) |
| 2019-23 | lan Bishop, PhD. Graduate School of Oceanography |
| 2019-21 | Samuel Gurr, PhD. Biological and Environmental Sciences |
| 2018-21 | Erin Borbee, PhD. Biological and Environmental Sciences |
| 2019 | Martin Hellwig, PhD. Computer Science. |
| 2018-19 | Evelyn Takyi, MS. Biological and Environmental Sciences |
| 2014-15 | Patricia M. Cockett, MS. Biology (Texas A&M Corpus Christi) |

Postdoctoral scholars

2024- Dr. Coline Caillon 2022-23 Dr. Alexandra Hooks

PROFESSIONAL SERVICE

Participant

| 2022- | The Evolving Seas RCN: Training and Integration Workshop (Website) Participant |
|-----------|--|
| 2022-2023 | Building Eelgrass Resiliency along the Mid-Atlantic and Southern New England Coast Workshop Series (Website) Participant |
| 2021-2023 | The Evolving Seas RCN: Virtual Lab Meeting Training Program |
| 2021- | Biological and Environmental Science Graduate Program Leadership Committee Coordinator of the Evolution and Marine Biology Graduate Specialization |
| 2021 | MarineOmics Panel Seminar on GBS/RADseq genotyping pipelines (Video) Invited Panelist |
| 2021- | MarineOmics Working Group (Website) Faculty advisor |
| 2021- | GSO Science Saturday Lab had an interactive research experience in Blount Aquarium |
| 2020-2021 | The Evolving Seas RCN: 2020 Virtual Lab Meeting Training Program Host and mentor for PhD student Camila Mac Loughlin Aleman, Centro de Investigaciones Biológicas del Noroeste |
| 2020- | Synthesis and outlook for future improvements in understanding and application of the biology of sea star wasting disease Working Group (Website) |

2018-Functional Re-annotation of Oyster Genomes with Epigenetic Resources (FROGER) working group **Participant** Moorea Coral Reef LTER Connectivity working group 2018-**Participant** Equity, inclusion, diversity committee, Biological Sciences 2018-Member 2018-Committee to establish a graduate Data Science Program at URI Member **Eastern Oyster Genome Consortium working group** 2017-Participant, leading genome description manuscript 2014dDocent RADseq Bioinformatics pipeline Developer, software has over 85,000 downloads and website with 95,000 visitors 2019 Faculty host at URI Welcome Day Ad hoc committee for Cruickshank Lecture, Biological Sciences 2018-2019 Chair 2018 The power of RNA: Broad application of RNA-based sequencing for transcriptome and genome analysis Science Webinar- AAAS (LINK) 2017 Faculty host at URI Fall Open House (both events)

Peer Reviewer

Proceedings of the Royal Society B Ecology and Evolution **Evolution MEPS** Biological Invasions **JEMBE** Marine Biology Molecular Ecology Journal of Heredity **BioScience** PLoS One Conservation Genetics Conservation Genetics Resources Heredity Transactions of AFS Genetica Aquatic Biology PeerJ PLoS Genetics Axios Biology Letters Open Science Journal of Fish Biology Nature Protocols Molecular Ecology Resources (X6) Genome Biology and Evolution Scientific Reports Methods in Ecology and Evolution Hydrobiologia Diversity and Distribution G3: GeneslGenomeslGenetics New Zeal J Mar Fres NSF: Bio. Oce. Ecological Applications Washington Sea Grant NSF Louisiana Sea Grant Graduate Women in Science

AWARDS AND HONORS

| 2015 | Top 300 Reviewers of Molecular Ecology |
|---------|--|
| 2009,11 | Best Paper Honorable Mention, Albert L. Tester Memorial Symposium |
| 2005-06 | National Science Foundation Graduate Research Fellowship Honorable Mention |

PUBLIC OUTREACH

2017- Skype a Scientist

The Skype a Scientist matches scientists with classrooms around the world! Scientists will skype into the classroom for 30-60 minute Q and A sessions that can cover the scientist's expertise or what it's like to be a scientist

2016- Scientist Pen pal

Working with Letters to a Pre-Scientist (http://www.prescientist.org) to connect with middle school students who want to learn more about being a scientist

2007- **Underwater Photographer**

Photo chosen for "Featured Image" for *PeerJ* (Nov 2013)

Photo chosen for "Image of the Week" for *Nature Communications* (March 2011)

Images have appeared in several publications including NOAA public reports, the Division of Land and Natural Resources Hawaii outreach posters, and several HIMB and UC Davis press releases.

2016 Nahant Coastal BioBlitz

Sponsored by the Ocean Genome Legacy and Northeastern University Marine Science Center- Volunteer Photographer and Scientist

2016 BLUE On Tour Film Festival-Corpus Christi

Sponsored by the Harte Research Institute-Volunteer Social Media Coordinator

2010-11 Participant in the Pacific Symposium for Science and Sustainability

Judged and moderated a high school science competition

2007-11 Scientific Blogger

2007

Created a blog to document the experience of being an NSF EAPSI fellow in Australia. jbpaustralia2011.wordpress.com

Created a blog to document field research experience aboard the NOAA R/V Hi'ialakai, Northwestern Hawaiian Islands Research cruise in September of 2007.

https://sites.google.com/site/jpuritz/cruise

Google changed this service and formatting for this webpage is no longer correct

TRAINING AND CERTIFICATIONS

NOAA Advanced Coxswain

| 2014-16 | Texas A&M Corpus Christi Scientific Diver (AAUS Reciprocity) |
|---------|--|
| 2005-12 | University of Hawaii Scientific Diver (AAUS Reciprocity) |