# JONATHAN B. PURITZ JR.

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## 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

2017- **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 12/1/2021)**

**1858 Citations**
**29 Peer reviewed journal articles and book chapters**
**19 h-index**

***Pre-prints, accepted manuscripts, and manuscripts under review***

2021 Schiebelhut, L., Giakoumis, M., Castilho, R., Duffin, P.J., Puritz, J.B., Wares, J.P., Wessel, G.M., Dawson, M.N. No short-term change in genetic estimates of effective population size despite massive widespread mortality in *Pisaster ochraceus*. ***Under Review at the Biological Bulletin.***

2021 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. ***Accepted at ICES Journal of Marine Science.***

2021 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***

2021 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

2021 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

2021 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

2018 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*

2018 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

2018 **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

2017 **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

2017 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

2016 **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

2016 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*

2015 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

2015 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

2014 **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

2014 **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

2013 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

2013 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

2013 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

2012 **Puritz, J. B.,** Keever, C. C., Addison, J. A., Byrne, M. Hart, M. W, Grosberg, R. K., Toonen, R. J. Extraordinarily rapid life history divergence between *Cryptasterina* sea star species. ***Proc. R. Soc. B*.** DOI: 10.1098/rspb.2012.1343

2012 **Puritz, J. B**., Addison, J. A., Toonen, R. J. Next-Generation Phylogeography: the application of 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. Iacchei**, 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

2005 Walker, A. N., Bush, P., **Puritz, J**. **B.**, Wilson, T., Chang, E. S., Miller, T., Holloway, K., 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 = $2,290,153**

2021 **USDA- Agriculture Research Service** **$250,000**
Population genomic evaluation of wild and cultured Eastern oyster populations from the Northeast region
(PI Puritz; Total to Puritz: 250,000; additional $50,000 in match Project Total: $300,000)

2021 **National Science Foundation** **$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)

2020 **National Science Foundation** **$298,978**
*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)

2019 **Rhode Island Sea Grant** **$249,192**
*How do multiple coastal stressors limit oyster recruitment?*
(PI Puritz; Total to Puritz: $249,192; additional $124,617 in match Project Total: $373,809)

2019 **US Fish and Wildlife Service** **$80,435**
*Assessing Horseshoe Crab Population Structure within Southern New England* (PIs: McManus, Ameral, and Puritz; Total to Puritz $80,435; additional match: $43,302 (URI) & $13,992 DEM; Project Total: $163,715

2019 **Atlantic Coast Marine Fisheries Council** **$690,000**
*From Sequence to Consequence: Genomic Selection to Expand And improve Selective Breeding for The Eastern Oyster* (Eastern Oyster Breeding Consortium; URI PIs: Puritz and Gomez-Chiarri;
Total to URI: $690,000; Project Total: $4,363,092)

2016 **Harte Research Institute Staff Innovations Grant (Declined)** **$5,000**
Collaborative proposal to investigate evolutionary impacts of barotrauma in red snapper

2015 **Texas Research Development Fund Program** **$20,000**
*Impacts of sewage effluent on genomic diversity and connectivity of marine intertidal communities*

2011 **NSF East Asian Pacific Science Institute Fellowship** **$8,000**
Fellowship for two months of research at the University of Sydney

2011 **University of Hawaii Graduate Student Organization** **$750**
Travel Grant for 2011 Annual Meeting of the Western Society of Naturalists

2010 **University of Hawaii Graduate Student Organization** **$900**
Travel Grant for 2010 Benthic Ecology Meeting

2009 **University of Hawaii Graduate Student Organization** **$750**
Research grant for sample collection trip to California

2008 **The Charles H. and Margaret B. Edmondson Research Fund** **$1,500**
Research on the population connectivity of *Linckia multifora*

2007 **Ecology, Evolution, and Conservation Biology Travel Grant** **$750**
Travel Grant for 2007 Annual Meeting of the Western Society of Naturalists

2007 **University of Hawaii Arts and Sciences Advisory Council Award** **$3,000**
Research expedition to the Chagos Archipelago

## TEACHING EXPERIENCE

### Classes

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: Invertebrate Zoology

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

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 diversity**
Department of Biology Seminar- University of Louisiana at Lafayette

2015 **Using next-generation sequencing to examine patterns of coastal genomic 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

## CONFERENCE PRESENTATIONS (LAST 5 YEARS)

^undergraduate mentee, # post-baccalaureate mentee, ***** graduate student mentee

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.

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.

2021 **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.

2016 **Mind the gap: the effects of INDELs and over-splitting on population genetic inference from RAD sequencing**
Puritz, J. B., Portnoy, D. S., Gold, J. R. Evolution. Raleigh, NC

2016 **Testing the Genomic Impacts of the DWH Oil Spill on Red Snapper**
Puritz, J. B., Portnoy, D. S., Gold, J. R. Gulf of Mexico Oil Spill and Ecosystem Science Conference. Tampa Bay, FL.

## CONFERENCE POSTERS (LAST 5 YEARS)

^undergraduate mentee, # post-baccalaureate mentee, * graduate student mentee

2021 **The effect of environmental parameters on *Crassostrea virginica* shell sizes**.
Satkowski^, S., Zyck*, A., Puritz, J.B. 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.

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**

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.

2015 **Fishing for selection, but only catching bias: examining library effects in double-digest RAD data in non-model marine species**
Puritz, J. B., Hollenbeck, C. M., Gold, J. R. Plant and Animal Genomes. San Diego, CA.

## MENTORING

### *Undergraduate*

| | |
|---|---|
| 2021- | 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*

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|---|---|---|---|
| 2021- | Gabriel Barrett | MS. | Biological and Environmental Sciences |
| 2020- | Megan Guidry | PhD. | Biological and Environmental Sciences |
| 2019- | 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*

| | |
|---|---|
| 2020- | Tyler Devos, M.S. Biological and Environmental Sciences |
| 2019- | Benjamin Ha, PhD. Ecology and Evolutionary Biology (UCLA) |
| 2019- | Ian 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) |

## PROFESSIONAL SERVICE

2021- **BES Leadership Committee**
Coordinator of the Evolution and Marine Biology Graduate Specialization

2021- **GSO Science Saturday**
Lab had an interactive research experience in Blount Aquarium

2021- **Sea Star Wasting Disease Working Group**
Participant

2018- **Functional Re-annotation of Oyster Genomes with Epigenetic Resources (FROGER) working group**
Participant

2018- **Moorea Coral Reef LTER Connectivity working group**
Participant

2018- **Equity, inclusion, diversity committee, Biological Sciences**
Member

2018- **Committee to establish a graduate Data Science Program at URI**
Member

2017- **Eastern Oyster Genome Consortium working group**
Participant, leading genome description manuscript

2014- **dDocent RADseq Bioinformatics pipeline**
Developer, software has over 50,000 downloads and website with 80,000 visitors

2019 **Faculty host at URI Welcome Day**

2018-2019 **Ad hoc committee for Cruickshank Lecture, Biological Sciences**
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)**

2015-16 **HRI Senior Research Staff Steering Committee Member**
Initiated the HRI Staff Innovations Grant Program

***Peer Reviewer***

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

## 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**
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

2014-16 **Texas A&M Corpus Christi Scientific Diver (AAUS Reciprocity)**

2005-12 **University of Hawaii Scientific Diver (AAUS Reciprocity)**

2007 **NOAA Advanced Coxswain**