

Scott A Pavey

List of Publications by Year in descending order

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Version: 2024-02-01

25
papers

979
citations

686830

13
h-index

642321

23
g-index

25
all docs

25
docs citations

25
times ranked

1668
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Transborder Gene Flow between Canada and the USA and Fine-Scale Population Structure of Atlantic Cod in the Broader Gulf of Maine Region. <i>Transactions of the American Fisheries Society</i> , 2021, 150, 560-577. | 0.6 | 1 |
| 2 | Genomic population structure of Striped Bass (<i>Morone saxatilis</i>) from the Gulf of St. Lawrence to Cape Fear River. <i>Evolutionary Applications</i> , 2020, 13, 1468-1486. | 1.5 | 13 |
| 3 | River-Specific Gene Expression Patterns Associated with Habitat Selection for Key Hormone-Coding Genes in Glass Eel-Stage American Eels. <i>Transactions of the American Fisheries Society</i> , 2018, 147, 855-868. | 0.6 | 0 |
| 4 | The evolution of the major histocompatibility complex in upstream versus downstream river populations of the longnose dace. <i>Ecology and Evolution</i> , 2017, 7, 3297-3311. | 0.8 | 4 |
| 5 | Draft genome of the American Eel (<i>Anguilla rostrata</i>). <i>Molecular Ecology Resources</i> , 2017, 17, 806-811. | 2.2 | 21 |
| 6 | RAD-Seq Reveals Patterns of Additive Polygenic Variation Caused by Spatially-Varying Selection in the American Eel (<i>Anguilla rostrata</i>). <i>Genome Biology and Evolution</i> , 2017, 9, 2974-2986. | 1.1 | 35 |
| 7 | Regional variation of gene regulation associated with storage lipid metabolism in American glass eels (<i>Anguilla rostrata</i>). <i>Comparative Biochemistry and Physiology Part A, Molecular & Integrative Physiology</i> , 2016, 196, 30-37. | 0.8 | 7 |
| 8 | RAD Sequencing Highlights Polygenic Discrimination of Habitat Ecotypes in the Panmictic American Eel. <i>Current Biology</i> , 2015, 25, 1666-1671. | 1.8 | 88 |
| 9 | Growth, Female Size, and Sex Ratio Variability in American Eel of Different Origins in Both Controlled Conditions and the Wild: Implications for Stocking Programs. <i>Transactions of the American Fisheries Society</i> , 2015, 144, 246-257. | 0.6 | 31 |
| 10 | Ecological release leads to novel ontogenetic diet shift in kokanee (<i>Oncorhynchus nerka</i>). <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2015, 72, 1718-1730. | 0.7 | 13 |
| 11 | Characterization of MHC class IIB for four endangered Australian freshwater fishes obtained from ecologically divergent populations. <i>Fish and Shellfish Immunology</i> , 2015, 46, 468-476. | 1.6 | 8 |
| 12 | Neutral and selective processes shape MHC gene diversity and expression in stocked brook charr populations (<i>Salvelinus fontinalis</i>). <i>Molecular Ecology</i> , 2014, 23, 1730-1748. | 2.0 | 21 |
| 13 | Nonparallelism in MHCII ² diversity accompanies nonparallelism in pathogen infection of lake whitefish (<i>Coregonus clupeaformis</i>) species pairs as revealed by next-generation sequencing. <i>Molecular Ecology</i> , 2013, 22, 3833-3849. | 2.0 | 38 |
| 14 | THE GENETIC ARCHITECTURE OF REPRODUCTIVE ISOLATION DURING SPECIATION-WITH-GENE-FLOW IN LAKE WHITEFISH SPECIES PAIRS ASSESSED BY RAD SEQUENCING. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 2483-2497. | 1.1 | 187 |
| 15 | Mapping phenotypic, expression and transmission ratio distortion QTL using RAD markers in the Lake Whitefish (<i>Coregonus clupeaformis</i>). <i>Molecular Ecology</i> , 2013, 22, 3036-3048. | 2.0 | 96 |
| 16 | Gene Coexpression Networks Reveal Key Drivers of Phenotypic Divergence in Lake Whitefish. <i>Molecular Biology and Evolution</i> , 2013, 30, 1384-1396. | 3.5 | 115 |
| 17 | What is needed for next-generation ecological and evolutionary genomics?. <i>Trends in Ecology and Evolution</i> , 2012, 27, 673-678. | 4.2 | 77 |
| 18 | A fast, highly sensitive double-nested PCR-based method to screen fish immunobiomes. <i>Molecular Ecology Resources</i> , 2012, 12, 1027-1039. | 2.2 | 11 |

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|----|---|-----|-----------|
| 19 | Ecological transcriptomics of lake-type and riverine sockeye salmon (<i>Oncorhynchus nerka</i>). <i>BMC Ecology</i> , 2011, 11, 31. | 3.0 | 12 |
| 20 | Full length MHC II ^β exon 2 primers for salmonids: a new resource for next generation sequencing. <i>Conservation Genetics Resources</i> , 2011, 3, 665-667. | 0.4 | 7 |
| 21 | RECENT ECOLOGICAL DIVERGENCE DESPITE MIGRATION IN SOCKEYE SALMON (<i>ONCORHYNCHUS NERKA</i>). <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 1773-1783. | 1.1 | 17 |
| 22 | The role of gene expression in ecological speciation. <i>Annals of the New York Academy of Sciences</i> , 2010, 1206, 110-129. | 1.8 | 134 |
| 23 | Perspectives: Gene expression in fisheries management. <i>Environmental Epigenetics</i> , 2010, 56, 157-156. | 0.9 | 13 |
| 24 | Contrasting Ecology Shapes Juvenile Lake- and Riverine Sockeye Salmon. <i>Transactions of the American Fisheries Society</i> , 2010, 139, 1584-1594. | 0.6 | 17 |
| 25 | Revisiting evolutionary dead ends in sockeye salmon (<i>Oncorhynchus nerka</i>) life history. <i>Canadian Journal of Fisheries and Aquatic Sciences</i> , 2007, 64, 1199-1208. | 0.7 | 13 |