

Mikhail V Matz

List of Publications by Year in descending order

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Version: 2024-02-01

132
papers

15,456
citations

28242

55
h-index

20343

116
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158
all docs

158
docs citations

158
times ranked

16206
citing authors

#	ARTICLE	IF	CITATIONS
1	Mechanisms and potential immune tradeoffs of accelerated coral growth induced by microfragmentation. <i>PeerJ</i> , 2022, 10, e13158.	0.9	4
2	Changes in gene body methylation do not correlate with changes in gene expression in Anthozoa or Hexapoda. <i>BMC Genomics</i> , 2022, 23, 234.	1.2	19
3	Coralâ€bleaching responses to climate change across biological scales. <i>Global Change Biology</i> , 2022, 28, 4229-4250.	4.2	44
4	Benchmarking DNA methylation assays in a reefâ€building coral. <i>Molecular Ecology Resources</i> , 2021, 21, 464-477.	2.2	12
5	Gene expression associated with disease resistance and long-term growth in a reef-building coral. <i>Royal Society Open Science</i> , 2021, 8, 210113.	1.1	10
6	Comparative neurotranscriptomics reveal widespread species differences associated with bonding. <i>BMC Genomics</i> , 2021, 22, 399.	1.2	7
7	Environmental specialization and cryptic genetic divergence in two massive coral species from the Florida Keys Reef Tract. <i>Molecular Ecology</i> , 2021, 30, 3468-3484.	2.0	27
8	Shuffling between <i>Cladocopium</i> and <i>Durusdinium</i> extensively modifies the physiology of each symbiont without stressing the coral host. <i>Molecular Ecology</i> , 2021, 30, 6585-6595.	2.0	10
9	How mitonuclear discordance and geographic variation have confounded species boundaries in a widely studied snake. <i>Molecular Phylogenetics and Evolution</i> , 2021, 162, 107194.	1.2	21
10	Six priorities to advance the science and practice of coral reef restoration worldwide. <i>Restoration Ecology</i> , 2021, 29, e13498.	1.4	36
11	A 2b-RAD parentage analysis pipeline for complex and mixed DNA samples. <i>Forensic Science International: Genetics</i> , 2021, 55, 102590.	1.6	3
12	Population genetics of the coral <i>Acropora millepora</i> : Toward genomic prediction of bleaching. <i>Science</i> , 2020, 369, .	6.0	167
13	<i>Cladocopium</i> community divergence in two <i>Acropora</i> coral hosts across multiple spatial scales. <i>Molecular Ecology</i> , 2020, 29, 4559-4572.	2.0	21
14	Metaâ€analysis of the coral environmental stress response: <i>Acropora</i> corals show opposing responses depending on stress intensity. <i>Molecular Ecology</i> , 2020, 29, 2855-2870.	2.0	60
15	Molecular tools for coral reef restoration: Beyond biomarker discovery. <i>Conservation Letters</i> , 2020, 13, e12687.	2.8	44
16	Estimating the potential for coral adaptation to global warming across the Indoâ€West Pacific. <i>Global Change Biology</i> , 2020, 26, 3473-3481.	4.2	54
17	Positive genetic associations among fitness traits support evolvability of a reefâ€building coral under multiple stressors. <i>Global Change Biology</i> , 2019, 25, 3294-3304.	4.2	50
18	Considerations for maximizing the adaptive potential of restored coral populations in the western Atlantic. <i>Ecological Applications</i> , 2019, 29, e01978.	1.8	163

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19	Role of host genetics and heat-tolerant algal symbionts in sustaining populations of the endangered coral <i>Orbicella faveolata</i> in the Florida Keys with ocean warming. <i>Global Change Biology</i> , 2019, 25, 1016-1031.	4.2	111
20	Comparative transcriptomics of sympatric species of coral reef fishes (genus: <i>Haemulon</i>). <i>PeerJ</i> , 2019, 7, e6541.	0.9	6
21	Effects of thermal stress on amount, composition, and antibacterial properties of coral mucus. <i>PeerJ</i> , 2019, 7, e6849.	0.9	23
22	CRISPR/Cas9-mediated genome editing in a reef-building coral. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 5235-5240.	3.3	110
23	Complex selection on a regulator of social cognition: Evidence of balancing selection, regulatory interactions and population differentiation in the prairie vole <i>Avpr1a</i> locus. <i>Molecular Ecology</i> , 2018, 27, 419-431.	2.0	3
24	Fantastic Beasts and How To Sequence Them: Ecological Genomics for Obscure Model Organisms. <i>Trends in Genetics</i> , 2018, 34, 121-132.	2.9	64
25	Role of gene body methylation in acclimatization and adaptation in a basal metazoan. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 13342-13346.	3.3	136
26	Potential and limits for rapid genetic adaptation to warming in a Great Barrier Reef coral. <i>PLoS Genetics</i> , 2018, 14, e1007220.	1.5	184
27	Contrasting effects of <i>Symbiodinium</i> identity on coral host transcriptional profiles across latitudes. <i>Molecular Ecology</i> , 2018, 27, 3103-3115.	2.0	23
28	Molecular characterization of larval development from fertilization to metamorphosis in a reef-building coral. <i>BMC Genomics</i> , 2018, 19, 17.	1.2	39
29	Relationship between <i>Acropora millepora</i> juvenile fluorescence and composition of newly established <i>Symbiodinium</i> assemblage. <i>PeerJ</i> , 2018, 6, e5022.	0.9	5
30	Transcriptome dynamics over a lunar month in a broadcast spawning acroporid coral. <i>Molecular Ecology</i> , 2017, 26, 2514-2526.	2.0	32
31	Modeled differences of coral life-history traits influence the refugium potential of a remote Caribbean reef. <i>Coral Reefs</i> , 2017, 36, 913-925.	0.9	30
32	Intraspecific differences in molecular stress responses and coral pathobiome contribute to mortality under bacterial challenge in <i>Acropora millepora</i> . <i>Scientific Reports</i> , 2017, 7, 2609.	1.6	78
33	Population structure and connectivity of the mountainous star coral, <i>Orbicella faveolata</i> , throughout the wider Caribbean region. <i>Ecology and Evolution</i> , 2017, 7, 9234-9246.	0.8	49
34	Rapid adaptive responses to climate change in corals. <i>Nature Climate Change</i> , 2017, 7, 627-636.	8.1	327
35	Gene expression plasticity as a mechanism of coral adaptation to a variable environment. <i>Nature Ecology and Evolution</i> , 2017, 1, 14.	3.4	306
36	Coral larvae for restoration and research: a large-scale method for rearing <i>Acropora millepora</i> larvae, inducing settlement, and establishing symbiosis. <i>PeerJ</i> , 2017, 5, e3732.	0.9	67

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37	Variation in heat shock protein expression at the latitudinal range limits of a widely distributed species, the <i>Ganville fritillaria</i> butterfly (<i>Melitaea cinxia</i>). <i>Physiological Entomology</i> , 2016, 41, 241-248.	0.6	15
38	Evolutionary Consequences of DNA Methylation in a Basal Metazoan. <i>Molecular Biology and Evolution</i> , 2016, 33, 2285-2293.	3.5	57
39	Red fluorescence in coral larvae is associated with a diapause-like state. <i>Molecular Ecology</i> , 2016, 25, 559-569.	2.0	28
40	Evolutionary origins of germline segregation in Metazoa: evidence for a germ stem cell lineage in the coral <i>Orbicella faveolata</i> (Cnidaria, Anthozoa). <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2016, 283, 20152128.	1.2	34
41	Differential responses of coral larvae to the colour of ambient light guide them to suitable settlement microhabitat. <i>Royal Society Open Science</i> , 2015, 2, 150358.	1.1	46
42	Fine-scale environmental specialization of reef-building corals might be limiting reef recovery in the Florida Keys. <i>Ecology</i> , 2015, 96, 3197-3212.	1.5	74
43	Exploring the role of Micronesian islands in the maintenance of coral genetic diversity in the Pacific Ocean. <i>Molecular Ecology</i> , 2015, 24, 70-82.	2.0	68
44	Estimating Trait Heritability in Highly Fecund Species. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 2639-2645.	0.8	17
45	A Hinge Migration Mechanism Unlocks the Evolution of Green-to-Red Photoconversion in GFP-like Proteins. <i>Structure</i> , 2015, 23, 34-43.	1.6	58
46	Heritable differences in fitness-related traits among populations of the mustard hill coral, <i>Porites astreoides</i> . <i>Heredity</i> , 2015, 115, 509-516.	1.2	74
47	Genomic determinants of coral heat tolerance across latitudes. <i>Science</i> , 2015, 348, 1460-1462.	6.0	473
48	Quantitative high resolution melting: two methods to determine SNP allele frequencies from pooled samples. <i>BMC Genetics</i> , 2015, 16, 62.	2.7	14
49	Gene expression associated with white syndromes in a reef building coral, <i>Acropora hyacinthus</i> . <i>BMC Genomics</i> , 2015, 16, 371.	1.2	271
50	Deep-Sequencing Method for Quantifying Background Abundances of Symbiodinium Types: Exploring the Rare Symbiodinium Biosphere in Reef-Building Corals. <i>PLoS ONE</i> , 2014, 9, e94297.	1.1	135
51	Quantifying cryptic <i>Symbiodinium</i> diversity within <i>Orbicella faveolata</i> and <i>Orbicella franksi</i> at the Flower Garden Banks, Gulf of Mexico. <i>PeerJ</i> , 2014, 2, e386.	0.9	55
52	A cross-ocean comparison of responses to settlement cues in reef-building corals. <i>PeerJ</i> , 2014, 2, e333.	0.9	22
53	Bimodal signatures of germline methylation are linked with gene expression plasticity in the coral <i>Acropora millepora</i> . <i>BMC Genomics</i> , 2014, 15, 1109.	1.2	89
54	Diagnostic gene expression biomarkers of coral thermal stress. <i>Molecular Ecology Resources</i> , 2014, 14, 667-678.	2.2	65

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55	Demystifying the <sc>RAD</sc> fad. <i>Molecular Ecology</i> , 2014, 23, 5937-5942.	2.0	199
56	So, you want to use next-generation sequencing in marine systems? Insight from the Pan-Pacific Advanced Studies Institute. <i>Bulletin of Marine Science</i> , 2014, 90, 79-122.	0.4	53
57	Novel polymorphic microsatellite markers for population genetics of the endangered Caribbean star coral, <i>Montastraea faveolata</i> . <i>Marine Biodiversity</i> , 2013, 43, 167-172.	0.3	47
58	Acidâ€Base Catalysis and Crystal Structures of a Least Evolved Ancestral GFP-like Protein Undergoing Green-to-Red Photoconversion. <i>Biochemistry</i> , 2013, 52, 8048-8059.	1.2	25
59	Deep relationships of Rhizaria revealed by phylogenomics: A farewell to Haeckelâ€™s Radiolaria. <i>Molecular Phylogenetics and Evolution</i> , 2013, 67, 53-59.	1.2	65
60	Evidence for a host role in thermotolerance divergence between populations of the mustard hill coral (<i><sc>P</sc>orites astreoides</i>) from different reef environments. <i>Molecular Ecology</i> , 2013, 22, 4335-4348.	2.0	158
61	Gene expression under chronic heat stress in populations of the mustard hill coral (<i><sc>P</sc>orites astreoides</i>) from different thermal environments. <i>Molecular Ecology</i> , 2013, 22, 4322-4334.	2.0	242
62	Gene Expression Signatures of Energetic Acclimatisation in the Reef Building Coral <i>Acropora millepora</i> . <i>PLoS ONE</i> , 2013, 8, e61736.	1.1	32
63	No Control Genes Required: Bayesian Analysis of qRT-PCR Data. <i>PLoS ONE</i> , 2013, 8, e71448.	1.1	137
64	Ecological Complexity of Coral Recruitment Processes: Effects of Invertebrate Herbivores on Coral Recruitment and Growth Depends Upon Substratum Properties and Coral Species. <i>PLoS ONE</i> , 2013, 8, e72830.	1.1	35
65	2b-RAD: a simple and flexible method for genome-wide genotyping. <i>Nature Methods</i> , 2012, 9, 808-810.	9.0	607
66	Multi-domain GFP-like proteins from two species of marine hydrozoans. <i>Photochemical and Photobiological Sciences</i> , 2012, 11, 637-644.	1.6	18
67	Multi-colored homologs of the green fluorescent protein from hydromedusa <i>Obelia</i> sp.. <i>Photochemical and Photobiological Sciences</i> , 2011, 10, 1303-1309.	1.6	14
68	Profiling gene expression responses of coral larvae (<i>Acropora millepora</i>) to elevated temperature and settlement inducers using a novel RNA-Seq procedure. <i>Molecular Ecology</i> , 2011, 20, no-no.	2.0	328
69	Fluorescence of coral larvae predicts their settlement response to crustose coralline algae and reflects stress. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2011, 278, 2691-2697.	1.2	53
70	Rapid Evolution of Coral Proteins Responsible for Interaction with the Environment. <i>PLoS ONE</i> , 2011, 6, e20392.	1.1	114
71	Development of Gene Expression Markers of Acute Heat-Light Stress in Reef-Building Corals of the Genus <i>Porites</i> . <i>PLoS ONE</i> , 2011, 6, e26914.	1.1	108
72	Evolution of Rhizaria: new insights from phylogenomic analysis of uncultivated protists. <i>BMC Evolutionary Biology</i> , 2010, 10, 377.	3.2	130

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73	Very Bright Green Fluorescent Proteins from the Pontellid Copepod <i>Pontella mimocerami</i> . <i>PLoS ONE</i> , 2010, 5, e11517.	1.1	30
74	Retracing Evolution of Red Fluorescence in GFP-Like Proteins from Faviina Corals. <i>Molecular Biology and Evolution</i> , 2010, 27, 225-233.	3.5	60
75	Fluorescent Proteins and Their Applications in Imaging Living Cells and Tissues. <i>Physiological Reviews</i> , 2010, 90, 1103-1163.	13.1	1,175
76	Characterization of a Group of MITEs with Unusual Features from Two Coral Genomes. <i>PLoS ONE</i> , 2010, 5, e10700.	1.1	14
77	Microsatellite Characterization and Marker Development from Public EST and WGS Databases in the Reef-Building Coral <i>Acropora millepora</i> (Cnidaria, Anthozoa, Scleractinia). <i>Journal of Heredity</i> , 2009, 100, 329-337.	1.0	42
78	Sequencing and de novo analysis of a coral larval transcriptome using 454 GSFlx. <i>BMC Genomics</i> , 2009, 10, 219.	1.2	405
79	Construction of a high-resolution genetic linkage map and comparative genome analysis for the reef-building coral <i>Acropora millepora</i> . <i>Genome Biology</i> , 2009, 10, R126.	13.9	55
80	Genetic variation in responses to a settlement cue and elevated temperature in the reef-building coral <i>Acropora millepora</i> . <i>Marine Ecology - Progress Series</i> , 2009, 392, 81-92.	0.9	102
81	Giant Deep-Sea Protist Produces Bilaterian-like Traces. <i>Current Biology</i> , 2008, 18, 1849-1854.	1.8	72
82	A Green Fluorescent Protein with Photoswitchable Emission from the Deep Sea. <i>PLoS ONE</i> , 2008, 3, e3766.	1.1	32
83	Diversity and Evolution of Coral Fluorescent Proteins. <i>PLoS ONE</i> , 2008, 3, e2680.	1.1	281
84	Blue light regulation of host pigment in reef-building corals. <i>Marine Ecology - Progress Series</i> , 2008, 364, 97-106.	0.9	110
85	Fluorescence lifetime imaging of coral fluorescent proteins. <i>Microscopy Research and Technique</i> , 2007, 70, 243-251.	1.2	18
86	Contributions of host and symbiont pigments to the coloration of reef corals. <i>FEBS Journal</i> , 2007, 274, 1102-1122.	2.2	101
87	It's cheap to be colorful. <i>FEBS Journal</i> , 2007, 274, 2496-2505.	2.2	64
88	Dealing with model uncertainty in reconstructing ancestral proteins in the laboratory: examples from archosaur visual pigments and coralfuorescent proteins. , 2007, , 164-180.		2
89	Ordered Differential Display. , 2006, 317, 059-074.		0
90	Statistical Approaches for DNA Barcoding. <i>Systematic Biology</i> , 2006, 55, 162-169.	2.7	122

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91	BOOK REVIEW Aglow in the Dark: The Revolutionary Science of Biofluorescence. Oceanography, 2006, 19, 155-157.	0.5	0
92	Photoinduced activation of GFP-like proteins in tissues of reef corals. , 2006, 6098, 64.		2
93	Are Corals Colorful?. Photochemistry and Photobiology, 2006, 82, 345.	1.3	79
94	Adaptive Evolution of Multicolored Fluorescent Proteins in Reef-Building Corals. Journal of Molecular Evolution, 2006, 62, 332-339.	0.8	90
95	Discovery and properties of GFP-like proteins from nonbioluminescent anthozoa. Methods of Biochemical Analysis, 2006, 47, 121-38.	0.2	4
96	Evolution of Function and Color in GFP-Like Proteins. Methods of Biochemical Analysis, 2005, 47, 139-161.	0.2	12
97	A likelihood ratio test for species membership based on DNA sequence data. Philosophical Transactions of the Royal Society B: Biological Sciences, 2005, 360, 1969-1974.	1.8	86
98	Applications of Ancestral Protein Reconstruction in Understanding Protein Function: GFP-Like Proteins. Methods in Enzymology, 2005, 395, 652-670.	0.4	20
99	Discovery and Properties of GFP-Like Proteins from Nonbioluminescent Anthozoa. Methods of Biochemical Analysis, 2005, , 121-138.	0.2	6
100	GFP-like Proteins as Ubiquitous Metazoan Superfamily: Evolution of Functional Features and Structural Complexity. Molecular Biology and Evolution, 2004, 21, 841-850.	3.5	394
101	Evolution of Coral Pigments Recreated. Science, 2004, 305, 1433-1433.	6.0	144
102	Simple cDNA normalization using kamchatka crab duplex-specific nuclease. Nucleic Acids Research, 2004, 32, 37e-37.	6.5	375
103	Amplification of cDNA Ends Using PCR Suppression Effect and Step-Out PCR. , 2003, 221, 41-50.		31
104	Molecular Basis and Evolutionary Origins of Color Diversity in Great Star Coral <i>Montastraea cavernosa</i> (Scleractinia: Faviida). Molecular Biology and Evolution, 2003, 20, 1125-1133.	3.5	102
105	Amplification of Representative cDNA Pools from Microscopic Amounts of Animal Tissue. , 2003, 221, 103-116.		21
106	Diversity and evolution of the green fluorescent protein family. Proceedings of the National Academy of Sciences of the United States of America, 2002, 99, 4256-4261.	3.3	340
107	Identification and characterization of a new family of C-type lectin-like genes from planaria <i>Girardia tigrina</i> . Glycobiology, 2002, 12, 463-472.	1.3	25
108	Amplification of Representative cDNA Samples from Microscopic Amounts of Invertebrate Tissue to Search for New Genes. , 2002, 183, 003-018.		32

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109	Family of the green fluorescent protein: Journey to the end of the rainbow. <i>BioEssays</i> , 2002, 24, 953-959.	1.2	131
110	Altering electrical connections in the nervous system of the pteropod mollusc <i>Clione limacina</i> by neuronal injections of gap junction mRNA. <i>European Journal of Neuroscience</i> , 2002, 16, 2475-2476.	1.2	20
111	GFP-like chromoproteins as a source of far-red fluorescent proteins. <i>FEBS Letters</i> , 2001, 507, 16-20.	1.3	240
112	Refined crystal structure of DsRed, a red fluorescent protein from coral, at 2.0-A resolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 462-467.	3.3	422
113	NATURAL ANIMAL COLORATION CAN BE DETERMINED BY A NON-FLUORESCENT GFP HOMOLOG. , 2001, , .		1
114	A ubiquitous family of putative gap junction molecules. <i>Current Biology</i> , 2000, 10, R473-R474.	1.8	485
115	Natural Animal Coloration Can Be Determined by a Nonfluorescent Green Fluorescent Protein Homolog. <i>Journal of Biological Chemistry</i> , 2000, 275, 25879-25882.	1.6	300
116	Novel fluorescent protein from <i>Discosoma</i> coral and its mutants possesses a unique far-red fluorescence. <i>FEBS Letters</i> , 2000, 479, 127-130.	1.3	136
117	"Fluorescent Timer": Protein That Changes Color with Time. , 2000, 290, 1585-1588.		347
118	Fluorescent proteins from nonbioluminescent Anthozoa species. <i>Nature Biotechnology</i> , 1999, 17, 969-973.	9.4	1,711
119	Amplification of cDNA ends based on template-switching effect and step-out PCR. <i>Nucleic Acids Research</i> , 1999, 27, 1558-1560.	6.5	381
120	Regulation of average length of complex PCR product. <i>Nucleic Acids Research</i> , 1999, 27, 23e-23.	6.5	57
121	Sequence-Independent Method for <i>In Vitro</i> Generation of Nested Deletions for Sequencing Large DNA Fragments. <i>Analytical Biochemistry</i> , 1998, 258, 138-141.	1.1	5
122	Inductive Interactions Regulating Body Patterning in Planarian, Revealed by Analysis of Expression of Novel Genes. <i>Developmental Biology</i> , 1998, 194, 172-181.	0.9	22
123	Different strategies of differential display: areas of application. <i>Nucleic Acids Research</i> , 1998, 26, 5537-5543.	6.5	71
124	Ordered differential display: a simple method for systematic comparison of gene expression profiles. <i>Nucleic Acids Research</i> , 1997, 25, 2541-2542.	6.5	96
125	Construction of cDNA Libraries from Small Amounts of Total RNA Using the Suppression PCR Effect. <i>Biochemical and Biophysical Research Communications</i> , 1997, 230, 285-288.	1.0	44
126	Whole mount <i>in situ</i> hybridization on freshwater planaria. <i>Technical Tips Online</i> , 1997, 2, 100-103.	0.2	0

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127	Molecule by molecule PCR amplification of complex DNA mixtures for direct sequencing: an approach to in vitro cloning. <i>Nucleic Acids Research</i> , 1996, 24, 2194-2195.	6.5	44
128	Primary structure of carboxypeptidase T: Delineation of functionally relevant features in Zn-carboxypeptidase family. <i>The Protein Journal</i> , 1992, 11, 561-570.	1.1	28
129	Crystal structure of carboxypeptidase T from <i>Thermoactinomyces vulgaris</i> . <i>FEBS Journal</i> , 1992, 208, 281-288.	0.2	65
130	Molecular cloning and primary structure of <i>Thermoactinomyces vulgaris</i> carboxypeptidase T A metalloenzyme endowed with dual substrate specificity. <i>FEBS Letters</i> , 1991, 291, 75-78.	1.3	27
131	Methods for Analysing mRNA Expression. , 0, , 163-407.		0
132	Novel fluorescent proteins: diversity, mutagenesis and applications. , 0, 2004, .		3