

# Mikhail V Matz

## List of Publications by Year in descending order

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132  
papers

15,456  
citations

28274

55  
h-index

19749

117  
g-index

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.	2.0	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.	2.8	19
3	Coralâ€bleaching responses to climate change across biological scales. <i>Global Change Biology</i> , 2022, 28, 4229-4250.	9.5	44
4	Benchmarking DNA methylation assays in a reefâ€building coral. <i>Molecular Ecology Resources</i> , 2021, 21, 464-477.	4.8	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.	2.4	10
6	Comparative neurotranscriptomics reveal widespread species differences associated with bonding. <i>BMC Genomics</i> , 2021, 22, 399.	2.8	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.	3.9	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.	3.9	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.	2.7	21
10	Six priorities to advance the science and practice of coral reef restoration worldwide. <i>Restoration Ecology</i> , 2021, 29, e13498.	2.9	36
11	A 2b-RAD parentage analysis pipeline for complex and mixed DNA samples. <i>Forensic Science International: Genetics</i> , 2021, 55, 102590.	3.1	3
12	Population genetics of the coral <i>Acropora millepora</i> : Toward genomic prediction of bleaching. <i>Science</i> , 2020, 369, .	12.6	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.	3.9	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.	3.9	60
15	Molecular tools for coral reef restoration: Beyond biomarker discovery. <i>Conservation Letters</i> , 2020, 13, e12687.	5.7	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.	9.5	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.	9.5	50
18	Considerations for maximizing the adaptive potential of restored coral populations in the western Atlantic. <i>Ecological Applications</i> , 2019, 29, e01978.	3.8	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.	9.5	111
20	Comparative transcriptomics of sympatric species of coral reef fishes (genus: <i>Haemulon</i> ). <i>PeerJ</i> , 2019, 7, e6541.	2.0	6
21	Effects of thermal stress on amount, composition, and antibacterial properties of coral mucus. <i>PeerJ</i> , 2019, 7, e6849.	2.0	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.	7.1	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.	3.9	3
24	Fantastic Beasts and How To Sequence Them: Ecological Genomics for Obscure Model Organisms. <i>Trends in Genetics</i> , 2018, 34, 121-132.	6.7	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.	7.1	136
26	Potential and limits for rapid genetic adaptation to warming in a Great Barrier Reef coral. <i>PLoS Genetics</i> , 2018, 14, e1007220.	3.5	184
27	Contrasting effects of <i>Symbiodinium</i> identity on coral host transcriptional profiles across latitudes. <i>Molecular Ecology</i> , 2018, 27, 3103-3115.	3.9	23
28	Molecular characterization of larval development from fertilization to metamorphosis in a reef-building coral. <i>BMC Genomics</i> , 2018, 19, 17.	2.8	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.	2.0	5
30	Transcriptome dynamics over a lunar month in a broadcast spawning acroporid coral. <i>Molecular Ecology</i> , 2017, 26, 2514-2526.	3.9	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.	2.2	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.	3.3	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.	1.9	49
34	Rapid adaptive responses to climate change in corals. <i>Nature Climate Change</i> , 2017, 7, 627-636.	18.8	327
35	Gene expression plasticity as a mechanism of coral adaptation to a variable environment. <i>Nature Ecology and Evolution</i> , 2017, 1, 14.	7.8	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.	2.0	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>Myiophobus cinctus</i> ). <i>Physiological Entomology</i> , 2016, 41, 241-248.	1.5	15
38	Evolutionary Consequences of DNA Methylation in a Basal Metazoan. <i>Molecular Biology and Evolution</i> , 2016, 33, 2285-2293.	8.9	57
39	Red fluorescence in coral larvae is associated with a diapause-like state. <i>Molecular Ecology</i> , 2016, 25, 559-569.	3.9	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.	2.6	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.	2.4	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.	3.2	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.	3.9	68
44	Estimating Trait Heritability in Highly Fecund Species. <i>G3: Genes, Genomes, Genetics</i> , 2015, 5, 2639-2645.	1.8	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.	3.3	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.	2.6	74
47	Genomic determinants of coral heat tolerance across latitudes. <i>Science</i> , 2015, 348, 1460-1462.	12.6	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.	2.8	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.	2.5	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.	2.0	55
52	A cross-ocean comparison of responses to settlement cues in reef-building corals. <i>PeerJ</i> , 2014, 2, e333.	2.0	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.	2.8	89
54	Diagnostic gene expression biomarkers of coral thermal stress. <i>Molecular Ecology Resources</i> , 2014, 14, 667-678.	4.8	65

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55	Demystifying the <sc>RAD</sc> fad. <i>Molecular Ecology</i> , 2014, 23, 5937-5942.	3.9	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.	0.8	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.	1.0	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.	2.5	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.	2.7	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.	3.9	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.	3.9	242
62	Gene Expression Signatures of Energetic Acclimatisation in the Reef Building Coral <i>Acropora millepora</i> . <i>PLoS ONE</i> , 2013, 8, e61736.	2.5	32
63	No Control Genes Required: Bayesian Analysis of qRT-PCR Data. <i>PLoS ONE</i> , 2013, 8, e71448.	2.5	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.	2.5	35
65	2b-RAD: a simple and flexible method for genome-wide genotyping. <i>Nature Methods</i> , 2012, 9, 808-810.	19.0	607
66	Multi-domain GFP-like proteins from two species of marine hydrozoans. <i>Photochemical and Photobiological Sciences</i> , 2012, 11, 637-644.	2.9	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.	2.9	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.	3.9	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.	2.6	53
70	Rapid Evolution of Coral Proteins Responsible for Interaction with the Environment. <i>PLoS ONE</i> , 2011, 6, e20392.	2.5	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.	2.5	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.	2.5	30
74	Retracing Evolution of Red Fluorescence in GFP-Like Proteins from Faviina Corals. <i>Molecular Biology and Evolution</i> , 2010, 27, 225-233.	8.9	60
75	Fluorescent Proteins and Their Applications in Imaging Living Cells and Tissues. <i>Physiological Reviews</i> , 2010, 90, 1103-1163.	28.8	1,175
76	Characterization of a Group of MITEs with Unusual Features from Two Coral Genomes. <i>PLoS ONE</i> , 2010, 5, e10700.	2.5	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.	2.4	42
78	Sequencing and de novo analysis of a coral larval transcriptome using 454 GSFlx. <i>BMC Genomics</i> , 2009, 10, 219.	2.8	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.	9.6	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.	1.9	102
81	Giant Deep-Sea Protist Produces Bilaterian-like Traces. <i>Current Biology</i> , 2008, 18, 1849-1854.	3.9	72
82	A Green Fluorescent Protein with Photoswitchable Emission from the Deep Sea. <i>PLoS ONE</i> , 2008, 3, e3766.	2.5	32
83	Diversity and Evolution of Coral Fluorescent Proteins. <i>PLoS ONE</i> , 2008, 3, e2680.	2.5	281
84	Blue light regulation of host pigment in reef-building corals. <i>Marine Ecology - Progress Series</i> , 2008, 364, 97-106.	1.9	110
85	Fluorescence lifetime imaging of coral fluorescent proteins. <i>Microscopy Research and Technique</i> , 2007, 70, 243-251.	2.2	18
86	Contributions of host and symbiont pigments to the coloration of reef corals. <i>FEBS Journal</i> , 2007, 274, 1102-1122.	4.7	101
87	It's cheap to be colorful. <i>FEBS Journal</i> , 2007, 274, 2496-2505.	4.7	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.	5.6	122

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91	BOOK REVIEW   Aglow in the Dark: The Revolutionary Science of Biofluorescence. Oceanography, 2006, 19, 155-157.	1.0	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.	2.5	79
94	Adaptive Evolution of Multicolored Fluorescent Proteins in Reef-Building Corals. Journal of Molecular Evolution, 2006, 62, 332-339.	1.8	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.	4.0	86
98	Applications of Ancestral Protein Reconstruction in Understanding Protein Function: GFP-Like Proteins. Methods in Enzymology, 2005, 395, 652-670.	1.0	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.	8.9	394
101	Evolution of Coral Pigments Recreated. Science, 2004, 305, 1433-1433.	12.6	144
102	Simple cDNA normalization using kamchatka crab duplex-specific nuclease. Nucleic Acids Research, 2004, 32, 37e-37.	14.5	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.	8.9	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.	7.1	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.	2.5	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.	2.5	131
110	Altering electrical connections in the nervous system of the pteropod mollusc <i>Clione limacin</i> by neuronal injections of gap junction mRNA. <i>European Journal of Neuroscience</i> , 2002, 16, 2475-2476.	2.6	20
111	GFP-like chromoproteins as a source of far-red fluorescent proteins. <i>FEBS Letters</i> , 2001, 507, 16-20.	2.8	240
112	Refined crystal structure of DsRed, a red fluorescent protein from coral, at 2.0-Å resolution. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 462-467.	7.1	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.	3.9	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.	3.4	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.	2.8	136
117	"Fluorescent Timer": Protein That Changes Color with Time. <i>Science</i> , 2000, 290, 1585-1588.	12.6	347
118	Fluorescent proteins from nonbioluminescent Anthozoa species. <i>Nature Biotechnology</i> , 1999, 17, 969-973.	17.5	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.	14.5	381
120	Regulation of average length of complex PCR product. <i>Nucleic Acids Research</i> , 1999, 27, 23e-23.	14.5	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.	2.4	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.	2.0	22
123	Different strategies of differential display: areas of application. <i>Nucleic Acids Research</i> , 1998, 26, 5537-5543.	14.5	71
124	Ordered differential display: a simple method for systematic comparison of gene expression profiles. <i>Nucleic Acids Research</i> , 1997, 25, 2541-2542.	14.5	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.	2.1	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.	14.5	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.	2.8	27
131	Methods for Analysing mRNA Expression. , 0, , 163-407.		0
132	Novel fluorescent proteins: diversity, mutagenesis and applications. , 0, 2004, .		3