

# Travis C Glenn

## List of Publications by Year in descending order

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223  
papers

13,938  
citations

46918

47  
h-index

25716

108  
g-index

245  
all docs

245  
docs citations

245  
times ranked

15577  
citing authors

#	ARTICLE	IF	CITATIONS
1	Estimating Movement Rates Between Eurasian and North American Birds That Are Vectors of Avian Influenza. <i>Avian Diseases</i> , 2022, 66, .	0.4	0
2	Comparison of Three Methods for Measuring Dietary Composition of Plains Hog-nosed Snakes. <i>Herpetologica</i> , 2022, 78, .	0.2	2
3	Tissue Distribution of Mercury in the Bodies of Wild American Alligators ( <i>Alligator mississippiensis</i> ) from a Coastal Marsh in Louisiana (USA). <i>Archives of Environmental Contamination and Toxicology</i> , 2022, 83, 13-20.	2.1	3
4	Population genetic divergence of bonnethead sharks <i>Sphyrna tiburo</i> in the western North Atlantic: Implications for conservation. <i>Aquatic Conservation: Marine and Freshwater Ecosystems</i> , 2021, 31, 83-98.	0.9	12
5	Whole genome genetic variation and linkage disequilibrium in a diverse collection of <i>Listeria monocytogenes</i> isolates. <i>PLoS ONE</i> , 2021, 16, e0242297.	1.1	0
6	Improved Microbial Community Characterization of 16S rRNA via Metagenome Hybridization Capture Enrichment. <i>Frontiers in Microbiology</i> , 2021, 12, 644662.	1.5	23
7	Ultraconserved elements reconstruct the evolution of Chagas disease vectoring kissing bugs ( <i>Reduviidae: Triatominae</i> ). <i>Systematic Entomology</i> , 2021, 46, 725-740.	1.7	24
8	Molecular Phylogeny and Evolution of Amazon Parrots in the Greater Antilles. <i>Genes</i> , 2021, 12, 608.	1.0	2
9	Unveiling the Gut Microbiota and Resistome of Wild Cotton Mice, <i>Peromyscus gossypinus</i> , from Heavy Metal- and Radionuclide-Contaminated Sites in the Southeastern United States. <i>Microbiology Spectrum</i> , 2021, 9, e0009721.	1.2	4
10	Escaping the fate of Sisyphus: assessing resistome hybridization baits for antimicrobial resistance gene capture. <i>Environmental Microbiology</i> , 2021, 23, 7523-7537.	1.8	3
11	A High-Quality Reference Genome Assembly of the Saltwater Crocodile, <i>Crocodylus porosus</i> , Reveals Patterns of Selection in Crocodylidae. <i>Genome Biology and Evolution</i> , 2020, 12, 3635-3646.	1.1	15
12	How microclimatic variables and blood meal sources influence <i>Rhodnius prolixus</i> abundance and <i>Trypanosoma cruzi</i> infection in <i>Attalea butyracea</i> and <i>Elaeis guineensis</i> palms?. <i>Acta Tropica</i> , 2020, 212, 105674.	0.9	4
13	An Open-Source Program (Haplo-ST) for Whole-Genome Sequence Typing Shows Extensive Diversity among <i>Listeria monocytogenes</i> Isolates in Outdoor Environments and Poultry Processing Plants. <i>Applied and Environmental Microbiology</i> , 2020, 87, .	1.4	5
14	Divergence, gene flow, and speciation in eight lineages of trans-Beringian birds. <i>Molecular Ecology</i> , 2020, 29, 3526-3542.	2.0	18
15	Agricultural pests consumed by common bat species in the United States corn belt: The importance of DNA primer choice. <i>Agriculture, Ecosystems and Environment</i> , 2020, 303, 107105.	2.5	17
16	Microbiota of Four Tissue Types in American Alligators ( <i>Alligator mississippiensis</i> ) Following Extended Dietary Selenomethionine Exposure. <i>Bulletin of Environmental Contamination and Toxicology</i> , 2020, 105, 381-386.	1.3	1
17	Co-occurrence of antibiotic, biocide, and heavy metal resistance genes in bacteria from metal and radionuclide contaminated soils at the Savannah River Site. <i>Microbial Biotechnology</i> , 2020, 13, 1179-1200.	2.0	89
18	Identification and characterization of microRNAs (miRNAs) and their transposable element origins in the saltwater crocodile, <i>Crocodylus porosus</i> . <i>Analytical Biochemistry</i> , 2020, 602, 113781.	1.1	6

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19	Integration of ecosystem science into radioecology: A consensus perspective. <i>Science of the Total Environment</i> , 2020, 740, 140031.	3.9	13
20	A High-Quality Genome Assembly of the North American Song Sparrow, <i>Melospiza melodia</i> . <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 1159-1166.	0.8	8
21	Genome comparison and transcriptome analysis of the invasive brown root rot pathogen, <i>Phellinus noxius</i> , from different geographic regions reveals potential enzymes associated with degradation of different wood substrates. <i>Fungal Biology</i> , 2020, 124, 144-154.	1.1	11
22	Comparison of the ruminal and fecal microbiotas in beef calves supplemented or not with concentrate. <i>PLoS ONE</i> , 2020, 15, e0231533.	1.1	56
23	Population genetics of two chromatic morphs of the Chagas disease vector <i>Rhodnius pallescens</i> Barber, 1932 in Panamá. <i>Infection, Genetics and Evolution</i> , 2020, 84, 104369.	1.0	2
24	Identification and characterization of a fast-neutron-induced mutant with elevated seed protein content in soybean. <i>Theoretical and Applied Genetics</i> , 2019, 132, 2965-2983.	1.8	7
25	Speciation despite gene flow in two owls ( <i>Aegolius</i> spp.): Evidence from 2,517 ultraconserved element loci. <i>Auk</i> , 2019, 136, .	0.7	8
26	Regional biogeography of microbiota composition in the Chagas disease vector <i>Rhodnius pallescens</i> . <i>Parasites and Vectors</i> , 2019, 12, 504.	1.0	17
27	Genomic mutations after multigenerational exposure of <i>Caenorhabditis elegans</i> to pristine and sulfidized silver nanoparticles. <i>Environmental Pollution</i> , 2019, 254, 113078.	3.7	31
28	Horizontal Gene Transfer and Acquired Antibiotic Resistance in <i>Salmonella enterica</i> Serovar Heidelberg following <i>In Vitro</i> Incubation in Broiler Ceca. <i>Applied and Environmental Microbiology</i> , 2019, 85, .	1.4	39
29	Bromate-induced Changes in p21 DNA Methylation and Histone Acetylation in Renal Cells. <i>Toxicological Sciences</i> , 2019, 168, 460-473.	1.4	7
30	Generalist host species drive <i>Trypanosoma cruzi</i> vector infection in oil palm plantations in the Orinoco region, Colombia. <i>Parasites and Vectors</i> , 2019, 12, 274.	1.0	16
31	Analysis of the Rumen Microbiota of Beef Calves Supplemented During the Suckling Phase. <i>Frontiers in Microbiology</i> , 2019, 10, 1131.	1.5	15
32	Examining the Effects of Chronic Selenium Exposure on Traditionally Used Stress Parameters in Juvenile American Alligators ( <i>Alligator mississippiensis</i> ). <i>Archives of Environmental Contamination and Toxicology</i> , 2019, 77, 14-21.	2.1	15
33	Earth history and the passerine superradiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 7916-7925.	3.3	238
34	Formation of a recent hybrid zone offers insight into the geographic puzzle and maintenance of species boundaries in musk turtles. <i>Molecular Ecology</i> , 2019, 28, 761-771.	2.0	17
35	Insight from an ultraconserved element bait set designed for hemipteran phylogenetics integrated with genomic resources. <i>Molecular Phylogenetics and Evolution</i> , 2019, 130, 297-303.	1.2	51
36	Adapterama III: Quadruple-indexed, double/triple-enzyme RADseq libraries (2RAD/3RAD). <i>PeerJ</i> , 2019, 7, e7724.	0.9	96

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37	Adapterama I: universal stubs and primers for 384 unique dual-indexed or 147,456 combinatorially-indexed Illumina libraries (iTru & iNext). PeerJ, 2019, 7, e7755.	0.9	243
38	Adapterama II: universal amplicon sequencing on Illumina platforms (TaggiMatrix). PeerJ, 2019, 7, e7786.	0.9	47
39	Long-term treatment with green tea polyphenols modifies the gut microbiome of female sprague-dawley rats. Journal of Nutritional Biochemistry, 2018, 56, 55-64.	1.9	64
40	45 Analysis of the Gastrointestinal Tract-Associated Microbiome of Calves Supplemented during the Suckling Phase.. Journal of Animal Science, 2018, 96, 24-24.	0.2	0
41	Complete mitochondrial genome of the yellowfin tuna ( <i>Thunnus albacares</i> ) and the blackfin tuna ( <i>Thunnus atlanticus</i> ): notes on mtDNA introgression and paraphyly on tunas. Conservation Genetics Resources, 2018, 10, 697-699.	0.4	3
42	Mitochondrial genomes of the Pacific sierra mackerel <i>Scomberomorus sierra</i> and the Monterey Spanish mackerel <i>Scomberomorus concolor</i> (Perciformes, Scombridae). Conservation Genetics Resources, 2018, 10, 471-474.	0.4	1
43	Conflicting Evolutionary Histories of the Mitochondrial and Nuclear Genomes in New World Myotis Bats. Systematic Biology, 2018, 67, 236-249.	2.7	56
44	Resolving taxonomic turbulence and uncovering cryptic diversity in the musk turtles ( <i>Sternotherus</i> ) using robust demographic modeling. Molecular Phylogenetics and Evolution, 2018, 120, 1-15.	1.2	23
45	95 Analysis Of The Gastrointestinal Tract-Associated Microbiome Of Calves Supplemented During The Suckling Phase.. Journal of Animal Science, 2018, 96, 408-408.	0.2	0
46	Isolation and characterization of microsatellite markers for conservation management of the endangered Great-billed Seed-finch, <i>Sporophila maximiliani</i> (Aves, Passeriformes), and cross-amplification in other congeners. Molecular Biology Reports, 2018, 45, 2815-2819.	1.0	4
47	Transcriptome Changes of <i>Escherichia coli</i> , <i>Enterococcus faecalis</i> , and <i>Escherichia coli</i> O157:H7 Laboratory Strains in Response to Photo-Degraded DOM. Frontiers in Microbiology, 2018, 9, 882.	1.5	6
48	A High-Quality Reference Genome for the Invasive Mosquitofish <i>Gambusia affinis</i> Using a Chicago Library. G3: Genes, Genomes, Genetics, 2018, 8, 1855-1861.	0.8	16
49	Dietary Selenomethionine Administration and Its Effects on the American Alligator ( <i>Alligator</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Contamination and Toxicology, 2018, 75, 37-44.	2.1	11
50	Influence of landscape heterogeneity on the functional connectivity of Allegheny woodrats ( <i>Neotoma magister</i> ) in Virginia. Conservation Genetics, 2018, 19, 1259-1268.	0.8	10
51	Ultraconserved elements (UCEs) illuminate the population genomics of a recent, high-latitude avian speciation event. PeerJ, 2018, 6, e5735.	0.9	31
52	Complete mitogenome sequences of the pacific red snapper ( <i>Lutjanus peru</i> ) and the spotted rose snapper ( <i>Lutjanus guttatus</i> ). Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis, 2017, 28, 223-224.	0.7	6
53	Dietary Selenomethionine Administration in the American Alligator ( <i>Alligator mississippiensis</i> ): Hepatic and Renal Se Accumulation and Its Effects on Growth and Body Condition. Archives of Environmental Contamination and Toxicology, 2017, 72, 439-448.	2.1	16
54	Genistein prevention of hyperglycemia and improvement of glucose tolerance in adult non-obese diabetic mice are associated with alterations of gut microbiome and immune homeostasis. Toxicology and Applied Pharmacology, 2017, 332, 138-148.	1.3	57

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55	Multiple Paternity Benefits Female Marbled Salamanders by Increasing Survival of Progeny to Metamorphosis. <i>Ethology</i> , 2017, 123, 307-315.	0.5	5
56	Blood Meal Source Characterization Using Illumina Sequencing in the Chagas Disease Vector <i>Rhodnius pallescens</i> (Hemiptera: Reduviidae) in Panamá. <i>Journal of Medical Entomology</i> , 2017, 54, 1786-1789.	0.9	36
57	Habitat predictors of genetic diversity for two sympatric wetland breeding amphibian species. <i>Ecology and Evolution</i> , 2017, 7, 6271-6283.	0.8	8
58	The Novel Evolution of the Sperm Whale Genome. <i>Genome Biology and Evolution</i> , 2017, 9, 3260-3264.	1.1	33
59	Rapid Microbiome Changes in Freshly Deposited Cow Feces under Field Conditions. <i>Frontiers in Microbiology</i> , 2016, 7, 500.	1.5	49
60	Use of sonic tomography to detect and quantify wood decay in living trees. <i>Applications in Plant Sciences</i> , 2016, 4, 1600060.	0.8	32
61	Addressing ecological effects of radiation on populations and ecosystems to improve protection of the environment against radiation: Agreed statements from a Consensus Symposium. <i>Journal of Environmental Radioactivity</i> , 2016, 158-159, 21-29.	0.9	75
62	Assessing the microbiomes of scald and chiller tank waters throughout a typical commercial poultry processing day. <i>Poultry Science</i> , 2016, 95, 2372-2382.	1.5	26
63	Chronic Ingestion of Coal Fly-Ash Contaminated Prey and Its Effects on Health and Immune Parameters in Juvenile American Alligators ( <i>Alligator mississippiensis</i> ). <i>Archives of Environmental Contamination and Toxicology</i> , 2016, 71, 347-358.	2.1	15
64	Capturing Darwin's dream. <i>Molecular Ecology Resources</i> , 2016, 16, 1051-1058.	2.2	22
65	Nephrotoxicity of epigenetic inhibitors used for the treatment of cancer. <i>Chemico-Biological Interactions</i> , 2016, 258, 21-29.	1.7	6
66	<scp>RAD</scp>cap: sequence capture of dual digest <scp>RAD</scp>seq libraries with identifiable duplicates and reduced missing data. <i>Molecular Ecology Resources</i> , 2016, 16, 1264-1278.	2.2	117
67	Sequence Capture versus Restriction Site Associated DNA Sequencing for Shallow Systematics. <i>Systematic Biology</i> , 2016, 65, 910-924.	2.7	220
68	Targeted DNA Region Re-sequencing. , 2016, , 43-68.		9
69	Detection of an Enigmatic Plethodontid Salamander Using Environmental DNA. <i>Copeia</i> , 2016, 104, 78-82.	1.4	19
70	Analysis of a Rapid Evolutionary Radiation Using Ultraconserved Elements: Evidence for a Bias in Some Multispecies Coalescent Methods. <i>Systematic Biology</i> , 2016, 65, 612-627.	2.7	137
71	Aflatoxin B <sub>1</sub> Induced Compositional Changes in Gut Microbial Communities of Male F344 Rats. <i>Toxicological Sciences</i> , 2016, 150, 54-63.	1.4	78
72	Avoiding Missing Data Biases in Phylogenomic Inference: An Empirical Study in the Landfowl (Aves: Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5	8.5	208

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73	Implementing and testing the multispecies coalescent model: A valuable paradigm for phylogenomics. <i>Molecular Phylogenetics and Evolution</i> , 2016, 94, 447-462.	1.2	321
74	Screening wild and semi-free ranging great apes for putative sexually transmitted diseases: Evidence of Trichomonadidae infections. <i>American Journal of Primatology</i> , 2015, 77, 1075-1085.	0.8	9
75	IN OVO AND IN VITRO SUSCEPTIBILITY OF AMERICAN ALLIGATORS (ALLIGATOR MISSISSIPPIENSIS) TO AVIAN INFLUENZA VIRUS INFECTION. <i>Journal of Wildlife Diseases</i> , 2015, 51, 187-198.	0.3	7
76	Development and characterization of microsatellite loci for common raven ( <i>Corvus corax</i> ) and cross species amplification in other Corvidae. <i>BMC Research Notes</i> , 2015, 8, 655.	0.6	2
77	Resolving phylogenetic relationships of the recently radiated carnivorous plant genus <i>Sarracenia</i> using target enrichment. <i>Molecular Phylogenetics and Evolution</i> , 2015, 85, 76-87.	1.2	108
78	Novel and cross-amplified microsatellite loci for the critically endangered São Paulo marsh antwren <i>Formicivora paludicola</i> (Aves: Thamnophilidae). <i>Conservation Genetics Resources</i> , 2015, 7, 129-131.	0.4	3
79	Development of 12 novel microsatellite loci for invasive Chinese privet ( <i>Ligustrum sinense</i> ) from its introduced range. <i>Conservation Genetics Resources</i> , 2015, 7, 467-469.	0.4	0
80	Development of 31 new microsatellite loci for two mole salamanders ( <i>Ambystoma laterale</i> and <i>A. tigrinum</i> ). <i>Conservation Genetics Resources</i> , 2015, 7, 470-472.	0.4	2
81	Characterization of 15 microsatellite loci in kudzu ( <i>Pueraria montana</i> var. <i>lobata</i> ) from the native and introduced ranges. <i>Conservation Genetics Resources</i> , 2015, 7, 403-405.	0.4	6
82	Impacts of degraded DNA on restriction enzyme associated DNA sequencing (RADSeq). <i>Molecular Ecology Resources</i> , 2015, 15, 1304-1315.	2.2	114
83	Eleven microsatellites in an emerging invader, <i>Phytolacca americana</i> (Phytolaccaceae), from its native and introduced ranges. <i>Applications in Plant Sciences</i> , 2015, 3, 1500002.	0.8	7
84	A phylogenomic analysis of turtles. <i>Molecular Phylogenetics and Evolution</i> , 2015, 83, 250-257.	1.2	244
85	Assessment of Environmental DNA for Detecting Presence of Imperiled Aquatic Amphibian Species in Isolated Wetlands. <i>Journal of Fish and Wildlife Management</i> , 2015, 6, 498-510.	0.4	29
86	Comparative Genome Analyses Reveal Distinct Structure in the Saltwater Crocodile MHC. <i>PLoS ONE</i> , 2014, 9, e114631.	1.1	22
87	Three crocodylian genomes reveal ancestral patterns of evolution among archosaurs. <i>Science</i> , 2014, 346, 1254449.	6.0	300
88	Whole-genome analyses resolve early branches in the tree of life of modern birds. <i>Science</i> , 2014, 346, 1320-1331.	6.0	1,583
89	A genetic map of <i>Peromyscus</i> with chromosomal assignment of linkage groups (a <i>Peromyscus</i> genetic map). <i>PLoS ONE</i> , 2014, 9, e114631.	1.0	24
90	Development and characterization of microsatellite loci for two species of Beringian birds, rock sandpiper ( <i>Calidris ptilocnemis</i> ) and Pacific wren ( <i>Troglodytes pacificus</i> ). <i>Conservation Genetics Resources</i> , 2014, 6, 175-177.	0.4	3

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91	Target Capture and Massively Parallel Sequencing of Ultraconserved Elements for Comparative Studies at Shallow Evolutionary Time Scales. <i>Systematic Biology</i> , 2014, 63, 83-95.	2.7	286
92	The drivers of tropical speciation. <i>Nature</i> , 2014, 515, 406-409.	13.7	452
93	The evolution of peafowl and other taxa with ocelli (eyespot): a phylogenomic approach. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2014, 281, 20140823.	1.2	47
94	Incongruence among different mitochondrial regions: A case study using complete mitogenomes. <i>Molecular Phylogenetics and Evolution</i> , 2014, 78, 314-323.	1.2	75
95	Expression profiling of lymph node cells from deer mice infected with Andes virus. <i>BMC Immunology</i> , 2013, 14, 18.	0.9	18
96	Significant variance in genetic diversity among populations of <i>Schistosoma haematobium</i> detected using microsatellite DNA loci from a genome-wide database. <i>Parasites and Vectors</i> , 2013, 6, 300.	1.0	26
97	Specialized stem cell niche enables repetitive renewal of alligator teeth. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, E2009-18.	3.3	97
98	Development and Characterization of Microsatellite Primers in <i>Geranium carolinianum</i> (Geraniaceae) with 454 Sequencing. <i>Applications in Plant Sciences</i> , 2013, 1, 1300006.	0.8	6
99	THE ROLE OF INBREEDING DEPRESSION AND MATING SYSTEM IN THE EVOLUTION OF HETEROSTYLY. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 2309-2322.	1.1	18
100	Microsatellite Markers in the Western Prairie Fringed Orchid, <i>Platanthera praeclara</i> (Orchidaceae). <i>Applications in Plant Sciences</i> , 2013, 1, 1200413.	0.8	9
101	STRAW: Species TRee Analysis Web server. <i>Nucleic Acids Research</i> , 2013, 41, W238-W241.	6.5	93
102	Using phytohaemagglutinin to determine immune responsiveness in saltwater crocodiles ( <i>Crocodylus</i> )	0.6	13
103	A Phylogeny of Birds Based on Over 1,500 Loci Collected by Target Enrichment and High-Throughput Sequencing. <i>PLoS ONE</i> , 2013, 8, e54848.	1.1	287
104	Transcriptome Analysis of a North American Songbird, <i>Melospiza melodia</i> . <i>DNA Research</i> , 2012, 19, 325-333.	1.5	16
105	Not All Sequence Tags Are Created Equal: Designing and Validating Sequence Identification Tags Robust to Indels. <i>PLoS ONE</i> , 2012, 7, e42543.	1.1	267
106	Microsatellite primers for the neotropical epiphyte <i>Epidendrum firmum</i> (Orchidaceae). <i>American Journal of Botany</i> , 2012, 99, e450-2.	0.8	5
107	Characterization of unstable microsatellites in mice: No evidence for germline mutation induction following gamma radiation exposure. <i>Environmental and Molecular Mutagenesis</i> , 2012, 53, 599-607.	0.9	8
108	Transcriptome Sequencing and Annotation for the Jamaican Fruit Bat ( <i>Artibeus jamaicensis</i> ). <i>PLoS ONE</i> , 2012, 7, e48472.	1.1	77

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109	More than 1000 ultraconserved elements provide evidence that turtles are the sister group of archosaurs. <i>Biology Letters</i> , 2012, 8, 783-786.	1.0	331
110	Ultraconserved Elements Anchor Thousands of Genetic Markers Spanning Multiple Evolutionary Timescales. <i>Systematic Biology</i> , 2012, 61, 717-726.	2.7	983
111	Ultraconserved elements are novel phylogenomic markers that resolve placental mammal phylogeny when combined with species-tree analysis. <i>Genome Research</i> , 2012, 22, 746-754.	2.4	349
112	Development and characterization of tetranucleotide microsatellite loci for the American alligator ( <i>Alligator mississippiensis</i> ). <i>Conservation Genetics Resources</i> , 2012, 4, 567-570.	0.4	4
113	Fourteen novel microsatellite loci in the Chinese alligator ( <i>Alligator sinensis</i> ) isolated via 454 pyrosequencing. <i>Conservation Genetics Resources</i> , 2012, 4, 729-732.	0.4	4
114	Whole genome sequencing for quantifying germline mutation frequency in humans and model species: Cautious optimism. <i>Mutation Research - Reviews in Mutation Research</i> , 2012, 750, 96-106.	2.4	25
115	Reproductive Effects from Chronic, Multigenerational, Low Dose Rate Exposures to Radiation. <i>NATO Science for Peace and Security Series C: Environmental Security</i> , 2012, , 219-232.	0.1	2
116	The genome of the green anole lizard and a comparative analysis with birds and mammals. <i>Nature</i> , 2011, 477, 587-591.	13.7	575
117	Mating system in a gopher tortoise population established through multiple translocations: Apparent advantage of prior residence. <i>Biological Conservation</i> , 2011, 144, 175-183.	1.9	27
118	Field guide to next-generation DNA sequencers. <i>Molecular Ecology Resources</i> , 2011, 11, 759-769.	2.2	940
119	Large sets of edit-metric sequence identification tags to facilitate large-scale multiplexing of reads from massively parallel sequencing. <i>Nature Precedings</i> , 2011, , .	0.1	1
120	Genetic status of the wood stork ( <i>Mycteria americana</i> ) from the southeastern United States and the Brazilian Pantanal as revealed by mitochondrial DNA analysis. <i>Genetics and Molecular Research</i> , 2011, 10, 1910-1922.	0.3	5
121	Dinucleotide microsatellite markers in the genus <i>Caulerpa</i> . <i>Journal of Applied Phycology</i> , 2011, 23, 715-719.	1.5	6
122	Microsatellite markers isolated from the Mexican banded spring snail <i>Mexipyrghus churinceanus</i> . <i>Conservation Genetics Resources</i> , 2011, 3, 29-31.	0.4	2
123	Microsatellites isolated from the North American ground skink ( <i>Scincella lateralis</i> ). <i>Conservation Genetics Resources</i> , 2011, 3, 95-97.	0.4	1
124	Development and characterization of 18 microsatellite loci for the Southern Leopard Frog, <i>Rana sphenocéphala</i> . <i>Conservation Genetics Resources</i> , 2011, 3, 267-269.	0.4	4
125	Development and characterization of 12 microsatellite loci for the Dwarf Salamander, <i>Eurycea quadridigitata</i> . <i>Conservation Genetics Resources</i> , 2011, 3, 633-635.	0.4	1
126	Developing a community-based genetic nomenclature for anole lizards. <i>BMC Genomics</i> , 2011, 12, 554.	1.2	23



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127	Evaluating the Utility of Microsatellites for Investigations of Autopolyploid Taxa. <i>Journal of Heredity</i> , 2011, 102, 473-478.	1.0	13
128	Isolation and characterization of 14 polymorphic microsatellite DNA loci for the endangered Whooping Crane ( <i>Grus americana</i> ) and their applicability to other crane species. <i>Conservation Genetics Resources</i> , 2010, 2, 251-254.	0.4	14
129	Five hundred microsatellite loci for <i>Peromyscus</i> . <i>Conservation Genetics</i> , 2010, 11, 1243-1246.	0.8	15
130	QTL mapping for two commercial traits in farmed saltwater crocodiles ( <i>Crocodylus porosus</i> ). <i>Animal Genetics</i> , 2010, 41, 142-149.	0.6	6
131	Geographic Variation in the Mitochondrial Control Region of Black-throated Blue Warblers ( <i>Dendroica caerulescens</i> ). <i>Auk</i> , 2009, 126, 198-210.	0.7	8
132	A genetic linkage map for the saltwater crocodile ( <i>Crocodylus porosus</i> ). <i>BMC Genomics</i> , 2009, 10, 339.	1.2	29
133	Ten microsatellite loci from Northern Bobwhite ( <i>Colinus virginianus</i> ). <i>Conservation Genetics</i> , 2009, 10, 535-538.	0.8	13
134	253 Novel polymorphic microsatellites for the saltwater crocodile ( <i>Crocodylus porosus</i> ). <i>Conservation Genetics</i> , 2009, 10, 963-980.	0.8	23
135	Cross-species amplification of microsatellites in crocodylians: assessment and applications for the future. <i>Conservation Genetics</i> , 2009, 10, 935-954.	0.8	21
136	Characterization of microsatellite loci from the Malagasy endemic, <i>Tina striata</i> Radlk. (Sapindaceae). <i>Conservation Genetics</i> , 2009, 10, 1113-1115.	0.8	1
137	Fifteen polymorphic microsatellite loci from Jamaican streamertail hummingbirds ( <i>Trochilus</i> ). <i>Conservation Genetics</i> , 2009, 10, 1195-1198.	0.8	10
138	Development and characterization of nineteen polymorphic microsatellite loci from seaside alder, <i>Alnus maritima</i> . <i>Conservation Genetics</i> , 2009, 10, 1907-1910.	0.8	9
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