

# Nicolas Perrin

## List of Publications by Year in descending order

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105  
papers

7,018  
citations

76326

40  
h-index

71685

76  
g-index

109  
all docs

109  
docs citations

109  
times ranked

6880  
citing authors

#	ARTICLE	IF	CITATIONS
1	ECOLOGICAL-NICHE FACTOR ANALYSIS: HOW TO COMPUTE HABITAT-SUITABILITY MAPS WITHOUT ABSENCE DATA?. <i>Ecology</i> , 2002, 83, 2027-2036.	3.2	1,142
2	Sex Determination: Why So Many Ways of Doing It?. <i>PLoS Biology</i> , 2014, 12, e1001899.	5.6	916
3	Advances in our understanding of mammalian sex-biased dispersal. <i>Molecular Ecology</i> , 2007, 16, 1559-1578.	3.9	533
4	SEX REVERSAL: A FOUNTAIN OF YOUTH FOR SEX CHROMOSOMES?. <i>Evolution; International Journal of Organic Evolution</i> , 2009, 63, 3043-3049.	2.3	204
5	Ever-Young Sex Chromosomes in European Tree Frogs. <i>PLoS Biology</i> , 2011, 9, e1001062.	5.6	172
6	A rapid rate of sex-chromosome turnover and non-random transitions in true frogs. <i>Nature Communications</i> , 2018, 9, 4088.	12.8	149
7	Convergent Genetic Architecture Underlies Social Organization in Ants. <i>Current Biology</i> , 2014, 24, 2728-2732.	3.9	131
8	POPULATION DEMOGRAPHY AND THE EVOLUTION OF HELPING BEHAVIORS. <i>Evolution; International Journal of Organic Evolution</i> , 2006, 60, 1137-1151.	2.3	121
9	Is Sociality Driven by the Costs of Dispersal or the Benefits of Philopatry? A Role for Kin Discrimination Mechanisms. <i>American Naturalist</i> , 2001, 158, 471-483.	2.1	115
10	High-density sex-specific linkage maps of a European tree frog ( <i>Hyla arborea</i> ) identify the sex chromosome without information on offspring sex. <i>Heredity</i> , 2016, 116, 177-181.	2.6	102
11	Sex-Chromosome Turnovers: The Hot-Potato Model. <i>American Naturalist</i> , 2014, 183, 140-146.	2.1	101
12	SEX-CHROMOSOME TURNOVERS INDUCED BY DELETERIOUS MUTATION LOAD. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 635-645.	2.3	99
13	Cryptic diversity among Western Palearctic tree frogs: Postglacial range expansion, range limits, and secondary contacts of three European tree frog lineages ( <i>Hyla arborea</i> group). <i>Molecular Phylogenetics and Evolution</i> , 2012, 65, 1-9.	2.7	97
14	Mitochondrial and nuclear phylogeny of circum-Mediterranean tree frogs from the <i>Hyla arborea</i> group. <i>Molecular Phylogenetics and Evolution</i> , 2008, 49, 1019-1024.	2.7	93
15	TEMPERATURE-DEPENDENT TURNOVERS IN SEX-DETERMINATION MECHANISMS: A QUANTITATIVE MODEL. <i>Evolution; International Journal of Organic Evolution</i> , 2011, 65, 64-78.	2.3	92
16	A Concentric Analysis of the Impact of Urbanization on the Threatened European Tree Frog in an Agricultural Landscape. <i>Conservation Biology</i> , 2004, 18, 1599-1606.	4.7	90
17	HOMOLOGOUS SEX CHROMOSOMES IN THREE DEEPLY DIVERGENT ANURAN SPECIES. <i>Evolution; International Journal of Organic Evolution</i> , 2013, 67, 2434-2440.	2.3	77
18	A VERTEBRATE REPRODUCTIVE SYSTEM INVOLVING THREE PLOIDY LEVELS: HYBRID ORIGIN OF TRIPLOIDS IN A CONTACT ZONE OF DIPLOID AND TETRAPLOID PALEARCTIC GREEN TOADS ( <i>BUFO VIRIDIS</i> SUBGROUP)*. <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 944-959.	2.3	63

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19	Sex-Chromosome Recombination in Common Frogs Brings Water to the Fountain-of-Youth. <i>Molecular Biology and Evolution</i> , 2018, 35, 942-948.	8.9	63
20	Are glacial refugia hotspots of speciation and cytonuclear discordances? Answers from the genomic phylogeography of Spanish common frogs. <i>Molecular Ecology</i> , 2020, 29, 986-1000.	3.9	63
21	WHAT USES ARE MATING TYPES? THE "DEVELOPMENTAL SWITCH" MODEL. <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 947-956.	2.3	60
22	Genetic Bottlenecks Driven by Population Disconnection. <i>Conservation Biology</i> , 2010, 24, 1596-1605.	4.7	59
23	Cryptic recombination in the ever-young sex chromosomes of <i>Hyla</i> frogs. <i>Journal of Evolutionary Biology</i> , 2012, 25, 1947-1954.	1.7	57
24	Low rates of X-Y recombination, not turnovers, account for homomorphic sex chromosomes in several diploid species of Palearctic green toads ( <i>Bufo viridis</i> subgroup). <i>Journal of Evolutionary Biology</i> , 2013, 26, 674-682.	1.7	57
25	Sex-Chromosome Homomorphy in Palearctic Tree Frogs Results from Both Turnovers and X-Y Recombination. <i>Molecular Biology and Evolution</i> , 2015, 32, 2328-2337.	8.9	57
26	Random sex determination: When developmental noise tips the sex balance. <i>BioEssays</i> , 2016, 38, 1218-1226.	2.5	57
27	Extreme heterochiasmy and nascent sex chromosomes in European tree frogs. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2008, 275, 1577-1585.	2.6	55
28	Strong reproductive barriers in a narrow hybrid zone of West-Mediterranean green toads ( <i>Bufo</i> ). <i>Evolution</i> , 2010, 64, 1070-1080.	3.2	53
29	Evolutionary melting pots: a biodiversity hotspot shaped by ring diversifications around the Black Sea in the Eastern tree frog ( <i>Hyla orientalis</i> ). <i>Molecular Ecology</i> , 2016, 25, 4285-4300.	3.9	53
30	Phylogeography of a cryptic speciation continuum in Eurasian spadefoot toads ( <i>Pelobates</i> ). <i>Molecular Ecology</i> , 2019, 28, 3257-3270.	3.9	50
31	Integrating hybrid zone analyses in species delimitation: lessons from two anuran radiations of the Western Mediterranean. <i>Heredity</i> , 2020, 124, 423-438.	2.6	50
32	High-density linkage maps fail to detect any genetic component to sex determination in a <i>Rana temporaria</i> family. <i>Journal of Evolutionary Biology</i> , 2016, 29, 220-225.	1.7	49
33	Simultaneous Mendelian and clonal genome transmission in a sexually reproducing, all-triploid vertebrate. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2012, 279, 1293-1299.	2.6	47
34	Within-population polymorphism of sex-determination systems in the common frog ( <i>Rana temporaria</i> ). <i>Journal of Evolutionary Biology</i> , 2013, 26, 1569-1577.	1.7	47
35	THE EVOLUTION OF XY RECOMBINATION: SEXUALLY ANTAGONISTIC SELECTION VERSUS DELETERIOUS MUTATION LOAD. <i>Evolution; International Journal of Organic Evolution</i> , 2012, 66, 3155-3166.	2.3	46
36	Conservation phylogeography: does historical diversity contribute to regional vulnerability in European tree frogs ( <i>Hyla arborea</i> )?. <i>Molecular Ecology</i> , 2013, 22, 5669-5684.	3.9	45

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37	Mass of genes rather than master genes underlie the genomic architecture of amphibian speciation. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118, .	7.1	45
38	Timeframe of speciation inferred from secondary contact zones in the European tree frog radiation ( <i>Hyla arborea</i> group). BMC Evolutionary Biology, 2015, 15, 155.	3.2	44
39	Identifying homomorphic sex chromosomes from wild-caught adults with limited genomic resources. Molecular Ecology Resources, 2017, 17, 752-759.	4.8	44
40	Geographic variation in sex-chromosome differentiation in the common frog ( <i>Rana lessonae</i> ). Heredity, 2019, 123, 107-115.	3.9	43
41	Expanding the classical paradigm: what we have learnt from vertebrates about sex chromosome evolution. Philosophical Transactions of the Royal Society B: Biological Sciences, 2021, 376, 20200097.	4.0	43
42	Multiscale determinants of tree frog ( <i>Hyla arborea</i> L.) calling ponds in western Switzerland. Biodiversity and Conservation, 2004, 13, 2227-2235.	2.6	42
43	Phylogeography reveals an ancient cryptic radiation in East-Asian tree frogs ( <i>Hyla japonica</i> group) and complex relationships between continental and island lineages. BMC Evolutionary Biology, 2016, 16, 253.	3.2	42
44	Individual reproductive success and effective population size in the greater white-toothed shrew <i>Crocodyra russula</i> . Proceedings of the Royal Society B: Biological Sciences, 2000, 267, 701-705.	2.6	41
45	Sex chromosome turnovers and genetic drift: a simulation study. Journal of Evolutionary Biology, 2018, 31, 1413-1419.	1.7	40
46	A Sex-Specific Marker Reveals Male Heterogamety in European Tree Frogs. Molecular Biology and Evolution, 2006, 23, 1104-1106.	8.9	39
47	Inferring the degree of incipient speciation in secondary contact zones of closely related lineages of Palearctic green toads ( <i>Bufo viridis</i> subgroup). Heredity, 2014, 113, 9-20.	2.6	38
48	Sex-chromosome differentiation and "sex races" in the common frog ( <i>Rana temporaria</i> ). Proceedings of the Royal Society B: Biological Sciences, 2015, 282, 20142726.	2.6	35
49	Effect of biogeographic history on population vulnerability in European amphibians. Conservation Biology, 2015, 29, 1235-1241.	4.7	35
50	Empirical evidence for large X-effects in animals with undifferentiated sex chromosomes. Scientific Reports, 2016, 6, 21029.	3.3	35
51	Multiple origins of invasive and "native" water frogs ( <i>Pelophylax</i> spp.) in Switzerland. Biological Journal of the Linnean Society, 2014, 112, 442-449.	1.6	34
52	OPPORTUNITY FOR SEXUAL SELECTION AND EFFECTIVE POPULATION SIZE IN THE LEK-BREEDING EUROPEAN TREEFROG ( <i>HYLA ARBOREA</i> ). Evolution; International Journal of Organic Evolution, 2009, 63, 674-683.	2.3	30
53	A reciprocal translocation radically reshapes sex-linked inheritance in the common frog. Molecular Ecology, 2019, 28, 1877-1889.	3.9	30
54	The genetic contribution to sex determination and number of sex chromosomes vary among populations of common frogs ( <i>Rana temporaria</i> ). Heredity, 2016, 117, 25-32.	2.6	29

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55	A cryptic heterogametic transition revealed by sex-linked DNA markers in Palearctic green toads. <i>Journal of Evolutionary Biology</i> , 2011, 24, 1064-1070.	1.7	27
56	Trans-species variation in <i>Dmrt1</i> is associated with sex determination in four European tree-frog species. <i>Evolution; International Journal of Organic Evolution</i> , 2016, 70, 840-847.	2.3	27
57	<i>Dmrt1</i> polymorphism and sex-chromosome differentiation in <i>Rana temporaria</i> . <i>Molecular Ecology</i> , 2017, 26, 4897-4905.	3.9	27
58	Stronger transferability but lower variability in transcriptomic than in anonymous microsatellites: evidence from <i>Hyla</i> frogs. <i>Molecular Ecology Resources</i> , 2014, 14, 716-725.	4.8	26
59	GOOD GENES DRIVE FEMALE CHOICE FOR MATING PARTNERS IN THE LEK-BREEDING EUROPEAN TREEFROG. <i>Evolution; International Journal of Organic Evolution</i> , 2010, 64, 108-115.	2.3	25
60	Sex-chromosome differentiation parallels postglacial range expansion in European tree frogs ( <i>Hyla</i> ). <i>Evolution</i> , 2010, 64, 108-115.	2.3	25
61	Heterozygosity-fitness correlations among wild populations of European tree frog ( <i>Hyla</i> ). <i>Evolution</i> , 2010, 64, 108-115.	3.9	24
62	Cryptic invasion of Italian pool frogs ( <i>Pelophylax bergeri</i> ) across Western Europe unraveled by multilocus phylogeography. <i>Biological Invasions</i> , 2017, 19, 1407-1420.	2.4	24
63	Evolutionary and developmental dynamics of sex-biased gene expression in common frogs with proto-Y chromosomes. <i>Genome Biology</i> , 2018, 19, 156.	8.8	24
64	Tissue Specificity and Dynamics of Sex-Biased Gene Expression in a Common Frog Population with Differentiated, Yet Homomorphic, Sex Chromosomes. <i>Genes</i> , 2018, 9, 294.	2.4	24
65	Sex-chromosome evolution in frogs: what role for sex-antagonistic genes?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2021, 376, 20200094.	4.0	24
66	Phylogeography of Aegean green toads ( <i>Bufo viridis</i> subgroup): continental hybrid swarm vs. insular diversification with discovery of a new island endemic. <i>BMC Evolutionary Biology</i> , 2018, 18, 67.	3.2	23
67	<i>Dmrt1</i> polymorphism covaries with sex-determination patterns in <i>Rana temporaria</i> . <i>Ecology and Evolution</i> , 2016, 6, 5107-5117.	1.9	22
68	Hybridization and introgression between toads with different sex chromosome systems. <i>Evolution Letters</i> , 2020, 4, 444-456.	3.3	22
69	The effect of phylogeographic history on species boundaries: a comparative framework in <i>Hyla</i> tree frogs. <i>Scientific Reports</i> , 2020, 10, 5502.	3.3	21
70	Massive genetic introgression in threatened northern crested newts ( <i>Triturus cristatus</i> ) by an invasive congener ( <i>T. carnifex</i> ) in Western Switzerland. <i>Conservation Genetics</i> , 2016, 17, 839-846.	1.5	20
71	Are invasive marsh frogs ( <i>Pelophylax ridibundus</i> ) replacing the native <i>P. lessonae</i> / <i>P. esculentus</i> hybridogenetic complex in Western Europe? Genetic evidence from a field study. <i>Conservation Genetics</i> , 2014, 15, 869-878.	1.5	19
72	Impact of deleterious mutations, sexually antagonistic selection, and mode of recombination suppression on transitions between male and female heterogamety. <i>Heredity</i> , 2019, 123, 419-428.	2.6	19

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73	Introgressive hybridization of threatened European tree frogs ( <i>Hyla arborea</i> ) by introduced <i>H. intermedia</i> in Western Switzerland. <i>Conservation Genetics</i> , 2015, 16, 1507-1513.	1.5	18
74	Origin and genome evolution of polyploid green toads in Central Asia: evidence from microsatellite markers. <i>Heredity</i> , 2015, 114, 300-308.	2.6	18
75	Profound genetic divergence and asymmetric parental genome contributions as hallmarks of hybrid speciation in polyploid toads. <i>Proceedings of the Royal Society B: Biological Sciences</i> , 2018, 285, 20172667.	2.6	18
76	Sex-antagonistic genes, $\langle \text{sc} \rangle \text{XY} \langle \text{sc} \rangle$ recombination and feminized Y chromosomes. <i>Journal of Evolutionary Biology</i> , 2018, 31, 416-427.	1.7	18
77	Recombination is suppressed and variability reduced in a nascent Y chromosome. <i>Journal of Evolutionary Biology</i> , 2007, 20, 1182-1188.	1.7	16
78	Isolation and characterization of microsatellite loci for the European tree frog ( <i>Hyla</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50,542 Td (ar	4.8	16
79	Maintenance of biodiversity in vineyard-dominated landscapes: a case study on larval salamanders. <i>Animal Conservation</i> , 2012, 15, 136-141.	2.9	16
80	Reaching the edge of the speciation continuum: hybridization between three sympatric species of <i>Hyla</i> tree frogs. <i>Biological Journal of the Linnean Society</i> , 2019, 126, 743-750.	1.6	16
81	When Sex Chromosomes Recombine Only in the Heterogametic Sex: Heterochiasmy and Heterogamety in <i>Hyla</i> Tree Frogs. <i>Molecular Biology and Evolution</i> , 2021, 38, 192-200.	8.9	15
82	Viability and Management of an Endangered Capercaillie ( <i>Tetrao urogallus</i> ) Metapopulation in the Jura Mountains, Western Switzerland. <i>Biodiversity and Conservation</i> , 2006, 15, 2017-2032.	2.6	14
83	ON METAPOPOPULATION RESISTANCE TO DRIFT AND EXTINCTION. <i>Ecology</i> , 2006, 87, 1844-1855.	3.2	13
84	Maintenance of Ancestral Sex Chromosomes in Palearctic Tree Frogs: Direct Evidence from $\langle \text{sc} \rangle \text{XY} \langle \text{sc} \rangle$ <i>Hyla orientalis</i> . <i>Sexual Development</i> , 2013, 7, 261-266.	2.0	13
85	Diversification and speciation in tree frogs from the Maghreb ( <i>Hyla meridionalis sensu lato</i> ), with description of a new African endemic. <i>Molecular Phylogenetics and Evolution</i> , 2019, 134, 291-299.	2.7	13
86	Meiosis reveals the early steps in the evolution of a neo-XY sex chromosome pair in the African pygmy mouse <i>Mus minutoides</i> . <i>PLoS Genetics</i> , 2020, 16, e1008959.	3.5	13
87	A Key Transcription Cofactor on the Nascent Sex Chromosomes of European Tree Frogs ( <i>Hyla</i> ) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50,542 Td (ar	2.9	12
88	Phylogeography, more than elevation, accounts for sex chromosome differentiation in Swiss populations of the common frog ( <i>Rana temporaria</i> )*. <i>Evolution; International Journal of Organic Evolution</i> , 2020, 74, 644-654.	2.3	11
89	No evidence that Y-chromosome differentiation affects male fitness in a Swiss population of common frogs. <i>Journal of Evolutionary Biology</i> , 2020, 33, 401-409.	1.7	11
90	The contribution of patch topology and demographic parameters to population viability analysis predictions: the case of the European tree frog. <i>Population Ecology</i> , 2006, 48, 353-361.	1.2	10

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91	Range-Wide Sex-Chromosome Sequence Similarity Supports Occasional XY Recombination in European Tree Frogs ( <i>Hyla arborea</i> ). PLoS ONE, 2014, 9, e97959.	2.5	10
92	Dobzhansky-Muller incompatibilities, dominance drive, and sex-chromosome introgression at secondary contact zones: A simulation study. Evolution; International Journal of Organic Evolution, 2018, 72, 1350-1361.	2.3	10
93	First-generation linkage map for the European tree frog ( <i>Hyla arborea</i> ) with utility in congeneric species. BMC Research Notes, 2014, 7, 850.	1.4	9
94	Thirteen polymorphic microsatellite markers for the European green toad <i>Bufo viridis viridis</i> , a declining amphibian species. Conservation Genetics Resources, 2011, 3, 311-313.	0.8	7
95	New polymorphic microsatellite markers and development of mitotyping primers for West Mediterranean green toad species ( <i>Bufo viridis</i> subgroup). Molecular Ecology Resources, 2009, 9, 1138-1140.	4.8	6
96	Social systems: demographic and genetic issues. Molecular Ecology, 2012, 21, 443-446.	3.9	6
97	Capture and return of sexual genomes by hybridogenetic frogs provides clonal genome enrichment in a sexual species. Scientific Reports, 2021, 11, 1633.	3.3	6
98	Identification of ancestral sex chromosomes in the frog <i>Glandirana rugosa</i> bearing XX <sup>XY</sup> and ZZ <sup>ZW</sup> sex-determining systems. Molecular Ecology, 2022, 31, 3859-3870.	3.9	6
99	Invasion and eradication of a competitively superior species in heterogeneous landscapes. Ecological Modelling, 2011, 222, 398-406.	2.5	5
100	Polymorphism at a Sex-Linked Transcription Cofactor in European Tree Frogs ( <i>Hyla arborea</i> ): Sex-Antagonistic Selection or Neutral Processes?. Evolutionary Biology, 2011, 38, 208-213.	1.1	5
101	Discovery of a <i>Pelophylax saharicus</i> (Anura, Ranidae) population in Southern France: a new potentially invasive species of water frogs in Europe. Amphibia - Reptilia, 2021, 42, 427-442.	0.5	5
102	On tree frog cryptozoology and systematics – response to Y. Werner. Molecular Phylogenetics and Evolution, 2010, 57, 957-958.	2.7	4
103	Development and cross-amplification of thirty microsatellite loci in five diploid and polyploid Central Asian species of Palearctic green toads ( <i>Bufo viridis</i> subgroup). Conservation Genetics Resources, 2013, 5, 243-249.	0.8	3
104	ECOLOGICAL-NICHE FACTOR ANALYSIS: HOW TO COMPUTE HABITAT-SUITABILITY MAPS WITHOUT ABSENCE DATA?. , 2002, 83, 2027.		3
105	Evolutionary correlates of sex-determination systems. , 2014, , 115-132.		2