## Sylvain Glémin

## List of Publications by Year in descending order

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94433 102487 5,272 67 37 66 citations h-index g-index papers 76 76 76 6611 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Hunting for Beneficial Mutations: Conditioning on SIFT Scores When Estimating the Distribution of Fitness Effect of New Mutations. Genome Biology and Evolution, 2022, 14, .	2.5	13
2	What does the distribution of fitness effects of new mutations reflect? Insights from plants. New Phytologist, 2022, 233, 1613-1619.	7.3	6
3	Competitive ability depends on mating system and ploidy level across <i>Capsella</i> species. Annals of Botany, 2022, 129, 697-708.	2.9	2
4	The relative role of plasticity and demographic history in <i>Capsella bursa-pastoris</i> : a common garden experiment in Asia and Europe. AoB PLANTS, 2022, 14, .	2.3	4
5	Dioecy Is Associated with High Genetic Diversity and Adaptation Rates in the Plant Genus (i) Silene (i). Molecular Biology and Evolution, 2021, 38, 805-818.	8.9	31
6	Balancing selection in selfâ€fertilizing populations. Evolution; International Journal of Organic Evolution, 2021, 75, 1011-1029.	2.3	14
7	The ecology of the genome and the dynamics of the biological dark matter. Journal of Theoretical Biology, 2021, 518, 110641.	1.7	O
8	From Drift to Draft: How Much Do Beneficial Mutations Actually Contribute to Predictions of Ohta's Slightly Deleterious Model of Molecular Evolution?. Genetics, 2020, 214, 1005-1018.	2.9	25
9	Evolution of putative barrier loci at an intermediate stage of speciation with gene flow in campions ( <i>Silene</i> ). Molecular Ecology, 2020, 29, 3511-3525.	3.9	10
10	Shift in ecological strategy helps marginal populations of shepherd's purse ( <i>Capsella) Tj ETQq0 0 0 rgBT /Ove Sciences, 2020, 287, 20200463.</i>	erlock 10 Tr 2.6	f 50 387 Td (t 12
11	Genome Evolution in Outcrossing vs. Selfing vs. Asexual Species. Methods in Molecular Biology, 2019,		
	1910, 331-369.	0.9	51
12	Evolutionary transcriptomics reveals the origins of olives and the genomic changes associated with their domestication. Plant Journal, 2019, 100, 143-157.	5.7	51 64
12 13	Evolutionary transcriptomics reveals the origins of olives and the genomic changes associated with		
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13	Evolutionary transcriptomics reveals the origins of olives and the genomic changes associated with their domestication. Plant Journal, 2019, 100, 143-157.  Towards the new normal: Transcriptomic convergence and genomic legacy of the two subgenomes of an allopolyploid weed (Capsella bursa-pastoris). PLoS Genetics, 2019, 15, e1008131.	5.7 3.5	64 27
13 14	Evolutionary transcriptomics reveals the origins of olives and the genomic changes associated with their domestication. Plant Journal, 2019, 100, 143-157.  Towards the new normal: Transcriptomic convergence and genomic legacy of the two subgenomes of an allopolyploid weed (Capsella bursa-pastoris). PLoS Genetics, 2019, 15, e1008131.  Pervasive hybridizations in the history of wheat relatives. Science Advances, 2019, 5, eaav9188.  Parental legacy, demography, and admixture influenced the evolution of the two subgenomes of the	5.7 3.5 10.3	64 27 79
13 14 15	Evolutionary transcriptomics reveals the origins of olives and the genomic changes associated with their domestication. Plant Journal, 2019, 100, 143-157.  Towards the new normal: Transcriptomic convergence and genomic legacy of the two subgenomes of an allopolyploid weed (Capsella bursa-pastoris). PLoS Genetics, 2019, 15, e1008131.  Pervasive hybridizations in the history of wheat relatives. Science Advances, 2019, 5, eaav9188.  Parental legacy, demography, and admixture influenced the evolution of the two subgenomes of the tetraploid Capsella bursa-pastoris (Brassicaceae). PLoS Genetics, 2019, 15, e1007949.  Estimating the Fitness Effect of Deleterious Mutations During the Two Phases of the Life Cycle: A New	5.7 3.5 10.3	64 27 79 42

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19	Codon Usage Bias in Animals: Disentangling the Effects of Natural Selection, Effective Population Size, and GC-Biased Gene Conversion. Molecular Biology and Evolution, 2018, 35, 1092-1103.	8.9	111
20	Experimental Evidence for the Negative Effects of Self-Fertilization on the Adaptive Potential of Populations. Current Biology, 2017, 27, 237-242.	3.9	38
21	Domestication rewired gene expression and nucleotide diversity patterns in tomato. Plant Journal, 2017, 91, 631-645.	5.7	34
22	The Evolutionary Interplay between Adaptation and Self-Fertilization. Trends in Genetics, 2017, 33, 420-431.	6.7	64
23	Using the Ornstein–Uhlenbeck process to model the evolution of interacting populations. Journal of Theoretical Biology, 2017, 429, 35-45.	1.7	18
24	Genetic Diversity and the Efficacy of Purifying Selection across Plant and Animal Species. Molecular Biology and Evolution, 2017, 34, 1417-1428.	8.9	142
25	Inference of Distribution of Fitness Effects and Proportion of Adaptive Substitutions from Polymorphism Data. Genetics, 2017, 207, 1103-1119.	2.9	134
26	A large set of 26 new reference transcriptomes dedicated to comparative population genomics in crops and wild relatives. Molecular Ecology Resources, 2017, 17, 565-580.	4.8	53
27	Evolutionary forces affecting synonymous variations in plant genomes. PLoS Genetics, 2017, 13, e1006799.	3.5	36
28	The influence of population structure on gene expression and flowering time variation in the ubiquitous weed ⟨i⟩⟨scp⟩apsella bursaâ€pastoris⟨/i⟩ (Brassicaceae). Molecular Ecology, 2016, 25, 1106-1121.	3.9	36
29	Limits to Adaptation in Partially Selfing Species. Genetics, 2016, 203, 959-974.	2.9	47
30	Introns Structure Patterns of Variation in Nucleotide Composition in <i>Arabidopsis thaliana</i> Rice Protein-Coding Genes. Genome Biology and Evolution, 2015, 7, 2913-2928.	2.5	17
31	Molecular Evolution of Freshwater Snails with Contrasting Mating Systems. Molecular Biology and Evolution, 2015, 32, 2403-2416.	8.9	54
32	Quantification of GC-biased gene conversion in the human genome. Genome Research, 2015, 25, 1215-1228.	5.5	127
33	Inference of Purifying and Positive Selection in Three Subspecies of Chimpanzees (Pan troglodytes) from Exome Sequencing. Genome Biology and Evolution, 2015, 7, 1122-1132.	2.5	33
34	Extreme Recombination Frequencies Shape Genome Variation and Evolution in the Honeybee, Apis mellifera. PLoS Genetics, 2015, 11, e1005189.	3.5	98
35	The Red Queen Model of Recombination Hotspots Evolution in the Light of Archaic and Modern Human Genomes. PLoS Genetics, 2014, 10, e1004790.	3.5	62
36	Hitchhiking of Deleterious Alleles and the Cost of Adaptation in Partially Selfing Species. Genetics, 2014, 196, 281-293.	2.9	40

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37	Transcriptome population genomics reveals severe bottleneck and domestication cost in the <scp>A</scp> frican rice ( <i><scp>O</scp>ryza glaberrima</i> ). Molecular Ecology, 2014, 23, 2210-2227.	3.9	<b>7</b> 5
38	GC content evolution in coding regions of angiosperm genomes: a unifying hypothesis. Trends in Genetics, 2014, 30, 263-270.	6.7	79
39	The Rate of Molecular Adaptation in a Changing Environment. Molecular Biology and Evolution, 2013, 30, 1292-1301.	8.9	51
40	ADAPTATION AND MALADAPTATION IN SELFING AND OUTCROSSING SPECIES: NEW MUTATIONS VERSUS STANDING VARIATION. Evolution; International Journal of Organic Evolution, 2013, 67, 225-240.	2.3	128
41	Maleâ€specific <scp>DNA</scp> markers provide genetic evidence of an <scp>XY</scp> chromosome system, a recombination arrest and allow the tracing of paternal lineages in date palm. New Phytologist, 2013, 197, 409-415.	7.3	88
42	Reference-Free Population Genomics from Next-Generation Transcriptome Data and the Vertebrate–Invertebrate Gap. PLoS Genetics, 2013, 9, e1003457.	3.5	157
43	Patterns and Evolution of Nucleotide Landscapes in Seed Plants. Plant Cell, 2012, 24, 1379-1397.	6.6	88
44	Extinction and fixation times with dominance and inbreeding. Theoretical Population Biology, 2012, 81, 310-316.	1.1	30
45	Multigenic phylogeny and analysis of tree incongruences in Triticeae (Poaceae). BMC Evolutionary Biology, 2011, 11, 181.	3.2	72
46	COMPLEXITY, PLEIOTROPY, AND THE FITNESS EFFECT OF MUTATIONS. Evolution; International Journal of Organic Evolution, 2011, 65, 1559-1571.	2.3	57
47	GC-Biased Gene Conversion Impacts Ribosomal DNA Evolution in Vertebrates, Angiosperms, and Other Eukaryotes. Molecular Biology and Evolution, 2011, 28, 2561-2575.	8.9	53
48	GC-Biased Gene Conversion and Selection Affect GC Content in the Oryza Genus (rice). Molecular Biology and Evolution, 2011, 28, 2695-2706.	8.9	83
49	Surprising Fitness Consequences of GC-Biased Gene Conversion. II. Heterosis. Genetics, 2011, 187, 217-227.	2.9	10
50	Interaction of climate, demography and genetics: a ten-year study of Brassica insularis, a narrow endemic Mediterranean species. Conservation Genetics, 2010, 11, 509-526.	1.5	12
51	Surprising Fitness Consequences of GC-Biased Gene Conversion: I. Mutation Load and Inbreeding Depression. Genetics, 2010, 185, 939-959.	2.9	49
52	Mitochondrial whims: metabolic rate, longevity and the rate of molecular evolution. Biology Letters, 2009, 5, 413-416.	2.3	90
53	GC-biased gene conversion promotes the fixation of deleterious amino acid changes in primates. Trends in Genetics, 2009, 25, $1$ -5.	6.7	167
54	A comparative view of the evolution of grasses under domestication. New Phytologist, 2009, 183, 273-290.	7.3	193

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55	Consequences of Low Mate Availability in the Rare Selfâ€Incompatible Species <i>Brassica insularis</i> Conservation Biology, 2008, 22, 216-221.	4.7	33
56	Determination of Mitochondrial Genetic Diversity in Mammals. Genetics, 2008, 178, 351-361.	2.9	107
57	Mating Systems and the Efficacy of Selection at the Molecular Level. Genetics, 2007, 177, 905-916.	2.9	98
58	Strong Variations of Mitochondrial Mutation Rate across Mammals—the Longevity Hypothesis. Molecular Biology and Evolution, 2007, 25, 120-130.	8.9	394
59	Population Size Does Not Influence Mitochondrial Genetic Diversity in Animals. Science, 2006, 312, 570-572.	12.6	773
60	Bio++: a set of C++ libraries for sequence analysis, phylogenetics, molecular evolution and population genetics. BMC Bioinformatics, 2006, 7, 188.	2.6	101
61	Impact of mating systems on patterns of sequence polymorphism in flowering plants. Proceedings of the Royal Society B: Biological Sciences, 2006, 273, 3011-3019.	2.6	249
62	Lethals in subdivided populations. Genetical Research, 2005, 86, 41-51.	0.9	13
63	Plant selfâ€incompatibility systems: a molecular evolutionary perspective. New Phytologist, 2005, 168, 61-69.	7.3	136
64	Balancing Selection in the Wild: Testing Population Genetics Theory of Self-Incompatibility in the Rare Species Brassica insularis. Genetics, 2005, 171, 279-289.	2.9	74
65	When Genes Go to Sleep: The Population Genetic Consequences of Seed Dormancy and Monocarpic Perenniality. American Naturalist, 2004, 163, 295-311.	2.1	87
66	Patterns of Inbreeding Depression and Architecture of the Load in Subdivided Populations. Genetics, 2003, 165, 2193-2212.	2.9	100
67	Inbreeding Depression in Small Populations of Self-Incompatible Plants. Genetics, 2001, 159, 1217-1229.	2.9	63