

# Robert Debuchy

## List of Publications by Year in descending order

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45  
papers

2,372  
citations

218677

26  
h-index

254184

43  
g-index

47  
all docs

47  
docs citations

47  
times ranked

1989  
citing authors

#	ARTICLE	IF	CITATIONS
1	Population genomics of apricots unravels domestication history and adaptive events. <i>Nature Communications</i> , 2021, 12, 3956.	12.8	45
2	RNAi-Related Dicer and Argonaute Proteins Play Critical Roles for Meicyte Formation, Chromosome-Axes Lengths and Crossover Patterning in the Fungus <i>Sordaria macrospora</i> . <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 684108.	3.7	5
3	Comparative genomics applied to <i>Mucor</i> species with different lifestyles. <i>BMC Genomics</i> , 2020, 21, 135.	2.8	23
4	The taxonomy of the model filamentous fungus <i>Podospora anserina</i> . <i>MycKeys</i> , 2020, 75, 51-69.	1.9	6
5	A RID-like putative cytosine methyltransferase homologue controls sexual development in the fungus <i>Podospora anserina</i> . <i>PLoS Genetics</i> , 2019, 15, e1008086.	3.5	16
6	The mitochondrial translocase of the inner membrane PaTim54 is involved in defense response and longevity in <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2019, 132, 103257.	2.1	4
7	Whole-genome sequencing reveals recent and frequent genetic recombination between clonal lineages of <i>Cryphonectria parasitica</i> in western Europe. <i>Fungal Genetics and Biology</i> , 2019, 130, 122-133.	2.1	16
8	Building bridges to move recombination complexes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 12400-12409.	7.1	39
9	Population Genome Sequencing of the Scab Fungal Species <i>Venturia inaequalis</i> , <i>Venturia pirina</i> , <i>Venturia aucupariae</i> and <i>Venturia asperata</i> . <i>G3: Genes, Genomes, Genetics</i> , 2019, 9, 2405-2414.	1.8	33
10	A gene graveyard in the genome of the fungus <i>Podospora comata</i> . <i>Molecular Genetics and Genomics</i> , 2019, 294, 177-190.	2.1	29
11	PaPro1 and IDC4, Two Genes Controlling Stationary Phase, Sexual Development and Cell Degeneration in <i>Podospora anserina</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2018, 4, 85.	3.5	19
12	Asy2/Mer2: an evolutionarily conserved mediator of meiotic recombination, pairing, and global chromosome compaction. <i>Genes and Development</i> , 2017, 31, 1880-1893.	5.9	62
13	Inositol-phosphate signaling as mediator for growth and sexual reproduction in <i>Podospora anserina</i> . <i>Developmental Biology</i> , 2017, 429, 285-305.	2.0	6
14	Adaptive Horizontal Gene Transfers between Multiple Cheese-Associated Fungi. <i>Current Biology</i> , 2015, 25, 2562-2569.	3.9	110
15	Maintaining Two Mating Types: Structure of the Mating Type Locus and Its Role in Heterokaryosis in <i>Podospora anserina</i> . <i>Genetics</i> , 2014, 197, 421-432.	2.9	69
16	Multiple recent horizontal transfers of a large genomic region in cheese making fungi. <i>Nature Communications</i> , 2014, 5, 2876.	12.8	195
17	<i>Dra2</i> transposition in <i>Deinococcus radiodurans</i> is downregulated by <i>TnpB</i> . <i>Molecular Microbiology</i> , 2013, 88, 443-455.	2.5	46
18	A Network of HMG-box Transcription Factors Regulates Sexual Cycle in the Fungus <i>Podospora anserina</i> . <i>PLoS Genetics</i> , 2013, 9, e1003642.	3.5	58

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19	Systematic Deletion of Homeobox Genes in <i>Podospora anserina</i> Uncovers Their Roles in Shaping the Fruiting Body. <i>PLoS ONE</i> , 2012, 7, e37488.	2.5	37
20	Genome-Wide Gene Expression Profiling of Fertilization Competent Mycelium in Opposite Mating Types in the Heterothallic Fungus <i>Podospora anserina</i> . <i>PLoS ONE</i> , 2011, 6, e21476.	2.5	51
21	The importomer peroxins are differentially required for peroxisome assembly and meiotic development in <i>Podospora anserina</i> : insights into a new peroxisome import pathway. <i>Molecular Microbiology</i> , 2011, 82, 365-377.	2.5	50
22	A general framework for optimization of probes for gene expression microarray and its application to the fungus <i>Podospora anserina</i> . <i>BMC Research Notes</i> , 2010, 3, 171.	1.4	16
23	Tracing the Origin of the Fungal $\hat{1}$ Domain Places Its Ancestor in the HMG-Box Superfamily: Implication for Fungal Mating-Type Evolution. <i>PLoS ONE</i> , 2010, 5, e15199.	2.5	93
24	Gene deletion and allelic replacement in the filamentous fungus <i>Podospora anserina</i> . <i>Current Genetics</i> , 2008, 53, 249-258.	1.7	102
25	The genome sequence of the model ascomycete fungus <i>Podospora anserina</i> . <i>Genome Biology</i> , 2008, 9, R77.	9.6	301
26	Mutations in mating-type genes greatly decrease repeat-induced point mutation process in the fungus <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2008, 45, 207-220.	2.1	9
27	IDC1, a Pezizomycotina-specific gene that belongs to the PaMpk1 MAP kinase transduction cascade of the filamentous fungus <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2007, 44, 1219-1230.	2.1	53
28	The Function of the Coding Sequences for the Putative Pheromone Precursors in <i>Podospora anserina</i> Is Restricted to Fertilization. <i>Eukaryotic Cell</i> , 2005, 4, 407-420.	3.4	70
29	Altering a Gene Involved in Nuclear Distribution Increases the Repeat-Induced Point Mutation Process in the Fungus <i>Podospora anserina</i> . <i>Genetics</i> , 2004, 167, 151-159.	2.9	29
30	Characterization of the genomic organization of the region bordering the centromere of chromosome V of <i>Podospora anserina</i> by direct sequencing. <i>Fungal Genetics and Biology</i> , 2003, 39, 250-263.	2.1	25
31	<i>pah1</i> : a homeobox gene involved in hyphal morphology and microconidiogenesis in the filamentous ascomycete <i>Podospora anserina</i> . <i>Molecular Microbiology</i> , 2001, 39, 54-64.	2.5	51
32	Mutations in Mating-Type Genes of the Heterothallic Fungus <i>Podospora anserina</i> Lead to Self-Fertility. <i>Genetics</i> , 2001, 159, 545-556.	2.9	32
33	Co-expression of the Mating-Type Genes Involved in Internuclear Recognition Is Lethal in <i>Podospora anserina</i> . <i>Genetics</i> , 2000, 155, 657-669.	2.9	73
34	Internuclear Recognition: A Possible Connection between Euascomycetes and Homobasidiomycetes. <i>Fungal Genetics and Biology</i> , 1999, 27, 218-223.	2.1	56
35	A homologue of the yeast SHE4 gene is essential for the transition between the syncytial and cellular stages during sexual reproduction of the fungus <i>Podospora anserina</i> . <i>EMBO Journal</i> , 1998, 17, 1248-1258.	7.8	56
36	What is a. <i>Molecular Genetics and Genomics</i> , 1997, 256, 169.	2.4	2

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37	The mat- allele of <i>Podospora anserina</i> contains three regulatory genes required for the development of fertilized female organs. <i>Molecular Genetics and Genomics</i> , 1993, 241-241, 667-673.	2.4	96
38	The mating types of <i>Podospora anserina</i> : functional analysis and sequence of the fertilization domains. <i>Molecular Genetics and Genomics</i> , 1992, 233, 113-121.	2.4	113
39	Studies on the maintenance and expression of cloned DNA fragments in the nuclear genome of the green alga <i>Chlamydomonas Reinhardtii</i> . <i>Physiologia Plantarum</i> , 1990, 78, 254-260.	5.2	37
40	Studies on the maintenance and expression of cloned DNA fragments in the nuclear genome of the green alga <i>Chlamydomonas reinhardtii</i> . <i>Physiologia Plantarum</i> , 1990, 78, 254-260.	5.2	29
41	Chromosome walking towards a centromere in the filamentous fungus <i>Podospora anserina</i> : cloning of a sequence lethal at a two-copy state. <i>Current Genetics</i> , 1988, 13, 105-111.	1.7	14
42	Transformation by integration in <i>Podospora anserina</i> . <i>Molecular Genetics and Genomics</i> , 1987, 210, 129-134.	2.4	32
43	Transformation by integration in <i>Podospora anserina</i> . <i>Molecular Genetics and Genomics</i> , 1985, 200, 128.	2.4	56
44	Mating Systems and Sexual Morphogenesis in Ascomycetes. , 0, , 499-535.		99
45	<i>Cochliobolus</i> and <i>Podospora</i> : Mechanisms of Sex Determination and the Evolution of Reproductive Lifestyle. , 0, , 91-121.		6