

Philippe Silar

List of Publications by Year in descending order

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97
papers

3,514
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172457

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docs citations

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times ranked

2734
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#	ARTICLE	IF	CITATIONS
1	A gene cluster with positive and negative elements controls bistability and hysteresis of the crippled versus normal growth in the fungus <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2022, 161, 103711.	2.1	0
2	Recombination suppression and evolutionary strata around mating-type loci in fungi: documenting patterns and understanding evolutionary and mechanistic causes. <i>New Phytologist</i> , 2021, 229, 2470-2491.	7.3	46
3	Lignin degradation by ascomycetes. <i>Advances in Botanical Research</i> , 2021, 99, 77-113.	1.1	11
4	OSIP1 is a self-assembling DUF3129 protein required to protect fungal cells from toxins and stressors. <i>Environmental Microbiology</i> , 2021, 23, 1594-1607.	3.8	3
5	Size Variation of the Nonrecombining Region on the Mating-Type Chromosomes in the Fungal <i>Podospora anserina</i> Species Complex. <i>Molecular Biology and Evolution</i> , 2021, 38, 2475-2492.	8.9	13
6	(2803) Proposal to change the conserved type of <i>Podospora</i> , nom. cons. (<i>Ascomycota</i>). <i>Taxon</i> , 2021, 70, 429-430.	0.7	3
7	Important role of melanin for fertility in the fungus <i>Podospora anserina</i> . <i>G3: Genes, Genomes, Genetics</i> , 2021, 11, .	1.8	2
8	Lignin Degradation and Its Use in Signaling Development by the Coprophilous Ascomycete <i>Podospora anserina</i> . <i>Journal of Fungi (Basel, Switzerland)</i> , 2020, 6, 278.	3.5	11
9	The taxonomy of the model filamentous fungus <i>Podospora anserina</i> . <i>MycKeys</i> , 2020, 75, 51-69.	1.9	6
10	Appressorium: The Breakthrough in Dikarya. <i>Journal of Fungi (Basel, Switzerland)</i> , 2019, 5, 72.	3.5	27
11	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
12	Phenotypic instability in fungi. <i>Advances in Applied Microbiology</i> , 2019, 107, 141-187.	2.4	10
13	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
14	Characterization of three multicopper oxidases in the filamentous fungus <i>Podospora anserina</i> : A new role of an ABR1-like protein in fungal development?. <i>Fungal Genetics and Biology</i> , 2018, 116, 1-13.	2.1	23
15	Biomolecules from olive pruning waste in Sierra Mágina “Engaging the energy transition by multi-actor and multidisciplinary analyses. <i>Journal of Environmental Management</i> , 2018, 216, 204-213.	7.8	9
16	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
17	Cyclooxygenases and lipoxygenases are used by the fungus <i>Podospora anserina</i> to repel nematodes. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2018, 1862, 2174-2182.	2.4	10
18	Identification and characterization of PDC1, a novel protein involved in the epigenetic cell degeneration Crippled Growth in <i>Podospora anserina</i> . <i>Molecular Microbiology</i> , 2018, 110, 499-512.	2.5	5

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19	IDC2 and IDC3 , two genes involved in cell non-autonomous signaling of fruiting body development in the model fungus <i>Podospora anserina</i> . <i>Developmental Biology</i> , 2017, 421, 126-138.	2.0	19
20	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
21	Inactivation of Cellobiose Dehydrogenases Modifies the Cellulose Degradation Mechanism of <i>Podospora anserina</i> . <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	13
22	SymB and SymC, two membrane associated proteins, are required for <i>Podospora anserina</i> hyphal cell-cell fusion and maintenance of a mutualistic interaction with <i>Aspergillus oryzae</i> . <i>Molecular Microbiology</i> , 2017, 103, 657-677.	2.5	23
23	Prions and Prion-Like Phenomena in Epigenetic Inheritance. , 2017, , 61-72.		4
24	Species Delimitation in the <i>Podospora anserina</i> / <i>p. pauciseta</i> / <i>p. comata</i> Species Complex (Sordariales). <i>Cryptogamie, Mycologie</i> , 2017, 38, 485-506.	1.0	21
25	Plant biomass degrading ability of the coprophilic ascomycete fungus <i>Podospora anserina</i> . <i>Biotechnology Advances</i> , 2016, 34, 976-983.	11.7	41
26	The PaPsr1 and PaWhi2 genes are members of the regulatory network that connect stationary phase to mycelium differentiation and reproduction in <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2016, 94, 1-10.	2.1	14
27	Étude de la dynamique de colonisation microbienne de produits de construction. <i>Materiaux Et Techniques</i> , 2016, 104, 507.	0.9	1
28	Maintaining heterokaryosis in pseudo-homothallic fungi. <i>Communicative and Integrative Biology</i> , 2015, 8, e994382.	1.4	14
29	Screen for soil fungi highly resistant to dichloroaniline uncovers mostly <i>Fusarium</i> species. <i>Fungal Genetics and Biology</i> , 2015, 81, 82-87.	2.1	6
30	Bilirubin oxidase-like proteins from <i>Podospora anserina</i> : promising thermostable enzymes for application in transformation of plant biomass. <i>Environmental Microbiology</i> , 2015, 17, 866-875.	3.8	26
31	Identification of <i>NoxD</i> / <i>P</i> ro41 as the homologue of the p22 ^{phox} NADPH oxidase subunit in fungi. <i>Molecular Microbiology</i> , 2015, 95, 1006-1024.	2.5	56
32	Gene replacement in <i>Penicillium roqueforti</i> . <i>Current Genetics</i> , 2015, 61, 203-210.	1.7	7
33	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
34	Genes That Bias Mendelian Segregation. <i>PLoS Genetics</i> , 2014, 10, e1004387.	3.5	80
35	Systematic gene deletions evidences that laccases are involved in several stages of wood degradation in the filamentous fungus <i>Podospora anserina</i> . <i>Environmental Microbiology</i> , 2014, 16, 141-161.	3.8	48
36	Multiple recent horizontal transfers of a large genomic region in cheese making fungi. <i>Nature Communications</i> , 2014, 5, 2876.	12.8	195

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37	Genetic control of anastomosis in <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2014, 70, 94-103.	2.1	23
38	<i>Greeneria saprophytica</i> sp. nov. on dead leaves of <i>Syzygium cumini</i> from Chiang Rai, Thailand. <i>Phytotaxa</i> , 2014, 184, 275.	0.3	4
39	Morphology and phylogeny of <i>Pseudorobillarda eucalypti</i> sp. nov., from Thailand. <i>Phytotaxa</i> , 2014, 176, 251.	0.3	15
40	Morphology and phylogeny of <i>Chaetospermum</i> (asexual coelomycetous Basidiomycota). <i>Phytotaxa</i> , 2014, 175, 61.	0.3	7
41	Simple Genetic Tools to Study Fruiting Body Development in Fungi. <i>The Open Mycology Journal</i> , 2014, 8, 148-155.	0.8	10
42	Biotransformation of <i>Trichoderma</i> spp. and Their Tolerance to Aromatic Amines, a Major Class of Pollutants. <i>Applied and Environmental Microbiology</i> , 2013, 79, 4719-4726.	3.1	29
43	Rab-GDI Complex Dissociation Factor Expressed through Translational Frameshifting in Filamentous Ascomycetes. <i>PLoS ONE</i> , 2013, 8, e73772.	2.5	11
44	<i>Podospora anserina</i> : From Laboratory to Biotechnology. <i>Soil Biology</i> , 2013, , 283-309.	0.8	24
45	A Non-Mendelian MAPK-Generated Hereditary Unit Controlled by a Second MAPK Pathway in <i>Podospora anserina</i> . <i>Genetics</i> , 2012, 191, 419-433.	2.9	55
46	The transcriptional response to the inactivation of the PaMpk1 and PaMpk2 MAP kinase pathways in <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2012, 49, 643-652.	2.1	26
47	The PaAlr1 magnesium transporter is required for ascospore development in <i>Podospora anserina</i> . <i>Fungal Biology</i> , 2012, 116, 1111-1118.	2.5	10
48	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
49	Hyphal Interference: Self Versus Non-self Fungal Recognition and Hyphal Death. , 2012, , 155-170.		4
50	Wood Utilization Is Dependent on Catalase Activities in the Filamentous Fungus <i>Podospora anserina</i> . <i>PLoS ONE</i> , 2012, 7, e29820.	2.5	46
51	Fungi as a promising tool for bioremediation of soils contaminated with aromatic amines, a major class of pollutants. <i>Nature Reviews Microbiology</i> , 2011, 9, 477-477.	28.6	16
52	Grafting as a method for studying development in the filamentous fungus <i>Podospora anserina</i> . <i>Fungal Biology</i> , 2011, 115, 793-802.	2.5	18
53	Prions and Prion-like Phenomena in Epigenetic Inheritance. , 2011, , 63-76.		2
54	Epigenetics of Eukaryotic Microbes. , 2011, , 185-201.		1

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55	Sme4 coiled-coil protein mediates synaptonemal complex assembly, recombinosome relocalization, and spindle pole body morphogenesis. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 10614-10619.	7.1	28
56	Insights into the Phylogeny of Arylamine N-Acetyltransferases in Fungi. Journal of Molecular Evolution, 2010, 71, 141-152.	1.8	23
57	A general framework for optimization of probes for gene expression microarray and its application to the fungus <i>Podospora anserina</i> . BMC Research Notes, 2010, 3, 171.	1.4	16
58	The Nox/Ferric reductase/Ferric reductase-like families of Eumycetes. Fungal Biology, 2010, 114, 766-777.	2.5	31
59	An Acetyltransferase Conferring Tolerance to Toxic Aromatic Amine Chemicals. Journal of Biological Chemistry, 2009, 284, 18726-18733.	3.4	44
60	Functions and regulation of the Nox family in the filamentous fungus <i>Podospora anserina</i> : a new role in cellulose degradation. Molecular Microbiology, 2009, 74, 480-496.	2.5	109
61	The genome sequence of the model ascomycete fungus <i>Podospora anserina</i> . Genome Biology, 2008, 9, R77.	9.6	301
62	The Crucial Role of the Pls1 Tetraspanin during Ascospore Germination in <i>Podospora anserina</i> Provides an Example of the Convergent Evolution of Morphogenetic Processes in Fungal Plant Pathogens and Saprobes. Eukaryotic Cell, 2008, 7, 1809-1818.	3.4	79
63	Convergent evolution of morphogenetic processes in fungi. Communicative and Integrative Biology, 2008, 1, 180-181.	1.4	16
64	PaTrx1 and PaTrx3, Two Cytosolic Thioredoxins of the Filamentous Ascomycete <i>Podospora anserina</i> Involved in Sexual Development and Cell Degeneration. Eukaryotic Cell, 2007, 6, 2323-2331.	3.4	23
65	IDC1, a Pezizomycotina-specific gene that belongs to the PaMpk1 MAP kinase transduction cascade of the filamentous fungus <i>Podospora anserina</i> . Fungal Genetics and Biology, 2007, 44, 1219-1230.	2.1	53
66	Identification of PaPKS1, a polyketide synthase involved in melanin formation and its use as a genetic tool in <i>Podospora anserina</i> . Mycological Research, 2007, 111, 901-908.	2.5	44
67	Regulation, Cell Differentiation and Protein-Based Inheritance. Cell Cycle, 2006, 5, 2584-2587.	2.6	9
68	A mitotically inheritable unit containing a MAP kinase module. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 13445-13450.	7.1	59
69	Peroxide accumulation and cell death in filamentous fungi induced by contact with a contestant. Mycological Research, 2005, 109, 137-149.	2.5	91
70	Genetic control of an epigenetic cell degeneration syndrome in <i>Podospora anserina</i> . Fungal Genetics and Biology, 2005, 42, 564-577.	2.1	52
71	Incomplete Penetrance and Variable Expressivity of a Growth Defect as a Consequence of Knocking Out Two K ⁺ Transporters in the Eumycete Fungus <i>Podospora anserina</i> . Genetics, 2004, 166, 125-133.	2.9	18
72	PaASK1, a Mitogen-Activated Protein Kinase Kinase Kinase That Controls Cell Degeneration and Cell Differentiation in <i>Podospora anserina</i> . Genetics, 2004, 166, 1241-1252.	2.9	54

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73	Bistability and hysteresis of the 'Secteur' differentiation are controlled by a two-gene locus in <i>Nectria haematococca</i> . <i>BMC Biology</i> , 2004, 2, 18.	3.8	34
74	Two NADPH oxidase isoforms are required for sexual reproduction and ascospore germination in the filamentous fungus <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2004, 41, 982-997.	2.1	239
75	Non-Mendelian determinants of morphology in fungi. <i>Current Opinion in Microbiology</i> , 2003, 6, 641-645.	5.1	13
76	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
77	NADPH oxidase: an enzyme for multicellularity?. <i>Trends in Microbiology</i> , 2003, 11, 9-12.	7.7	97
78	Cell degeneration in the model system <i>Podospora anserina</i> . <i>Biogerontology</i> , 2001, 2, 1-17.	3.9	27
79	eEF1A Controls Ascospore Differentiation Through Elevated Accuracy, but Controls Longevity and Fruiting Body Formation Through Another Mechanism in <i>Podospora anserina</i> . <i>Genetics</i> , 2001, 158, 1477-1489.	2.9	10
80	In vivo labelling of functional ribosomes reveals spatial regulation during starvation in <i>Podospora anserina</i> . <i>BMC Genetics</i> , 2000, 1, 3.	2.7	6
81	Deletion and dosage modulation of the eEF1A gene in <i>Podospora anserina</i> : effect on the life cycle. <i>Biogerontology</i> , 2000, 1, 47-54.	3.9	12
82	Non-conventional infectious elements in filamentous fungi. <i>Trends in Genetics</i> , 1999, 15, 141-145.	6.7	30
83	What Triggers Senescence in <i>Podospora anserina</i> ?. <i>Fungal Genetics and Biology</i> , 1999, 27, 26-35.	2.1	21
84	Propagation of a Novel Cytoplasmic, Infectious and Deleterious Determinant Is Controlled by Translational Accuracy in <i>Podospora anserina</i> . <i>Genetics</i> , 1999, 151, 87-95.	2.9	59
85	Identification of the Genes Encoding the Cytosolic Translation Release Factors from <i>Podospora anserina</i> and Analysis of Their Role During the Life Cycle. <i>Genetics</i> , 1998, 149, 1763-1775.	2.9	26
86	Cloning, Sequencing, and Transgenic Expression of <i>Podospora curvicolla</i> and <i>Sordaria macrospora</i> eEF1A Genes: Relationship between Cytosolic Translation and Longevity in Filamentous Fungi. <i>Fungal Genetics and Biology</i> , 1997, 22, 191-198.	2.1	14
87	Contribution of various classes of defective mitochondrial DNA molecules to senescence in <i>Podospora anserina</i> . <i>Current Genetics</i> , 1997, 31, 171-178.	1.7	26
88	Cytosolic Ribosomal Mutations That Abolish Accumulation of Circular Intron in the Mitochondria Without Preventing Senescence of <i>Podospora anserina</i> . <i>Genetics</i> , 1997, 145, 697-705.	2.9	55
89	Genes that control longevity in <i>Podospora anserina</i> . <i>Mechanisms of Ageing and Development</i> , 1996, 90, 183-193.	4.6	17
90	Response of <i>Drosophila</i> metallothionein promoters to metallic, heat shock and oxidative stresses. <i>FEBS Letters</i> , 1996, 380, 33-38.	2.8	41

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91	Rapid methods for nucleic acids extraction from Petri dish-grown mycelia. <i>Current Genetics</i> , 1994, 25, 122-123.	1.7	217
92	Is translational accuracy an out-dated topic?. <i>Trends in Genetics</i> , 1994, 10, 71-72.	6.7	8
93	Increased longevity of EF-1 \pm high-fidelity mutants in <i>Podospora anserina</i> . <i>Journal of Molecular Biology</i> , 1994, 235, 231-236.	4.2	89
94	New shuttle vectors for direct cloning in <i>Saccharomyces cerevisiae</i> . <i>Gene</i> , 1991, 104, 99-102.	2.2	8
95	Metallothionein Mto gene of <i>Drosophila melanogaster</i> : Structure and regulation. <i>Journal of Molecular Biology</i> , 1990, 215, 217-224.	4.2	52
96	Expression of the <i>Drosophila melanogaster</i> metallothionein genes in yeast. <i>FEBS Letters</i> , 1990, 269, 273-276.	2.8	14
97	Mating Systems and Sexual Morphogenesis in <i>Ascomycetes</i> . , 0, , 499-535.		99