

# Philippe Silar

## List of Publications by Year in descending order

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97  
papers

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all docs

100  
docs citations

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times ranked

2734  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | The genome sequence of the model ascomycete fungus <i>Podospora anserina</i> . <i>Genome Biology</i> , 2008, 9, R77.  | 9.6  | 301       |
| 2  | 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       |
| 3  | Rapid methods for nucleic acids extraction from Petri dish-grown mycelia. <i>Current Genetics</i> , 1994, 25, 122-123.  | 1.7  | 217       |
| 4  | Multiple recent horizontal transfers of a large genomic region in cheese making fungi. <i>Nature Communications</i> , 2014, 5, 2876.  | 12.8 | 195       |
| 5  | Functions and regulation of the Nox family in the filamentous fungus <i>Podospora anserina</i> : a new role in cellulose degradation. <i>Molecular Microbiology</i> , 2009, 74, 480-496.  | 2.5  | 109       |
| 6  | Mating Systems and Sexual Morphogenesis in Ascomycetes. , 0, , 499-535.   |      | 99        |
| 7  | NADPH oxidase: an enzyme for multicellularity?. <i>Trends in Microbiology</i> , 2003, 11, 9-12.   | 7.7  | 97        |
| 8  | Peroxide accumulation and cell death in filamentous fungi induced by contact with a contestant. <i>Mycological Research</i> , 2005, 109, 137-149.   | 2.5  | 91        |
| 9  | 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        |
| 10 | Genes That Bias Mendelian Segregation. <i>PLoS Genetics</i> , 2014, 10, e1004387.   | 3.5  | 80        |
| 11 | 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 Saprobies. <i>Eukaryotic Cell</i> , 2008, 7, 1809-1818. | 3.4  | 79        |
| 12 | 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        |
| 13 | A mitotically inheritable unit containing a MAP kinase module. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2006, 103, 13445-13450.   | 7.1  | 59        |
| 14 | 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        |
| 15 | Identification of NoxD<sup>P</sup>ro41 as the homologue of the p22<sup>phox</sup> NADPH oxidase subunit in fungi. <i>Molecular Microbiology</i> , 2015, 95, 1006-1024.  | 2.5  | 56        |
| 16 | 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        |
| 17 | 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        |
| 18 | PaASK1, a Mitogen-Activated Protein Kinase Kinase Kinase That Controls Cell Degeneration and Cell Differentiation in <i>Podospora anserina</i> . <i>Genetics</i> , 2004, 166, 1241-1252.  | 2.9  | 54        |

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|----|--|------|-----------|
| 19 | 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        |
| 20 | Metallothionein Mto gene of <i>Drosophila melanogaster</i> : Structure and regulation. <i>Journal of Molecular Biology</i> , 1990, 215, 217-224.   | 4.2  | 52        |
| 21 | Genetic control of an epigenetic cell degeneration syndrome in <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2005, 42, 564-577.   | 2.1  | 52        |
| 22 | 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        |
| 23 | 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        |
| 24 | 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        |
| 25 | Identification of PaPKS1, a polyketide synthase involved in melanin formation and its use as a genetic tool in <i>Podospora anserina</i> . <i>Mycological Research</i> , 2007, 111, 901-908.   | 2.5  | 44        |
| 26 | An Acetyltransferase Conferring Tolerance to Toxic Aromatic Amine Chemicals. <i>Journal of Biological Chemistry</i> , 2009, 284, 18726-18733.  | 3.4  | 44        |
| 27 | 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        |
| 28 | Plant biomass degrading ability of the coprophilic ascomycete fungus <i>Podospora anserina</i> . <i>Biotechnology Advances</i> , 2016, 34, 976-983.  | 11.7 | 41        |
| 29 | 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        |
| 30 | 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        |
| 31 | The Nox/Ferric reductase/Ferric reductase-like families of Eumycetes. <i>Fungal Biology</i> , 2010, 114, 766-777.  | 2.5  | 31        |
| 32 | Non-conventional infectious elements in filamentous fungi. <i>Trends in Genetics</i> , 1999, 15, 141-145.  | 6.7  | 30        |
| 33 | 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        |
| 34 | 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        |
| 35 | Sme4 coiled-coil protein mediates synaptonemal complex assembly, recombinosome relocalization, and spindle pole body morphogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011, 108, 10614-10619. | 7.1  | 28        |
| 36 | Cell degeneration in the model system <i>Podospora anserina</i> . <i>Biogerontology</i> , 2001, 2, 1-17.   | 3.9  | 27        |

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|----|---|-----|-----------|
| 37 | Appressorium: The Breakthrough in Dikarya. <i>Journal of Fungi</i> (Basel, Switzerland), 2019, 5, 72.   | 3.5 | 27        |
| 38 | 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        |
| 39 | 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        |
| 40 | 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        |
| 41 | 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        |
| 42 | 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        |
| 43 | <i>Podospora anserina</i> : From Laboratory to Biotechnology. <i>Soil Biology</i> , 2013, , 283-309.  | 0.8 | 24        |
| 44 | PaTrx1 and PaTrx3, Two Cytosolic Thioredoxins of the Filamentous Ascomycete <i>Podospora anserina</i> Involved in Sexual Development and Cell Degeneration. <i>Eukaryotic Cell</i> , 2007, 6, 2323-2331.  | 3.4 | 23        |
| 45 | Insights into the Phylogeny of Arylamine N-Acetyltransferases in Fungi. <i>Journal of Molecular Evolution</i> , 2010, 71, 141-152.  | 1.8 | 23        |
| 46 | Genetic control of anastomosis in <i>Podospora anserina</i> . <i>Fungal Genetics and Biology</i> , 2014, 70, 94-103.  | 2.1 | 23        |
| 47 | SymB and SymC, two membrane associated proteins, are required for <i>Podospora anserina</i> hyphal cell fusion and maintenance of a mutualistic interaction with <i>Trichostema album</i> . <i>Molecular Microbiology</i> , 2017, 103, 657-677. | 2.5 | 23        |
| 48 | 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        |
| 49 | What Triggers Senescence in <i>Podospora anserina</i> ?. <i>Fungal Genetics and Biology</i> , 1999, 27, 26-35.  | 2.1 | 21        |
| 50 | Species Delimitation in the <i>Podospora anserina</i> / <i>p. paucisetata</i> / <i>p. comata</i> Species Complex (Sordariales). <i>Cryptogamie, Mycologie</i> , 2017, 38, 485-506.  | 1.0 | 21        |
| 51 | 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        |
| 52 | PaPro1 and IDC4, Two Genes Controlling Stationary Phase, Sexual Development and Cell Degeneration in <i>Podospora anserina</i> . <i>Journal of Fungi</i> (Basel, Switzerland), 2018, 4, 85.   | 3.5 | 19        |
| 53 | Incomplete Penetrance and Variable Expressivity of a Growth Defect as a Consequence of Knocking Out Two K <sup>+</sup> Transporters in the Eukaryotic Fungus <i>Podospora anserina</i> . <i>Genetics</i> , 2004, 166, 125-133.                  | 2.9 | 18        |
| 54 | 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        |

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|----|--|------|-----------|
| 55 | Genes that control longevity in <i>Podospora anserina</i> . <i>Mechanisms of Ageing and Development</i> , 1996, 90, 183-193.   | 4.6  | 17        |
| 56 | Convergent evolution of morphogenetic processes in fungi. <i>Communicative and Integrative Biology</i> , 2008, 1, 180-181.   | 1.4  | 16        |
| 57 | 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        |
| 58 | 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        |
| 59 | Morphology and phylogeny of <i>Pseudorobillarda eucalypti</i> sp. nov., from Thailand. <i>Phytotaxa</i> , 2014, 176, 251.  | 0.3  | 15        |
| 60 | Expression of the <i>Drosophila melanogaster</i> metallothionein genes in yeast. <i>FEBS Letters</i> , 1990, 269, 273-276.   | 2.8  | 14        |
| 61 | 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        |
| 62 | Maintaining heterokaryosis in pseudo-homothallic fungi. <i>Communicative and Integrative Biology</i> , 2015, 8, e994382.   | 1.4  | 14        |
| 63 | 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        |
| 64 | Non-Mendelian determinants of morphology in fungi. <i>Current Opinion in Microbiology</i> , 2003, 6, 641-645.  | 5.1  | 13        |
| 65 | 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        |
| 66 | 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        |
| 67 | 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        |
| 68 | Rab-GDI Complex Dissociation Factor Expressed through Translational Frameshifting in Filamentous Ascomycetes. <i>PLoS ONE</i> , 2013, 8, e73772.   | 2.5  | 11        |
| 69 | 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        |
| 70 | Lignin degradation by ascomycetes. <i>Advances in Botanical Research</i> , 2021, 99, 77-113.   | 1.1  | 11        |
| 71 | 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        |
| 72 | 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        |

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|----|---|-----|-----------|
| 73 | Phenotypic instability in fungi. <i>Advances in Applied Microbiology</i> , 2019, 107, 141-187.  | 2.4 | 10        |
| 74 | Simple Genetic Tools to Study Fruiting Body Development in Fungi. <i>The Open Mycology Journal</i> , 2014, 8, 148-155.  | 0.8 | 10        |
| 75 | 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        |
| 76 | Regulation, Cell Differentiation and Protein-Based Inheritance. <i>Cell Cycle</i> , 2006, 5, 2584-2587.   | 2.6 | 9         |
| 77 | 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         |
| 78 | New shuttle vectors for direct cloning in <i>Saccharomyces cerevisiae</i> . <i>Gene</i> , 1991, 104, 99-102.  | 2.2 | 8         |
| 79 | Is translational accuracy an out-dated topic?. <i>Trends in Genetics</i> , 1994, 10, 71-72.   | 6.7 | 8         |
| 80 | Morphology and phylogeny of <i>Chaetospermum</i> (asexual coelomycetous Basidiomycota). <i>Phytotaxa</i> , 2014, 175, 61.   | 0.3 | 7         |
| 81 | Gene replacement in <i>Penicillium roqueforti</i> . <i>Current Genetics</i> , 2015, 61, 203-210.  | 1.7 | 7         |
| 82 | 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         |
| 83 | 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         |
| 84 | 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         |
| 85 | The taxonomy of the model filamentous fungus <i>Podospora anserina</i> . <i>MycKeys</i> , 2020, 75, 51-69.  | 1.9 | 6         |
| 86 | 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         |
| 87 | <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         |
| 88 | Prions and Prion-Like Phenomena in Epigenetic Inheritance. , 2017, , 61-72.   |     | 4         |
| 89 | 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         |
| 90 | Hyphal Interference: Self Versus Non-self Fungal Recognition and Hyphal Death. , 2012, , 155-170.   |     | 4         |

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|----|---|-----|-----------|
| 91 | OSIP1 is a self-assembling DUF3129 protein required to protect fungal cells from toxins and stressors. Environmental Microbiology, 2021, 23, 1594-1607.   | 3.8 | 3         |
| 92 | (2803) Proposal to change the conserved type of <i>Podospora</i> , nom. cons. ( <i>Ascomycota</i> ). Taxon, 2021, 70, 429-430.  | 0.7 | 3         |
| 93 | Prions and Prion-like Phenomena in Epigenetic Inheritance. , 2011, , 63-76.   |     | 2         |
| 94 | Important role of melanin for fertility in the fungus <i>Podospora anserina</i> . G3: Genes, Genomes, Genetics, 2021, 11, .   | 1.8 | 2         |
| 95 | Epigenetics of Eukaryotic Microbes. , 2011, , 185-201.  |     | 1         |
| 96 | Étude de la dynamique de colonisation microbienne de produits de construction. Matériaux Et Techniques, 2016, 104, 507.   | 0.9 | 1         |
| 97 | 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> . Fungal Genetics and Biology, 2022, 161, 103711. | 2.1 | 0         |