

# Daniel G Panaccione

## List of Publications by Year in descending order

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78  
papers

4,285  
citations

109321

35  
h-index

110387

64  
g-index

81  
all docs

81  
docs citations

81  
times ranked

3430  
citing authors

#	ARTICLE	IF	CITATIONS
1	Contribution of a novel gene to lysergic acid amide synthesis in <i>Metarhizium brunneum</i> . <i>BMC Research Notes</i> , 2022, 15, 183.	1.4	4
2	A Baeyer-Villiger Monooxygenase Gene Involved in the Synthesis of Lysergic Acid Amides Affects the Interaction of the Fungus <i>Metarhizium brunneum</i> with Insects. <i>Applied and Environmental Microbiology</i> , 2021, 87, e0074821.	3.1	7
3	Independent Evolution of a Lysergic Acid Amide in <i>Aspergillus</i> Species. <i>Applied and Environmental Microbiology</i> , 2021, 87, e0180121.	3.1	6
4	Diversification of ergot alkaloids and heritable fungal symbionts in morning glories. <i>Communications Biology</i> , 2021, 4, 1362.	4.4	12
5	Genetic Reprogramming of the Ergot Alkaloid Pathway of <i>Metarhizium brunneum</i> . <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	15
6	Several <i>Metarhizium</i> Species Produce Ergot Alkaloids in a Condition-Specific Manner. <i>Applied and Environmental Microbiology</i> , 2020, 86, .	3.1	23
7	Endophytes matter: Variation of dung beetle performance across different endophyte-infected tall fescue cultivars. <i>Applied Soil Ecology</i> , 2020, 152, 103561.	4.3	5
8	Diversity and function of fungi associated with the fungivorous millipede, <i>Brachycybe lecontii</i> . <i>Fungal Ecology</i> , 2019, 41, 187-197.	1.6	17
9	Biodiversity of Convolvulaceous species that contain ergot alkaloids, indole diterpene alkaloids, and swainsonine. <i>Biochemical Systematics and Ecology</i> , 2019, 86, 103921.	1.3	10
10	Decreased Root-Knot Nematode Gall Formation in Roots of the Morning Glory <i>Ipomoea tricolor</i> Symbiotic with Ergot Alkaloid-Producing Fungal <i>Periglandula</i> Sp.. <i>Journal of Chemical Ecology</i> , 2019, 45, 879-887.	1.8	8
11	Psychoactive plant- and mushroom-associated alkaloids from two behavior modifying cicada pathogens. <i>Fungal Ecology</i> , 2019, 41, 147-164.	1.6	55
12	Molecular identification and characterization of endophytes from uncultivated barley. <i>Mycologia</i> , 2018, 110, 453-472.	1.9	7
13	Ergot Alkaloid Synthesis Capacity of <i>Penicillium camemberti</i> . <i>Applied and Environmental Microbiology</i> , 2018, 84, .	3.1	10
14	Biological activity of <i>Claviceps gigantea</i> in juvenile New Zealand rabbits. <i>Mycotoxin Research</i> , 2018, 34, 297-305.	2.3	0
15	Ergot Alkaloids of the Family Clavicipitaceae. <i>Phytopathology</i> , 2017, 107, 504-518.	2.2	76
16	Biosynthesis of the Pharmaceutically Important Fungal Ergot Alkaloid Dihydrolysergic Acid Requires a Specialized Allele of <i>cloA</i> . <i>Applied and Environmental Microbiology</i> , 2017, 83, .	3.1	14
17	Ergot alkaloids contribute to virulence in an insect model of invasive aspergillosis. <i>Scientific Reports</i> , 2017, 7, 8930.	3.3	36
18	Ergot Alkaloid Biosynthesis in the Maize ( <i>Zea mays</i> ) Ergot Fungus <i>Claviceps gigantea</i> . <i>Journal of Agricultural and Food Chemistry</i> , 2017, 65, 10703-10710.	5.2	9

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19	Toxin-producing <i>Epichloa bromicola</i> strains symbiotic with the forage grass <i>Elymus dahuricus</i> in China. <i>Mycologia</i> , 2017, 109, 847-859.	1.9	12
20	Chromosome-End Knockoff Strategy to Reshape Alkaloid Profiles of a Fungal Endophyte. <i>G3: Genes, Genomes, Genetics</i> , 2016, 6, 2601-2610.	1.8	19
21	Modulation of Ergot Alkaloids in a Grass-Endophyte Symbiosis by Alteration of mRNA Concentrations of an Ergot Alkaloid Synthesis Gene. <i>Journal of Agricultural and Food Chemistry</i> , 2016, 64, 4982-4989.	5.2	8
22	Functional analysis of the gene controlling hydroxylation of festuclavine in the ergot alkaloid pathway of <i>Neosartorya fumigata</i> . <i>Current Genetics</i> , 2016, 62, 853-860.	1.7	20
23	The role of fungi and invertebrates in litter decomposition in mitigated and reference wetlands. <i>Limnologica</i> , 2015, 54, 23-32.	1.5	18
24	Genetics, Genomics and Evolution of Ergot Alkaloid Diversity. <i>Toxins</i> , 2015, 7, 1273-1302.	3.4	83
25	Diversification of Ergot Alkaloids in Natural and Modified Fungi. <i>Toxins</i> , 2015, 7, 201-218.	3.4	49
26	Phylogenetic and chemotypic diversity of <i>Periglandula</i> species in eight new morning glory hosts (Convolvulaceae). <i>Mycologia</i> , 2015, 107, 667-678.	1.9	25
27	Identification and Structural Elucidation of Ergotryptamine, a New Ergot Alkaloid Produced by Genetically Modified <i>Aspergillus nidulans</i> and Natural Isolates of <i>Epichloa</i> Species. <i>Journal of Agricultural and Food Chemistry</i> , 2015, 63, 61-67.	5.2	18
28	Bioactive alkaloids in vertically transmitted fungal endophytes. <i>Functional Ecology</i> , 2014, 28, 299-314.	3.6	154
29	Accumulation of Ergot Alkaloids During Conidiophore Development in <i>Aspergillus fumigatus</i> . <i>Current Microbiology</i> , 2014, 68, 1-5.	2.2	17
30	Potential for Industrial Application of Microbes in Symbioses that Influence Plant Productivity and Sustainability in Agricultural, Natural, or Restored Ecosystems. <i>Industrial Biotechnology</i> , 2014, 10, 347-353.	0.8	1
31	Heterologous Expression of Lysergic Acid and Novel Ergot Alkaloids in <i>Aspergillus fumigatus</i> . <i>Applied and Environmental Microbiology</i> , 2014, 80, 6465-6472.	3.1	37
32	Differential Allocation of Seed-Borne Ergot Alkaloids During Early Ontogeny of Morning Glories (Convolvulaceae). <i>Journal of Chemical Ecology</i> , 2013, 39, 919-930.	1.8	26
33	Currencies of Mutualisms: Sources of Alkaloid Genes in Vertically Transmitted <i>Epichloae</i> . <i>Toxins</i> , 2013, 5, 1064-1088.	3.4	109
34	Plant-Symbiotic Fungi as Chemical Engineers: Multi-Genome Analysis of the Clavicipitaceae Reveals Dynamics of Alkaloid Loci. <i>PLoS Genetics</i> , 2013, 9, e1003323.	3.5	344
35	Partial Reconstruction of the Ergot Alkaloid Pathway by Heterologous Gene Expression in <i>Aspergillus nidulans</i> . <i>Toxins</i> , 2013, 5, 445-455.	3.4	46
36	Analysis and Modification of Ergot Alkaloid Profiles in Fungi. <i>Methods in Enzymology</i> , 2012, 515, 267-290.	1.0	42

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37	Lifestyle transitions in plant pathogenic <i>Colletotrichum</i> fungi deciphered by genome and transcriptome analyses. <i>Nature Genetics</i> , 2012, 44, 1060-1065.	21.4	840
38	Chemotypic and genotypic diversity in the ergot alkaloid pathway of <i>Aspergillus fumigatus</i> . <i>Mycologia</i> , 2012, 104, 804-812.	1.9	18
39	Ergot Alkaloids. , 2011, , 195-214.		1
40	Ergot cluster-encoded catalase is required for synthesis of chanoclavine-I in <i>Aspergillus fumigatus</i> . <i>Current Genetics</i> , 2011, 57, 201-211.	1.7	48
41	An Old Yellow Enzyme Gene Controls the Branch Point between <i>Aspergillus fumigatus</i> and <i>Claviceps purpurea</i> Ergot Alkaloid Pathways. <i>Applied and Environmental Microbiology</i> , 2010, 76, 3898-3903.	3.1	67
42	Controlling a Structural Branch Point in Ergot Alkaloid Biosynthesis. <i>Journal of the American Chemical Society</i> , 2010, 132, 12835-12837.	13.7	56
43	A Role for Old Yellow Enzyme in Ergot Alkaloid Biosynthesis. <i>Journal of the American Chemical Society</i> , 2010, 132, 1776-1777.	13.7	54
44	Contribution of ergot alkaloids to suppression of a grass-feeding caterpillar assessed with gene knockout endophytes in perennial ryegrass. <i>Entomologia Experimentalis Et Applicata</i> , 2008, 126, 138-147.	1.4	67
45	Association of ergot alkaloids with conidiation in <i>Aspergillus fumigatus</i> . <i>Mycologia</i> , 2007, 99, 804-811.	1.9	45
46	Association of ergot alkaloids with conidiation in <i>Aspergillus fumigatus</i> . <i>Mycologia</i> , 2007, 99, 804-811.	1.9	48
47	Chapter 2 Ergot Alkaloids " Biology and Molecular Biology. <i>The Alkaloids Chemistry and Biology</i> , 2006, 63, 45-86.	2.0	184
48	Effects of Ergot Alkaloids on Food Preference and Satiety in Rabbits, As Assessed with Gene-Knockout Endophytes in Perennial Ryegrass ( <i>Lolium perenne</i> ). <i>Journal of Agricultural and Food Chemistry</i> , 2006, 54, 4582-4587.	5.2	76
49	Pathways to Diverse Ergot Alkaloid Profiles in Fungi. <i>Recent Advances in Phytochemistry</i> , 2006, , 23-52.	0.5	3
50	Ergot alkaloids are not essential for endophytic fungus-associated population suppression of the lesion nematode, <i>Pratylenchus scribneri</i> , on perennial ryegrass. <i>Nematology</i> , 2006, 8, 583-590.	0.6	37
51	Origins and significance of ergot alkaloid diversity in fungi. <i>FEMS Microbiology Letters</i> , 2005, 251, 9-17.	1.8	89
52	The ergot alkaloid gene cluster in <i>Claviceps purpurea</i> : Extension of the cluster sequence and intra species evolution. <i>Phytochemistry</i> , 2005, 66, 1312-1320.	2.9	122
53	Abundant Respirable Ergot Alkaloids from the Common Airborne Fungus <i>Aspergillus fumigatus</i> . <i>Applied and Environmental Microbiology</i> , 2005, 71, 3106-3111.	3.1	85
54	Structural analysis of a peptide synthetase gene required for ergopeptide production in the endophytic fungus <i>Neotyphodium lolii</i> . <i>DNA Sequence</i> , 2005, 16, 379-385.	0.7	14

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55	An Ergot Alkaloid Biosynthesis Gene and Clustered Hypothetical Genes from <i>Aspergillus fumigatus</i> . <i>Applied and Environmental Microbiology</i> , 2005, 71, 3112-3118.	3.1	103
56	The determinant step in ergot alkaloid biosynthesis by an endophyte of perennial ryegrass. <i>Fungal Genetics and Biology</i> , 2004, 41, 189-198.	2.1	105
57	Biochemical Outcome of Blocking the Ergot Alkaloid Pathway of a Grass Endophyte. <i>Journal of Agricultural and Food Chemistry</i> , 2003, 51, 6429-6437.	5.2	53
58	Identification of differentially expressed genes in the mutualistic association of tall fescue with <i>Neotyphodium coenophialum</i> . <i>Physiological and Molecular Plant Pathology</i> , 2003, 63, 305-317.	2.5	69
59	Characterization of dilution enrichment cultures obtained from size-fractionated soil bacteria by BIOLOG <sup>®</sup> community-level physiological profiles and restriction analysis of 16S rRNA genes. <i>Soil Biology and Biochemistry</i> , 2001, 33, 1555-1562.	8.8	25
60	Diversity of <i>Cenococcum geophilum</i> isolates from serpentine and non-serpentine soils. <i>Mycologia</i> , 2001, 93, 645-652.	1.9	42
61	Elimination of ergovaline from a grass- <i>Neotyphodium</i> endophyte symbiosis by genetic modification of the endophyte. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 12820-12825.	7.1	164
62	Diversity of <i>Cenococcum geophilum</i> Isolates from Serpentine and Non-Serpentine Soils. <i>Mycologia</i> , 2001, 93, 645.	1.9	39
63	Significance of fungal peptide secondary metabolites in the agri-food industry. <i>Applied Mycology and Biotechnology</i> , 2001, 1, 115-143.	0.3	2
64	Organic acid eÅ—udation by <i>Laccaria bicolor</i> and <i>Pisolithus tinctorius</i> eÅ—posed to aluminum in vitro. <i>Canadian Journal of Forest Research</i> , 2001, 31, 703-710.	1.7	17
65	Presence of peptide synthetase gene transcripts and accumulation of ergopeptines in <i>Claviceps purpurea</i> and <i>Neotyphodium coenophialum</i> . <i>Canadian Journal of Microbiology</i> , 1998, 44, 80-86.	1.7	22
66	Presence of peptide synthetase gene transcripts and accumulation of ergopeptines in <i>Claviceps purpurea</i> and <i>Neotyphodium coenophialum</i> . <i>Canadian Journal of Microbiology</i> , 1998, 44, 80-86.	1.7	2
67	Metalaxyl stimulation of growth of isolates of <i>Phytophthora infestans</i> . <i>Mycologia</i> , 1997, 89, 289-292.	1.9	25
68	Transposon-like sequences at the TOX2 locus of the plant-pathogenic fungus <i>Cochliobolus carbonum</i> . <i>Gene</i> , 1996, 176, 103-109.	2.2	35
69	Multiple families of peptide synthetase genes from ergopeptide-producing fungi. <i>Mycological Research</i> , 1996, 100, 429-436.	2.5	30
70	Identification of peptide synthetase-encoding genes from filamentous fungi producing host-selective phytotoxins or analogs. <i>Gene</i> , 1995, 165, 207-211.	2.2	49
71	The fungal genus <i>Cochliobolus</i> and toxin-mediated plant disease. <i>Trends in Microbiology</i> , 1993, 1, 14-20.	7.7	17
72	Host-Selective Toxins and Disease Specificity: Perspectives and Progress. <i>Annual Review of Phytopathology</i> , 1993, 31, 275-303.	7.8	80

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73	The PYR1 gene of the plant pathogenic fungus <i>Colletotrichum graminicola</i> : selection by intraspecific complementation and sequence analysis. <i>Molecular Genetics and Genomics</i> , 1992, 235, 74-80.	2.4	18
74	Gene expression associated with light-induced conidiation in <i>Colletotrichum graminicola</i> . <i>Canadian Journal of Microbiology</i> , 1991, 37, 165-167.	1.7	3
75	Endopolygalacturonase Is Not Required for Pathogenicity of <i>Cochliobolus carbonum</i> on Maize. <i>Plant Cell</i> , 1990, 2, 1191.	6.6	53
76	Characterization of two divergent $\beta$ -tubulin genes from <i>Colletotrichum graminicola</i> . <i>Gene</i> , 1990, 86, 163-170.	2.2	61
77	Conidial Dimorphism in <i>Colletotrichum graminicola</i> . <i>Mycologia</i> , 1989, 81, 876.	1.9	21
78	<i>Colletotrichum graminicola</i> Transformed with Homologous and Heterologous Benomyl-Resistance Genes Retains Expected Pathogenicity to Corn. <i>Molecular Plant-Microbe Interactions</i> , 1988, 1, 113.	2.6	53