

Jan Stenlid

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

280
papers

15,657
citations

60
h-index

115
g-index

286
ext. papers

18,221
ext. citations

4.8
avg, IF

6.41
L-index

#	Paper	IF	Citations
280	The Paleozoic origin of enzymatic lignin decomposition reconstructed from 31 fungal genomes. <i>Science</i> , 2012 , 336, 1715-9	33.3	1129
279	New primers to amplify the fungal ITS2 region--evaluation by 454-sequencing of artificial and natural communities. <i>FEMS Microbiology Ecology</i> , 2012 , 82, 666-77	4.3	985
278	Roots and associated fungi drive long-term carbon sequestration in boreal forest. <i>Science</i> , 2013 , 339, 1615-8	33.3	866
277	Spatial separation of litter decomposition and mycorrhizal nitrogen uptake in a boreal forest. <i>New Phytologist</i> , 2007 , 173, 611-620	9.8	658
276	Fungal community analysis by high-throughput sequencing of amplified markers--a user's guide. <i>New Phytologist</i> , 2013 , 199, 288-299	9.8	601
275	The plant cell wall-decomposing machinery underlies the functional diversity of forest fungi. <i>Science</i> , 2011 , 333, 762-5	33.3	417
274	Biogeographical patterns and determinants of invasion by forest pathogens in Europe. <i>New Phytologist</i> , 2013 , 197, 238-250	9.8	351
273	Carbon sequestration is related to mycorrhizal fungal community shifts during long-term succession in boreal forests. <i>New Phytologist</i> , 2015 , 205, 1525-1536	9.8	339
272	Population structure of <i>Heterobasidion annosum</i> as determined by somatic incompatibility, sexual incompatibility, and isoenzyme patterns. <i>Canadian Journal of Botany</i> , 1985 , 63, 2268-2273		289
271	Comparative genomics of <i>Ceriporiopsis subvermispora</i> and <i>Phanerochaete chrysosporium</i> provide insight into selective ligninolysis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012 , 109, 5458-63	11.5	225
270	Widespread <i>Phytophthora</i> infestations in European nurseries put forest, semi-natural and horticultural ecosystems at high risk of <i>Phytophthora</i> diseases. <i>Forest Pathology</i> , 2016 , 46, 134-163	1.2	187
269	Conifer root and butt rot caused by <i>Heterobasidion annosum</i> (Fr.) Bref. s.l. <i>Molecular Plant Pathology</i> , 2005 , 6, 395-409	5.7	183
268	Intraspecific genetic variation in <i>Heterobasidion annosum</i> revealed by amplification of minisatellite DNA. <i>Mycological Research</i> , 1994 , 98, 57-63		171
267	Insight into trade-off between wood decay and parasitism from the genome of a fungal forest pathogen. <i>New Phytologist</i> , 2012 , 194, 1001-1013	9.8	168
266	Size, distribution and biomass of genets in populations of <i>Suillus bovinus</i> (L.: Fr.) Roussel revealed by somatic incompatibility. <i>New Phytologist</i> , 1994 , 128, 225-234	9.8	158
265	Population structure and dynamics in <i>Suillus bovinus</i> as indicated by spatial distribution of fungal clones. <i>New Phytologist</i> , 1990 , 115, 487-493	9.8	150
264	Biodiversity and ecosystem functioning relations in European forests depend on environmental context. <i>Ecology Letters</i> , 2017 , 20, 1414-1426	10	149

263	Diversity and abundance of resupinate theleporoid fungi as ectomycorrhizal symbionts in Swedish boreal forests. <i>Molecular Ecology</i> , 2000 , 9, 1985-96	5.7	148
262	A novel comparative research platform designed to determine the functional significance of tree species diversity in European forests. <i>Perspectives in Plant Ecology, Evolution and Systematics</i> , 2013 , 15, 281-291	3	143
261	Biotic homogenization can decrease landscape-scale forest multifunctionality. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 3557-62	11.5	134
260	Emerging pathogens: fungal host jumps following anthropogenic introduction. <i>Trends in Ecology and Evolution</i> , 2005 , 20, 420-1	10.9	129
259	Translocation of 32P between interacting mycelia of a wood-decomposing fungus and ectomycorrhizal fungi in microcosm systems. <i>New Phytologist</i> , 1999 , 144, 183-193	9.8	129
258	Patterns of fungal communities among and within decaying logs, revealed by 454 sequencing. <i>Molecular Ecology</i> , 2012 , 21, 4514-32	5.7	128
257	Replacing monocultures with mixed-species stands: Ecosystem service implications of two production forest alternatives in Sweden. <i>Ambio</i> , 2016 , 45 Suppl 2, 124-39	6.5	125
256	Jack-of-all-trades effects drive biodiversity-ecosystem multifunctionality relationships in European forests. <i>Nature Communications</i> , 2016 , 7, 11109	17.4	120
255	The effect of fungal pathogens on the water and carbon economy of trees: implications for drought-induced mortality. <i>New Phytologist</i> , 2014 , 203, 1028-1035	9.8	117
254	Fungal communities in mycorrhizal roots of conifer seedlings in forest nurseries under different cultivation systems, assessed by morphotyping, direct sequencing and mycelial isolation. <i>Mycorrhiza</i> , 2005 , 16, 33-41	3.9	115
253	Investigations concerning the role of <i>Chalara fraxinea</i> in declining <i>Fraxinus excelsior</i> . <i>Plant Pathology</i> , 2009 , 58, 284-292	2.8	110
252	Controlling and predicting the spread of heterobasidion annosum from infected stumps and trees of picea abies. <i>Scandinavian Journal of Forest Research</i> , 1987 , 2, 187-198	1.7	106
251	Ecology and molecular characterization of dark septate fungi from roots, living stems, coarse and fine woody debris. <i>Mycological Research</i> , 2004 , 108, 965-73		100
250	Occurrence and pathogenicity of fungi in necrotic and non-symptomatic shoots of declining common ash (<i>Fraxinus excelsior</i>) in Sweden. <i>European Journal of Forest Research</i> , 2009 , 128, 51-60	2.7	84
249	Retracing the routes of introduction of invasive species: the case of the <i>Sirex noctilio</i> woodwasp. <i>Molecular Ecology</i> , 2012 , 21, 5728-44	5.7	83
248	Spatiotemporal patterns in ectomycorrhizal populations. <i>Canadian Journal of Botany</i> , 1995 , 73, 1222-1230		83
247	Population Dynamics of the Root Rot Fungus <i>Heterobasidion annosum</i> Following Thinning of <i>Picea abies</i> . <i>Oikos</i> , 1993 , 66, 247	4	83
246	The importance of inoculum size for the competitive ability of wood decomposing fungi. <i>FEMS Microbiology Ecology</i> , 1993 , 12, 169-176	4.3	83

245	Wood-inhabiting fungi in stems of <i>Fraxinus excelsior</i> in declining ash stands of northern Lithuania, with particular reference to <i>Armillaria cepistipes</i> . <i>Scandinavian Journal of Forest Research</i> , 2005 , 20, 337-346	1.7	80
244	Partial intersterility in <i>Heterobasidion annosum</i> . <i>Mycological Research</i> , 1991 , 95, 1153-1159		79
243	Plant pathogens. Mitochondrial control of fungal hybrid virulence. <i>Nature</i> , 2001 , 411, 438	50.4	78
242	Towards standardization of the description and publication of next-generation sequencing datasets of fungal communities. <i>New Phytologist</i> , 2011 , 191, 314-318	9.8	77
241	Evolution of family 18 glycoside hydrolases: diversity, domain structures and phylogenetic relationships. <i>Journal of Molecular Microbiology and Biotechnology</i> , 2009 , 16, 208-23	0.9	75
240	Effects of resource availability on mycelial interactions and 32P transfer between a saprotrophic and an ectomycorrhizal fungus in soil microcosms. <i>FEMS Microbiology Ecology</i> , 2001 , 38, 43-52	4.3	75
239	Species associations during the succession of wood-inhabiting fungal communities. <i>Fungal Ecology</i> , 2014 , 11, 17-28	4.1	73
238	Pathogenic fungal species hybrids infecting plants. <i>Microbes and Infection</i> , 2002 , 4, 1353-9	9.3	73
237	Initial fungal colonizer affects mass loss and fungal community development in <i>Picea abies</i> logs 6yr after inoculation. <i>Fungal Ecology</i> , 2011 , 4, 449-460	4.1	71
236	Emerging Diseases in European Forest Ecosystems and Responses in Society. <i>Forests</i> , 2011 , 2, 486-504	2.8	71
235	Wood-inhabiting fungal communities in woody debris of Norway spruce (<i>Picea abies</i> (L.) Karst.), as reflected by sporocarps, mycelial isolations and T-RFLP identification. <i>FEMS Microbiology Ecology</i> , 2006 , 55, 57-67	4.3	69
234	Spore deposition of wood-decaying fungi: importance of landscape composition. <i>Ecography</i> , 2004 , 27, 103-111	6.5	68
233	Transmission of double-stranded RNA in <i>Heterobasidion annosum</i> . <i>Fungal Genetics and Biology</i> , 2002 , 36, 147-54	3.9	68
232	Evolutionary history of the conifer root rot fungus <i>Heterobasidion annosum sensu lato</i> . <i>Molecular Ecology</i> , 2010 , 19, 4979-93	5.7	66
231	Evolutionary significance of imbalanced nuclear ratios within heterokaryons of the basidiomycete fungus <i>Heterobasidion parviporum</i> . <i>Evolution; International Journal of Organic Evolution</i> , 2008 , 62, 2279-96	3.8	65
230	Functional analysis of glycoside hydrolase family 18 and 20 genes in <i>Neurospora crassa</i> . <i>Fungal Genetics and Biology</i> , 2012 , 49, 717-30	3.9	64
229	Molecular markers reveal genetic isolation and phylogeography of the S and F intersterility groups of the wood-decay fungus <i>Heterobasidion annosum</i> . <i>Molecular Phylogenetics and Evolution</i> , 2003 , 29, 94-101	4.1	64
228	Transcriptional analysis of <i>Pinus sylvestris</i> roots challenged with the ectomycorrhizal fungus <i>Laccaria bicolor</i> . <i>BMC Plant Biology</i> , 2008 , 8, 19	5.3	63

227	Global diversity and taxonomy of the <i>Inonotus linteus</i> complex (Hymenochaetales, Basidiomycota): <i>Sanghuangporus</i> gen. nov., <i>Tropicoporus excentrodendri</i> and <i>T. guanacastensis</i> gen. et spp. nov., and 17 new combinations. <i>Fungal Diversity</i> , 2016 , 77, 335-347	17.6	62
226	Chemical and transcriptional responses of Norway spruce genotypes with different susceptibility to <i>Heterobasidion</i> spp. infection. <i>BMC Plant Biology</i> , 2011 , 11, 154	5.3	62
225	Competitive Hierarchies of Wood Decomposing Basidiomycetes in Artificial Systems Based on Variable Inoculum Sizes. <i>Oikos</i> , 1997 , 79, 77	4	62
224	Abundance and viability of fungal spores along a forestry gradient [Responses to habitat loss and isolation?]. <i>Oikos</i> , 2004 , 104, 35-42	4	62
223	Global geographic distribution and host range of <i>Dothistroma</i> species: a comprehensive review. <i>Forest Pathology</i> , 2016 , 46, 408-442	1.2	61
222	Wood-decay fungi in fine living roots of conifer seedlings. <i>New Phytologist</i> , 2007 , 174, 441-446	9.8	61
221	Nuclear reassortment between vegetative mycelia in natural populations of the basidiomycete <i>Heterobasidion annosum</i> . <i>Fungal Genetics and Biology</i> , 2004 , 41, 563-70	3.9	61
220	Variable response of different functional groups of saproxylic beetles to substrate manipulation and forest management: Implications for conservation strategies. <i>Forest Ecology and Management</i> , 2007 , 242, 496-510	3.9	59
219	Habitat generalists and specialists in microbial communities across a terrestrial-freshwater gradient. <i>Scientific Reports</i> , 2016 , 6, 37719	4.9	58
218	Molecular identification of wood-inhabiting fungi in an unmanaged <i>Picea abies</i> forest in Sweden. <i>Forest Ecology and Management</i> , 1999 , 115, 203-211	3.9	58
217	Differential growth of S- and P-isolates of <i>Heterobasidion annosum</i> in <i>Picea abies</i> and <i>Pinus sylvestris</i> . <i>Transactions of the British Mycological Society</i> , 1988 , 90, 209-213		58
216	Utilizing ITS1 and ITS2 to study environmental fungal diversity using pyrosequencing. <i>FEMS Microbiology Ecology</i> , 2013 , 84, 165-75	4.3	57
215	Stump removal to control root disease in Canada and Scandinavia: A synthesis of results from long-term trials. <i>Forest Ecology and Management</i> , 2013 , 290, 5-14	3.9	57
214	Selective replacement between species of wood-rotting basidiomycetes, a laboratory study. <i>Mycological Research</i> , 1997 , 101, 714-720		57
213	Fungi in decayed roots of conifer seedlings in forest nurseries, afforested clear-cuts and abandoned farmland. <i>Plant Pathology</i> , 2006 , 55, 117-129	2.8	57
212	Comparative evolutionary histories of the fungal chitinase gene family reveal non-random size expansions and contractions due to adaptive natural selection. <i>Evolutionary Bioinformatics</i> , 2008 , 4, 47-60 ⁹	1.9	56
211	A genome-wide association study identifies genomic regions for virulence in the non-model organism <i>Heterobasidion annosum</i> s.s. <i>PLoS ONE</i> , 2013 , 8, e53525	3.7	56
210	Friend or foe? Biological and ecological traits of the European ash dieback pathogen <i>Hymenoscyphus fraxineus</i> in its native environment. <i>Scientific Reports</i> , 2016 , 6, 21895	4.9	55

209	Population structure of <i>Hymenoscyphus pseudoalbidus</i> and its genetic relationship to <i>Hymenoscyphus albidus</i> . <i>Fungal Ecology</i> , 2012 , 5, 147-153	4.1	54
208	Transcript profiling of a conifer pathosystem: response of <i>Pinus sylvestris</i> root tissues to pathogen (<i>Heterobasidion annosum</i>) invasion. <i>Tree Physiology</i> , 2007 , 27, 1441-58	4.2	54
207	Pectic isozyme profiles of intersterility groups in <i>Heterobasidion annosum</i> . <i>Mycological Research</i> , 1991 , 95, 531-536		51
206	Socio-ecological implications of modifying rotation lengths in forestry. <i>Ambio</i> , 2016 , 45 Suppl 2, 109-23	6.5	50
205	Estimating the frequency of stem rot in <i>Picea abies</i> using an increment borer. <i>Scandinavian Journal of Forest Research</i> , 1986 , 1, 303-308	1.7	50
204	Susceptibility of conifer and broadleaf seedlings to Swedish S and P strains of <i>Heterobasidion annosum</i> . <i>Plant Pathology</i> , 1995 , 44, 73-79	2.8	49
203	Genetic control of somatic incompatibility in the root-rotting basidiomycete <i>Heterobasidion annosum</i> . <i>Mycological Research</i> , 1993 , 97, 1229-1233		49
202	Stump removal to control root rot in forest stands. A literature study. <i>Silva Fennica</i> , 2008 , 42,	1.9	49
201	Nitrogen and carbon reallocation in fungal mycelia during decomposition of boreal forest litter. <i>PLoS ONE</i> , 2014 , 9, e92897	3.7	48
200	Comparative analysis of transcript abundance in <i>Pinus sylvestris</i> after challenge with a saprotrophic, pathogenic or mutualistic fungus. <i>Tree Physiology</i> , 2008 , 28, 885-97	4.2	48
199	Expressed sequences from the basidiomycetous tree pathogen <i>Heterobasidion annosum</i> during early infection of scots pine. <i>Fungal Genetics and Biology</i> , 2003 , 39, 51-9	3.9	47
198	Continental mapping of forest ecosystem functions reveals a high but unrealised potential for forest multifunctionality. <i>Ecology Letters</i> , 2018 , 21, 31-42	10	47
197	Diverse ecological roles within fungal communities in decomposing logs of <i>Picea abies</i> . <i>FEMS Microbiology Ecology</i> , 2015 , 91,	4.3	46
196	Light and scanning electron microscopy studies of the early infection stages of <i>Hymenoscyphus pseudoalbidus</i> on <i>Fraxinus excelsior</i> . <i>Plant Pathology</i> , 2013 , 62, 1294-1301	2.8	46
195	Comparative molecular evolution of trichoderma chitinases in response to mycoparasitic interactions. <i>Evolutionary Bioinformatics</i> , 2010 , 6, 1-26	1.9	46
194	Fungi inhabiting stems of <i>Picea abies</i> in a managed stand in Lithuania. <i>Forest Ecology and Management</i> , 1998 , 109, 119-126	3.9	46
193	Modeling infection and spread of <i>Heterobasidion annosum</i> in even-aged Fennoscandian conifer stands. <i>Canadian Journal of Forest Research</i> , 2005 , 35, 74-84	1.9	46
192	Growing evidence for facultative biotrophy in saprotrophic fungi: data from microcosm tests with 201 species of wood-decay basidiomycetes. <i>New Phytologist</i> , 2017 , 215, 747-755	9.8	45

191	Modelling root rot incidence in Sweden using tree, site and stand variables. <i>Scandinavian Journal of Forest Research</i> , 2005 , 20, 165-176	1.7	45
190	Persistence and long-term impact of Rotstop biological control agent on mycodiversity in <i>Picea abies</i> stumps. <i>Biological Control</i> , 2005 , 32, 295-304	3.8	45
189	Planting <i>Betula pendula</i> on pine sites infested by <i>Heterobasidion annosum</i> : disease transfer, silvicultural evaluation, and community of wood-inhabiting fungi. <i>Canadian Journal of Forest Research</i> , 2004 , 34, 120-130	1.9	44
188	<i>Heterobasidion annosum</i> infection of <i>Picea abies</i> following manual or mechanized stump treatment. <i>Scandinavian Journal of Forest Research</i> , 2005 , 20, 154-164	1.7	44
187	Long-term reduction in the diameter growth of butt rot affected Norway spruce, <i>Picea abies</i> . <i>Forest Ecology and Management</i> , 1995 , 74, 239-243	3.9	44
186	Somatic incompatibility and nuclear reassortment in <i>Heterobasidion annosum</i> . <i>Mycological Research</i> , 1993 , 97, 1223-1228		44
185	Changes in fungal community of Scots pine (<i>Pinus sylvestris</i>) needles along a latitudinal gradient in Sweden. <i>Fungal Ecology</i> , 2015 , 17, 126-139	4.1	43
184	Glucose and ammonium additions affect needle decomposition and carbon allocation by the litter degrading fungus <i>Mycena epipterygia</i> . <i>Soil Biology and Biochemistry</i> , 2008 , 40, 995-999	7.5	43
183	Population structure of the wood decay fungus <i>Fomitopsis pinicola</i> . <i>Heredity</i> , 1999 , 83 (Pt 3), 354-60	3.6	43
182	Fungi vectored by the bark beetle <i>Ips typographus</i> following hibernation under the bark of standing trees and in the forest litter. <i>Microbial Ecology</i> , 2009 , 58, 651-9	4.4	41
181	Impact of biological (Rotstop) and chemical (urea) treatments on fungal community structure in freshly cut <i>Picea abies</i> stumps. <i>Biological Control</i> , 2004 , 31, 405-413	3.8	41
180	Population structure and responses to disturbance of the basidiomycete <i>Resinicium bicolor</i> . <i>Oecologia</i> , 1990 , 85, 178-184	2.9	40
179	A <i>Picea abies</i> linkage map based on SNP markers identifies QTLs for four aspects of resistance to <i>Heterobasidion parviporum</i> infection. <i>PLoS ONE</i> , 2014 , 9, e101049	3.7	39
178	Population structure and mating system in <i>Marasmius androsaceus</i> Fr. <i>New Phytologist</i> , 1991 , 119, 307-318	3.4	39
177	Afforestation of abandoned farmland with conifer seedlings inoculated with three ectomycorrhizal fungi - impact on plant performance and ectomycorrhizal community. <i>Mycorrhiza</i> , 2007 , 17, 337-348	3.9	38
176	Effects of clear-cutting, thinning, and wood moisture content on the susceptibility of Norway spruce stumps to <i>Heterobasidion annosum</i> . <i>Canadian Journal of Forest Research</i> , 1998 , 28, 759-765	1.9	38
175	Phenotypic interactions between tree hosts and invasive forest pathogens in the light of globalization and climate change. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2016 , 371,	5.8	38
174	Identification of Norway Spruce MYB-bHLH-WDR Transcription Factor Complex Members Linked to Regulation of the Flavonoid Pathway. <i>Frontiers in Plant Science</i> , 2017 , 8, 305	6.2	35

173	Intronic and plasmid-derived regions contribute to the large mitochondrial genome sizes of Agaricomycetes. <i>Current Genetics</i> , 2014 , 60, 303-13	2.9	35
172	Clonality and genetic variation in <i>Amylostereum areolatum</i> and <i>A. chailletii</i> from northern Europe. <i>New Phytologist</i> , 1998 , 139, 751-758	9.8	35
171	Urea treatment reduced <i>Heterobasidion annosum</i> s.l. root rot in <i>Picea abies</i> after 15 years. <i>Forest Ecology and Management</i> , 2008 , 255, 2876-2882	3.9	35
170	Identifying the tree species compositions that maximize ecosystem functioning in European forests. <i>Journal of Applied Ecology</i> , 2019 , 56, 733-744	5.8	35
169	Bark beetles have a decisive impact on fungal communities in Norway spruce stem sections. <i>Fungal Ecology</i> , 2014 , 7, 47-58	4.1	34
168	Do bark beetles facilitate the establishment of rot fungi in Norway spruce?. <i>Fungal Ecology</i> , 2011 , 4, 262-269	4.1	34
167	Fungal C translocation restricts N-mineralization in heterogeneous environments. <i>Functional Ecology</i> , 2010 , 24, 454-459	5.6	34
166	Calcium concentrations of soil affect suppressiveness against <i>Aphanomyces</i> root rot of pea. <i>Soil Biology and Biochemistry</i> , 2007 , 39, 2222-2229	7.5	34
165	Molecular and morphological investigation of <i>Daldinia</i> in northern Europe. <i>Mycological Research</i> , 2000 , 104, 275-280		34
164	Population genetics of <i>Fomitopsis rosea</i> , a wood-decay fungus of the old-growth European taiga. <i>Molecular Ecology</i> , 1999 , 8, 703-710	5.7	34
163	Root rot, associated fungi and their impact on health condition of declining <i>Fraxinus excelsior</i> stands in Lithuania. <i>Scandinavian Journal of Forest Research</i> , 2011 , 26, 128-135	1.7	33
162	Differential gene expression during interactions between <i>Heterobasidion annosum</i> and <i>Physisporinus sanguinolentus</i> . <i>FEMS Microbiology Letters</i> , 2004 , 241, 79-85	2.9	33
161	Diffuse competition for heterogeneous substrate in soil among six species of wood-decomposing basidiomycetes. <i>Oecologia</i> , 1996 , 106, 531-538	2.9	33
160	Butt rot incidence, causal fungi, and related yield loss in <i>Picea abies</i> stands of Latvia. <i>Canadian Journal of Forest Research</i> , 2011 , 41, 2337-2345	1.9	32
159	Fungal disease incidence along tree diversity gradients depends on latitude in European forests. <i>Ecology and Evolution</i> , 2016 , 6, 2426-38	2.8	32
158	Biosynthesis of fomannoxin in the root rotting pathogen <i>Heterobasidion occidentale</i> . <i>Phytochemistry</i> , 2012 , 84, 31-9	4	31
157	Double-stranded RNA transmission through basidiospores of <i>Heterobasidion annosum</i> . <i>Mycological Research</i> , 2004 , 108, 149-53		31
156	Silvicultural and pathological evaluation of Scots pine afforestations mixed with deciduous trees to reduce the infections by <i>Heterobasidion annosum</i> s.s.. <i>Forest Ecology and Management</i> , 2004 , 201, 275-283	2.9	30

155	Diplodia Tip Blight on Its Way to the North: Drivers of Disease Emergence in Northern Europe. <i>Frontiers in Plant Science</i> , 2018 , 9, 1818	6.2	29
154	Understanding the role of sapwood loss and reaction zone formation on radial growth of Norway spruce (<i>Picea abies</i>) trees decayed by <i>Heterobasidion annosum</i> s.l.. <i>Forest Ecology and Management</i> , 2012 , 274, 201-209	3.9	29
153	Identification of quantitative trait loci affecting virulence in the basidiomycete <i>Heterobasidion annosum</i> s.l. <i>Current Genetics</i> , 2007 , 52, 35-44	2.9	29
152	An AFLP-markers based genetic linkage map of <i>Heterobasidion annosum</i> locating intersterility genes. <i>Fungal Genetics and Biology</i> , 2005 , 42, 519-27	3.9	29
151	Enniatins of <i>Fusarium</i> sp. strain F31 and their inhibition of <i>Botrytis cinerea</i> spore germination. <i>Journal of Natural Products</i> , 2004 , 67, 851-7	4.9	29
150	Spread of S and P group isolates of <i>Heterobasidion annosum</i> within and among <i>Picea abies</i> trees in central Lithuania. <i>Canadian Journal of Forest Research</i> , 1998 , 28, 961-966	1.9	29
149	Optimized metabarcoding with Pacific biosciences enables semi-quantitative analysis of fungal communities. <i>New Phytologist</i> , 2020 , 228,	9.8	28
148	Fungal communities in Norway spruce stumps along a latitudinal gradient in Sweden. <i>Forest Ecology and Management</i> , 2016 , 371, 50-58	3.9	28
147	The primary module in Norway spruce defence signalling against <i>H. annosum</i> s.l. seems to be jasmonate-mediated signalling without antagonism of salicylate-mediated signalling. <i>Planta</i> , 2013 , 237, 1037-45	4.7	28
146	Fungi in foliage and shoots of <i>Fraxinus excelsior</i> in eastern Ukraine: a first report on <i>Hymenoscyphus pseudoalbidus</i> . <i>Forest Pathology</i> , 2013 , 43, 462-467	1.2	28
145	QTL mapping of resistance to leaf rust in <i>Salix</i> . <i>Tree Genetics and Genomes</i> , 2011 , 7, 1219-1235	2.1	28
144	<i>Heterobasidion annosum</i> root rot in <i>Picea abies</i> : Modelling economic outcomes of stump treatment in Scandinavian coniferous forests. <i>Scandinavian Journal of Forest Research</i> , 2006 , 21, 414-423 ¹⁻⁷		28
143	Variation in spread of <i>Heterobasidion annosum</i> in clones of <i>Picea abies</i> grown at different vegetation phases under greenhouse conditions. <i>Scandinavian Journal of Forest Research</i> , 1996 , 11, 137-144		28
142	Population genetics of the wood-decay fungus <i>Phlebia centrifuga</i> P. Karst. in fragmented and continuous habitats. <i>Molecular Ecology</i> , 2007 , 16, 3326-33	5.7	27
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1	Population dynamics of forest tree pathogens 2022 , 131-143		