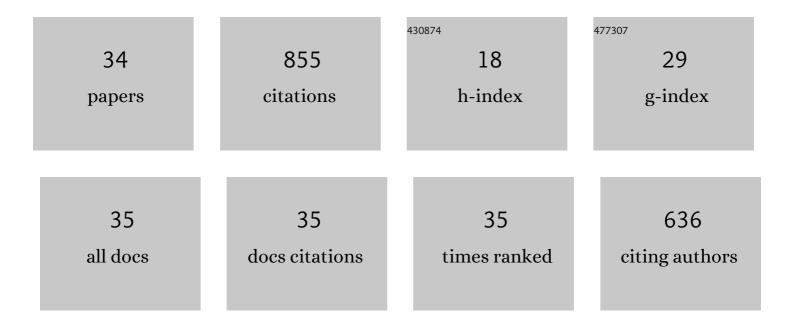
Martin Schnittler

List of Publications by Year in descending order

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MADTIN SCHNITTIED

| # | Article | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Myxomycete diversity and distribution from the fossil record to the present. Biodiversity and Conservation, 2008, 17, 285-301. | 2.6 | 91 |
| 2 | Towards a phylogenetic classification of the Myxomycetes. Phytotaxa, 2019, 399, 209. | 0.3 | 61 |
| 3 | Two-Gene Phylogeny of Bright-Spored Myxomycetes (Slime Moulds, Superorder Lucisporidia). PLoS ONE, 2013, 8, e62586. | 2.5 | 58 |
| 4 | Size matters—a comparison of three methods to assess age- and size-dependent climate sensitivity of trees. Trees - Structure and Function, 2019, 33, 183-192. | 1.9 | 54 |
| 5 | Tuning the Voices of a Choir: Detecting Ecological Gradients in Time-Series Populations. PLoS ONE, 2016, 11, e0158346. | 2.5 | 50 |
| 6 | Root suckering patterns in Populus euphratica (Euphrates poplar, Salicaceae). Trees - Structure and Function, 2009, 23, 991-1001. | 1.9 | 47 |
| 7 | Sex or no sex? Group I introns and independent marker genes reveal the existence of three sexual but reproductively isolated biospecies in Trichia varia (Myxomycetes). Organisms Diversity and Evolution, 2015, 15, 631-650. | 1.6 | 43 |
| 8 | Barcoding myxomycetes with molecular markers: challenges and opportunities. Nova Hedwigia, 2017, 104, 323-341. | 0.4 | 39 |
| 9 | The genus <i>Alwisia</i> (Myxomycetes) revalidated, with two species new to science. Mycologia, 2014, 106, 936-948. | 1.9 | 33 |
| 10 | A critical revision of the <i>Tubifera ferruginosa</i> complex. Mycologia, 2015, 107, 959-985. | 1.9 | 33 |
| 11 | What an Intron May Tell: Several Sexual Biospecies Coexist in Meriderma spp. (Myxomycetes). Protist, 2016, 167, 234-253. | 1.5 | 33 |
| 12 | A habitat colonisation model for spore-dispersed organisms—Does it work with eumycetozoans?. Mycological Research, 2008, 112, 697-707. | 2.5 | 29 |
| 13 | Genetic barcoding of darkâ€spored myxomycetes (Amoebozoa)—Identification, evaluation and application of a sequence similarity threshold for species differentiation in <scp>NGS</scp> studies. Molecular Ecology Resources, 2018, 18, 306-318. | 4.8 | 27 |
| 14 | Speciation in progress? A phylogeographic study among populations of Hemitrichia serpula (Myxomycetes). PLoS ONE, 2017, 12, e0174825. | 2.5 | 27 |
| 15 | Phylogenetic position of the enigmatic myxomycete genus Kelleromyxa revealed by SSU rDNA sequences. Mycological Progress, 2013, 12, 599-608. | 1.4 | 24 |
| 16 | First insight into dead wood protistan diversity: a molecular sampling of bright-spored Myxomycetes (Amoebozoa, slime-moulds) in decaying beech logs. FEMS Microbiology Ecology, 2015, 91, . | 2.7 | 23 |
| 17 | Myxomycete diversity in the Tarim basin and eastern Tian-Shan, Xinjiang Prov., China. Fungal Diversity, 2013, 59, 91-108. | 12.3 | 21 |
| 18 | Biogeographical assessment of myxomycete assemblages from Neotropical and Asian Palaeotropical forests. Journal of Biogeography, 2017, 44, 1524-1536. | 3.0 | 21 |

MARTIN SCHNITTLER

| # | Article | IF | CITATIONS |
|----|--|---------------------|--------------------|
| 19 | A four year survey reveals a coherent pattern between occurrence of fruit bodies and soil amoebae populations for nivicolous myxomycetes. Scientific Reports, 2018, 8, 11662. | 3.3 | 18 |
| 20 | Different Degrees of Niche Differentiation for Bacteria, Fungi, and Myxomycetes Within an Elevational Transect in the German Alps. Microbial Ecology, 2019, 78, 764-780. | 2.8 | 16 |
| 21 | A new species of <i>Alwisia</i> (Myxomycetes) from New South Wales and Tasmania. Mycologia, 2014, 106, 1212-1219. | 1.9 | 14 |
| 22 | Development of two microsatellite multiplex PCR systems for high throughput genotyping in Populus euphratica. Journal of Forestry Research, 2009, 20, 195-198. | 3.6 | 12 |
| 23 | Morphological and molecular investigations of Gagea (Liliaceae) in southeastern Kazakhstan with special reference to putative altitudinal hybrid zones. Plant Systematics and Evolution, 2016, 302, 985-1007. | 0.9 | 11 |
| 24 | Species-specific effects of thermal stress on the expression of genetic variation across a diverse group of plant and animal taxa under experimental conditions. Heredity, 2021, 126, 23-37. | 2.6 | 11 |
| 25 | Pseudocapillitium or true capillitium? A study of capillitial structures in Alwisia bombarda (Myxomycetes). Nova Hedwigia, 2014, 99, 441-451. | 0.4 | 9 |
| 26 | Systematic revision of the <i>Tubifera casparyi–T</i> . <i>dictyoderma</i> complex: Resurrection of the genus <i>Siphoptychium</i> and introduction of the new genus <i>Thecotubifera</i> . Mycologia, 2019, 111, 981-997. | 1.9 | 9 |
| 27 | Genetic diversity and hybrid formation in Central European club-mosses (Diphasiastrum,) Tj ETQq1 1 0.784314 Phylogenetics and Evolution, 2019, 131, 181-192. | 1 rgBT /Ovei 2.7 | lock 10 Tf 50 8 |
| 28 | Population structure and the influence of microenvironment and genetic similarity on individual growth at Alaskan white spruce treelines. Science of the Total Environment, 2021, 798, 149267. | 8.0 | 8 |
| 29 | Contributions to "E-Taxonomyâ€⊷ A virtual approach to the flora of Mongolia (FloraGREIF). Feddes Repertorium, 2013, 123, n/a-n/a. | 0.5 | 7 |
| 30 | Genetic structure of the protist <i>Physarum albescens</i> (Amoebozoa) revealed by multiple markers and genotyping by sequencing. Molecular Ecology, 2022, 31, 372-390. | 3.9 | 7 |
| 31 | Studies of life history of Gagea graeca (Liliaceae) based on morphological and molecular methods. , 2017, 58, 40. | | 3 |
| 32 | The phylogeny and phylogenetically based classification of myxomycetes. , 2022, , 97-124. | | 3 |
| | | | |
| 33 | A workflow for low-cost automated image analysis of myxomycete spore numbers, size and shape. PeerJ, 2021, 9, e12471. | 2.0 | 3 |