

# Natascha Dorothea Wagner

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/777035/publications.pdf>

Version: 2024-02-01

10  
papers

257  
citations

1163117

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1372567

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#	ARTICLE	IF	CITATIONS
1	Untying Gordian knots: unraveling reticulate polyploid plant evolution by genomic data using the large <i>Ranunculus auricomus</i> species complex. <i>New Phytologist</i> , 2022, 235, 2081-2098.	7.3	19
2	Ancient DNA extraction methods for herbarium specimens: When is it worth the effort?. <i>Applications in Plant Sciences</i> , 2022, 10, .	2.1	9
3	Restriction-site associated DNA sequencing data reveal a radiation of willow species ( <i>Salix</i> L.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 59, 44-57.	3.1	27
4	Conservation in the face of hybridisation: genome-wide study to evaluate taxonomic delimitation and conservation status of a threatened orchid species. <i>Conservation Genetics</i> , 2021, 22, 151-168.	1.5	8
5	Chromosome-scale assembly of the genome of <i>Salix dunnii</i> reveals a male-heterogametic sex determination system on chromosome 7. <i>Molecular Ecology Resources</i> , 2021, 21, 1966-1982.	4.8	28
6	The Evolutionary History, Diversity, and Ecology of Willows ( <i>Salix</i> L.) in the European Alps. <i>Diversity</i> , 2021, 13, 146.	1.7	17
7	Highly Diverse Shrub Willows ( <i>Salix</i> L.) Share Highly Similar Plastomes. <i>Frontiers in Plant Science</i> , 2021, 12, 662715.	3.6	12
8	Phylogenomic Relationships and Evolution of Polyploid <i>Salix</i> Species Revealed by RAD Sequencing Data. <i>Frontiers in Plant Science</i> , 2020, 11, 1077.	3.6	54
9	RAD-seq reveals genetic structure of the F2-generation of natural willow hybrids ( <i>Salix</i> L.) and a great potential for interspecific introgression. <i>BMC Plant Biology</i> , 2018, 18, 317.	3.6	31
10	RAD sequencing resolved phylogenetic relationships in European shrub willows ( <i>Salix</i> L. subg.) Tj ETQq0 0 0 rgBT /Overlock 10 Tf and Evolution, 2018, 8, 8243-8255.	1.9	39