

# ValÃ©rie Geffroy

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8239234/publications.pdf>

Version: 2024-02-01

10  
papers

1,691  
citations

1163117

8  
h-index

1372567

10  
g-index

10  
all docs

10  
docs citations

10  
times ranked

2190  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-Wide Identification of Key Components of RNA Silencing in Two Phaseolus vulgaris Genotypes of Contrasting Origin and Their Expression Analyses in Response to Fungal Infection. <i>Genes</i> , 2022, 13, 64.	2.4	4
2	A common bean truncated CRINKLY4 kinase controls gene-for-gene resistance to the fungus <i>Colletotrichum lindemuthianum</i> . <i>Journal of Experimental Botany</i> , 2021, 72, 3569-3581.	4.8	21
3	R-BPMV-Mediated Resistance to Bean pod mottle virus in Phaseolus vulgaris L. Is Heat-Stable but Elevated Temperatures Boost Viral Infection in Susceptible Genotypes. <i>Viruses</i> , 2021, 13, 1239.	3.3	4
4	The INCREASE project: Intelligent Collections of food&#x2013;legume genetic resources for European agrofood systems. <i>Plant Journal</i> , 2021, 108, 646-660.	5.7	29
5	Development of molecular markers linked to disease resistance genes in common bean based on whole genome sequence. <i>Plant Science</i> , 2016, 242, 351-357.	3.6	118
6	Fine mapping of Co-x, an anthracnose resistance gene to a highly virulent strain of <i>Colletotrichum lindemuthianum</i> in common bean. <i>Theoretical and Applied Genetics</i> , 2014, 127, 1653-1666.	3.6	59
7	A reference genome for common bean and genome-wide analysis of dual domestications. <i>Nature Genetics</i> , 2014, 46, 707-713.	21.4	1,159
8	Evolution of a Complex Disease Resistance Gene Cluster in Diploid <i>Phaseolus</i> and Tetraploid <i>Glycine</i> . <i>Plant Physiology</i> , 2012, 159, 336-354.	4.8	76
9	A Nomadic Subtelomeric Disease Resistance Gene Cluster in Common Bean. <i>Plant Physiology</i> , 2009, 151, 1048-1065.	4.8	121
10	Resistance to <i>Colletotrichum lindemuthianum</i> in <i>Phaseolus vulgaris</i> : a case study for mapping two independent genes. <i>Theoretical and Applied Genetics</i> , 2008, 116, 407-415.	3.6	100