

Jean-Luc Gallois

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

2,409
citations

279487

23
h-index

329751

37
g-index

40
all docs

40
docs citations

40
times ranked

2522
citing authors

#	ARTICLE	IF	CITATIONS
1	Prime Editing in the model plant <i>Physcomitrium patens</i> and its potential in the tetraploid potato. <i>Plant Science</i> , 2022, 316, 111162.	1.7	32
2	CRISPR-based knock-out of eIF4E2 in a cherry tomato background successfully recapitulates resistance to pepper veinal mottle virus. <i>Plant Science</i> , 2022, 316, 111160.	1.7	18
3	Exploring New Routes for Genetic Resistances to Potyviruses: The Case of the <i>Arabidopsis thaliana</i> Phosphoglycerates Kinases (PGK) Metabolic Enzymes. <i>Viruses</i> , 2022, 14, 1245.	1.5	4
4	Gene Editing in Potato Using CRISPR-Cas9 Technology. <i>Methods in Molecular Biology</i> , 2021, 2354, 331-351.	0.4	4
5	A blueprint for gene function analysis through Base Editing in the model plant <i>Physcomitrium (Physcomitrella) patens</i> . <i>New Phytologist</i> , 2021, 230, 1258-1272.	3.5	18
6	When a knockout is an Achilles™ heel: Resistance to one potyvirus species triggers hypersusceptibility to another one in <i>Arabidopsis thaliana</i> . <i>Molecular Plant Pathology</i> , 2021, 22, 334-347.	2.0	18
7	Knock-out mutation of eukaryotic initiation factor 4E2 (eIF4E2) confers resistance to pepper veinal mottle virus in tomato. <i>Virology</i> , 2020, 539, 11-17.	1.1	19
8	Precision Breeding Made Real with CRISPR: Illustration through Genetic Resistance to Pathogens. <i>Plant Communications</i> , 2020, 1, 100102.	3.6	32
9	Expanding the CRISPR Toolbox in <i>P. patens</i> Using SpCas9-NG Variant and Application for Gene and Base Editing in Solanaceae Crops. <i>International Journal of Molecular Sciences</i> , 2020, 21, 1024.	1.8	44
10	Analysis of tomato spotted wilt virus RNA-dependent RNA polymerase adaptative evolution and constrained domains using homology protein structure modelling. <i>Journal of General Virology</i> , 2020, 101, 334-346.	1.3	5
11	Host Plant Resistance to Pests and Pathogens, the Genetic Leverage in Integrated Pest and Disease Management. , 2020, , 259-283.		3
12	A complex <i>eIF4E</i> locus impacts the durability of <i>va</i> resistance to <i>Potato virus Y</i> in tobacco. <i>Molecular Plant Pathology</i> , 2019, 20, 1051-1066.	2.0	18
13	The <i>Solanum tuberosum</i> GBSSI gene: a target for assessing gene and base editing in tetraploid potato. <i>Plant Cell Reports</i> , 2019, 38, 1065-1080.	2.8	78
14	Mimicking natural polymorphism in <i>eIF4E</i> by CRISPR-Cas9 base editing is associated with resistance to potyviruses. <i>Plant Biotechnology Journal</i> , 2019, 17, 1736-1750.	4.1	129
15	Trans-species synthetic gene design allows resistance pyramiding and broad-spectrum engineering of virus resistance in plants. <i>Plant Biotechnology Journal</i> , 2018, 16, 1569-1581.	4.1	64
16	Role of the Genetic Background in Resistance to Plant Viruses. <i>International Journal of Molecular Sciences</i> , 2018, 19, 2856.	1.8	62
17	PUX10 Is a CDC48A Adaptor Protein That Regulates the Extraction of Ubiquitinated Oleosins from Seed Lipid Droplets in <i>Arabidopsis</i> . <i>Plant Cell</i> , 2018, 30, 2116-2136.	3.1	64
18	eIF4E Resistance: Natural Variation Should Guide Gene Editing. <i>Trends in Plant Science</i> , 2017, 22, 411-419.	4.3	98

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19	Identification of major loci and genomic regions controlling acid and volatile content in tomato fruit: implications for flavor improvement. <i>New Phytologist</i> , 2017, 215, 624-641.	3.5	65
20	A <i>TILLING</i> approach to generate broad-spectrum resistance to potyviruses in tomato is hampered by <i>elf4E</i> gene redundancy. <i>Plant Journal</i> , 2016, 85, 717-729.	2.8	76
21	A new <i>elf4E1</i> allele characterized by RNAseq data mining is associated with resistance to potato virus Y in tomato albeit with a low durability. <i>Journal of General Virology</i> , 2016, 97, 3063-3072.	1.3	23
22	Human Management of a Wild Plant Modulates the Evolutionary Dynamics of a Gene Determining Recessive Resistance to Virus Infection. <i>PLoS Genetics</i> , 2016, 12, e1006214.	1.5	20
23	Pyramiding resistances based on translation initiation factors in <i>Arabidopsis</i> is impaired by male gametophyte lethality. <i>Plant Signaling and Behavior</i> , 2014, 9, e27940.	1.2	29
24	Cloning of the <i>Arabidopsis rwm1</i> gene for resistance to <i>Watermelon mosaic virus</i> points to a new function for natural virus resistance genes. <i>Plant Journal</i> , 2014, 79, 705-716.	2.8	62
25	Specific requirement for translation initiation factor 4E or its isoform drives plant host susceptibility to Tobacco etch virus. <i>BMC Plant Biology</i> , 2014, 14, 67.	1.6	24
26	Functional characterization of the plant ubiquitin regulatory X (UBX) domain-containing protein AtPUX7 in <i>Arabidopsis thaliana</i> . <i>Gene</i> , 2013, 526, 299-308.	1.0	29
27	A SNP associated with alternative splicing of RPT5b causes unequal redundancy between RPT5a and RPT5b among <i>Arabidopsis thaliana</i> natural variation. <i>BMC Plant Biology</i> , 2010, 10, 158.	1.6	9
28	Single amino acid changes in the turnip mosaic virus viral genome-linked protein (VPg) confer virulence towards <i>Arabidopsis thaliana</i> mutants knocked out for eukaryotic initiation factors <i>elf(iso)4E</i> and <i>elf(iso)4G</i> . <i>Journal of General Virology</i> , 2010, 91, 288-293.	1.3	69
29	The <i>Arabidopsis</i> Proteasome RPT5 Subunits Are Essential for Gametophyte Development and Show Accession-Dependent Redundancy. <i>Plant Cell</i> , 2009, 21, 442-459.	3.1	65
30	Natural variation and functional analyses provide evidence for co-evolution between plant <i>elf4E</i> and potyviral VPg. <i>Plant Journal</i> , 2008, 54, 56-68.	2.8	207
31	Coordinated and selective recruitment of <i>elf4E</i> and <i>elf4G</i> factors for potyvirus infection in <i>Arabidopsis thaliana</i> . <i>FEBS Letters</i> , 2007, 581, 1041-1046.	1.3	109
32	AtPRD1 is required for meiotic double strand break formation in <i>Arabidopsis thaliana</i> . <i>EMBO Journal</i> , 2007, 26, 4126-4137.	3.5	102
33	Simultaneous mutations in translation initiation factors <i>elf4E</i> and <i>elf(iso)4E</i> are required to prevent pepper vein mottle virus infection of pepper. <i>Journal of General Virology</i> , 2006, 87, 2089-2098.	1.3	140
34	WUSCHEL induces shoot stem cell activity and developmental plasticity in the root meristem. <i>Genes and Development</i> , 2004, 18, 375-380.	2.7	229
35	The <i>Arabidopsis</i> nuclear DAL gene encodes a chloroplast protein which is required for the maturation of the plastid ribosomal RNAs and is essential for chloroplast differentiation. <i>Plant Molecular Biology</i> , 2003, 51, 651-663.	2.0	62
36	Combined SHOOT MERISTEMLESS and WUSCHEL trigger ectopic organogenesis in <i>Arabidopsis</i> . <i>Development (Cambridge)</i> , 2002, 129, 3207-3217.	1.2	221

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37	Combined SHOOT MERISTEMLESS and WUSCHEL trigger ectopic organogenesis in Arabidopsis. <i>Development (Cambridge)</i> , 2002, 129, 3207-17.	1.2	110
38	The Arabidopsis chloroplast ribosomal protein L21 is encoded by a nuclear gene of mitochondrial origin. <i>Gene</i> , 2001, 274, 179-185.	1.0	20