

Christophe DÃ©lye

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8755805/publications.pdf>

Version: 2024-02-01

64
papers

4,644
citations

94433

37
h-index

110387

64
g-index

65
all docs

65
docs citations

65
times ranked

2686
citing authors

#	ARTICLE	IF	CITATIONS
1	Deciphering the evolution of herbicide resistance in weeds. <i>Trends in Genetics</i> , 2013, 29, 649-658.	6.7	462
2	Unravelling the genetic bases of non-target-site-based resistance (NTSR) to herbicides: a major challenge for weed science in the forthcoming decade. <i>Pest Management Science</i> , 2013, 69, 176-187.	3.4	364
3	High-throughput microsatellite isolation through 454 GS-FLX Titanium pyrosequencing of enriched DNA libraries. <i>Molecular Ecology Resources</i> , 2011, 11, 638-644.	4.8	276
4	Weed resistance to acetyl coenzyme A carboxylase inhibitors: an update. <i>Weed Science</i> , 2005, 53, 728-746.	1.5	241
5	A mutation in the 14 alpha-demethylase gene of <i>Uncinula necator</i> that correlates with resistance to a sterol biosynthesis inhibitor. <i>Applied and Environmental Microbiology</i> , 1997, 63, 2966-2970.	3.1	194
6	Molecular Bases for Sensitivity to Acetyl-Coenzyme A Carboxylase Inhibitors in Black-Grass. <i>Plant Physiology</i> , 2005, 137, 794-806.	4.8	176
7	PCR cloning and detection of point mutations in the eburicol 14 a -demethylase (CYP51) gene from <i>Erysiphe graminis</i> f. sp. <i>hordei</i> , a "recalcitrant" fungus. <i>Current Genetics</i> , 1998, 34, 399-403.	1.7	125
8	An Isoleucine Residue within the Carboxyl-Transferase Domain of Multidomain Acetyl-Coenzyme A Carboxylase Is a Major Determinant of Sensitivity to Aryloxyphenoxypropionate But Not to Cyclohexanedione Inhibitors. <i>Plant Physiology</i> , 2003, 132, 1716-1723.	4.8	122
9	Prevalence of cross or multiple resistance to the acetyl-coenzyme A carboxylase inhibitors fenoxaprop, clodinafop and pinoxaden in black-grass (<i>Alopecurus myosuroides</i> Huds.) in France. <i>Pest Management Science</i> , 2010, 66, 168-177.	3.4	120
10	RNA-Seq analysis of rye-grass transcriptomic response to an herbicide inhibiting acetolactate-synthase identifies transcripts linked to non-target-site-based resistance. <i>Plant Molecular Biology</i> , 2015, 87, 473-487.	3.9	115
11	An isoleucine-leucine substitution in chloroplastic acetyl-CoA carboxylase from green foxtail (<i>Setaria viridis</i> L. Beauv.) is responsible for resistance to the cyclohexanedione herbicide sethoxydim. <i>Planta</i> , 2002, 214, 421-427.	3.2	106
12	Geographical variation in resistance to acetyl-coenzyme A carboxylase-inhibiting herbicides across the range of the arable weed <i>Alopecurus myosuroides</i> (black-grass). <i>New Phytologist</i> , 2010, 186, 1005-1017.	7.3	103
13	Fitness costs associated with three mutant acetyl-coenzyme A carboxylase alleles endowing herbicide resistance in black-grass <i>Alopecurus myosuroides</i> . <i>Journal of Applied Ecology</i> , 2008, 45, 939-947.	4.0	99
14	PCR-based detection of resistance to acetyl-CoA carboxylase-inhibiting herbicides in black-grass (<i>Alopecurus myosuroides</i> Huds) and ryegrass (<i>Lolium rigidum</i> Gaud). <i>Pest Management Science</i> , 2002, 58, 474-478.	3.4	94
15	Complex genetic control of non-target-site-based resistance to herbicides inhibiting acetyl-coenzyme A carboxylase and acetolactate-synthase in <i>Alopecurus myosuroides</i> Huds.. <i>Plant Science</i> , 2010, 178, 501-509.	3.6	88
16	Non-target-site-based resistance should be the centre of attention for herbicide resistance research: <i>Alopecurus myosuroides</i> as an illustration. <i>Weed Research</i> , 2011, 51, 433-437.	1.7	87
17	Molecular Bases for Sensitivity to Tubulin-Binding Herbicides in Green Foxtail. <i>Plant Physiology</i> , 2004, 136, 3920-3932.	4.8	85
18	Genetic Diversity and Pathogenic Variability Among Isolates of <i>Colletotrichum</i> Species from Strawberry. <i>Phytopathology</i> , 2003, 93, 219-228.	2.2	80

#	ARTICLE	IF	CITATIONS
19	Cross-resistance patterns to ACCase-inhibiting herbicides conferred by mutant ACCase isoforms in <i>Alopecurus myosuroides</i> Huds. (black-grass), re-examined at the recommended herbicide field rate. <i>Pest Management Science</i> , 2008, 64, 1179-1186.	3.4	76
20	Molecular Mechanisms of Herbicide Resistance. <i>Weed Science</i> , 2015, 63, 91-115.	1.5	73
21	Weed response to herbicides: regional-scale distribution of herbicide resistance alleles in the grass weed <i>Alopecurus myosuroides</i> . <i>New Phytologist</i> , 2006, 171, 861-874.	7.3	72
22	A new insight into arable weed adaptive evolution: mutations endowing herbicide resistance also affect germination dynamics and seedling emergence. <i>Annals of Botany</i> , 2013, 111, 681-691.	2.9	72
23	Genetic variation and population structure in black-grass (<i>Alopecurus myosuroides</i> Huds.), a successful, herbicide-resistant, annual grass weed of winter cereal fields. <i>Molecular Ecology</i> , 2007, 16, 3161-3172.	3.9	67
24	Status of black grass (<i>Alopecurus myosuroides</i>) resistance to acetyl-coenzyme A carboxylase inhibitors in France. <i>Weed Research</i> , 2007, 47, 95-105.	1.7	66
25	ALOMYbase, a resource to investigate non-target-site-based resistance to herbicides inhibiting acetolactate-synthase (ALS) in the major grass weed <i>Alopecurus myosuroides</i> (black-grass). <i>BMC Genomics</i> , 2015, 16, 590.	2.8	66
26	SNP markers for black-grass (<i>Alopecurus myosuroides</i> Huds.) genotypes resistant to acetyl CoA-carboxylase inhibiting herbicides. <i>Theoretical and Applied Genetics</i> , 2002, 104, 1114-1120.	3.6	65
27	Evolution and diversity of the mechanisms endowing resistance to herbicides inhibiting acetolactate-synthase (ALS) in corn poppy (<i>Papaver rhoeas</i> L.). <i>Plant Science</i> , 2011, 180, 333-342.	3.6	62
28	RAPD Analysis Provides Insight into the Biology and Epidemiology of <i>Uncinula necator</i> . <i>Phytopathology</i> , 1997, 87, 670-677.	2.2	60
29	PCR Assays That Identify the Grapevine Dieback Fungus <i>Eutypa lata</i> . <i>Applied and Environmental Microbiology</i> , 2000, 66, 4475-4480.	3.1	58
30	Reference Genes to Study Herbicide Stress Response in <i>Lolium</i> sp.: Up-Regulation of P450 Genes in Plants Resistant to Acetolactate-Synthase Inhibitors. <i>PLoS ONE</i> , 2013, 8, e63576.	2.5	58
31	DNA Analysis of Herbarium Specimens of the Grass Weed <i>Alopecurus myosuroides</i> Reveals Herbicide Resistance Pre-Dated Herbicides. <i>PLoS ONE</i> , 2013, 8, e75117.	2.5	55
32	'Universal' primers for PCR-sequencing of grass chloroplastic acetyl-CoA carboxylase domains involved in resistance to herbicides. <i>Weed Research</i> , 2005, 45, 323-330.	1.7	54
33	'Universal'™ PCR assays detecting mutations in acetyl-coenzyme A carboxylase or acetolactate synthase that endow herbicide resistance in grass weeds. <i>Weed Research</i> , 2011, 51, 353-362.	1.7	54
34	Adaptive introgression from maize has facilitated the establishment of teosinte as a noxious weed in Europe. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 25618-25627.	7.1	54
35	Origin of primary infections of grape by <i>Uncinula necator</i> : RAPD analysis discriminates two biotypes. <i>Mycological Research</i> , 1998, 102, 283-288.	2.5	51
36	A molecular assay for the proactive detection of target site-based resistance to herbicides inhibiting acetolactate synthase in <i>Alopecurus myosuroides</i> . <i>Weed Research</i> , 2008, 48, 97-101.	1.7	42

#	ARTICLE	IF	CITATIONS
37	Transcriptional markers enable identification of rye-grass (<i>Lolium</i> sp.) plants with non-target-site-based resistance to herbicides inhibiting acetolactate-synthase. <i>Plant Science</i> , 2017, 257, 22-36.	3.6	42
38	Occurrence, genetic control and evolution of non-target-site based resistance to herbicides inhibiting acetolactate synthase (ALS) in the dicot weed <i>Papaver rhoeas</i> . <i>Plant Science</i> , 2015, 238, 158-169.	3.6	40
39	Nucleotide Variability at the Acetyl Coenzyme A Carboxylase Gene and the Signature of Herbicide Selection in the Grass Weed <i>Alopecurus myosuroides</i> (Huds.). <i>Molecular Biology and Evolution</i> , 2004, 21, 884-892.	8.9	39
40	High gene flow promotes the genetic homogeneity of arable weed populations at the landscape level. <i>Basic and Applied Ecology</i> , 2010, 11, 504-512.	2.7	37
41	Variation in the gene encoding acetolactate synthase in <i>Lolium</i> species and proactive detection of mutant, herbicide-resistant alleles. <i>Weed Research</i> , 2009, 49, 326-336.	1.7	36
42	Validation of a set of reference genes to study response to herbicide stress in grasses. <i>BMC Research Notes</i> , 2012, 5, 18.	1.4	35
43	Fitness cost due to herbicide resistance may trigger genetic background evolution. <i>Evolution; International Journal of Organic Evolution</i> , 2015, 69, 271-278.	2.3	35
44	Nested Allele-Specific PCR Primers Distinguish Genetic Groups of <i>Uncinula necator</i> . <i>Applied and Environmental Microbiology</i> , 1999, 65, 3950-3954.	3.1	34
45	Cloning and sequence analysis of the eburicol 14 α -demethylase gene of the obligate biotrophic grape powdery mildew fungus. <i>Gene</i> , 1997, 195, 29-33.	2.2	31
46	Herbicide Safeners Decrease Sensitivity to Herbicides Inhibiting Acetolactate-Synthase and Likely Activate Non-Target-Site-Based Resistance Pathways in the Major Grass Weed <i>Lolium</i> sp. (Rye-Grass). <i>Frontiers in Plant Science</i> , 2017, 8, 1310.	3.6	27
47	A RAPD Assay for Strain Typing of the Biotrophic Grape Powdery Mildew Fungus <i>Uncinula necator</i> Using DNA Extracted from the Mycelium. <i>Experimental Mycology</i> , 1995, 19, 234-237.	1.6	25
48	Gene flow increases the initial frequency of herbicide resistance alleles in unselected <i>Lolium rigidum</i> populations. <i>Agriculture, Ecosystems and Environment</i> , 2011, 142, 403-409.	5.3	24
49	New gSSR and EST-SSR markers reveal high genetic diversity in the invasive plant <i>Ambrosia artemisiifolia</i> L. and can be transferred to other invasive <i>Ambrosia</i> species. <i>PLoS ONE</i> , 2017, 12, e0176197.	2.5	23
50	Multiple origins for black-grass (<i>Alopecurus myosuroides</i> Huds) target-site-based resistance to herbicides inhibiting acetyl-CoA carboxylase. <i>Pest Management Science</i> , 2004, 60, 35-41.	3.4	22
51	Genetic basis, evolutionary origin and spread of resistance to herbicides inhibiting acetolactate synthase in common groundsel (<i>Senecio vulgaris</i>). <i>Pest Management Science</i> , 2016, 72, 89-102.	3.4	19
52	Choosing the best cropping systems to target pleiotropic effects when managing single gene herbicide resistance in grass weeds. A blackgrass simulation study. <i>Pest Management Science</i> , 2016, 72, 1910-1925.	3.4	18
53	New tools for studying epidemiology and resistance of grape powdery mildew to DMI fungicides. <i>Pest Management Science</i> , 1997, 51, 309-314.	0.4	17
54	Using next-generation sequencing to detect mutations endowing resistance to pesticides: application to acetolactate synthase (ALS)-based resistance in barnyard grass, a polyploid grass weed. <i>Pest Management Science</i> , 2015, 71, 675-685.	3.4	14

#	ARTICLE	IF	CITATIONS
55	Harnessing the power of next-generation sequencing technologies to the purpose of high-throughput pesticide resistance diagnosis. <i>Pest Management Science</i> , 2020, 76, 543-552.	3.4	14
56	Isolation and Characterisation of 11 Polymorphic Microsatellite Markers in <i>Papaver rhoeas</i> L. (Corn) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 5 Sciences, 2013, 14, 470-479.	4.1	11
57	Multiple resistance of <i>Papaver rhoeas</i> L. to 2,4-D and acetolactate synthase inhibitors in four European countries. <i>Weed Research</i> , 2019, 59, 367-376.	1.7	11
58	A high diversity of mechanisms endows ALS-inhibiting herbicide resistance in the invasive common ragweed (<i>Ambrosia artemisiifolia</i> L.). <i>Scientific Reports</i> , 2021, 11, 19904.	3.3	11
59	A single polymerase chain reaction-based assay for simultaneous detection of two mutations conferring resistance to tubulin-binding herbicides in <i>Setaria viridis</i> . <i>Weed Research</i> , 2005, 45, 228-235.	1.7	9
60	Molecular evidence of biased inheritance of trifluralin herbicide resistance in foxtail millet. <i>Plant Breeding</i> , 2006, 125, 254-258.	1.9	9
61	Rapid isolation of both double-stranded RNA and PCR-suitable DNA from the obligate biotrophic phytopathogenic fungus <i>Uncinula necator</i> using a commercially available reagent. <i>Journal of Virological Methods</i> , 1998, 74, 149-153.	2.1	8
62	High conservation of the transcriptional response to acetolactate synthase-inhibiting herbicides across plant species. <i>Weed Research</i> , 2018, 58, 2-7.	1.7	8
63	Lab meets field: Accelerated selection and field monitoring concur that non-target-site-based resistance evolves first in the dicotyledonous, allergenic weed <i>Ambrosia artemisiifolia</i> . <i>Plant Science</i> , 2022, 317, 111202.	3.6	2
64	Herbicide Resistance in <i>Setaria</i> . <i>Plant Genetics and Genomics: Crops and Models</i> , 2017, , 251-266.	0.3	1