

# Henri Darmency

## List of Publications by Year in descending order

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Version: 2024-02-01

39  
papers

813  
citations

471509

17  
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501196

28  
g-index

40  
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40  
docs citations

40  
times ranked

654  
citing authors

#	ARTICLE	IF	CITATIONS
1	Estimated effects of cornflower presence on winter wheat. <i>Biological Agriculture and Horticulture</i> , 2022, 38, 113-123.	1.0	2
2	Relative success of frost-resistant variants of <i>Avena fatua</i> : a field experiment. <i>Journal of Agricultural Science</i> , 2020, 158, 558-563.	1.3	0
3	Morphological differences among <i>Raphanus raphanistrum</i> populations and their relationship to related crops. <i>Plant Breeding</i> , 2019, 138, 907-915.	1.9	2
4	Does genetic variability in weeds respond to non-chemical selection pressure in arable fields?. <i>Weed Research</i> , 2019, 59, 260-264.	1.7	8
5	A quantitative genetic examination of non-target-site resistance applied to <i>Avena</i> species. <i>Weed Research</i> , 2018, 58, 69-75.	1.7	5
6	Variation of seed dormancy and longevity in <i>Raphanus raphanistrum</i> L.. <i>Seed Science Research</i> , 2018, 28, 34-40.	1.7	5
7	Relationship between weed dormancy and herbicide rotations: implications in resistance evolution. <i>Pest Management Science</i> , 2017, 73, 1994-1999.	3.4	25
8	Gene Introgression in Weeds Depends on Initial Gene Location in the Crop: <i>Brassica napus</i> <i>Raphanus raphanistrum</i> Model. <i>Genetics</i> , 2017, 206, 1361-1372.	2.9	9
9	Simulating changes in cropping practises in conventional and glyphosate-tolerant maize. I. Effects on weeds. <i>Environmental Science and Pollution Research</i> , 2017, 24, 11582-11600.	5.3	23
10	Longevity, dormancy and germination of <i>Cyanus segetum</i> . <i>Weed Research</i> , 2017, 57, 361-371.	1.7	8
11	Assessing fitness parameters of hybrids between weed beets and transgenic sugar beets. <i>Plant Breeding</i> , 2017, 136, 969-976.	1.9	2
12	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
13	Consequences of Gene Flow between Transgenic, Insect-Resistant Crops and Their Wild Relatives. , 2016, , 423-430.		0
14	Effects of fragmentation and population size on the genetic diversity of <i>Centaurea cyanus</i> (Asteraceae) populations. <i>Plant Ecology and Evolution</i> , 2015, 148, 191-198.	0.7	8
15	Does gall midge larvae cause pre-dispersal seed mortality and limit cornflower population growth?. <i>Acta Oecologica</i> , 2015, 69, 167-172.	1.1	7
16	Variation of inbreeding depression in <i>Centaurea cyanus</i> L., a self-incompatible species. <i>Flora: Morphology, Distribution, Functional Ecology of Plants</i> , 2015, 212, 24-29.	1.2	12
17	Genetic diversity of the declining arable plant <i>Centaurea cyanus</i> : population fragmentation within an agricultural landscape is not associated with enhanced spatial genetic structure. <i>Weed Research</i> , 2014, 54, 436-444.	1.7	13
18	Pseudo-self-compatibility in <i>Centaurea cyanus</i> L. <i>Flora: Morphology, Distribution, Functional Ecology of Plants</i> , 2014, 209, 325-331.	1.2	14

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19	Spread of introgressed insect-resistance genes in wild populations of Brassica juncea: a simulated in-vivo approach. Transgenic Research, 2013, 22, 747-756.	2.4	23
20	Consequences of gene flow between oilseed rape (Brassica napus) and its relatives. Plant Science, 2013, 211, 42-51.	3.6	60
21	Pleiotropic effects of herbicide-resistance genes on crop yield: a review. Pest Management Science, 2013, 69, 897-904.	3.4	29
22	<i>Centaurea cyanus</i> as a biological indicator of segetal species richness in arable fields. Weed Research, 2012, 52, 551-563.	1.7	21
23	Identifying key life-traits for the dynamics and gene flow in a weedy crop relative: Sensitivity analysis of the GeneSys simulation model for weed beet ( <i>Beta vulgaris</i> ssp. <i>vulgaris</i> ). Ecological Modelling, 2010, 221, 225-237.	2.5	9
24	Backcrosses to Brassica napus of hybrids between B. juncea and B. napus as a source of herbicide-resistant volunteer-like feral populations. Plant Science, 2010, 179, 459-465.	3.6	34
25	Identifying key components of weed beet management using sensitivity analyses of the GeneSys model in GM sugar beet. Weed Research, 2009, 49, 581-591.	1.7	13
26	GeneSys-Beet: A model of the effects of cropping systems on gene flow between sugar beet and weed beet. Field Crops Research, 2008, 107, 245-256.	5.1	20
27	Transgene escape in sugar beet production fields: data from six years farm scale monitoring. Environmental Biosafety Research, 2007, 6, 197-206.	1.1	31
28	Crop and density effects on weed beet growth and reproduction. Weed Research, 2004, 44, 50-59.	1.7	21
29	Emergence and growth of hybrids between Brassica napus and Raphanus raphanistrum. New Phytologist, 2003, 158, 561-567.	7.3	23
30	Polymorphism for interspecific hybridisation within a population of wild radish ( <i>Raphanus</i> ) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 307 Td 169-172.	2.2	20
31	Assessment of interspecific hybridization between transgenic oilseed rape and wild radish under normal agronomic conditions. Theoretical and Applied Genetics, 2000, 100, 1233-1239.	3.6	79
32	Spontaneous hybridizations between oilseed rape and wild radish. Molecular Ecology, 1998, 7, 1467-1473.	3.9	73
33	The impact of hybrids between genetically modified crop plants and their related species: introgression and weediness. Molecular Ecology, 1994, 3, 37-40.	3.9	84
34	Genetic Diversity for Competitive and Reproductive Ability in Wild Oats ( <i>Avena fatua</i> ). Weed Science, 1992, 40, 215-219.	1.5	7
35	Variation of Transhexadecenoic Acid Content in Two Triazine Resistant Mutants of <i>Chenopodium album</i> and Their Susceptible Progenitor. Plant Physiology, 1988, 86, 967-970.	4.8	18
36	Abscisic-acid in triazine-resistant and susceptible poa annua. Plant Science, 1987, 49, 81-83.	3.6	6

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37	Character inheritance and polymorphism in a wild oat ( <i>Avena fatua</i> ) population. Canadian Journal of Botany, 1987, 65, 2352-2356.	1.1	6
38	Breeding foxtail millet ( <i>Setaria italica</i> ) for quantitative traits after interspecific hybridization and polyploidization. Genome, 1987, 29, 453-456.	2.0	33
39	INTERPRETING THE EVOLUTION OF A TRIAZINE-RESISTANT POPULATION OF <i>POA ANNUA</i> L.. New Phytologist, 1983, 95, 299-304.	7.3	26