

Marie-Hélène Dizier

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1658209/publications.pdf>

Version: 2024-02-01

63
papers

3,483
citations

361413

20
h-index

138484

58
g-index

64
all docs

64
docs citations

64
times ranked

5514
citing authors

#	ARTICLE	IF	CITATIONS
1	A Large-Scale, Consortium-Based Genomewide Association Study of Asthma. <i>New England Journal of Medicine</i> , 2010, 363, 1211-1221.	27.0	1,762
2	Effect of 17q21 Variants and Smoking Exposure in Early-Onset Asthma. <i>New England Journal of Medicine</i> , 2008, 359, 1985-1994.	27.0	351
3	Genome Screen for Asthma and Related Phenotypes in the French EGEA Study. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2000, 162, 1812-1818.	5.6	217
4	Meta-analysis identifies seven susceptibility loci involved in the atopic march. <i>Nature Communications</i> , 2015, 6, 8804.	12.8	148
5	Epidemiological Study of the Genetics and Environment of Asthma, Bronchial Hyperresponsiveness, and Atopy. <i>American Journal of Respiratory and Critical Care Medicine</i> , 1997, 156, S123-S129.	5.6	126
6	Meta-analysis of 20 genome-wide linkage studies evidenced new regions linked to asthma and atopy. <i>European Journal of Human Genetics</i> , 2010, 18, 700-706.	2.8	54
7	Detection of a recessive major gene for high IgE levels acting independently of specific response to allergens. <i>Genetic Epidemiology</i> , 1995, 12, 93-105.	1.3	52
8	EGEA (Epidemiological study on the Genetics and Environment of Asthma, bronchial) Tj ETQq0 0 0 rgBT /Overlock 10 Tf 50 462 Td (hyp	2.9	44
9	Clustering patterns of LOD scores for asthma-related phenotypes revealed by a genome-wide screen in 295 French EGEA families. <i>Human Molecular Genetics</i> , 2004, 13, 3103-3113.	2.9	36
10	Indication of linkage of serum IgE levels to the interleukin-4 gene and exclusion of the contribution of the (-590 C to T) interleukin-4 promoter polymorphism to IgE variation. <i>Genetic Epidemiology</i> , 1999, 16, 84-94.	1.3	35
11	Familial Resemblance of Asthma Severity in the EGEA* Study. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2002, 165, 185-189.	5.6	35
12	Evidence for gene Ã— smoking exposure interactions in a genome-wide linkage screen of asthma and bronchial hyper-responsiveness in EGEA families. <i>European Journal of Human Genetics</i> , 2007, 15, 810-815.	2.8	35
13	Associations between Nitric Oxide Synthase Genes and Exhaled NO-Related Phenotypes according to Asthma Status. <i>PLoS ONE</i> , 2012, 7, e36672.	2.5	33
14	Genome screen in the French EGEA study: detection of linked regions shared or not shared by allergic rhinitis and asthma. <i>Genes and Immunity</i> , 2005, 6, 95-102.	4.1	31
15	Sex-specific effect of IL9 polymorphisms on lung function and polysensitization. <i>Genes and Immunity</i> , 2009, 10, 559-565.	4.1	26
16	DNA methylation within melatonin receptor 1A (MTNR1A) mediates paternally transmitted genetic variant effect on asthma plus rhinitis. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 138, 748-753.	2.9	25
17	Identification of a new locus at 16q12 associated with time to asthma onset. <i>Journal of Allergy and Clinical Immunology</i> , 2016, 138, 1071-1080.	2.9	25
18	Scores of asthma and asthma severity reveal new regions of linkage in EGEA study families. <i>European Respiratory Journal</i> , 2007, 30, 253-259.	6.7	24

#	ARTICLE	IF	CITATIONS
19	Evidence for linkage of a new region (11p14) to eczema and allergic diseases. <i>Human Genetics</i> , 2008, 122, 605-614.	3.8	24
20	Genes Interacting with Occupational Exposures to Low Molecular Weight Agents and Irritants on Adult-Onset Asthma in Three European Studies. <i>Environmental Health Perspectives</i> , 2017, 125, 207-214.	6.0	23
21	Two-disease locus model: Sib pair method using information on both HLA and Gm. <i>Genetic Epidemiology</i> , 1986, 3, 343-356.	1.3	20
22	Confirmation of Psoriasis Susceptibility Loci on Chromosome 6p21 and 20p13 in French Families. <i>Journal of Investigative Dermatology</i> , 2007, 127, 1403-1409.	0.7	20
23	The triangle test statistic (TTS): a test of genetic homogeneity using departure from the triangle constraints in IBD distribution among affected sib pairs. <i>Annals of Human Genetics</i> , 2000, 64, 433-442.	0.8	19
24	The ANO3/MUC15 locus is associated with eczema in families ascertained through asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2012, 129, 1547-1553.e3.	2.9	18
25	Evidence for a pleiotropic QTL on chromosome 5q13 influencing both time to asthma onset and asthma score in French EGEA families. <i>Human Genetics</i> , 2007, 121, 711-719.	3.8	17
26	The nuclear factor I/A (NFIA) gene is associated with the asthma plus rhinitis phenotype. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 134, 576-582.e1.	2.9	17
27	Interaction between the DNAH9 gene and early smoke exposure in bronchial hyperresponsiveness. <i>European Respiratory Journal</i> , 2016, 47, 1072-1081.	6.7	17
28	European collaborative study of early-onset bipolar disorder: Evidence for genetic heterogeneity on 2q14 according to age at onset. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2010, 153B, 1425-1433.	1.7	16
29	Network-assisted analysis of GWAS data identifies a functionally-relevant gene module for childhood-onset asthma. <i>Scientific Reports</i> , 2017, 7, 938.	3.3	14
30	Evidence for a Locus in 1p31 Region Specifically Linked to the Co-Morbidity of Asthma and Allergic Rhinitis in the EGEA Study. <i>Human Heredity</i> , 2007, 63, 162-167.	0.8	13
31	Genetic heterogeneity according to age at onset in bipolar disorder: A combined positional cloning and candidate gene approach. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2012, 159B, 653-659.	1.7	13
32	Indication of linkage and genetic heterogeneity of asthma according to age at onset on chromosome 7q in 107 French EGEA families. <i>European Journal of Human Genetics</i> , 2001, 9, 867-872.	2.8	12
33	Replication of Association between ADAM33 Polymorphisms and Psoriasis. <i>PLoS ONE</i> , 2008, 3, e2448.	2.5	12
34	Segregation analysis of IgE levels in 335 French families (EGEA) using different strategies to correct for the ascertainment through a correlated trait (asthma). , 2000, 18, 128-142.		11
35	Familial correlations and inter-relationships of four asthma-associated quantitative phenotypes in 320 French EGEA families ascertained through asthmatic probands. <i>European Journal of Human Genetics</i> , 2004, 12, 955-963.	2.8	11
36	A common variant in RAB27A gene is associated with fractional exhaled nitric oxide levels in adults. <i>Clinical and Experimental Allergy</i> , 2015, 45, 797-806.	2.9	11

#	ARTICLE	IF	CITATIONS
37	Epidemiologic Study of the Genetics and Environment of Asthma, Bronchial Hyperresponsiveness, and Atopy. <i>Chest</i> , 2002, 121, 27S.	0.8	9
38	Testing linkage and Gene Ã— Environment interaction: Comparison of different affected sib-pair methods. <i>Genetic Epidemiology</i> , 2003, 25, 73-79.	1.3	9
39	A novel role for ciliary function in atopy: ADGRV1 and DNAH5 interactions. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 1659-1667.e11.	2.9	9
40	The <i>COL5A3</i> and <i>MMP9</i> genes interact in eczema susceptibility. <i>Clinical and Experimental Allergy</i> , 2018, 48, 297-305.	2.9	9
41	Genome-wide interaction study of early-life smoking exposure on time-to-asthma onset in childhood. <i>Clinical and Experimental Allergy</i> , 2019, 49, 1342-1351.	2.9	9
42	Genetics of emotional reactivity in bipolar disorders. <i>Journal of Affective Disorders</i> , 2015, 188, 101-106.	4.1	8
43	Gain of power of the general regression model compared to Cochran-Armitage Trend tests: simulation study and application to bipolar disorder. <i>BMC Genetics</i> , 2017, 18, 24.	2.7	8
44	A Gm Haplotype Study in Relation with HLA-DR in 155 Insulin-Dependent Diabetic Patients and Their Affected and Non Affected Siblings. <i>Experimental and Clinical Endocrinology and Diabetes</i> , 1987, 89, 325-332.	1.2	7
45	A New Correction for Multiple Testing in Gene-Gene Interaction Studies. <i>Annals of Human Genetics</i> , 2015, 79, 380-384.	0.8	7
46	Segregation analysis of two genetic markers in IDDM families under two-locus models. <i>Genetic Epidemiology</i> , 1989, 6, 71-75.	1.3	6
47	Genes Involved in Interleukin-1 Receptor Type II Activities Are Associated With Asthmatic Phenotypes. <i>Allergy, Asthma and Immunology Research</i> , 2016, 8, 466.	2.9	5
48	Two-disease-locus model: segregation analysis using information on two markers in nuclear families. Application to IDDM *. <i>Tissue Antigens</i> , 1990, 36, 1-7.	1.0	4
49	Comparative Power of Family-Based Association Strategies to Detect Disease-Causing Variants Under Two-Locus Models. <i>Genetic Epidemiology</i> , 2012, 36, 848-855.	1.3	4
50	Interactive effect between ATPase-related genes and early-life tobacco smoke exposure on bronchial hyper-responsiveness detected in asthma-ascertained families. <i>Thorax</i> , 2019, 74, 254-260.	5.6	4
51	Departure from the triangle constraints in discordant sib pairs: A test for genetic heterogeneity. <i>Genetic Epidemiology</i> , 1999, 17, S685-S689.	1.3	3
52	Indication of linkage and genetic heterogeneity for asthma and atopy on chromosomes 8p and 12q in 107 French EGEA families. <i>European Journal of Human Genetics</i> , 2003, 11, 590-596.	2.8	3
53	Identification of OCA2 as a novel locus for the comorbidity of asthma plus eczema. <i>Clinical and Experimental Allergy</i> , 2021, , .	2.9	3
54	PID1 is associated to a respiratory endotype related to occupational exposures to irritants. <i>Free Radical Biology and Medicine</i> , 2021, 172, 503-507.	2.9	3

#	ARTICLE	IF	CITATIONS
55	Genome-Wide Association Study of Fluorescent Oxidation Products Accounting for Tobacco Smoking Status in Adults from the French EGEA Study. <i>Antioxidants</i> , 2022, 11, 802.	5.1	3
56	Modeling the role of two susceptibility loci by the MASC method. <i>Genetic Epidemiology</i> , 1995, 12, 589-593.	1.3	1
57	Triangle Test Statistic in Discordant Sib Pairs: Test of Genetic Heterogeneity of Asthma and Atopy in CSCA Families. <i>Genetic Epidemiology</i> , 2001, 21, S192-7.	1.3	1
58	Homogeneity of Asthma Genome Scan Results. <i>Genetic Epidemiology</i> , 2001, 21, S44-8.	1.3	1
59	Impact of the diagnosis definition on linkage detection. <i>BMC Genetics</i> , 2005, 6, S140.	2.7	1
60	Genome-wide interaction study of environmental tobacco smoke exposure on time-to-asthma onset in childhood. , 2018, , .		1
61	General regression model: A "model-free" association test for quantitative traits allowing to test for the underlying genetic model. <i>Annals of Human Genetics</i> , 2020, 84, 280-290.	0.8	0
62	Génétique de l'asthme. <i>Medecine/Sciences</i> , 1991, 7, 1041.	0.2	0
63	Segregation analysis of IgE levels in 335 French families (EGEA) using different strategies to correct for the ascertainment through a correlated trait (asthma). <i>Genetic Epidemiology</i> , 2000, 18, 128.	1.3	0