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List of Publications by Year in descending order

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471509 395702 2,461 34 17 33 citations h-index g-index papers 36 36 36 5713 citing authors docs citations times ranked all docs

#	Article	IF	CITATIONS
1	Prediction and evaluation of the effect of pre-centrifugation sample management on the measurable untargeted LC-MS plasma metabolome. Analytica Chimica Acta, 2021, 1182, 338968.	5.4	9
2	A familyâ€based genomeâ€wide association study of recurrent aphthous stomatitis. Oral Diseases, 2020, 26, 1696-1705.	3.0	6
3	Radiation of the urinary bladder attenuates the development of lipopolysaccharide-induced cystitis. International Immunopharmacology, 2020, 83, 106334.	3.8	3
4	Altered peripheral amino acid profile indicate a systemic impact of active celiac disease and a possible role of amino acids in disease pathogenesis. PLoS ONE, 2018, 13, e0193764.	2.5	17
5	Different <i>>scp>DRB1*03:01â€<scp>DQB1</scp>*02:01</i> haplotypes confer different risk for celiac disease. Hla, 2017, 90, 95-101.	0.6	19
6	A family-based genome-wide association study of chronic rhinosinusitis with nasal polyps implicates several genes in the disease pathogenesis. PLoS ONE, 2017, 12, e0185244.	2.5	19
7	The angiotensin-converting enzyme gene insertion polymorphism: a higher risk for psoriasis in male patients. British Journal of Dermatology, 2016, 175, 824-826.	1.5	6
8	The influence of heredity versus environment on coeliac disease. Gut, 2016, 65, 1779-1780.	12.1	1
9	Role of proneurotensin as marker of paediatric coeliac disease. Clinical and Experimental Immunology, 2016, 186, 387-392.	2.6	5
10	Validity of histology for the diagnosis of paediatric coeliac disease: a Swedish multicentre study. Scandinavian Journal of Gastroenterology, 2016, 51, 427-433.	1.5	25
11	Shared Genetic Factors Involved in Celiac Disease, Type 2 Diabetes and Anorexia Nervosa Suggest Common Molecular Pathways for Chronic Diseases. PLoS ONE, 2016, 11, e0159593.	2.5	14
12	Single Nucleotide Polymorphisms in the FADS Gene Cluster but not the ELOVL2 Gene are Associated with Serum Polyunsaturated Fatty Acid Composition and Development of Allergy (in a Swedish Birth) Tj ETQq0 C	0 4gB T /O	veøløck 10 Tf
13	Characterisation of a Swedish cohort with orofacial granulomatosis with or without Crohn's disease. Oral Diseases, 2015, 21, e98-104.	3.0	28
14	Genes involved in muscle contractility and nutrient signaling pathways within celiac disease risk loci show differential mRNA expression. BMC Medical Genetics, 2015, 16, 44.	2.1	13
15	Association analysis of GWAS and candidate gene loci in a Pakistani population with psoriasis. Molecular Immunology, 2015, 64, 190-194.	2.2	30
16	Heredity of nasal polyps. Rhinology, 2015, 53, 25-28.	1.3	10
17	Study Designs for Exploring the Non-HLA Genetics in Celiac Disease. Methods in Molecular Biology, 2015, 1326, 35-44.	0.9	0
18	Haplotypes of the inducible nitric oxide synthase gene are strongly associated with exhaled nitric oxide levels in adults: a population-based study. Journal of Medical Genetics, 2014, 51, 449-454.	3.2	9

#	Article	IF	Citations
19	Genetic association study of QT interval highlights role for calcium signaling pathways in myocardial repolarization. Nature Genetics, 2014, 46, 826-836.	21.4	281
20	A Possible Mechanism behind Autoimmune Disorders Discovered By Genome-Wide Linkage and Association Analysis in Celiac Disease. PLoS ONE, 2013, 8, e70174.	2.5	51
21	Single nucleotide polymorphisms in the <i>NOS2</i> and <i>NOS3</i> genes are associated with exhaled nitric oxide. Journal of Medical Genetics, 2012, 49, 200-205.	3.2	24
22	Effect of Five Genetic Variants Associated with Lung Function on the Risk of Chronic Obstructive Lung Disease, and Their Joint Effects on Lung Function. American Journal of Respiratory and Critical Care Medicine, 2011, 184, 786-795.	5.6	128
23	Genome-wide association study identifies five loci associated with lung function. Nature Genetics, 2010, 42, 36-44.	21.4	518
24	A genome-wide association study identifies new psoriasis susceptibility loci and an interaction between HLA-C and ERAP1. Nature Genetics, 2010, 42, 985-990.	21.4	918
25	Meta-Analysis of Genome-Wide Linkage Studies in Celiac Disease. Human Heredity, 2009, 68, 223-230.	0.8	10
26	Association Between Genotypes and Phenotypes in Coeliac Disease. Journal of Pediatric Gastroenterology and Nutrition, 2009, 49, 165-169.	1.8	14
27	Association study of IL2/IL21 and FcgRIIa: significant association with the IL2/IL21 region in Scandinavian coeliac disease families. Genes and Immunity, 2008, 9, 364-367.	4.1	35
28	Searching for genes influencing a complex disease: the case of coeliac disease. European Journal of Human Genetics, 2008, 16, 542-553.	2.8	7
29	A comprehensive screen for SNP associations on chromosome region 5q31–33 in Swedish/Norwegian celiac disease families. European Journal of Human Genetics, 2007, 15, 980-987.	2.8	20
30	Meta and pooled analysis of European coeliac disease data. European Journal of Human Genetics, 2003, 11, 828-834.	2.8	79
31	A collaborative European search for non-DQA1*05-DQB1*02 celiac disease loci on HLA-DR3 haplotypes: analysis of transmission from homozygous parents. Human Immunology, 2003, 64, 350-358.	2.4	27
32	The IL12B gene does not confer susceptibility to coeliac disease. Tissue Antigens, 2002, 59, 70-72.	1.0	17
33	Variation in the CTLA4/CD28 gene region confers an increased risk of coeliac disease. Annals of Human Genetics, 2002, 66, 125-37.	0.8	11
34	Genome-wide linkage analysis of Scandinavian affected sib-pairs supports presence of susceptibility loci for celiac disease on chromosomes 5 and 11. European Journal of Human Genetics, 2001, 9, 938-944.	2.8	80