

Eva Ellinghaus

List of Publications by Year in descending order

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Version: 2024-02-01

56
papers

7,679
citations

81839

39
h-index

138417

58
g-index

63
all docs

63
docs citations

63
times ranked

13211
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Identification of 15 new psoriasis susceptibility loci highlights the role of innate immunity. <i>Nature Genetics</i> , 2012, 44, 1341-1348. | 9.4 | 848 |
| 2 | Analysis of five chronic inflammatory diseases identifies 27 new associations and highlights disease-specific patterns at shared loci. <i>Nature Genetics</i> , 2016, 48, 510-518. | 9.4 | 617 |
| 3 | Dense genotyping of immune-related disease regions identifies nine new risk loci for primary sclerosing cholangitis. <i>Nature Genetics</i> , 2013, 45, 670-675. | 9.4 | 339 |
| 4 | Genome-wide association study identifies a psoriasis susceptibility locus at TRAF3IP2. <i>Nature Genetics</i> , 2010, 42, 991-995. | 9.4 | 331 |
| 5 | Genome-wide association analysis identifies three psoriasis susceptibility loci. <i>Nature Genetics</i> , 2010, 42, 1000-1004. | 9.4 | 313 |
| 6 | Combined Analysis of Genome-wide Association Studies for Crohn Disease and Psoriasis Identifies Seven Shared Susceptibility Loci. <i>American Journal of Human Genetics</i> , 2012, 90, 636-647. | 2.6 | 290 |
| 7 | Association analyses identify six new psoriasis susceptibility loci in the Chinese population. <i>Nature Genetics</i> , 2010, 42, 1005-1009. | 9.4 | 287 |
| 8 | High-density mapping of the MHC identifies a shared role for HLA-DRB1*01:03 in inflammatory bowel diseases and heterozygous advantage in ulcerative colitis. <i>Nature Genetics</i> , 2015, 47, 172-179. | 9.4 | 280 |
| 9 | Large scale meta-analysis characterizes genetic architecture for common psoriasis associated variants. <i>Nature Communications</i> , 2017, 8, 15382. | 5.8 | 251 |
| 10 | Genome-wide Association Analysis of Psoriatic Arthritis and Cutaneous Psoriasis Reveals Differences in Their Genetic Architecture. <i>American Journal of Human Genetics</i> , 2015, 97, 816-836. | 2.6 | 245 |
| 11 | Genome-wide association analysis in primary sclerosing cholangitis identifies two non-HLA susceptibility loci. <i>Nature Genetics</i> , 2011, 43, 17-19. | 9.4 | 221 |
| 12 | Extended analysis of a genome-wide association study in primary sclerosing cholangitis detects multiple novel risk loci. <i>Journal of Hepatology</i> , 2012, 57, 366-375. | 1.8 | 196 |
| 13 | Genomics and drug profiling of fatal TCF3-HLF ^{hi} positive acute lymphoblastic leukemia identifies recurrent mutation patterns and therapeutic options. <i>Nature Genetics</i> , 2015, 47, 1020-1029. | 9.4 | 190 |
| 14 | Fine Mapping Major Histocompatibility Complex Associations in Psoriasis and Its Clinical Subtypes. <i>American Journal of Human Genetics</i> , 2014, 95, 162-172. | 2.6 | 182 |
| 15 | Genome-wide association study for ulcerative colitis identifies risk loci at 7q22 and 22q13 (IL17REL). <i>Nature Genetics</i> , 2010, 42, 292-294. | 9.4 | 177 |
| 16 | Gene Expression in Skin and Lymphoblastoid Cells: Refined Statistical Method Reveals Extensive Overlap in cis-eQTL Signals. <i>American Journal of Human Genetics</i> , 2010, 87, 779-789. | 2.6 | 169 |
| 17 | Genome-wide Comparative Analysis of Atopic Dermatitis and Psoriasis Gives Insight into Opposing Genetic Mechanisms. <i>American Journal of Human Genetics</i> , 2015, 96, 104-120. | 2.6 | 163 |
| 18 | Genetic correlations among psychiatric and immune-related phenotypes based on genome-wide association data. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2018, 177, 641-657. | 1.1 | 158 |

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|----|---|------|-----------|
| 19 | Enhanced meta-analysis and replication studies identify five new psoriasis susceptibility loci. <i>Nature Communications</i> , 2015, 6, 7001. | 5.8 | 156 |
| 20 | Genome-wide meta-analysis identifies multiple novel associations and ethnic heterogeneity of psoriasis susceptibility. <i>Nature Communications</i> , 2015, 6, 6916. | 5.8 | 154 |
| 21 | Genome-wide association analysis in Primary sclerosing cholangitis and ulcerative colitis identifies risk loci at <i>GPR35</i> and <i>TCF4</i> . <i>Hepatology</i> , 2013, 58, 1074-1083. | 3.6 | 150 |
| 22 | Association Between Variants of PRDM1 and NDP52 and Crohn's Disease, Based on Exome Sequencing and Functional Studies. <i>Gastroenterology</i> , 2013, 145, 339-347. | 0.6 | 149 |
| 23 | Whole-genome sequencing of a sporadic primary immunodeficiency cohort. <i>Nature</i> , 2020, 583, 90-95. | 13.7 | 148 |
| 24 | Epidermal lipid composition, barrier integrity, and eczematous inflammation are associated with skin microbiome configuration. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 1668-1676.e16. | 1.5 | 131 |
| 25 | Three ulcerative colitis susceptibility loci are associated with primary sclerosing cholangitis and indicate a role for <i>IL2</i> , <i>REL</i> , and <i>CARD9</i> . <i>Hepatology</i> , 2011, 53, 1977-1985. | 3.6 | 110 |
| 26 | Mutational Characterization of the Bile Acid Receptor TGR5 in Primary Sclerosing Cholangitis. <i>PLoS ONE</i> , 2010, 5, e12403. | 1.1 | 106 |
| 27 | Identification of germline susceptibility loci in ETV6-RUNX1-rearranged childhood acute lymphoblastic leukemia. <i>Leukemia</i> , 2012, 26, 902-909. | 3.3 | 106 |
| 28 | Genome-Wide Meta-Analysis of Psoriatic Arthritis Identifies Susceptibility Locus at REL. <i>Journal of Investigative Dermatology</i> , 2012, 132, 1133-1140. | 0.3 | 99 |
| 29 | Genetic signature to provide robust risk assessment of psoriatic arthritis development in psoriasis patients. <i>Nature Communications</i> , 2018, 9, 4178. | 5.8 | 95 |
| 30 | Identification of Immune-Relevant Factors Conferring Sarcoidosis Genetic Risk. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2015, 192, 727-736. | 2.5 | 94 |
| 31 | Reduced FOXP3+ regulatory T cells in patients with primary sclerosing cholangitis are associated with IL2RA gene polymorphisms. <i>Journal of Hepatology</i> , 2014, 60, 1010-1016. | 1.8 | 90 |
| 32 | Development of a high-resolution NGS-based HLA-typing and analysis pipeline. <i>Nucleic Acids Research</i> , 2015, 43, e70-e70. | 6.5 | 77 |
| 33 | Imputation of KIR Types from SNP Variation Data. <i>American Journal of Human Genetics</i> , 2015, 97, 593-607. | 2.6 | 73 |
| 34 | Association of CLEC16A with human common variable immunodeficiency disorder and role in murine B cells. <i>Nature Communications</i> , 2015, 6, 6804. | 5.8 | 63 |
| 35 | Conditional analysis identifies three novel major histocompatibility complex loci associated with psoriasis. <i>Human Molecular Genetics</i> , 2012, 21, 5185-5192. | 1.4 | 58 |
| 36 | Genome-wide association analysis for chronic venous disease identifies EFEMP1 and KCNH8 as susceptibility loci. <i>Scientific Reports</i> , 2017, 7, 45652. | 1.6 | 48 |

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|----|--|-----|-----------|
| 37 | Construction and benchmarking of a multi-ethnic reference panel for the imputation of HLA class I and II alleles. <i>Human Molecular Genetics</i> , 2019, 28, 2078-2092. | 1.4 | 48 |
| 38 | Fine mapping and replication of genetic risk loci in primary sclerosing cholangitis. <i>Scandinavian Journal of Gastroenterology</i> , 2012, 47, 820-826. | 0.6 | 47 |
| 39 | Microbiomarkers in inflammatory bowel diseases: caveats come with caviar. <i>Gut</i> , 2017, 66, 1734-1738. | 6.1 | 47 |
| 40 | Predicting Humoral Alloimmunity from Differences in Donor and Recipient HLA Surface Electrostatic Potential. <i>Journal of Immunology</i> , 2018, 201, 3780-3792. | 0.4 | 47 |
| 41 | A variant at 9p21.3 functionally implicates CDKN2B in paediatric B-cell precursor acute lymphoblastic leukaemia aetiology. <i>Nature Communications</i> , 2016, 7, 10635. | 5.8 | 44 |
| 42 | Exome-wide association study reveals novel psoriasis susceptibility locus at TNFSF15 and rare protective alleles in genes contributing to type I IFN signalling. <i>Human Molecular Genetics</i> , 2017, 26, 4301-4313. | 1.4 | 41 |
| 43 | Abundant Genetic Overlap between Blood Lipids and Immune-Mediated Diseases Indicates Shared Molecular Genetic Mechanisms. <i>PLoS ONE</i> , 2015, 10, e0123057. | 1.1 | 40 |
| 44 | Protein-coding variants contribute to the risk of atopic dermatitis and skin-specific gene expression. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 145, 1208-1218. | 1.5 | 29 |
| 45 | Association to the Glypican-5 gene in multiple sclerosis. <i>Journal of Neuroimmunology</i> , 2010, 226, 194-197. | 1.1 | 20 |
| 46 | Nonsyndromic cleft lip with or without cleft palate and cancer: Evaluation of a possible common genetic background through the analysis of GWAS data. <i>Genomics Data</i> , 2016, 10, 22-29. | 1.3 | 19 |
| 47 | Transethnic analysis of the human leukocyte antigen region for ulcerative colitis reveals not only shared but also ethnicity-specific disease associations. <i>Human Molecular Genetics</i> , 2021, 30, 356-369. | 1.4 | 19 |
| 48 | CD4+ T cells from patients with primary sclerosing cholangitis exhibit reduced apoptosis and down-regulation of proapoptotic Bim in peripheral blood. <i>Journal of Leukocyte Biology</i> , 2017, 101, 589-597. | 1.5 | 15 |
| 49 | Genetic association with B-cell acute lymphoblastic leukemia in allogeneic transplant patients differs by age and sex. <i>Blood Advances</i> , 2017, 1, 1717-1728. | 2.5 | 15 |
| 50 | Genome-wide association study of psoriasis in an Egyptian population. <i>Experimental Dermatology</i> , 2019, 28, 623-627. | 1.4 | 15 |
| 51 | Frequent and sex-biased deletion of SLX4IP by illegitimate V(D)J-mediated recombination in childhood acute lymphoblastic leukemia. <i>Human Molecular Genetics</i> , 2014, 23, 590-601. | 1.4 | 13 |
| 52 | Replication Study of Ulcerative Colitis Risk Loci in a Lithuanian/Latvian Case-Control Sample. <i>Inflammatory Bowel Diseases</i> , 2013, 19, 2349-2355. | 0.9 | 10 |
| 53 | Childhood acute lymphoblastic leukemia-associated risk-loci KZF1, ARID5B and CEBPE and risk of pediatric non-Hodgkin lymphoma: a report from the Berlin-Frankfurt-Münster Study Group. <i>Leukemia and Lymphoma</i> , 2015, 56, 814-816. | 0.6 | 9 |
| 54 | Genetic and transcriptional analysis of inflammatory bowel disease-associated pathways in patients with GUCY2C-linked familial diarrhea. <i>Scandinavian Journal of Gastroenterology</i> , 2018, 53, 1264-1273. | 0.6 | 9 |

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|----|--|-----|-----------|
| 55 | Large-Scale Imputation of KIR Copy Number and HLA Alleles in North American and European Psoriasis Case-Control Cohorts Reveals Association of Inhibitory KIR2DL2 With Psoriasis. <i>Frontiers in Immunology</i> , 2021, 12, 684326. | 2.2 | 7 |
| 56 | Genetic Variation in ABCC4 and CFTR and Acute Pancreatitis during Treatment of Pediatric Acute Lymphoblastic Leukemia. <i>Journal of Clinical Medicine</i> , 2021, 10, 4815. | 1.0 | 2 |