

Annette Hastie

List of Publications by Year in descending order

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49
papers

4,565
citations

147566

31
h-index

197535

49
g-index

49
all docs

49
docs citations

49
times ranked

6086
citing authors

#	ARTICLE	IF	CITATIONS
1	Sputum neutrophil counts are associated with more severe asthma phenotypes using cluster analysis. <i>Journal of Allergy and Clinical Immunology</i> , 2014, 133, 1557-1563.e5.	1.5	488
2	Analyses of asthma severity phenotypes and inflammatory proteins in subjects stratified by sputum granulocytes. <i>Journal of Allergy and Clinical Immunology</i> , 2010, 125, 1028-1036.e13.	1.5	405
3	Plasma interleukin-6 concentrations, metabolic dysfunction, and asthma severity: a cross-sectional analysis of two cohorts. <i>Lancet Respiratory Medicine</i> , 2016, 4, 574-584.	5.2	375
4	COVID-19-related Genes in Sputum Cells in Asthma. Relationship to Demographic Features and Corticosteroids. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 202, 83-90.	2.5	370
5	Airway Mucin Concentration as a Marker of Chronic Bronchitis. <i>New England Journal of Medicine</i> , 2017, 377, 911-922.	13.9	279
6	Biomarker surrogates do not accurately predict sputum eosinophil and neutrophil percentages in asthmatic subjects. <i>Journal of Allergy and Clinical Immunology</i> , 2013, 132, 72-80.e12.	1.5	224
7	Association of sputum and blood eosinophil concentrations with clinical measures of COPD severity: an analysis of the SPIROMICS cohort. <i>Lancet Respiratory Medicine</i> , 2017, 5, 956-967.	5.2	211
8	Baseline Features of the Severe Asthma Research Program (SARP III) Cohort: Differences with Age. <i>Journal of Allergy and Clinical Immunology: in Practice</i> , 2018, 6, 545-554.e4.	2.0	210
9	Extracellular DNA, Neutrophil Extracellular Traps, and Inflammasome Activation in Severe Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019, 199, 1076-1085.	2.5	165
10	Refractory airway type 2 inflammation in a large subgroup of asthmatic patients treated with inhaled corticosteroids. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, 104-113.e14.	1.5	135
11	Evidence for Exacerbation-Prone Asthma and Predictive Biomarkers of Exacerbation Frequency. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2020, 202, 973-982.	2.5	105
12	Comparison of Proteomic Assessment Methods in Multiple Cohort Studies. <i>Proteomics</i> , 2020, 20, e1900278.	1.3	103
13	Effect of rare variants in ADRB2 on risk of severe exacerbations and symptom control during longacting β_2 agonist treatment in a multiethnic asthma population: a genetic study. <i>Lancet Respiratory Medicine</i> , 2014, 2, 204-213.	5.2	100
14	Genome-wide association study identifies TH1 pathway genes associated with lung function in asthmatic patients. <i>Journal of Allergy and Clinical Immunology</i> , 2013, 132, 313-320.e15.	1.5	98
15	Obesity alters pathology and treatment response in inflammatory disease. <i>Nature</i> , 2022, 604, 337-342.	13.7	93
16	Multiview Cluster Analysis Identifies Variable Corticosteroid Response Phenotypes in Severe Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019, 199, 1358-1367.	2.5	91
17	Effects of Age and Disease Severity on Systemic Corticosteroid Responses in Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2017, 195, 1439-1448.	2.5	87
18	eQTL of bronchial epithelial cells and bronchial alveolar lavage deciphers GWAS-identified asthma genes. <i>Allergy: European Journal of Allergy and Clinical Immunology</i> , 2015, 70, 1309-1318.	2.7	82

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19	Airway mucin MUC5AC and MUC5B concentrations and the initiation and progression of chronic obstructive pulmonary disease: an analysis of the SPIROMICS cohort. <i>Lancet Respiratory Medicine</i> , 2021, 9, 1241-1254.	5.2	80
20	Differential Expression of TRAIL and TRAIL Receptors in Allergic Asthmatics Following Segmental Antigen Challenge: Evidence for a Role of TRAIL in Eosinophil Survival. <i>Journal of Immunology</i> , 2002, 169, 5986-5996.	0.4	76
21	Natural killer cell-mediated inflammation resolution is disabled in severe asthma. <i>Science Immunology</i> , 2017, 2, .	5.6	76
22	Mucus Plugs and Emphysema in the Pathophysiology of Airflow Obstruction and Hypoxemia in Smokers. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021, 203, 957-968.	2.5	71
23	Genetic analyses identify GSDMB associated with asthma severity, exacerbations, and antiviral pathways. <i>Journal of Allergy and Clinical Immunology</i> , 2021, 147, 894-909.	1.5	50
24	Complex association patterns for inflammatory mediators in induced sputum from subjects with asthma. <i>Clinical and Experimental Allergy</i> , 2018, 48, 787-797.	1.4	49
25	IL-6 trans-signaling increases expression of airways disease genes in airway smooth muscle. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2015, 309, L129-L138.	1.3	42
26	Design of a multi-center immunophenotyping analysis of peripheral blood, sputum and bronchoalveolar lavage fluid in the Subpopulations and Intermediate Outcome Measures in COPD Study (SPIROMICS). <i>Journal of Translational Medicine</i> , 2015, 13, 19.	1.8	41
27	Mixed Sputum Granulocyte Longitudinal Impact on Lung Function in the Severe Asthma Research Program. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021, 203, 882-892.	2.5	39
28	Investigation of the relationship between IL-6 and type 2 biomarkers in patients with severe asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 145, 430-433.	1.5	38
29	BAL Cell Gene Expression in Severe Asthma Reveals Mechanisms of Severe Disease and Influences of Medications. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2019, 200, 837-856.	2.5	37
30	Association of urine mitochondrial DNA with clinical measures of COPD in the SPIROMICS cohort. <i>JCI Insight</i> , 2020, 5, .	2.3	37
31	Lung microbiota associations with clinical features of COPD in the SPIROMICS cohort. <i>Npj Biofilms and Microbiomes</i> , 2021, 7, 14.	2.9	33
32	ALX receptor ligands define a biochemical endotype for severe asthma. <i>JCI Insight</i> , 2017, 2, .	2.3	29
33	The Impact of Insulin Resistance on Loss of Lung Function and Response to Treatment in Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2022, 206, 1096-1106.	2.5	28
34	Baseline sputum eosinophil ⁺ neutrophil subgroups [™] clinical characteristics and longitudinal trajectories for NHLBI Severe Asthma Research Program (SARP 3) cohort. <i>Journal of Allergy and Clinical Immunology</i> , 2020, 146, 222-226.	1.5	25
35	The Precision Interventions for Severe and/or Exacerbation-Prone (PreciSE) Asthma Network: An overview of Network organization, procedures, and interventions. <i>Journal of Allergy and Clinical Immunology</i> , 2022, 149, 488-516.e9.	1.5	24
36	Expression of asthma susceptibility genes in bronchial epithelial cells and bronchial alveolar lavage in the Severe Asthma Research Program (SARP) cohort. <i>Journal of Asthma</i> , 2016, 53, 775-782.	0.9	23

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37	IL-1RA regulates immunopathogenesis during fungal-associated allergic airway inflammation. <i>JCI Insight</i> , 2019, 4, .	2.3	21
38	The common \hat{I}^3 -chain cytokine IL-7 promotes immunopathogenesis during fungal asthma. <i>Mucosal Immunology</i> , 2018, 11, 1352-1362.	2.7	20
39	Identification of Sputum Biomarkers Predictive of Pulmonary Exacerbations in COPD. <i>Chest</i> , 2022, 161, 1239-1249.	0.4	20
40	Systemic Markers of Inflammation in Smokers With Symptoms Despite Preserved Spirometry in SPIROMICS. <i>Chest</i> , 2019, 155, 908-917.	0.4	18
41	Responsiveness to Parenteral Corticosteroids and Lung Function Trajectory in Adults with Moderate-to-Severe Asthma. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2021, 203, 841-852.	2.5	14
42	Alveolar eosinophilia in current smokers with chronic obstructive pulmonary disease in the SPIROMICS cohort. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 429-432.	1.5	12
43	The chemokine CX3CL1/fractalkine regulates immunopathogenesis during fungal-associated allergic airway inflammation. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2021, 320, L393-L404.	1.3	9
44	Alterations in vasodilator-stimulated phosphoprotein (VASP) phosphorylation: associations with asthmatic phenotype, airway inflammation and \hat{I}^2 -agonist use. <i>Respiratory Research</i> , 2006, 7, 25.	1.4	8
45	A Metabolomic Severity Score for Airflow Obstruction and Emphysema. <i>Metabolites</i> , 2022, 12, 368.	1.3	8
46	Chitinase 3-like-1 protects airway function despite promoting type 2 inflammation during fungal-associated allergic airway inflammation. <i>American Journal of Physiology - Lung Cellular and Molecular Physiology</i> , 2021, 320, L615-L626.	1.3	7
47	Effects of bronchoscopy on lung function in asthmatics. <i>Journal of Asthma</i> , 2017, 54, 866-871.	0.9	3
48	Tumour necrosis factor-related apoptosis-inducing ligand (TRAIL) and receptors in type 1, type 2 and type 17 inflammation in cross-sectional asthma study. <i>Thorax</i> , 2020, 75, 808-811.	2.7	3
49	Lung function, airway and peripheral basophils and eosinophils are associated with molecular pharmacogenomic endotypes of steroid response in severe asthma. <i>Thorax</i> , 2022, 77, 452-460.	2.7	3