

# Anthony Rowe

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8056780/publications.pdf>

Version: 2024-02-01

17  
papers

1,909  
citations

687363

13  
h-index

888059

17  
g-index

17  
all docs

17  
docs citations

17  
times ranked

3175  
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical and inflammatory characteristics of the European U-BIOPRED adult severe asthma cohort. <i>European Respiratory Journal</i> , 2015, 46, 1308-1321.	6.7	434
2	U-BIOPRED clinical adult asthma clusters linked to a subset of sputum omics. <i>Journal of Allergy and Clinical Immunology</i> , 2017, 139, 1797-1807.	2.9	236
3	Application of omics technologies to biomarker discovery in inflammatory lung diseases. <i>European Respiratory Journal</i> , 2013, 42, 802-825.	6.7	234
4	The burden of severe asthma in childhood and adolescence: results from the paediatric U-BIOPRED cohorts. <i>European Respiratory Journal</i> , 2015, 46, 1322-1333.	6.7	179
5	Sputum transcriptomics reveal upregulation of IL-1 receptor family members in patients with severe asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 560-570.	2.9	166
6	A Severe Asthma Disease Signature from Gene Expression Profiling of Peripheral Blood from U-BIOPRED Cohorts. <i>American Journal of Respiratory and Critical Care Medicine</i> , 2017, 195, 1311-1320.	5.6	152
7	Epithelial IL-6 trans-signaling defines a new asthma phenotype with increased airway inflammation. <i>Journal of Allergy and Clinical Immunology</i> , 2019, 143, 577-590.	2.9	140
8	Pathway discovery using transcriptomic profiles in adult-onset severe asthma. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 1280-1290.	2.9	105
9	The discovery net system for high throughput bioinformatics. <i>Bioinformatics</i> , 2003, 19, i225-i231.	4.1	67
10	Gene expression analysis in RA: towards personalized medicine. <i>Pharmacogenomics Journal</i> , 2014, 14, 93-106.	2.0	65
11	Improving data and knowledge management to better integrate health care and research. <i>Journal of Internal Medicine</i> , 2013, 274, 321-328.	6.0	44
12	A computational framework for complex disease stratification from multiple large-scale datasets. <i>BMC Systems Biology</i> , 2018, 12, 60.	3.0	43
13	The RA-MAP Consortium: a working model for academia-industry collaboration. <i>Nature Reviews Rheumatology</i> , 2018, 14, 53-60.	8.0	15
14	U-BIOPRED: evaluation of the value of a public-private partnership to industry. <i>Drug Discovery Today</i> , 2018, 23, 1622-1634.	6.4	14
15	Characterization of disease course and remission in early seropositive rheumatoid arthritis: results from the TACERA longitudinal cohort study. <i>Therapeutic Advances in Musculoskeletal Disease</i> , 2021, 13, 1759720X21110439.	2.7	6
16	Data and knowledge management in translational research: implementation of the eTRIKS platform for the IMI OncoTrack consortium. <i>BMC Bioinformatics</i> , 2019, 20, 164.	2.6	5
17	RA-MAP, molecular immunological landscapes in early rheumatoid arthritis and healthy vaccine recipients. <i>Scientific Data</i> , 2022, 9, 196.	5.3	4