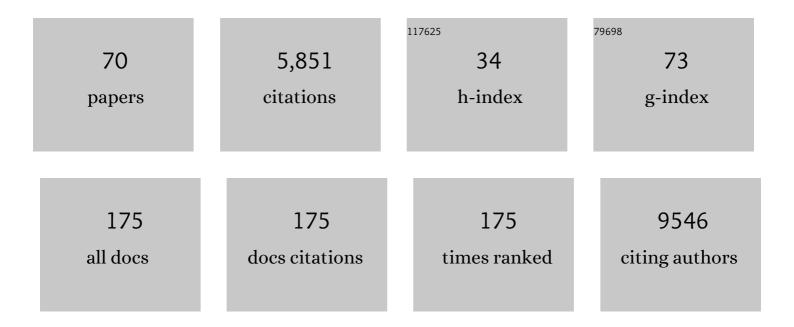
## **Charles Auffray**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8247154/publications.pdf Version: 2024-02-01



| #  | Article   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | DNA Methylation in Newborns and Maternal Smoking in Pregnancy: Genome-wide Consortium<br>Meta-analysis. American Journal of Human Genetics, 2016, 98, 680-696.  | 6.2  | 717       |
| 2  | Clinical and inflammatory characteristics of the European U-BIOPRED adult severe asthma cohort.<br>European Respiratory Journal, 2015, 46, 1308-1321.   | 6.7  | 434       |
| 3  | Systems medicine: the future of medical genomics and healthcare. Genome Medicine, 2009, 1, 2.   | 8.2  | 333       |
| 4  | U-BIOPRED clinical adult asthma clusters linked to a subset of sputum omics. Journal of Allergy and Clinical Immunology, 2017, 139, 1797-1807.  | 2.9  | 236       |
| 5  | Application of 'omics technologies to biomarker discovery in inflammatory lung diseases. European<br>Respiratory Journal, 2013, 42, 802-825.  | 6.7  | 234       |
| 6  | Revolutionizing medicine in the 21 <sup>st</sup> century through systems approaches. Biotechnology<br>Journal, 2012, 7, 992-1001.   | 3.5  | 225       |
| 7  | Making sense of big data in health research: Towards an EU action plan. Genome Medicine, 2016, 8, 71.   | 8.2  | 190       |
| 8  | The P4 Health Spectrum – A Predictive, Preventive, Personalized and Participatory Continuum for<br>Promoting Healthspan. Progress in Cardiovascular Diseases, 2017, 59, 506-521.                            | 3.1  | 178       |
| 9  | DNA methylation in childhood asthma: an epigenome-wide meta-analysis. Lancet Respiratory<br>Medicine,the, 2018, 6, 379-388.   | 10.7 | 170       |
| 10 | Sputum transcriptomics reveal upregulation of IL-1 receptor family members in patients with severe asthma. Journal of Allergy and Clinical Immunology, 2018, 141, 560-570.                                  | 2.9  | 166       |
| 11 | A Transcriptome-driven Analysis of Epithelial Brushings and Bronchial Biopsies to Define Asthma<br>Phenotypes in U-BIOPRED. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 443-455. | 5.6  | 165       |
| 12 | A Severe Asthma Disease Signature from Gene Expression Profiling of Peripheral Blood from<br>U-BIOPRED Cohorts. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 1311-1320.           | 5.6  | 152       |
| 13 | Mechanisms of the Development of Allergy (MeDALL): Introducing novel concepts in allergy phenotypes. Journal of Allergy and Clinical Immunology, 2017, 139, 388-399.  | 2.9  | 145       |
| 14 | Predictive, preventive, personalized and participatory medicine: back to the future. Genome Medicine, 2010, 2, 57.  | 8.2  | 144       |
| 15 | Participatory medicine: a driving force for revolutionizing healthcare. Genome Medicine, 2013, 5, 110.  | 8.2  | 137       |
| 16 | An Integrative Systems Biology Approach to Understanding Pulmonary Diseases. Chest, 2010, 137,<br>1410-1416.  | 0.8  | 135       |
| 17 | Genome-Wide Interaction Analysis of Air Pollution Exposure and Childhood Asthma with Functional<br>Follow-up. American Journal of Respiratory and Critical Care Medicine, 2017, 195, 1373-1383.             | 5.6  | 107       |
| 18 | Pathway discovery using transcriptomic profiles in adult-onset severe asthma. Journal of Allergy and<br>Clinical Immunology, 2018, 141, 1280-1290.  | 2.9  | 105       |

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|----|---|-----|-----------|
| 19 | "T2-high―in severe asthma related to blood eosinophil, exhaled nitric oxide andÂserum periostin.<br>European Respiratory Journal, 2019, 53, 1800938.                                | 6.7 | 104       |
| 20 | From functional genomics to systems biology: concepts and practices. Comptes Rendus - Biologies, 2003, 326, 879-892.  | 0.2 | 103       |
| 21 | COVID-19 Disease Map, building a computational repository of SARS-CoV-2 virus-host interaction mechanisms. Scientific Data, 2020, 7, 136.   | 5.3 | 99        |
| 22 | Systems medicine disease maps: community-driven comprehensive representation of disease mechanisms. Npj Systems Biology and Applications, 2018, 4, 21.                              | 3.0 | 84        |
| 23 | IL-17–high asthma with features of a psoriasis immunophenotype. Journal of Allergy and Clinical<br>Immunology, 2019, 144, 1198-1213.  | 2.9 | 80        |
| 24 | The road from systems biology to systems medicine. Pediatric Research, 2013, 73, 502-507.   | 2.3 | 78        |
| 25 | Enabling multiscale modeling in systems medicine. Genome Medicine, 2014, 6, 21.   | 8.2 | 76        |
| 26 | Scale relativity theory and integrative systems biology: 1. Progress in Biophysics and Molecular<br>Biology, 2008, 97, 79-114.  | 2.9 | 75        |
| 27 | Representing and querying disease networks using graph databases. BioData Mining, 2016, 9, 23.  | 4.0 | 75        |
| 28 | Exposure to Traffic-Related Air Pollution and Serum Inflammatory Cytokines in Children.<br>Environmental Health Perspectives, 2017, 125, 067007.                                    | 6.0 | 71        |
| 29 | Stratification of asthma phenotypes by airway proteomic signatures. Journal of Allergy and Clinical<br>Immunology, 2019, 144, 70-82.  | 2.9 | 59        |
| 30 | Scale relativity theory and integrative systems biology: 2 Macroscopic quantum-type mechanics.<br>Progress in Biophysics and Molecular Biology, 2008, 97, 115-157.                  | 2.9 | 56        |
| 31 | COVID19 Disease Map, a computational knowledge repository of virus–host interaction mechanisms.<br>Molecular Systems Biology, 2021, 17, e10387.                                     | 7.2 | 53        |
| 32 | The Genexpress IMAGE Knowledge Base of the Human Brain Transcriptome: A Prototype Integrated Resource for Functional and Computational Genomics. Genome Research, 1999, 9, 195-209. | 5.5 | 52        |
| 33 | Community-driven roadmap for integrated disease maps. Briefings in Bioinformatics, 2019, 20, 659-670.   | 6.5 | 48        |
| 34 | Towards a European health research and innovation cloud (HRIC). Genome Medicine, 2020, 12, 18.  | 8.2 | 46        |
| 35 | Transcriptomic gene signatures associated with persistent airflow limitation in patients with severe asthma. European Respiratory Journal, 2017, 50, 1602298.                       | 6.7 | 44        |
| 36 | A computational framework for complex disease stratification from multiple large-scale datasets.<br>BMC Systems Biology, 2018, 12, 60.  | 3.0 | 43        |

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|----|--|-----|-----------|
| 37 | Blood eosinophil count and airway epithelial transcriptome relationships in COPD versus asthma.<br>Allergy: European Journal of Allergy and Clinical Immunology, 2020, 75, 370-380.  | 5.7 | 37        |
| 38 | P4 Medicine Needs P4 Education. Current Pharmaceutical Design, 2014, 20, 6071-6072.  | 1.9 | 37        |
| 39 | Origins of Systems Biology in William Harvey's Masterpiece on the Movement of the Heart and the<br>Blood in Animals. International Journal of Molecular Sciences, 2009, 10, 1658-1669.   | 4.1 | 33        |
| 40 | Computational analysis of multimorbidity between asthma, eczema and rhinitis. PLoS ONE, 2017, 12, e0179125.  | 2.5 | 33        |
| 41 | From genomic medicine to precision medicine: highlights of 2015. Genome Medicine, 2016, 8, 12.   | 8.2 | 32        |
| 42 | Contribution of airway eosinophils in airway wall remodeling in asthma: Role of<br><i><scp>MMP</scp>â€10</i> and <i><scp>MET</scp></i> . Allergy: European Journal of Allergy and<br>Clinical Immunology, 2019, 74, 1102-1112.             | 5.7 | 32        |
| 43 | Longitudinally Stable, Clinically Defined Clusters of Patients with Asthma Independently Identified in the ADEPT and U-BIOPRED Asthma Studies. Annals of the American Thoracic Society, 2016, 13, S102-S103.                               | 3.2 | 30        |
| 44 | Self–organized living systems: conjunction of a stable organization with chaotic fluctuations in<br>biological space–time. Philosophical Transactions Series A, Mathematical, Physical, and Engineering<br>Sciences, 2003, 361, 1125-1139. | 3.4 | 29        |
| 45 | Systems Medicine: The Future of Medical Genomics, Healthcare, and Wellness. Methods in Molecular<br>Biology, 2016, 1386, 43-60.  | 0.9 | 29        |
| 46 | Editorial: Systems biology and personalized medicine – the future is now. Biotechnology Journal, 2012, 7, 938-939.   | 3.5 | 28        |
| 47 | Recon2Neo4j: applying graph database technologies for managing comprehensive genome-scale networks. Bioinformatics, 2017, 33, 1096-1098.   | 4.1 | 25        |
| 48 | Newt: a comprehensive web-based tool for viewing, constructing and analyzing biological maps.<br>Bioinformatics, 2021, 37, 1475-1477.  | 4.1 | 24        |
| 49 | AsthmaMap: An expertâ€driven computational representation of disease mechanisms. Clinical and Experimental Allergy, 2018, 48, 916-918.   | 2.9 | 21        |
| 50 | Prediction of chronic lung allograft dysfunction: a systems medicine challenge. European Respiratory<br>Journal, 2014, 43, 689-693.  | 6.7 | 20        |
| 51 | STON: exploring biological pathways using the SBCN standard and graph databases. BMC<br>Bioinformatics, 2016, 17, 494.   | 2.6 | 19        |
| 52 | Introducing PIONEER: a project to harness big data in prostate cancer research. Nature Reviews<br>Urology, 2020, 17, 351-362.  | 3.8 | 18        |
| 53 | EpiGeNet: A Graph Database of Interdependencies Between Genetic and Epigenetic Events in Colorectal<br>Cancer. Journal of Computational Biology, 2017, 24, 969-980.  | 1.6 | 16        |
| 54 | Epithelial dysregulation in obese severe asthmatics with gastro-oesophageal reflux. European<br>Respiratory Journal, 2019, 53, 1900453.  | 6.7 | 15        |

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|----|--|------------------|-------------|
| 55 | Instability of sputum molecular phenotypes in U-BIOPRED severe asthma. European Respiratory Journal,<br>2021, 57, 2001836.                                 | 6.7              | 13          |
| 56 | Ten years of Genome Medicine. Genome Medicine, 2019, 11, 7.  | 8.2              | 11          |
| 57 | Progress in integrative systems biology, physiology and medicine: towards a scale-relative biology.<br>European Physical Journal A, 2020, 56, 1.           | 2.5              | 11          |
| 58 | Looking back at genomic medicine in 2011. Genome Medicine, 2012, 4, 9.   | 8.2              | 10          |
| 59 | Sharing knowledge: a new frontier for public-private partnerships in medicine. Genome Medicine, 2009,<br>1, 29.  | 8.2              | 8           |
| 60 | Human-like layout algorithms for signalling hypergraphs: outlining requirements. Briefings in Bioinformatics, 2018, , .                                    | 6.5              | 8           |
| 61 | 2012 highlights in translational 'omics. Genome Medicine, 2013, 5, 10.   | 8.2              | 7           |
| 62 | Viva Europa, a Land of Excellence in Research and Innovation for Health and Wellbeing. Progress in<br>Preventive Medicine (New York, N Y ), 2017, 2, e006. | 0.7              | 6           |
| 63 | AsthmaMap: An interactive knowledge repository for mechanisms of asthma. Journal of Allergy and<br>Clinical Immunology, 2021, 147, 853-856.                | 2.9              | 6           |
| 64 | The Key Role of Patient Involvement in the Development of Core Outcome Sets in Prostate Cancer.<br>European Urology Focus, 2021, 7, 943-946.               | 3.1              | 6           |
| 65 | Foreword: from the Transcriptome conferences to the Systemoscope International Consortium.<br>Comptes Rendus - Biologies, 2003, 326, 867-875.              | 0.2              | 3           |
| 66 | COVID-19 and beyond:Âa call for action andÂaudacious solidarity to all the citizens and nations,Âit is<br>humanity's fight. F1000Research, 0, 9, 1130.     | 1.6              | 3           |
| 67 | Interview with a Thought Leader on Systems Medicine—Charles Auffray, PhD. Systems Medicine (New) Tj ETQq   | 1 1 0.784<br>1.1 | 314 rgBT /O |
| 68 | Computational Infrastructures for Data and Knowledge Management in Systems Biology. , 2013, ,<br>377-397.  |                  | 2           |
| 69 | A multi-omics data integration approach to identify a predictive molecular signature of CLAD. , 2015, , .  |                  | 2           |
| 70 | Informing epidemic (research) responses in a timely fashion by knowledge management - a Zika virus<br>use case. Biology Open, 2020, 9, .                   | 1.2              | 1           |