

Virginia M Pascual

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

13,405
citations

117625

34
h-index

289244

40
g-index

45
all docs

45
docs citations

45
times ranked

17055
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | Interferon and Granulopoiesis Signatures in Systemic Lupus Erythematosus Blood. <i>Journal of Experimental Medicine</i> , 2003, 197, 711-723. | 8.5 | 1,760 |
| 2 | An interferon-inducible neutrophil-driven blood transcriptional signature in human tuberculosis. <i>Nature</i> , 2010, 466, 973-977. | 27.8 | 1,632 |
| 3 | Human Blood CXCR5+CD4+ T Cells Are Counterparts of T Follicular Cells and Contain Specific Subsets that Differentially Support Antibody Secretion. <i>Immunity</i> , 2011, 34, 108-121. | 14.3 | 1,376 |
| 4 | Induction of Dendritic Cell Differentiation by IFN- α in Systemic Lupus Erythematosus. <i>Science</i> , 2001, 294, 1540-1543. | 12.6 | 1,160 |
| 5 | Plasmacytoid Dendritic Cells Induce Plasma Cell Differentiation through Type I Interferon and Interleukin 6. <i>Immunity</i> , 2003, 19, 225-234. | 14.3 | 929 |
| 6 | Type I Interferon in Systemic Lupus Erythematosus and Other Autoimmune Diseases. <i>Immunity</i> , 2006, 25, 383-392. | 14.3 | 840 |
| 7 | Extracellular Vesicle and Particle Biomarkers Define Multiple Human Cancers. <i>Cell</i> , 2020, 182, 1044-1061.e18. | 28.9 | 691 |
| 8 | A Modular Analysis Framework for Blood Genomics Studies: Application to Systemic Lupus Erythematosus. <i>Immunity</i> , 2008, 29, 150-164. | 14.3 | 623 |
| 9 | Personalized Immunomonitoring Uncovers Molecular Networks that Stratify Lupus Patients. <i>Cell</i> , 2016, 165, 551-565. | 28.9 | 524 |
| 10 | Oxidized mitochondrial nucleoids released by neutrophils drive type I interferon production in human lupus. <i>Journal of Experimental Medicine</i> , 2016, 213, 697-713. | 8.5 | 363 |
| 11 | TLR recognition of self nucleic acids hampers glucocorticoid activity in lupus. <i>Nature</i> , 2010, 465, 937-941. | 27.8 | 320 |
| 12 | Modular Transcriptional Repertoire Analyses of Adults With Systemic Lupus Erythematosus Reveal Distinct Type I and Type II Interferon Signatures. <i>Arthritis and Rheumatology</i> , 2014, 66, 1583-1595. | 5.6 | 302 |
| 13 | Systems Scale Interactive Exploration Reveals Quantitative and Qualitative Differences in Response to Influenza and Pneumococcal Vaccines. <i>Immunity</i> , 2013, 38, 831-844. | 14.3 | 284 |
| 14 | Systemic lupus erythematosus: all roads lead to type I interferons. <i>Current Opinion in Immunology</i> , 2006, 18, 676-682. | 5.5 | 254 |
| 15 | Increased Frequency of Pre-germinal Center B Cells and Plasma Cell Precursors in the Blood of Children with Systemic Lupus Erythematosus. <i>Journal of Immunology</i> , 2001, 167, 2361-2369. | 0.8 | 231 |
| 16 | Blood leukocyte microarrays to diagnose systemic onset juvenile idiopathic arthritis and follow the response to IL-1 blockade. <i>Journal of Experimental Medicine</i> , 2007, 204, 2131-2144. | 8.5 | 215 |
| 17 | Mapping systemic lupus erythematosus heterogeneity at the single-cell level. <i>Nature Immunology</i> , 2020, 21, 1094-1106. | 14.5 | 212 |
| 18 | TLR7 gain-of-function genetic variation causes human lupus. <i>Nature</i> , 2022, 605, 349-356. | 27.8 | 208 |

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|----|--|------|-----------|
| 19 | A CD4+ T cell population expanded in lupus blood provides B cell help through interleukin-10 and succinate. <i>Nature Medicine</i> , 2019, 25, 75-81. | 30.7 | 189 |
| 20 | A Genomic Approach to Human Autoimmune Diseases. <i>Annual Review of Immunology</i> , 2010, 28, 535-571. | 21.8 | 137 |
| 21 | IL1 Receptor Antagonist Controls Transcriptional Signature of Inflammation in Patients with Metastatic Breast Cancer. <i>Cancer Research</i> , 2018, 78, 5243-5258. | 0.9 | 119 |
| 22 | Host Immune Transcriptional Profiles Reflect the Variability in Clinical Disease Manifestations in Patients with <i>Staphylococcus aureus</i> Infections. <i>PLoS ONE</i> , 2012, 7, e34390. | 2.5 | 100 |
| 23 | Blood dendritic cells and DC-poietins in systemic lupus erythematosus. <i>Human Immunology</i> , 2002, 63, 1172-1180. | 2.4 | 92 |
| 24 | Erythroid mitochondrial retention triggers myeloid-dependent type I interferon in human SLE. <i>Cell</i> , 2021, 184, 4464-4479.e19. | 28.9 | 90 |
| 25 | The Transcriptional Signature of Active Tuberculosis Reflects Symptom Status in Extra-Pulmonary and Pulmonary Tuberculosis. <i>PLoS ONE</i> , 2016, 11, e0162220. | 2.5 | 81 |
| 26 | A narrow repertoire of transcriptional modules responsive to pyogenic bacteria is impaired in patients carrying loss-of-function mutations in MYD88 or IRAK4. <i>Nature Immunology</i> , 2014, 15, 1134-1142. | 14.5 | 75 |
| 27 | Functional rare and low frequency variants in BLK and BANK1 contribute to human lupus. <i>Nature Communications</i> , 2019, 10, 2201. | 12.8 | 73 |
| 28 | Differences in Antibody Responses Between Trivalent Inactivated Influenza Vaccine and Live Attenuated Influenza Vaccine Correlate With the Kinetics and Magnitude of Interferon Signaling in Children. <i>Journal of Infectious Diseases</i> , 2014, 210, 224-233. | 4.0 | 69 |
| 29 | Transcriptional profiling unveils type I and II interferon networks in blood and tissues across diseases. <i>Nature Communications</i> , 2019, 10, 2887. | 12.8 | 65 |
| 30 | Longitudinal profiling of human blood transcriptome in healthy and lupus pregnancy. <i>Journal of Experimental Medicine</i> , 2019, 216, 1154-1169. | 8.5 | 56 |
| 31 | Transcriptional specialization of human dendritic cell subsets in response to microbial vaccines. <i>Nature Communications</i> , 2014, 5, 5283. | 12.8 | 51 |
| 32 | Extracellular vesicle and particle-mediated communication shapes innate and adaptive immune responses. <i>Journal of Experimental Medicine</i> , 2021, 218, . | 8.5 | 47 |
| 33 | IFN Priming Is Necessary but Not Sufficient To Turn on a Migratory Dendritic Cell Program in Lupus Monocytes. <i>Journal of Immunology</i> , 2014, 192, 5586-5598. | 0.8 | 40 |
| 34 | The E3 ubiquitin ligase Itch inhibits p38 signaling and skin inflammation through the ubiquitylation of Tab1. <i>Science Signaling</i> , 2015, 8, ra22. | 3.6 | 37 |
| 35 | The immune roadmap for understanding multi-system inflammatory syndrome in children: opportunities and challenges. <i>Nature Medicine</i> , 2020, 26, 1819-1824. | 30.7 | 32 |
| 36 | Single Cell Analysis of Blood Mononuclear Cells Stimulated Through Either LPS or Anti-CD3 and Anti-CD28. <i>Frontiers in Immunology</i> , 2021, 12, 636720. | 4.8 | 32 |

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 37 | Development of a fixed module repertoire for the analysis and interpretation of blood transcriptome data. <i>Nature Communications</i> , 2021, 12, 4385. | 12.8 | 29 |
| 38 | The Blood Transcriptome of Experimental Melioidosis Reflects Disease Severity and Shows Considerable Similarity with the Human Disease. <i>Journal of Immunology</i> , 2015, 195, 3248-3261. | 0.8 | 20 |
| 39 | Analysis of Transcriptional Signatures in Response to <i>Listeria monocytogenes</i> Infection Reveals Temporal Changes That Result from Type I Interferon Signaling. <i>PLoS ONE</i> , 2016, 11, e0150251. | 2.5 | 10 |
| 40 | Mass Cytometry Defines Virus-Specific CD4+ T Cells in Influenza Vaccination. <i>ImmunoHorizons</i> , 2020, 4, 774-788. | 1.8 | 3 |
| 41 | Breaching self-tolerance by targeting the gatekeeper. <i>Journal of Experimental Medicine</i> , 2021, 218, . | 8.5 | 1 |