## David S Deluca

List of Publications by Year in descending order

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DAVID S DELLICA

#	Article	IF	CITATIONS
1	The Genotype-Tissue Expression (GTEx) project. Nature Genetics, 2013, 45, 580-585.	21.4	6,815
2	The Genotype-Tissue Expression (GTEx) pilot analysis: Multitissue gene regulation in humans. Science, 2015, 348, 648-660.	12.6	4,659
3	The human transcriptome across tissues and individuals. Science, 2015, 348, 660-665.	12.6	1,127
4	<i>SF3B1</i> and Other Novel Cancer Genes in Chronic Lymphocytic Leukemia. New England Journal of Medicine, 2011, 365, 2497-2506.	27.0	1,021
5	A Novel Approach to High-Quality Postmortem Tissue Procurement: The GTEx Project. Biopreservation and Biobanking, 2015, 13, 311-319.	1.0	674
6	Comparative analysis of RNA sequencing methods for degraded or low-input samples. Nature Methods, 2013, 10, 623-629.	19.0	419
7	Complementary genomic approaches highlight the PI3K/mTOR pathway as a common vulnerability in osteosarcoma. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E5564-73.	7.1	355
8	Effect of predicted protein-truncating genetic variants on the human transcriptome. Science, 2015, 348, 666-669.	12.6	252
9	Transcriptomic Characterization of SF3B1 Mutation Reveals Its Pleiotropic Effects in Chronic Lymphocytic Leukemia. Cancer Cell, 2016, 30, 750-763.	16.8	173
10	A Pan-Cancer Analysis of Transcriptome Changes Associated with Somatic Mutations in U2AF1 Reveals Commonly Altered Splicing Events. PLoS ONE, 2014, 9, e87361.	2.5	168
11	SOX2 and p63 colocalize at genetic loci in squamous cell carcinomas. Journal of Clinical Investigation, 2014, 124, 1636-1645.	8.2	151
12	Mutated <i>BCR-ABL</i> Generates Immunogenic T-cell Epitopes in CML Patients. Clinical Cancer Research, 2012, 18, 5761-5772.	7.0	57
13	MULTIPRED2: A computational system for large-scale identification of peptides predicted to bind to HLA supertypes and alleles. Journal of Immunological Methods, 2011, 374, 53-61.	1.4	55
14	Clinical Significance of SERPINA1 Gene and Its Encoded Alpha1-antitrypsin Protein in NSCLC. Cancers, 2019, 11, 1306.	3.7	52
15	Novel sequence feature variant type analysis of the HLA genetic association in systemic sclerosis. Human Molecular Genetics, 2010, 19, 707-719.	2.9	37
16	Amino acid 95 causes strong alteration of peptide position PÎ $^{\circ}$ in HLA-B*41 variants. Immunogenetics, 2007, 59, 253-259.	2.4	25
17	Complications and risk factors in pediatric bronchoscopy in a tertiary pediatric respiratory center. Pediatric Pulmonology, 2018, 53, 619-627.	2.0	24
18	The FMS-like tyrosine kinase-3 ligand/lung dendritic cell axis contributes to regulation of pulmonary fibrosis. Thorax, 2019, 74, 947-957.	5.6	24

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19	A modular concept of HLA for comprehensive peptide binding prediction. Immunogenetics, 2006, 59, 25-35.	2.4	22
20	Serum Levels of Alpha1-antitrypsin and Their Relationship With COPD in the General Spanish Population. Archivos De Bronconeumologia, 2020, 56, 76-83.	0.8	22
21	Filtration and Normalization of Sequencing Read Data in Whole-Metagenome Shotgun Samples. PLoS ONE, 2016, 11, e0165015.	2.5	22
22	ILâ€17 regulates DC migration to the peribronchial LNs and allergen presentation in experimental allergic asthma. European Journal of Immunology, 2020, 50, 1019-1033.	2.9	14
23	High-throughput minor histocompatibility antigen prediction. Bioinformatics, 2009, 25, 2411-2417.	4.1	11
24	MaHCO: an ontology of the major histocompatibility complex for immunoinformatic applications and text mining. Bioinformatics, 2009, 25, 2064-2070.	4.1	9
25	The Delivery of α1-Antitrypsin Therapy Through Transepidermal Route: Worthwhile to Explore. Frontiers in Pharmacology, 2020, 11, 983.	3.5	9
26	SERPINA1 gene polymorphisms in a populationâ€based ALSPAC cohort. Pediatric Pulmonology, 2019, 54, 1474-1478.	2.0	6
27	Single cell versus single nucleus: transcriptome differences in the murine kidney after ischemia-reperfusion injury. American Journal of Physiology - Renal Physiology, 2022, 323, F171-F181.	2.7	5
28	Smokers with COPD Show a Shift in Energy and Nitrogen Metabolism at Rest and During Exercise. International Journal of COPD, 2020, Volume 15, 1-13.	2.3	4
29	Polymerization of misfolded Z alpha-1 antitrypsin protein lowers CX3CR1 expression in human PBMCs. ELife, 2021, 10, .	6.0	4
30	Implementing the Modular MHC Model for Predicting Peptide Binding. Methods in Molecular Biology, 2007, 409, 261-271.	0.9	4
31	A comparison of curated gene sets versus transcriptomics-derived gene signatures for detecting pathway activation in immune cells. BMC Bioinformatics, 2020, 21, 28.	2.6	4
32	The Replacement Mutation in HLA-DRB1*1211 Affects a Likely Keystone Position. Human Immunology, 2005, 66, 1254-1257.	2.4	3
33	Human Leukocyte Antigen Typing Using a Knowledge Base Coupled with a High-Throughput Oligonucleotide Probe Array Analysis. Frontiers in Immunology, 2014, 5, 597.	4.8	3
34	SF3B1 Mutation Alters The Selection Of 3' RNA Splice Sites In Chronic Lymphocytic Leukemia. Blood, 2013, 122, 117-117.	1.4	2
35	Peptides Derived From Mutated BCR-ABL Elicit T Cell Immunity In CML Patients. Blood, 2010, 116, 887-887.	1.4	1
36	MicroRNA-449a Inhibits Triple Negative Breast Cancer by Disturbing DNA Repair and Chromatid Separation. International Journal of Molecular Sciences, 2022, 23, 5131.	4.1	1

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37	Factor VIII Is a Potential Autosomal B Cell Minor Histocompatibility Antigen in Chronic Gvhd Blood, 2009, 114, 1167-1167.	1.4	0
38	Systematic Identification of Personal Mutated Tumor-Specific Neoantigens in CLL. Blood, 2012, 120, 954-954.	1.4	0
39	Tumor Neoantigens Are Abundant Across Cancers. Blood, 2013, 122, 3265-3265.	1.4	0