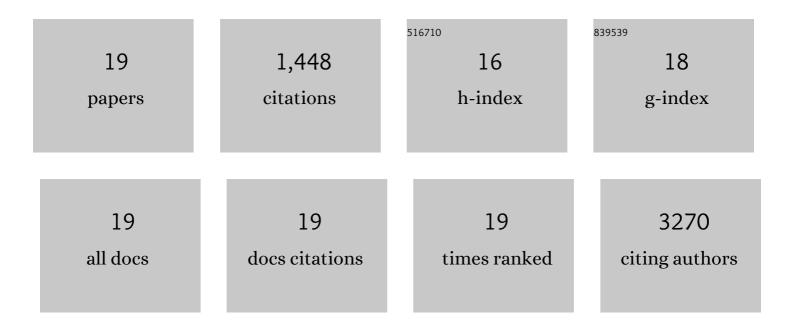
Alice M Walsh

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Integration of the Transcriptome and Genome-Wide Landscape of BRD2 and BRD4 Binding Motifs Identifies Key Superenhancer Genes and Reveals the Mechanism of Bet Inhibitor Action in Rheumatoid Arthritis Synovial Fibroblasts. Journal of Immunology, 2021, 206, 422-431.	0.8	23
2	Molecular correlates of response to nivolumab at baseline and on treatment in patients with RCC. , 2021, 9, e001506.		23
3	Elevated serum interleukin-8 is associated with enhanced intratumor neutrophils and reduced clinical benefit of immune-checkpoint inhibitors. Nature Medicine, 2020, 26, 688-692.	30.7	296
4	STK11 and KEAP1 mutations as prognostic biomarkers in an observational real-world lung adenocarcinoma cohort. ESMO Open, 2020, 5, e000706.	4.5	139
5	Clinical Validation of <i>PBRM1</i> Alterations as a Marker of Immune Checkpoint Inhibitor Response in Renal Cell Carcinoma. JAMA Oncology, 2019, 5, 1631.	7.1	166
6	Integrative analysis reveals CD38 as a therapeutic target for plasma cell-rich pre-disease and established rheumatoid arthritis and systemic lupus erythematosus. Arthritis Research and Therapy, 2018, 20, 85.	3.5	83
7	Immune checkpoint inhibitor PD-1 pathway is down-regulated in synovium at various stages of rheumatoid arthritis disease progression. PLoS ONE, 2018, 13, e0192704.	2.5	82
8	Enriched Cd141+ DCs in the joint are transcriptionally distinct, activated, and contribute to joint pathogenesis. JCI Insight, 2018, 3, .	5.0	30
9	Serum interleukin 8 (IL-8) may serve as a biomarker of response to immuno-oncology (I-O) therapy Journal of Clinical Oncology, 2018, 36, 3025-3025.	1.6	6
10	CD40L-Dependent Pathway Is Active at Various Stages of Rheumatoid Arthritis Disease Progression. Journal of Immunology, 2017, 198, 4490-4501.	0.8	73
11	Triple DMARD treatment in early rheumatoid arthritis modulates synovial T cell activation and plasmablast/plasma cell differentiation pathways. PLoS ONE, 2017, 12, e0183928.	2.5	33
12	Joint-specific DNA methylation and transcriptome signatures in rheumatoid arthritis identify distinct pathogenic processes. Nature Communications, 2016, 7, 11849.	12.8	104
13	Integrative genomic deconvolution of rheumatoid arthritis GWAS loci into gene and cell type associations. Genome Biology, 2016, 17, 79.	8.8	70
14	Sprouty2 Drives Drug Resistance and Proliferation in Glioblastoma. Molecular Cancer Research, 2015, 13, 1227-1237.	3.4	29
15	p53Î [°] is a transcriptionally inactive p53 isoform able to reprogram cells toward a metastatic-like state. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E3287-96.	7.1	73
16	Differential parsing of EGFR endocytic flux among parallel internalization pathways in lung cancer cells with EGFR-activating mutations. Integrative Biology (United Kingdom), 2014, 6, 312-323.	1.3	5
17	Regulation of EGFR trafficking and cell signaling by Sprouty2 and MIG6 in lung cancer cells. Journal of Cell Science, 2013, 126, 4339-4348.	2.0	26
18	Kinetic model for lamellipodal actin-integrin 'clutch' dynamics. Cell Adhesion and Migration, 2008, 2, 95-105.	2.7	39

#	Article	IF	CITATIONS
19	Cell patterning chip for controlling the stem cell microenvironment. Biomaterials, 2007, 28, 3208-3216.	11.4	148