## Willie June Brickey

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5507999/publications.pdf

Version: 2024-02-01

1040056 1281871 1,170 12 9 11 citations h-index g-index papers 12 12 12 2064 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Multi-omics analyses of radiation survivors identify radioprotective microbes and metabolites. Science, 2020, 370, .	12.6	260
2	Inflammasome-independent role of AIM2 in suppressing colon tumorigenesis via DNA-PK and Akt. Nature Medicine, 2015, 21, 906-913.	30.7	230
3	NLRP12 attenuates colon inflammation by maintaining colonic microbial diversity and promoting protective commensal bacterial growth. Nature Immunology, 2017, 18, 541-551.	14.5	225
4	The Inhibitory Innate Immune Sensor NLRP12 Maintains a Threshold against Obesity by Regulating Gut Microbiota Homeostasis. Cell Host and Microbe, 2018, 24, 364-378.e6.	11.0	158
5	GVHD-associated, inflammasome-mediated loss of function in adoptively transferred myeloid-derived suppressor cells. Blood, 2015, 126, 1621-1628.	1.4	104
6	NLRP12 Regulates Anti-viral RIG-I Activation via Interaction with TRIM25. Cell Host and Microbe, 2019, 25, 602-616.e7.	11.0	66
7	H2-DMαâ~'/â^' Mice Show the Importance of Major Histocompatibility Complexâ€"Bound Peptide in Cardiac Allograft Rejection. Journal of Experimental Medicine, 2000, 192, 31-40.	8.5	65
8	TRAF3IP3 negatively regulates cytosolic RNA induced anti-viral signaling by promoting TBK1 K48 ubiquitination. Nature Communications, 2020, 11, 2193.	12.8	33
9	The Toll–Like Receptor 2/6 Agonist, FSL–1 Lipopeptide, Therapeutically Mitigates Acute Radiation Syndrome. Scientific Reports, 2017, 7, 17355.	3.3	24
10	Re-Examination of the Exacerbating Effect of Inflammasome Components during Radiation Injury. Radiation Research, 2021, , .	1.5	4
11	Isolates of Salmonella typhimurium circumvent NLRP3 inflammasome recognition in macrophages during the chronic phase of infection. Journal of Biological Chemistry, 2022, 298, 101461.	3.4	1
12	Expression Profiling to Identify Candidate Genes Associated with Allergic Phenotypes. Methods in Molecular Biology, 2013, 1032, 287-295.	0.9	0