Naama Geva-Zatorsky

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/5890923/naama-geva-zatorsky-publications-by-year.pdf

Version: 2024-04-03

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

34	4,652	22	35
papers	citations	h-index	g-index
35	5,609	2 0.8 avg, IF	5.23
ext. papers	ext. citations		L-index

#	Paper	IF	Citations
34	Oral Capsulized Fecal Microbiota Transplantation for Eradication of Carbapenemase-producing Enterobacteriaceae Colonization With a Metagenomic Perspective. <i>Clinical Infectious Diseases</i> , 2021 , 73, e166-e175	11.6	15
33	Metagenomic analysis reveals the signature of gut microbiota associated with human chronotypes. <i>FASEB Journal</i> , 2021 , 35, e22011	0.9	3
32	Identification of bacteria-derived HLA-bound peptides in melanoma. <i>Nature</i> , 2021 , 592, 138-143	50.4	52
31	Strain-level immunomodulatory variation of gut bacteria. FEBS Letters, 2021, 595, 1322-1327	3.8	1
30	The hygiene hypothesis, the COVID pandemic, and consequences for the human microbiome. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021 , 118,	11.5	45
29	Analysis of a phase-variable restriction modification system of the human gut symbiont Bacteroides fragilis. <i>Nucleic Acids Research</i> , 2020 , 48, 11040-11053	20.1	2
28	Gut Bacteria-Not for the Faint of Heart. <i>Cell Host and Microbe</i> , 2020 , 27, 1-3	23.4	28
27	Microbial bile acid metabolites modulate gut RORl regulatory T cell homeostasis. <i>Nature</i> , 2020 , 577, 410-415	50.4	278
26	Gut microbiota - host interactions now also brain-immune axis. <i>Current Opinion in Neurobiology</i> , 2020 , 62, 53-59	7.6	12
25	Direct on-the-spot detection of SARS-CoV-2 in patients. <i>Experimental Biology and Medicine</i> , 2020 , 245, 1187-1193	3.7	23
24	Phage-Bacteria Associations: Analyze. Match. Develop Therapies. <i>Cell Host and Microbe</i> , 2020 , 28, 353-3	35 2 53.4	O
23	When Cultures Meet: The Landscape of "Social" Interactions between the Host and Its Indigenous Microbes. <i>BioEssays</i> , 2019 , 41, e1900002	4.1	3
22	What Came First: The Microbiota or the Tr(egg) Cells?. <i>Immunity</i> , 2018 , 48, 1072-1074	32.3	8
21	Gut microbes as a therapeutic armory. <i>Drug Discovery Today: Disease Models</i> , 2018 , 28, 51-59	1.3	2
20	Mining the Human Gut Microbiota for Immunomodulatory Organisms. <i>Cell</i> , 2017 , 168, 928-943.e11	56.2	356
19	Identifying species of symbiont bacteria from the human gut that, alone, can induce intestinal Th17 cells in mice. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, E8141-E8150	11.5	230
18	Bacteroides fragilis type VI secretion systems use novel effector and immunity proteins to antagonize human gut Bacteroidales species. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016 , 113, 3627-32	11.5	123

LIST OF PUBLICATIONS

17	MUCOSAL IMMUNOLOGY. Individual intestinal symbionts induce a distinct population of ROR⊞ regulatory T cells. <i>Science</i> , 2015 , 349, 993-7	33.3	487
16	In vivo imaging and tracking of host-microbiota interactions via metabolic labeling of gut anaerobic bacteria. <i>Nature Medicine</i> , 2015 , 21, 1091-100	50.5	129
15	Noise genetics: inferring protein function by correlating phenotype with protein levels and localization in individual human cells. <i>PLoS Genetics</i> , 2014 , 10, e1004176	6	15
14	Using bleach-chase to measure protein half-lives in living cells. <i>Nature Protocols</i> , 2012 , 7, 801-11	18.8	12
13	Dynamic proteomics of human protein level and localization across the cell cycle. <i>PLoS ONE</i> , 2012 , 7, e48722	3.7	15
12	Proteome half-life dynamics in living human cells. <i>Science</i> , 2011 , 331, 764-8	33.3	236
11	Dynamic Proteomics: a database for dynamics and localizations of endogenous fluorescently-tagged proteins in living human cells. <i>Nucleic Acids Research</i> , 2010 , 38, D508-12	20.1	22
10	Fourier analysis and systems identification of the p53 feedback loop. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010 , 107, 13550-5	11.5	73
9	Protein dynamics in drug combinations: a linear superposition of individual-drug responses. <i>Cell</i> , 2010 , 140, 643-51	56.2	80
8	Protein dynamics in individual human cells: experiment and theory. <i>PLoS ONE</i> , 2009 , 4, e4901	3.7	51
7	Dynamic proteomics of individual cancer cells in response to a drug. <i>Science</i> , 2008 , 322, 1511-6	33.3	467
6	Generation of a fluorescently labeled endogenous protein library in living human cells. <i>Nature Protocols</i> , 2007 , 2, 1515-27	18.8	58
5	Oscillations and variability in the p53 system. <i>Molecular Systems Biology</i> , 2006 , 2, 2006.0033	12.2	446
4	Laser autofocusing system for high-resolution cell biological imaging. <i>Journal of Microscopy</i> , 2006 , 221, 145-51	1.9	51
3	Dynamic proteomics in individual human cells uncovers widespread cell-cycle dependence of nuclear proteins. <i>Nature Methods</i> , 2006 , 3, 525-31	21.6	117
2	Variability and memory of protein levels in human cells. <i>Nature</i> , 2006 , 444, 643-6	50.4	440
1	Dynamics of the p53-Mdm2 feedback loop in individual cells. <i>Nature Genetics</i> , 2004 , 36, 147-50	36.3	772