## **Gautier Stoll**

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2807827/publications.pdf

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840585 1125617 14 816 11 13 citations h-index g-index papers 16 16 16 1216 all docs docs citations times ranked citing authors

#	Article	IF	Citations
1	A TLR3 Ligand Reestablishes Chemotherapeutic Responses in the Context of FPR1 Deficiency. Cancer Discovery, 2021, 11, 408-423.	7.7	28
2	Metabolic features of cancer cells impact immunosurveillance. , 2021, 9, e002362.		11
3	Autoimmunity affecting the biliary tract fuels the immunosurveillance of cholangiocarcinoma. Journal of Experimental Medicine, 2021, 218, .	4.2	20
4	Paradoxical implication of BAX/BAK in the persistence of tetraploid cells. Cell Death and Disease, 2021, 12, 1039.	2.7	7
5	Oxidative phosphorylation as a potential therapeutic target for cancer therapy. International Journal of Cancer, 2020, 146, 10-17.	2.3	125
6	Dynamical Boolean Modeling of Immunogenic Cell Death. Frontiers in Physiology, 2020, 11, 590479.	1.3	13
7	Immunoprophylactic and immunotherapeutic control of hormone receptor-positive breast cancer. Nature Communications, 2020, 11, 3819.	5.8	71
8	AIF-regulated oxidative phosphorylation supports lung cancer development. Cell Research, 2019, 29, 579-591.	5.7	58
9	PhysiBoSS: a multi-scale agent-based modelling framework integrating physical dimension and cell signalling. Bioinformatics, 2019, 35, 1188-1196.	1.8	88
10	Preclinical evaluation of a MAGE-A3 vaccination utilizing the oncolytic Maraba virus currently in first-in-human trials. Oncolmmunology, 2019, 8, e1512329.	2.1	53
11	Immune effectors responsible for the elimination of hyperploid cancer cells. OncoImmunology, 2018, 7, e1463947.	2.1	14
12	The CoLoMoTo Interactive Notebook: Accessible and Reproducible Computational Analyses for Qualitative Biological Networks. Frontiers in Physiology, 2018, 9, 680.	1.3	67
13	MaBoSS 2.0: an environment for stochastic Boolean modeling. Bioinformatics, 2017, 33, 2226-2228.	1.8	118
14	Continuous time boolean modeling for biological signaling: application of Gillespie algorithm. BMC Systems Biology, 2012, 6, 116.	3.0	134