## Michael Schuster

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	<scp>PDâ€L1</scp> overexpression correlates with <scp><i>JAK2</i>â€V617F</scp> mutational burden and is associated with 9p uniparental disomy in myeloproliferative neoplasms. American Journal of Hematology, 2022, 97, 390-400.	2.0	8
2	RANK links thymic regulatory T cells to fetal loss and gestational diabetes in pregnancy. Nature, 2021, 589, 442-447.	13.7	52
3	Acute BAF perturbation causes immediate changes in chromatin accessibility. Nature Genetics, 2021, 53, 269-278.	9.4	103
4	SARS-CoV-2 infects and replicates in cells of the human endocrine and exocrine pancreas. Nature Metabolism, 2021, 3, 149-165.	5.1	378
5	EZH2 inactivation in RAS-driven myeloid neoplasms hyperactivates RAS-signaling and increases MEK inhibitor sensitivity. Leukemia, 2021, 35, 1521-1526.	3.3	3
6	High-throughput drug screening identifies the ATR-CHK1 pathway as a therapeutic vulnerability of CALR mutated hematopoietic cells. Blood Cancer Journal, 2021, 11, 137.	2.8	1
7	Mutations and variants of ONECUT1 in diabetes. Nature Medicine, 2021, 27, 1928-1940.	15.2	24
8	Genomic epidemiology of superspreading events in Austria reveals mutational dynamics and transmission properties of SARS-CoV-2. Science Translational Medicine, 2020, 12, .	5.8	203
9	MTHFD1 interaction with BRD4 links folate metabolism to transcriptional regulation. Nature Genetics, 2019, 51, 990-998.	9.4	61
10	Mutational landscape of the transcriptome offers putative targets for immunotherapy of myeloproliferative neoplasms. Blood, 2019, 134, 199-210.	0.6	54
11	Mapping the chemical chromatin reactivation landscape identifies BRD4-TAF1 cross-talk. Nature Chemical Biology, 2016, 12, 504-510.	3.9	43
12	Whole-exome sequencing identifies novel MPL and JAK2 mutations in triple-negative myeloproliferative neoplasms. Blood, 2016, 127, 325-332.	0.6	228