Micah A Luftig

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Single-cell RNA-seq reveals transcriptomic heterogeneity mediated by host–pathogen dynamics in lymphoblastoid cell lines. ELife, 2021, 10, .	6.0	26
2	Evidence of Epsteinâ€Barr virus heterogeneous gene expression in adult lung transplant recipients with posttransplant lymphoproliferative disorder. Journal of Medical Virology, 2021, 93, 5040-5047.	5.0	2
3	Highly recurrent CBS epimutations in gastric cancer CpG island methylator phenotypes and inflammation. Genome Biology, 2021, 22, 167.	8.8	10
4	Monocarboxylate transporter antagonism reveals metabolic vulnerabilities of viral-driven lymphomas. Proceedings of the National Academy of Sciences of the United States of America, 2021, 118,	7.1	34
5	Massively parallel quantification of phenotypic heterogeneity in single-cell drug responses. Science Advances, 2021, 7, eabf9840.	10.3	9
6	Epstein-Barr Virus Genomes Reveal Population Structure and Type 1 Association with Endemic Burkitt Lymphoma. Journal of Virology, 2020, 94, .	3.4	20
7	The Role of EBV-Induced Hypermethylation in Gastric Cancer Tumorigenesis. Viruses, 2020, 12, 1222.	3.3	33
8	Identification of Host Biomarkers of Epstein-Barr Virus Latency IIb and Latency III. MBio, 2019, 10, .	4.1	20
9	The whole-genome landscape of Burkitt lymphoma subtypes. Blood, 2019, 134, 1598-1607.	1.4	113
10	Reprogramming of cellular metabolic pathways by human oncogenic viruses. Current Opinion in Virology, 2019, 39, 60-69.	5.4	20
11	Intracellular BH3 Profiling Reveals Shifts in Antiapoptotic Dependency in Human B Cell Maturation and Mitogen-Stimulated Proliferation. Journal of Immunology, 2018, 200, 1727-1736.	0.8	6
12	c-Myc Represses Transcription of Epstein-Barr Virus Latent Membrane Protein 1 Early after Primary B Cell Infection. Journal of Virology, 2018, 92, .	3.4	33
13	Molecular features and translational outlook for Epstein–Barr virus-associated gastric cancer. Future Virology, 2018, 13, 803-818.	1.8	3
14	DNA Tumour Viruses and the Host DNA Damage Response. , 2018, , .		0
15	Limited nucleotide pools restrict Epstein–Barr virus-mediated B-cell immortalization. Oncogenesis, 2017, 6, e349-e349.	4.9	26
16	The Epstein-Barr virus miR-BHRF1 microRNAs regulate viral gene expression in cis. Virology, 2017, 512, 113-123.	2.4	24
17	Epstein-Barr Virus Induces Adhesion Receptor CD226 (DNAM-1) Expression during Primary B-Cell Transformation into Lymphoblastoid Cell Lines. MSphere, 2017, 2,	2.9	8
18	Characterization of the EBV-Induced Persistent DNA Damage Response. Viruses, 2017, 9, 366.	3.3	17

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19	Epstein-Barr virus ensures B cell survival by uniquely modulating apoptosis at early and late times after infection. ELife, 2017, 6, .	6.0	54
20	Recent advances in understanding Epstein-Barr virus. F1000Research, 2017, 6, 386.	1.6	55
21	Metabolic stress is a barrier to Epstein–Barr virus-mediated B-cell immortalization. Proceedings of the United States of America, 2016, 113, E782-90.	7.1	94
22	To Be or Not IIb: A Multi-Step Process for Epstein-Barr Virus Latency Establishment and Consequences for B Cell Tumorigenesis. PLoS Pathogens, 2015, 11, e1004656.	4.7	121
23	Dynamic Epstein–Barr Virus Gene Expression on the Path to B-Cell Transformation. Advances in Virus Research, 2014, 88, 279-313.	2.1	73
24	Viruses and the DNA Damage Response: Activation and Antagonism. Annual Review of Virology, 2014, 1, 605-625.	6.7	124
25	Mitogen-Induced B-Cell Proliferation Activates Chk2-Dependent G1/S Cell Cycle Arrest. PLoS ONE, 2014, 9, e87299.	2.5	32
26	Interplay Between DNA Tumor Viruses and the Host DNA Damage Response. Current Topics in Microbiology and Immunology, 2013, 371, 229-257.	1.1	39
27	Use of Viral Systems to Study miRNA-Mediated Regulation of Gene Expression in Human Cells. Methods in Molecular Biology, 2013, 936, 143-156.	0.9	1
28	Epstein-Barr Virus Induces Global Changes in Cellular mRNA Isoform Usage That Are Important for the Maintenance of Latency. Journal of Virology, 2013, 87, 12291-12301.	3.4	12
29	Genetic heterogeneity of diffuse large B-cell lymphoma. Proceedings of the National Academy of Sciences of the United States of America, 2013, 110, 1398-1403.	7.1	494
30	Heavy LIFting: tumor promotion and radioresistance in NPC. Journal of Clinical Investigation, 2013, 123, 4999-5001.	8.2	22
31	A component of the mir-17-92 polycistronic oncomir promotes oncogene-dependent apoptosis. ELife, 2013, 2, e00822.	6.0	75
32	The Viral and Cellular MicroRNA Targetome in Lymphoblastoid Cell Lines. PLoS Pathogens, 2012, 8, e1002484.	4.7	321
33	Analysis of Epstein-Barr Virus-Regulated Host Gene Expression Changes through Primary B-Cell Outgrowth Reveals Delayed Kinetics of Latent Membrane Protein 1-Mediated NF-I®B Activation. Journal of Virology, 2012, 86, 11096-11106.	3.4	85
34	The Epstein-Barr Virus (EBV)-Induced Tumor Suppressor MicroRNA MiR-34a Is Growth Promoting in EBV-Infected B Cells. Journal of Virology, 2012, 86, 6889-6898.	3.4	81
35	The DNA damage response in viral-induced cellular transformation. British Journal of Cancer, 2012, 106, 429-435.	6.4	36
36	SplicerEX: A tool for the automated detection and classification of mRNA changes from conventional and splice-sensitive microarray expression data. Rna, 2012, 18, 1435-1445.	3.5	2

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37	Enhanced outgrowth of EBV-transformed chronic lymphocytic leukemia B cells mediated by coculture with macrophage feeder cells. Blood, 2012, 119, e35-e44.	1.4	12
38	The role of microRNAs in Epstein-Barr virus latency and lytic reactivation. Microbes and Infection, 2011, 13, 1156-1167.	1.9	56
39	Structure of Herpes Simplex Virus Glycoprotein D Bound to the Human Receptor Nectin-1. PLoS Pathogens, 2011, 7, e1002277.	4.7	154
40	At a crossroads: human DNA tumor viruses and the host DNA damage response. Future Virology, 2011, 6, 813-830.	1.8	23
41	Deep sequencing of the small RNA transcriptome of normal and malignant human B cells identifies hundreds of novel microRNAs. Blood, 2010, 116, e118-e127.	1.4	188
42	Virally Induced Cellular MicroRNA miR-155 Plays a Key Role in B-Cell Immortalization by Epstein-Barr Virus. Journal of Virology, 2010, 84, 11670-11678.	3.4	182
43	An ATM/Chk2-Mediated DNA Damage-Responsive Signaling Pathway Suppresses Epstein-Barr Virus Transformation of Primary Human B Cells. Cell Host and Microbe, 2010, 8, 510-522.	11.0	211
44	Alternative Splicing Is a Major Mechanism of Gene Regulation In Diffuse Large B Cell Lymphoma. Blood, 2010, 116, 803-803.	1.4	0
45	MDM2-Dependent Inhibition of p53 Is Required for Epstein-Barr Virus B-Cell Growth Transformation and Infected-Cell Survival. Journal of Virology, 2009, 83, 2491-2499.	3.4	53
46	Affinity maturation and characterization of a human monoclonal antibody against HIV-1 gp41. MAbs, 2009, 1, 462-474.	5.2	20
47	A Comprehensive Identification of the Microrna Transcriptome and Its Application in B Cell Malignancies Blood, 2009, 114, 2403-2403.	1.4	0
48	Structural basis for HIV-1 neutralization by a gp41 fusion intermediate–directed antibody. Nature Structural and Molecular Biology, 2006, 13, 740-747.	8.2	122
49	Epstein-Barr virus latent infection membrane protein 1 TRAF-binding site induces NIK/IKKÂ-dependent noncanonical NF-ÂB activation. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 141-146.	7.1	161
50	Latent infection membrane protein transmembrane FWLY is critical for intermolecular interaction, raft localization, and signaling. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 278-283.	7.1	62
51	Proteins of purified Epstein-Barr virus. Proceedings of the National Academy of Sciences of the United States of America, 2004, 101, 16286-16291.	7.1	383
52	Epstein-Barr virus latent membrane protein 1 activation of NF-ÂB through IRAK1 and TRAF6. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 15595-15600.	7.1	120
53	Effects of the NIK aly Mutation on NF-κB Activation by the Epstein-Barr Virus Latent Infection Membrane Protein, Lymphotoxin β Receptor, and CD40. Journal of Biological Chemistry, 2001, 276, 14602-14606.	3.4	36
54	Glycoprotein B of Human Herpesvirus 8 Is a Component of the Virion in a Cleaved Form Composed of Amino- and Carboxyl-Terminal Fragments. Virology, 2000, 269, 18-25.	2.4	41