## Romy Kerber

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Unique human immune signature of Ebola virus disease in Guinea. Nature, 2016, 533, 100-104.	13.7	170
2	The N-Terminal Domain of the Arenavirus L Protein Is an RNA Endonuclease Essential in mRNA Transcription. PLoS Pathogens, 2010, 6, e1001038.	2.1	145
3	Transcriptomic signatures differentiate survival from fatal outcomes in humans infected with Ebola virus. Genome Biology, 2017, 18, 4.	3.8	115
4	Structure of the Lassa Virus Nucleoprotein Revealed by X-ray Crystallography, Small-angle X-ray Scattering, and Electron Microscopy. Journal of Biological Chemistry, 2011, 286, 38748-38756.	1.6	47
5	Chimeric Mice with Competent Hematopoietic Immunity Reproduce Key Features of Severe Lassa Fever. PLoS Pathogens, 2016, 12, e1005656.	2.1	41
6	Laboratory Findings, Compassionate Use of Favipiravir, and Outcome in Patients With Ebola Virus Disease, Guinea, 2015—A Retrospective Observational Study. Journal of Infectious Diseases, 2019, 220, 195-202.	1.9	38
7	Evaluation of RealStar Reverse Transcription–Polymerase Chain Reaction Kits for Filovirus Detection in the Laboratory and Field. Journal of Infectious Diseases, 2016, 214, S243-S249.	1.9	33
8	Development of an antibody capture ELISA using inactivated Ebola Zaire Makona virus. Medical Microbiology and Immunology, 2016, 205, 173-183.	2.6	32
9	Ebola Virus Disease Is Characterized by Poor Activation and Reduced Levels of Circulating CD16 <sup>+</sup> Monocytes. Journal of Infectious Diseases, 2016, 214, S275-S280.	1.9	31
10	Deep Sequencing of RNA from Blood and Oral Swab Samples Reveals the Presence of Nucleic Acid from a Number of Pathogens in Patients with Acute Ebola Virus Disease and Is Consistent with Bacterial Translocation across the Gut. MSphere, 2017, 2, .	1.3	30
11	Cross-Species Analysis of the Replication Complex of Old World Arenaviruses Reveals Two Nucleoprotein Sites Involved in L Protein Function. Journal of Virology, 2011, 85, 12518-12528.	1.5	29
12	Kinetics of Soluble Mediators of the Host Response in Ebola Virus Disease. Journal of Infectious Diseases, 2018, 218, S496-S503.	1.9	25
13	Variation around the dominant viral genome sequence contributes to viral load and outcome in patients with Ebola virus disease. Genome Biology, 2020, 21, 238.	3.8	18
14	Heterologous arenavirus vector prime-boost overrules self-tolerance for efficient tumor-specific CD8 TÂcell attack. Cell Reports Medicine, 2021, 2, 100209.	3.3	16
15	Rift Valley fever virus minigenome system for investigating the role of L protein residues in viral transcription and replication. Journal of General Virology, 2019, 100, 1093-1098.	1.3	13