Aare Abroi

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/308429/publications.pdf

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		1039406	839053
17	319	9	18
papers	citations	h-index	g-index
20 all docs	20 docs citations	20 times ranked	397 citing authors

#	Article	IF	CITATIONS
1	Transcriptional and replicational activation functions in the bovine papillomavirus type 1 E2 protein are encoded by different structural determinants. Journal of Virology, 1996, 70, 6169-6179.	1.5	70
2	Analysis of Chromatin Attachment and Partitioning Functions of Bovine Papillomavirus Type 1 E2 Protein. Journal of Virology, 2004, 78, 2100-2113.	1.5	43
3	Are viruses a source of new protein folds for organisms? – Virosphere structure space and evolution. BioEssays, 2011, 33, 626-635.	1.2	37
4	Effect of Bovine Papillomavirus E2 Protein-Specific Monoclonal Antibodies on Papillomavirus DNA Replication. Journal of Virology, 1999, 73, 4670-4677.	1.5	28
5	Episomal Maintenance of Plasmids with Hybrid Origins in Mouse Cells. Journal of Virology, 2005, 79, 15277-15288.	1.5	25
6	Characterization of the Functional Activities of the Bovine Papillomavirus Type 1 E2 Protein Single-Chain Heterodimers. Journal of Virology, 2006, 80, 11218-11225.	1.5	22
7	Bovine papillomavirus type 1 E2 protein heterodimer is functional in papillomavirus DNA replication in vivo. Virology, 2009, 386, 353-359.	1.1	13
8	Did Viruses Evolve As a Distinct Supergroup from Common Ancestors of Cells?. Genome Biology and Evolution, 2016, 8, 2474-2481.	1.1	12
9	Conservation of the E8 CDS of the E8^E2 protein among mammalian papillomaviruses. Journal of General Virology, 2016, 97, 2333-2345.	1.3	11
10	Nuclear myosin 1 associates with papillomavirus E2 regulatory protein and influences viral replication. Virology, 2018, 514, 142-155.	1.1	9
11	The Common Partner of Several Methyltransferases TRMT112 Regulates the Expression of N6AMT1 Isoforms in Mammalian Cells. Biomolecules, 2019, 9, 422.	1.8	9
12	Human TRMT112-Methyltransferase Network Consists of Seven Partners Interacting with a Common Co-Factor. International Journal of Molecular Sciences, 2021, 22, 13593.	1.8	8
13	A protein domain-based view of the virosphere–host relationship. Biochimie, 2015, 119, 231-243.	1.3	7
14	Protein Structure-Guided Hidden Markov Models (HMMs) as A Powerful Method in the Detection of Ancestral Endogenous Viral Elements. Viruses, 2019, 11, 320.	1.5	7
15	The Enigmatic Origin of Papillomavirus Protein Domains. Viruses, 2017, 9, 240.	1.5	6
16	Characterization of the nuclear matrix targeting sequence (NMTS) of the BPV1 E8/E2 protein — the shortest known NMTS. Nucleus, 2015, 6, 289-300.	0.6	5
17	cRegions—a tool for detecting conserved cis-elements in multiple sequence alignment of diverged coding sequences. PeerJ, 2019, 6, e6176.	0.9	2