

David Forgacs

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4097003/publications.pdf>

Version: 2024-02-01

14
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1040056

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#	ARTICLE	IF	CITATIONS
1	Immunisation of ferrets and mice with recombinant SARS-CoV-2 spike protein formulated with Advax-SM adjuvant protects against COVID-19 infection. <i>Vaccine</i> , 2021, 39, 5940-5953.	3.8	44
2	PARIS and SPARTA: Finding the Achillesâ€™ Heel of SARS-CoV-2. <i>MSphere</i> , 2022, 7, e0017922.	2.9	25
3	SARS-CoV-2 mRNA Vaccines Elicit Different Responses in Immunologically Naïve and Pre-Immune Humans. <i>Frontiers in Immunology</i> , 2021, 12, 728021.	4.8	20
4	Convergent antibody evolution and clonotype expansion following influenza virus vaccination. <i>PLoS ONE</i> , 2021, 16, e0247253.	2.5	19
5	Antibody affinity maturation and cross-variant activity following SARS-CoV-2 mRNA vaccination: Impact of prior exposure and sex. <i>EBioMedicine</i> , 2021, 74, 103748.	6.1	17
6	Genomic evaluation of hybridization in historic and modern North American Bison (<i>Bison bison</i>). <i>Scientific Reports</i> , 2022, 12, 6397.	3.3	17
7	The spatial genetic differentiation of the legume pod borer, <i>Maruca vitrata</i> F. (Lepidoptera: Tj ETQq1 1 0.784314 rgBT /Overloc 12	1.0	12
8	Mitochondrial Genome Analysis Reveals Historical Lineages in Yellowstone Bison. <i>PLoS ONE</i> , 2016, 11, e0166081.	2.5	11
9	The Effect of Waning on Antibody Levels and Memory B Cell Recall following SARS-CoV-2 Infection or Vaccination. <i>Vaccines</i> , 2022, 10, 696.	4.4	11
10	Development of SNP-Based Genomic Tools for the Canadian Bison Industry: Parentage Verification and Subspecies Composition. <i>Frontiers in Genetics</i> , 2020, 11, 585999.	2.3	10
11	The Expectations and Challenges of Wildlife Disease Research in the Era of Genomics: Forecasting with a Horizon Scan-like Exercise. <i>Journal of Heredity</i> , 2019, 110, 261-274.	2.4	9
12	Evaluation of fecal samples as a valid source of DNA by comparing paired blood and fecal samples from American bison (<i>Bison bison</i>). <i>BMC Genetics</i> , 2019, 20, 22.	2.7	7
13	Presence of diverse <i>Rickettsia</i> spp. and absence of <i>Borrelia burgdorferi</i> sensu lato in ticks in an East Texas forest with reduced tick density associated with controlled burns. <i>Ticks and Tick-borne Diseases</i> , 2020, 11, 101310.	2.7	6
14	An Expanded Reverse Line Blot Hybridization Protocol for the Simultaneous Detection of Numerous Tick-Borne Pathogens in North America. <i>Journal of Medical Entomology</i> , 2016, 53, 721-726.	1.8	1