

Margaret Littlejohn

List of Publications by Year in descending order

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Version: 2024-02-01

48
papers

1,930
citations

304368

22
h-index

264894

42
g-index

50
all docs

50
docs citations

50
times ranked

2355
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|-----|-----------|
| 1 | Serum hepatitis B surface antigen and hepatitis B e antigen titers: Disease phase influences correlation with viral load and intrahepatic hepatitis B virus markers. <i>Hepatology</i> , 2010, 51, 1933-1944. | 3.6 | 366 |
| 2 | The phenylpropenamide derivative AT-130 blocks HBV replication at the level of viral RNA packaging. <i>Antiviral Research</i> , 2007, 76, 168-177. | 1.9 | 141 |
| 3 | Long-Term Entecavir Treatment Results in Sustained Antiviral Efficacy and Prolonged Life Span in the Woodchuck Model of Chronic Hepatitis Infection. <i>Journal of Infectious Diseases</i> , 2001, 184, 1236-1245. | 1.9 | 140 |
| 4 | The evolution and clinical impact of hepatitis B virus genome diversity. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2020, 17, 618-634. | 8.2 | 107 |
| 5 | Insights From Deep Sequencing of the HBV Genome—Unique, Tiny, and Misunderstood. <i>Gastroenterology</i> , 2019, 156, 384-399. | 0.6 | 92 |
| 6 | Origins and Evolution of Hepatitis B Virus and Hepatitis D Virus. <i>Cold Spring Harbor Perspectives in Medicine</i> , 2016, 6, a021360. | 2.9 | 84 |
| 7 | Possible origins and evolution of the hepatitis B virus (HBV). <i>Seminars in Cancer Biology</i> , 2013, 23, 561-575. | 4.3 | 76 |
| 8 | Patterns and Causes of Suboptimal Response to Tenofovir-Based Therapy in Individuals Coinfected With HIV and Hepatitis B Virus. <i>Clinical Infectious Diseases</i> , 2013, 56, e87-e94. | 2.9 | 69 |
| 9 | A Divergent Hepatitis D-Like Agent in Birds. <i>Viruses</i> , 2018, 10, 720. | 1.5 | 69 |
| 10 | SEQHEPB: A sequence analysis program and relational database system for chronic hepatitis B. <i>Antiviral Research</i> , 2007, 75, 64-74. | 1.9 | 68 |
| 11 | <i>In Vitro</i> Studies Show that Sequence Variability Contributes to Marked Variation in Hepatitis B Virus Replication, Protein Expression, and Function Observed across Genotypes. <i>Journal of Virology</i> , 2016, 90, 10054-10064. | 1.5 | 49 |
| 12 | The molecular epidemiology of hepatitis B in the Indigenous people of northern Australia. <i>Journal of Gastroenterology and Hepatology (Australia)</i> , 2013, 28, 1234-1241. | 1.4 | 47 |
| 13 | Deep sequencing shows that HBV basal core promoter and precore variants reduce the likelihood of HBsAg loss following tenofovir disoproxil fumarate therapy in HBeAg-positive chronic hepatitis B. <i>Gut</i> , 2017, 66, 2013-2023. | 6.1 | 42 |
| 14 | Quantitative HBsAg and HBeAg Predict Hepatitis B Seroconversion after Initiation of HAART in HIV-HBV Coinfected Individuals. <i>PLoS ONE</i> , 2013, 8, e61297. | 1.1 | 41 |
| 15 | Molecular virology of hepatitis B virus, subgenotype C4 in northern Australian Indigenous populations. <i>Journal of Medical Virology</i> , 2014, 86, 695-706. | 2.5 | 38 |
| 16 | Analysis of genomic-length HBV sequences to determine genotype and subgenotype reference sequences. <i>Journal of General Virology</i> , 2020, 101, 271-283. | 1.3 | 38 |
| 17 | Identification of a novel hepatitis B virus precore/core deletion mutant in HIV/hepatitis B virus co-infected individuals. <i>Aids</i> , 2007, 21, 1701-1710. | 1.0 | 36 |
| 18 | Quantification of HBV Covalently Closed Circular DNA from Liver Tissue by Real-Time PCR. , 2004, 95, 41-50. | | 31 |

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|----|--|-----|-----------|
| 19 | Molecular and clinical characteristics of hepatitis B virus in Korea. <i>Journal of Medical Virology</i> , 2010, 82, 1126-1134. | 2.5 | 31 |
| 20 | HBV mutations in untreated HIV-HBV co-infection using genomic length sequencing. <i>Virology</i> , 2010, 405, 539-547. | 1.1 | 28 |
| 21 | <scp>ALT</scp> flares during nucleotide analogue therapy are associated with <scp>HB</scp>sAg loss in genotype A <scp>HB</scp>eAgâ€positive chronic hepatitis B. <i>Liver International</i> , 2018, 38, 1760-1769. | 1.9 | 26 |
| 22 | Tracing Ancient Human Migrations into Sahul Using Hepatitis B Virus Genomes. <i>Molecular Biology and Evolution</i> , 2019, 36, 942-954. | 3.5 | 26 |
| 23 | Defective hepatitis B virus DNA is not associated with disease status but is reduced by polymerase mutations associated with drug resistance. <i>Hepatology</i> , 2008, 48, 741-749. | 3.6 | 24 |
| 24 | Molecular epidemiology of hepatitis delta virus in the Western Pacific region. <i>Journal of Clinical Virology</i> , 2014, 61, 34-39. | 1.6 | 23 |
| 25 | Hepatitis Delta Virus (HDV) and Delta-Like Agents: Insights Into Their Origin. <i>Frontiers in Microbiology</i> , 2021, 12, 652962. | 1.5 | 22 |
| 26 | Long-Term TDF-Inclusive ART and Progressive Rates of HBsAg Loss in HIV-HBV Coinfectionâ€Lessons for Functional HBV Cure?. <i>Journal of Acquired Immune Deficiency Syndromes (1999)</i> , 2020, 84, 527-533. | 0.9 | 20 |
| 27 | Quantitative analysis of the splice variants expressed by the major hepatitis B virus genotypes. <i>Microbial Genomics</i> , 2021, 7, . | 1.0 | 16 |
| 28 | Stopping nucleot(s)ide analogues in nonâ€cirrhotic <scp>HBeAg</scp>â€negative chronic hepatitis B patients: <scp>HBsAg</scp> loss at 96â€weeks is associated with low baseline <scp>HBsAg</scp> levels. <i>Alimentary Pharmacology and Therapeutics</i> , 2022, 56, 310-320. | 1.9 | 16 |
| 29 | Effects of long-term tenofovir-based combination antiretroviral therapy in HIV-hepatitis B virus coinfection on persistent hepatitis B virus viremia and the role of hepatitis B virus quasispecies diversity. <i>Aids</i> , 2016, 30, 1597-1606. | 1.0 | 15 |
| 30 | Evaluation of dried blood spots for hepatitis B and D serology and nucleic acid testing. <i>Journal of Medical Virology</i> , 2022, 94, 642-648. | 2.5 | 14 |
| 31 | Origins and Evolution of the Primate Hepatitis B Virus. <i>Frontiers in Microbiology</i> , 2021, 12, 653684. | 1.5 | 14 |
| 32 | Molecular characterization of hepatitis B virus (HBV) in African children living in Australia identifies genotypes and variants associated with poor clinical outcome. <i>Journal of General Virology</i> , 2018, 99, 1103-1114. | 1.3 | 12 |
| 33 | Hepatitis B and D in the Pacific Islands of Kiribati. <i>Journal of Clinical Virology</i> , 2020, 129, 104527. | 1.6 | 11 |
| 34 | In vitro replication phenotype of a novel (âˆ1G) hepatitis B virus variant associated with HIV coâ€infection. <i>Journal of Medical Virology</i> , 2012, 84, 1166-1176. | 2.5 | 10 |
| 35 | Sub-optimal protection against past hepatitis B virus infection where subtype mismatch exists between vaccine and circulating viral genotype in northern Australia. <i>Vaccine</i> , 2018, 36, 3533-3540. | 1.7 | 10 |
| 36 | Factors Associated with Elevated ALT in an International HIV/HBV Co-Infected Cohort on Long-Term HAART. <i>PLoS ONE</i> , 2011, 6, e26482. | 1.1 | 10 |

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|----|---|-----|-----------|
| 37 | Analysis of Hepatitis B Virus Haplotype Diversity Detects Striking Sequence Conservation Across Genotypes and Chronic Disease Phase. <i>Hepatology</i> , 2021, 73, 1652-1670. | 3.6 | 9 |
| 38 | HBV variants are common in the "immune-tolerant" phase of chronic hepatitis B. <i>Journal of Viral Hepatitis</i> , 2020, 27, 1061-1070. | 1.0 | 8 |
| 39 | Genotypic profiles of hepatitis B in African immigrants and their clinical relevance. <i>Journal of Medical Virology</i> , 2017, 89, 1000-1007. | 2.5 | 7 |
| 40 | Hepatitis B among immigrants from Myanmar: Genotypes and their clinical relevance. <i>Journal of Medical Virology</i> , 2018, 90, 271-276. | 2.5 | 7 |
| 41 | Epidemiology and phylogenetic analysis of hepatitis D virus infection in Australia. <i>Internal Medicine Journal</i> , 2018, 48, 1308-1317. | 0.5 | 7 |
| 42 | Secreted hepatitis B virus splice variants differ by HBV genotype and across phases of chronic hepatitis B infection. <i>Journal of Viral Hepatitis</i> , 2022, 29, 604-615. | 1.0 | 6 |
| 43 | Towards Genotype-Specific Care for Chronic Hepatitis B: The First 6 Years Follow Up From the CHARM Cohort Study. <i>Open Forum Infectious Diseases</i> , 2019, 6, ofz469. | 0.4 | 5 |
| 44 | Point of care and oral fluid hepatitis B testing in remote Indigenous communities of northern Australia. <i>Journal of Viral Hepatitis</i> , 2020, 27, 407-414. | 1.0 | 5 |
| 45 | Clinical stage drugs targeting inhibitor of apoptosis proteins purge episomal Hepatitis B viral genome in preclinical models. <i>Cell Death and Disease</i> , 2021, 12, 641. | 2.7 | 4 |
| 46 | A novel hepatitis B virus S gene insertion associated with reduced humoral immunity and diagnostic escape. <i>Internal Medicine Journal</i> , 2014, 44, 709-710. | 0.5 | 2 |
| 47 | Antiviral drug resistance and hepatitis B: a continuing public health problem. <i>Antiviral Therapy</i> , 2017, 22, 643-644. | 0.6 | 1 |
| 48 | Molecular Phylogenetics of Hepatitis D Virus in New Zealand and the Implications for Pacific Island Countries. <i>Intervirology</i> , 2021, 64, 102-107. | 1.2 | 0 |