

Julietta Trink's

List of Publications by Year in descending order

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Version: 2024-02-01

14
papers

113
citations

1477746

6
h-index

1372195

10
g-index

15
all docs

15
docs citations

15
times ranked

228
citing authors

#	ARTICLE	IF	CITATIONS
1	Influence of ethnicity on the distribution of genetic polymorphisms associated with risk of chronic liver disease in South American populations. BMC Genetics, 2015, 16, 93.	2.7	25
2	Role of HLA-DP and HLA-DQ on the clearance of hepatitis B virus and the risk of chronic infection in a multiethnic population. Liver International, 2017, 37, 1476-1487.	1.9	23
3	Prevalence and Factors Related to Natural Resistance-Associated Substitutions to Direct-Acting Antivirals in Patients with Genotype 1 Hepatitis C Virus Infection. Viruses, 2019, 11, 3.	1.5	15
4	Distribution of genetic polymorphisms associated with hepatitis C virus (HCV) antiviral response in a multiethnic and admixed population. Pharmacogenomics Journal, 2014, 14, 549-554.	0.9	11
5	Plasma and stool metabolomics to identify microbiota derived-biomarkers of metabolic dysfunction-associated fatty liver disease: effect of PNPLA3 genotype. Metabolomics, 2021, 17, 58.	1.4	10
6	Increased prevalence of human herpesvirus type 8 (HHV-8) genome among blood donors from North-Western Argentina. Journal of Medical Virology, 2017, 89, 518-527.	2.5	7
7	Pre-treatment prediction of response to peginterferon plus ribavirin in chronic hepatitis C genotype 3. World Journal of Hepatology, 2015, 7, 703.	0.8	5
8	Evolving Trends in the Hepatitis C Virus Molecular Epidemiology Studies: From the Viral Sequences to the Human Genome. Epidemiology Research International, 2012, 2012, 1-10.	0.2	4
9	Clinical utility of pharmacogenomics in the management of hepatitis C. Pharmacogenomics and Personalized Medicine, 2014, 7, 339.	0.4	4
10	Hepatitis C virus pharmacogenomics in Latin American populations: implications in the era of direct-acting antivirals. Pharmacogenomics and Personalized Medicine, 2017, Volume10, 79-91.	0.4	3
11	Genetic diversity and phylogeographic analysis of human herpesvirus type 8 (HHV-8) in two distant regions of Argentina: Association with the genetic ancestry of the population. Infection, Genetics and Evolution, 2020, 85, 104523.	1.0	3
12	The genetic variability of hepatitis B virus subgenotype F1b precore/core gene is related to the outcome of the acute infection. Virus Research, 2020, 277, 197840.	1.1	2
13	A simple and rapid approach for human herpesvirus type 8 subtype characterization using single base extension. Letters in Applied Microbiology, 2021, 73, 308-317.	1.0	1
14	P9: Distribution of genetic polymorphisms associated to hepatitis C virus (HCV) antiviral response in a multiethnic and admixed population. Journal of Viral Hepatitis, 2013, 20, 20-20.	1.0	0