## Umbertina Villano

## List of Publications by Year

 in descending orderSource: https:|/exaly.com/author-pdf/112949/publications.pdf
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Molecular Epidemiology of Hepatitis C Virus Cenotype 4 Isolates in Egypt and Analysis of the Variability
3 of Envelope Proteins E1 and E2 in Patients with Chronic Hepatitis. Journal of Clinical Microbiology,
2005, 43, 1902-1909.

Seroprevalence and anti-HEV persistence in the general population of the Republic of San Marino. ,

6 An integrated approach identifies IFN-regulated microRNAs and targeted mRNAs modulated by different HCV replicon clones. BMC Genomics, 2011, 12, 485.

8 Hepatitis E virus genotypes and subgenotypes causing acute hepatitis, Bulgaria, 2013â€"2015. PLoS ONE, 2018, 13, e0198045.Journal of Gastroenterology and Hepatology, 1997, 9, 1081-1084.

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11. Evolutionary dynamics of HBVâ€D1 genotype epidemic in Turkey. Journal of Medical Virology, 2014, 86,
    109-116.
11 109-116.
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$5.0 \quad 20$

20

12 microRNA levels in paraffin-embedded indolent B-cell non-Hodgkin lymphoma tissues from patients chronically infected with hepatitis B or C virus. BMC Infectious Diseases, 2014, 14, S6.
2.9

14
13 Hepatitis E Outbreak in the Central Part of Italy Sustained by Multiple HEV Genotype 3 Strains,
Juneâ€"December 2019. Viruses, 2021, 13, 1159.
3.3

14

Lack of WHV integration nearby N-myc2 and in the downstream b3n and win loci in a considerable
14 fraction of liver tumors with activated N-myc2 from naturally infected wild woodchucks. Virology,
2.4

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2006, 345, 258-269.

The win locus involved in activation of the distal N -myc2 gene upon WHV integration in woodchuck
15 liver tumors harbors S/MAR elements. Virology, 2004, 329, 1-10.
$2.4 \quad 12$

Molecular epidemiology and phylogenetic analysis of Hepatitis B virus in a group of migrants in Italy.
BMC Infectious Diseases, 2015, 15, 287.
2.9

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Naturally Occurring Surface Antigen Variants of Hepatitis B Virus in Tunisian Patients. Intervirology,
2016, 59, 36-47.
2.8

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19

Population-based survey of hepatitis A virus infection in the Republic of San Marino. European Journal of Epidemiology, 1997, 13, 687-689.
5.7

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A computational approach to identify point mutations associated with occult hepatitis B : significant mutations affect coding regions but not regulative elements of HBV. Virology Journal, 2011, 8, 394.
$3.4 \quad 9$

Human hepatitis E virus circulation in Bulgaria: Deep Bayesian phylogenetic analysis for viral spread
5.0
control in the country. Journal of Medical Virology, 2019, 91, 132-138.

In vivo transmission and dynamics of deleted genomes after experimental infection of woodchuck hepatitis B virus in adult animals. Virus Genes, 2002, 25, 147-157.
$1.6 \quad 7$

Prevalence and risk factors for hepatitis $E$ virus infection in blood donors: a nationwide survey in Italy, 2017 to 2019. Eurosurveillance, 2022, 27, .
7.0

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24 Evolutionary dynamics of HBVâ€Đ7 subgenotype in Tunisia. Journal of Medical Virology, 2017, 89, 469-475.
5.0

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25 An Evaluation of Hepatitis E Virus Molecular Typing Methods. Clinical Chemistry, 2021, 68, 181-191.

Hepatitis E virus infection prevalence among men who have sex with men involved in a hepatitis $A$ virus outbreak in Italy. Blood Transfusion, 2019, , 1-5.

