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D Ge, J Fellay, AJ Thompson, JS Simon, KV Shianna... - Nature, 2009 - nature.com

... The IFN- proteins, encoded by the IL28A/B and IL29 genes, were first described in 2003 (refs 13, 14). ... Population genetic structure of variable drug response. Nature Genet. ... RESEARCH. ITPA gene variants protect against anaemia in patients treated for chronic hepatitis C ...

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... Numerous investigations have documented the genetic variability of HBV in HBsAg-positive patients (reviewed in Carman et al. ... In vitro expression of the S gene led to the synthesis of HBsAg, which ... tent with this view, HBV genomes isolated either during an acute hepatitis B or 30 ...

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Genetic variation in IL28B and spontaneous clearance of hepatitis C virus

DL Thomas, CL Thio, MP Martin, Y Qi, D Ge... - Nature, 2009 - nature.com

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VE Buckwold, Z Xu, M Chen, TS Yen, J Ou - Journal of virology, 1996 - Am Soc Microbiol

... 1993. Viral genetic variation: hepatitis B virus as a clinical example ... Cloning and characterization of a human ribosomal protein gene with enhanced expression in fetal and neoplastic cells ... sequence of the hepatitis B viral genome and the identification of the major genes, p. 57 ...

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Genetic variation in IL28B is associated with chronic hepatitis C and treatment failure: a genome-wide association study

..., B Hirschel, R Malinverni, D Moradpour, B Müllhaupt... - Gastroenterology, 2010 - Elsevier

... like receptors, 20 chemokines, and interleukins as well as IFN-stimulated genes 21, 22 ... for the purpose of mapping the candidate causal variant or genetic region using ... recombination profile indicated that the signal encompassed the whole IL28B gene (Supplementary Figures 1 ...

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Genetic variation of hepatitis B virus and its significance for pathogenesis

Zhen-Hua Zhang, Chun-Chen Wu, Xin-Wen Chen, Xu Li, Jun Li, Meng-Ji Lu

Abstract

Hepatitis B virus (HBV) has a worldwide distribution and is endemic in many populations. Due to its unique life cycle which requires an error-prone reverse transcriptase for replication, it constantly evolves, resulting in tremendous genetic variation in the forms of genotypes, subgenotypes, and mutations. In the recent years, much effort has been invested into studies on the relationship between HBV genetic variation and HBV-related pathogenesis, which has profound implications in the natural history of HBV infection. viral detection.

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