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Retrospective Cohort Study

Hepatitis B virus persistent infection-related single nucleotide polymorphisms in HLA regions are associated with viral load in hepatoma families

Hsieh A-R et al. HBV viral load in HCC relatives

Abstract

BACKGROUND

Genome-wide association studies from Asia indicate that HLA-DP and HLA-DQ loci are important in persistent HBV infections. One of the key elements for HBV-related carcinogenesis is persistent viral replication and inflammation.

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Recently, a great number of genetic studies based on single nucleotide polymorphisms (SNPs) and genome-wide association studies have been performed to search for host determinants of the development of chronic HBV infection, clinical outcomes, therapeutic efficacy, and responses to hepatitis B vaccines, with a focus on human leukocyte antigens (HLA), cytokine genes, and toll-like ...

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Jul 10, 2019 · Those subjects who demonstrate direct and strong immune responses may die of cytokine storm in fulminant hepatitis, SARS, influenza, or other infections. Therefore, the persistent HBV infection-related single nucleotide polymorphisms (SNPs) on HLA-DP and -DQ loci could be an adaptive evolutionary response to the local environment.

Author: Dar-In Tai Publish Year: 2020