



Alcohol consumption leads to loss of healthy life, but the *ADH1B*2* allele may still protect from NASH

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Nonalcoholic fatty liver disease (NAFLD) is now the most common cause of chronic liver disease worldwide, affecting almost a quarter of the world population (1,2), and is a main cause of severe hepatic complications such as cirrhosis and liver cancer. In histological assessment NAFLD and alcoholic liver disease (ALD) are difficult to distinguish; NAFLD diagnosis thus requires a limit of alcohol intake. However, NAFLD and ALD very frequently coexist in the same patient, making it difficult to identify the exact cause of liver complications. Almost 5–6% of all worldwide deaths (~3 million) are caused by the harmful use of alcohol (<https://www.who.int/publications/i/item/9789241565639>), therefore alcohol represents a major health problem at a global scale. There is an association between the amount of alcohol consumption and ALD, however, only 30% of chronic drinkers develop alcoholic hepatitis, and 10–20% progress to advanced fibrosis or cirrhosis (3), underscoring the role of genetic factors involved in disease severity and progression. Any level of alcohol consumption, regardless of the amount, leads to loss of healthy life. The recommendation in clinical practice should be to avoid alcohol intake, particularly in the presence of any liver disease. Moderate alcohol consumption has been reported by many studies to be associated with less severe NAFLD, although some of these cross-sectional studies may be affected by selection bias (4–6). In contrast a longitudinal analysis of liver biopsies from patients with NAFLD showed that modest alcohol consumption was associated with lower

NASH resolution in comparison with nondrinkers (7). Recently, Chang *et al.* showed that the risk of liver steatosis in low and moderate alcohol consumers (MACs) was lower in comparison with nondrinkers (8); however, the proportion of hepatic steatosis plus fibrosis was higher in drinkers. The exact mechanism of the deleterious effect of alcohol consumption on NAFLD in obese patients is still unclear. When combined with free fatty acids the polyphenol resveratrol resulted in the stimulation of profibrogenic effects in hepatic stellate cells (key cells for induction and propagation of hepatic fibrosis) instead of any protective role (9). Therefore, it is important to shed light on the exact role of alcohol intake in addition to NAFLD to solve the currently rather conflicting data on this issue. The work published by Vilar-Gomez *et al.* in *Gastroenterology* (10) clarifies some knowledge gaps about alcohol metabolism and consumption, and the severity of NAFLD by studying the role of alcohol dehydrogenase (ADH)-1B, in particular *ADH1B*2*, in this context. The authors started from a finding generated in a previous work by Sookoian *et al.*, where they observed that patients with low alcohol intake and the *ADH1B*2* allele showed a less severe NAFLD by histology compared to other patients studied (11). Vilar-Gomez *et al.* studied 1,697 patients enrolled into various studies conducted by the Nonalcoholic Steatohepatitis Clinical Research Network (NASH CRN) over a period of 10 years [2009–2019]; the NASH CRN Pathology committee reviewed all liver

biopsies, and comprehensive alcohol consumption was obtained by AUDIT and LDH questionnaire, binge and heavy drinkers were excluded (170 patients). Remarkably, the frequency of *ADH1B*2* carriage varied across race being high in Asians/Pacific Islanders/Hawaiians (86%) and low in non-Hispanic whites (8%), Hispanics (14%), and Blacks (4%), but the study was focused on the 1,153 non-Hispanic whites, which were mainly female, obese, and hypertensive. Among the 1,153 patients, 30% had advanced fibrosis, and 60% had definite NASH. The cohort included 720 non-drinkers and 433 moderate drinkers, but no heavy alcohol consumers. *ADH1B*2* carriers were more likely to be male, and moderate alcohol consumption was similar between *ADH1B*1* and *ADH1B*2* carriers.

ADH and aldehyde dehydrogenase (ALDH) are the primary enzymes involved in hepatic alcohol metabolism and there are variants that encode enzymes with different activity, vary in ethnic distribution, influencing the level of alcohol oxidation to the less toxic acetaldehyde, the level of alcohol consumption, and the risk of alcoholism or alcohol dependence (12). For example, the presence of the *ADH1B*2* and *ADH1B*3* alleles would be associated with a higher oxidative capacity leading to faster ethanol oxidation. Therefore, the presence of *ADH1B*2* allele has shown a protective effect against alcohol dependence in Asian patients (13,14). Vilar-Gomez *et al.* showed that, in comparison with non-drinkers, the prevalence of definite NASH (defined as NAFLD activity score ≥ 4), and advanced fibrosis was lower in patients with MAC; moreover, they identified an inverse dose-dependent relationship between the amount of alcohol intake and risk of definite NASH up to a daily dose of 28 g alcohol. Although this was observed for both the *ADH1B** and the *ADH1B*2* allele, the NASH risk was significantly lower in patients carrying the *ADH1B*2* allele.

In summary, this study demonstrated a protective effect for patients carrying the *ADH1B*2* allele on the risk of severe histological damage of NAFLD, including fibrosis. Patients with MAC have lower risk of definite NASH in a dose-dependent manner. The *ADH1B* polymorphism *ADH1B*2* not only exerts a protective effect on NAFLD patients with MAC but also may reduce the calories gained by alcohol metabolism as patients had lower BMI in other studies (15). It seems that the beneficial effect of a highly metabolic ADH activity is diminished in patients with higher BMI with a limit of 37 kg/m² in the study of Vilar-Gomez *et al.* Further studies are needed to clarify the cause for this BMI-dependent effect on NASH risk of this allele.

The limitations of the study as disclosed by the authors included possible misclassification of alcohol consumption by self-assessment, the lower number of Asians/Pacific Islanders/Hawaiians individuals included in the trial, lack of information on socio-economic patient status, physical activity and other possible confounding factors. Despite these limitations the present work demonstrates an unequivocal protective effect against NASH and NASH fibrosis by the *ADH1B*2* allele, which is independent of MAC but offset by very high BMI in a Caucasian population. This study opens the door for further studies in specific settings to unravel the exact mechanisms of the protection conferred by the *ADH1B*2* allele.

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