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SPARC inhibition accelerates NAFLD-associated hepatocellular carcinoma development by dysregulating hepatic lipid metabolism

Agostina M Onorato¹, Esteban Fiore¹, Juan Bayo¹, Cecilia Casali²³, María Fernandez-Tomé²³, Marcelo Rodríguez¹, Luciana Domínguez¹, Josepmaría Argemi⁴, Florencia Hidalgo⁵, Cristian Favre⁵, Mariana García¹, Catalina Atorrasagasti¹, Guillermo D Mazzolini¹⁶

Affiliations PMID: 33641248 DOI: 10.1111/liv.14857

Abstract

Background and aims: Non-alcoholic fatty liver (NAFLD) and its more serious form non-alcoholic steatohepatitis increase risk of hepatocellular carcinoma (HCC). Lipid metabolic alterations and its role in HCC development remain unclear. SPARC (Secreted Protein, Acidic and Rich in Cysteine) is involved in lipid metabolism, NAFLD and diabetes, but the effects on hepatic lipid metabolism and HCC development is unknown. The aim of this study was to evaluate the role of SPARC in HCC development in the context of NAFLD.

Methods: Primary hepatocyte cultures from knockout (SPARC^{-/-}) or wild-type (SPARC^{+/+}) mice, and HepG2 cells were used to assess the effects of free fatty acids on lipid accumulation, expression of lipogenic genes and de novo triglyceride (TG) synthesis. A NAFLD-HCC model was stabilized on SPARC^{-/-} or SPARC^{+/+} mice. Correlations among SPARC, lipid metabolism-related gene expression patterns and clinical prognosis were studied using HCC gene expression dataset.

Results: SPARC^{-/-} mice increases hepatic lipid deposits over time. Hepatocytes from SPARC^{-/-} mice or inhibition of SPARC by an antisense adenovirus in HepG2 cells resulted in increased TG deposit, expression of lipid-related genes and nuclear translocation of SREBP1c. Human HCC database analysis revealed that SPARC negatively correlated with genes involved in lipid metabolism, and with poor survival. In NAFLD-HCC murine model, the absence of SPARC accelerates HCC development. RNA-seq study revealed that pathways related to lipid metabolism, cellular detoxification and proliferation were upregulated in SPARC^{-/-} tumour-bearing mice.

Conclusions: The absence of SPARC is associated with an altered hepatic lipid metabolism, and an accelerated NAFLD-related HCC development.

Keywords: hepatocellular carcinoma; lipid deposits; metabolism; non-alcoholic fatty liver disease; secreted protein acidic and rich in cysteine; sterol regulatory element-binding protein1c.

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