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► **To cite this version:**

Jérôme Eeckhoutte. Non-Alcoholic Steatohepatitis (NASH) and Liver Fibrosis: Molecular and Multicellular Control of Evolving Diseased States. *Cells*, 2022, 11 (16), pp.2551. 10.3390/cells11162551 . hal-03752605

**HAL Id: hal-03752605**

**<https://hal.science/hal-03752605v1>**

Submitted on 17 Aug 2022

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Editorial

# Non-Alcoholic Steatohepatitis (NASH) and Liver Fibrosis: Molecular and Multicellular Control of Evolving Diseased States

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Non-alcoholic fatty liver disease (NAFLD), the most common chronic liver disease, has emerged as a major threat to public health [1]. Indeed, obesity and type 2 diabetes are independent risk factors of NAFLD for which increases contribute to the devastating consequences of the metabolic disease pandemic [2]. Importantly, NAFLD presents with a range of conditions initiated by relatively benign hepatic steatosis, which can evolve to non-alcoholic steatohepatitis (NASH); fibrosis; and ultimately, cirrhosis [3]. Advanced stages promote hepatocellular carcinoma and/or liver failure [1]. In this context, NASH has been the fastest growing indication for liver transplantation in Western countries and is expected to become the leading cause of liver transplant in the United States [4,5].

NASH is characterized by an escalation in the disease complexity, where suffering hepatocytes trigger inflammation and, secondarily, liver fibrosis [1–3]. These processes are increasingly recognized as being driven by dynamic and coordinated changes in the complexity and activities of the various liver cell populations [6]. Indeed, in addition to hepatocytes and immune cells, resident or recruited, the role exerted by cholangiocytes, (myo)fibroblasts, and endothelial cells is also now being analyzed in detail (e.g., [7–9]). In this context, our understanding of the molecular and multicellular mechanisms underlying the various NAFLD stages is still limited and defining how/why the disease evolves to more advanced stages has remained challenging. For instance, the kinetic, spatial organization, and functional hierarchy of changes in cellular states and activities largely remain to be described together with the identity and integration of intercellular signals involved. At the individual cell-type level, molecular (mal)adaptations accompanying the disease progression, including transcriptomic dysregulation, also need to be further characterized. Indeed, diagnosis and therapeutic treatment of NASH and fibrosis remain controversial and of limited efficacy [10–12] and would greatly benefit from a greater understanding of the molecular mechanisms at play in the multicellular processes defining advanced NAFLD stages. The purpose of this Special Issue is to highlight recent findings in those areas, enlightening how NASH and its associated liver fibrosis develop through coordinated modulation of the activities of different liver cell-types.

**Funding:** Work at Inserm U1011 is supported by the Agence Nationale de la Recherche (ANR) grants “HSCreg” (ANR-21-CE14-0032), “PreciNASH” (ANR-16-RHUS-0006-PreciNASH), and “European Genomic Institute for Diabetes” E.G.I.D (ANR-10-LABX-0046), a French State fund managed by ANR under the frame program Investissements d’Avenir I-SITE ULNE/ANR-16-IDEX-0004 ULNE and by Fondation Recherche Médicale (FRM grants EQU202203014645 and EQU202203014650).

**Conflicts of Interest:** The author declares no conflict of interest.



**Citation:** Eeckhoutte, J. Non-Alcoholic Steatohepatitis (NASH) and Liver Fibrosis: Molecular and Multicellular Control of Evolving Diseased States. *Cells* **2022**, *11*, 2551. <https://doi.org/10.3390/cells11162551>

Received: 10 August 2022

Accepted: 15 August 2022

Published: 17 August 2022

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