Pathogenesis of HCV and Liver Fibrosis

Abstract

Despite the oppressive research, hepatitis C virus (HCV) pathogenesis is still evolving. A new prospective assembling pathogenic theories of HCV and its related liver fibrosis, would be an additional advocate. Recurrent HCV-induced inflammation, hepatic stellate cells activation, along with hepatic progenitor cells proliferation and apoptosis; all together might be portraying the most probable HCV pathogenic pathways.

Mini Review

Chronic infection with hepatitis C virus (HCV) affected more than 170 million people worldwide with up to 90% of these will progress to chronic liver disease [1]. In the era of direct antiviral drugs, the interest in HCV pathophysiological scripts is still roaming in every hepatologist’s mind. Efforts are exerted to coalesce the scattered parts of this unresolved story.

The interplay between inflammation, apoptosis, necrosis, and fibrosis is the fundamental knob. However, which of which the on button is the dilemma to be concealed. Cytokines are key mediators, playing an important role in the defense against viral infections, determining the pattern of host immune response and inhibiting viral replication [2]. Th1 cytokines such as IL-2 and interferon gamma are required for host anti viral immune responses, while Th2 cytokines (IL-4, IL-10) can inhibit the development of these effect or mechanisms [3]. The positive connection between interferon gamma, interleukin 2 producing cells with the biochemical and necroinflammatory and fibrotic activities of chronic HCV infection had been documented (Figure 1) [4]. Proinflammatory cytokines (promoting inflammation) such as tumor necrosis factor alpha (TNF-a) and interferon gamma (Inf G) were reported to be overexpressed in chronic HCV directly correlated to the necroinflammatory state and state of fibrosis. While anti-inflammatory cytokines such as IL-28 demonstrated an association with low HAI Ishak score of necro inflammation and low AST level (p=0.002) (Figure 2 & 3) [5,6].

Unveiling the other part of the story: HCV related Liver fibrosis is the next step. It is often initiated by core protein activation of the major sensing cells; hepatic stellate cells (HSC). On activation a series of events take place; starting from secreting extracellular matrix compounds alpha-smooth muscle actin expression (alpha SMA) matrix metalproteases (MMP), secreting several growth factors, and ending by enhanced neoangiogenesis which is a common pathological feature of chronic HCV infection [7]. The intense expression of the regenerative marker hepatocyte growth factor activator (HGFA) in liver tissues of chronic HCV patients which was documented to be in line with the expression of transforming growth factor beta1 (TGFβ1) and alpha smooth muscle actin (α-SMA), known as fibrogenic markers. A parallel expression of HGFA with the potent immune cytokine TGFβ1 with the grade of activity in CHC patients make them as activity markers. The coparallel expression of HGFA, TGFβ1 and α-SMA with advanced stage of fibrosis and cirrhosis in HCV related chronic hepatitis might illustrate a pleiotropic complex relationship between regeneration and fibrosis [8].

Figure 1: TNF-a expression was in favor of higher grade of portal tract inflammation compared to the group lacking TNF-a expression.

In chronic HCV, activation of hepatic progenitor cells (HPCs) as a potential reservoir of cholangiocyte/hepatocyte differentiation [9], has been coupled to the severity of inflammation, stage of fibrosis along with the concomitant clinical status of liver disease (Figure 4) [10].
Apoptosis as an important form of host immune response against viral infections, its role as a pathogenic integrator in chronic HCV infection is validated in most studies [11]. Eminently enhanced hepatocyte apoptosis, was found to be interconnected to the inflammation severity, stages of fibrosis, transaminases levels and viral load in CHC patients [12]. Such studies had demonstrated the Fas antigen and the Fas ligand bonding on T cells [13]. However the antiapoptotic marker bcl2 protein as an antiapoptotic mediators participating substantially in the apoptotic process; have not been systematically and thoroughly ascribed in liver tissue of chronic CHC (Figure 5) [14]. A complementary harmonized converse connection between activation of hepatic progenitor cells and antiapoptotic mediators is suggested for an evolving research in the horizon with rewarding results. It might be potentially incriminated in chronic HCV pathogenesis. Suppositions might open new horizons in HCV understanding. New roles for antiapoptotic factor along with HPC antagonists might be advocated, paving the way for more future studies.

References


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