Dear editors,

Thank you for your reply and for the reviewers' comments concerning our manuscript entitled "Multiple inflammatory mediators crosstalk networks in the plasticity of liver fibrosis" that we submitted to *World Journal of Gastroenterology*. Those comments on our study are insightful and we feel they have led to significant improvement of our paper.

We have revised our manuscript according to the reviewers' suggestions (Shown in Blue color within manuscript). The following is a point-by-point response to the reviewers' comments and questions.

Reviewers' comments:

Reviewer #1 (Remarks to the Author):

Well written manuscript dealing with important issue in the field of hepatology but some comments to be mentioned - As regard autophagy , abreife description of the role of autophagy in liver fibrosis must be mentioned -As regard MRNA 200b was missed in liver fibrosis (see Besheer et al 2019 Diffusion-weighted magnetic resonance imaging and micro-RNA in the diagnosis of hepatic fibrosis in chronic hepatitis C virus. World J Gastroenterol 2019;25(11): 1289-1431 -Most of the reference in the study focusing on HBVe.g TLRs although there are many studies on HCV. Fakhir et al. Genetic variations in toll-like receptors 7 and 8 modulate natural hepatitis C outcomes and liver disease progression. Liver Int. 2018;38(3):432–442. El-Bendary et al (2018) The association of single nucleotide polymorphisms of Toll-like receptor 3, Toll-like receptor 7 and Toll-like receptor 8 genes with the susceptibility to HCV infection, British Journal of Biomedical Science, 75:4, 175-181, -few grammatical errors Page6 line 114 ;Collagen III Page8 line 174: neutrophils instead of neutrophills.

[Response] We are grateful for your comments and we have revised our manuscript according to your suggestions. Additionally, we substantially revised references and corrected grammar errors.

Reviewer #2 (Remarks to the Author):

This review article describes about multiple inflammatory mediator cross-talking networks. The epigenetic crosstalk between histone acetylation and miRNAs may be described more in detail in terms of the inhibition of HSC activation.

[Response] We are grateful for your comments and we have revised our manuscript according to your suggestions.