

Stimulation of EPOR/CD131 in ethanol-exposed rats results in changed expression of neural regeneration-, autophagy, apoptosis- and neuroinflammation-related genes

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Due to the intricacy of ethanol pharmacodynamics, the neurobiological basis of ethanol-induced neurodegeneration is still poorly understood. Nonetheless, long-term effects of known pathogenic factors that cause neuronal damage, such as oxidative stress, neuroinflammation and glutamate excitotoxicity, result in the activation of proapoptotic cascades and a decrease in neurotrophin synthesis. Some of the defense mechanisms targeted at responding to the neurotoxic effects of ethanol include autophagy activation and apoptosis inhibition; thus, pharmacological regulation of apoptosis and autophagy may have potential therapeutic benefits in alcohol-induced neurodegeneration. Erythropoietin, an endogenous regulator of erythropoiesis (EPO) possesses both the properties of an apoptosis inhibitor and an autophagy activator [Zubareva, 2016]. These properties are mediated by the EPOR/CD131 receptor agonist, which has been linked to tissue protection, anti-inflammatory, and antioxidant effects, even in nerve tissue damage [Zhong, 2020].

Aim: To analyze the expression patterns of genes associated with autophagy, apoptosis, neuroregeneration and neuroinflammation in ethanol-induced neurodegeneration.

Materials and methods: The study included 48 male wistar rats, (age 20 weeks, weight; 270-300g) grouped into 4 according to ethanol sensitivity. 1) Ethanol; 2) Ethanol + PHBSP; 3) Ethanol + EPO; and 4) Control.

To simulate ethanol-induced neurodegeneration, Rats were given 5%, 10%, 20% ethanol in this specific order in the span of 20 weeks. PHBSP and recombinant erythropoietin were administered at doses of 5 μ g/kg and 10 μ g/kg subcutaneously twice a week. Ethanol and control groups received apyrogenic water in an equivalent volume. Quantitative PCR was carried out to analyze the expression of autophagy, neuroinflammation and apoptosis genes in the rats.

Results: The qPCR analysis showed increase in the expression of *Atg7*, *Becn1* and *LC31* genes, associated with autophagy, which is consistent with results from other studies [Li, 2018]. Furthermore, this study revealed that EPO and PHBSP significantly attenuated the expression of *Il1b*, *Tnf* and *Ccl2* proinflammatory genes, reduced proapoptotic gene expression and restored the impaired expression of the *Bdnf* and *Ngf* neurotrophic factor genes.

Conclusion: Results confirm the significant roles of therapeutic regulation of apoptosis and autophagy in ethanol-induced neurodegeneration.

References

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Illustrations

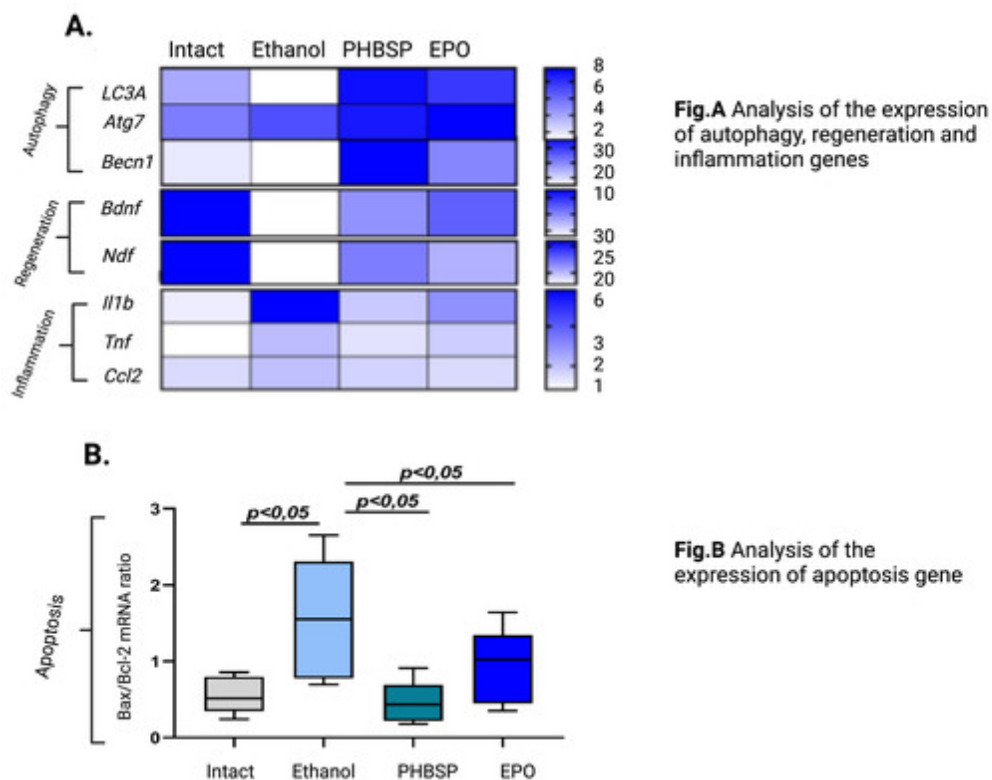


Рис. 1. Analysis of the expression of autophagy, regeneration and inflammation genes