

DEREGULATION OF AUTOPHAGY FLUX AND GENE EXPRESSION INDUCED BY TOBACCO SMOKE AMONG IRAQI SMOKERS

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(Received 4 April 2021, Revised 5 June 2021, Accepted 12 June 2021)

ABSTRACT : Cigarette smoking is a lifestyle behavior that causes significant adverse health effects. Cigarette smoke contains chemicals, many of which are lead to the production of reactive oxygen species (ROS), which can lead to apoptosis and autophagy. To estimate the association of Cigarette smoking with the autophagy and immunity, technology of real time polymerase chain reaction (RTPCR) for gene expression of (LC3A, LC3B, LC3C, myd88) was used. Enzyme-linked immunosorbent assay (ELISA) technique was utilized to measurement the amount of TNF- α protein. The ratios of LC3A/LC3B and LC3B/LC3C were calculated to estimate the autophagy flux. The results indicate the expression of LC3B, LC3C and Myd88 genes in smokers is increased significantly ($p \leq 0.001$) compared with non-smokers. While the expression of LC3A gene is decreased significantly. Findings show that the serum mean concentration of (TNF- α) for smokers group is increased significantly ($p \leq 0.001$). The ratio of LC3A/LC3B is increased highly significant ($p \leq 0.001$) in smokers individuals. While there is no significant differences in the ratio of LC3B/LC3C between smokers and non-smokers.

Key words : Smokers, free radical production, autophagy flux, LC3 gene, gene expression.

How to cite : Miyada Kh Hassan, Adel M. Rabee and Haider F. Ghazi (2022) Deregulation of autophagy flux and gene expression induced by tobacco smoke among Iraqi smokers. *Biochem. Cell. Arch.* **22**, 2415-2422. DocID: https://connectjournals.com/03896.2022.22.2415

INTRODUCTION

Autophagy is a process that removes damaged organelles and proteins by lysosomal degradation which greatly contributes to cell homeostasis and the prevention of various diseases. Autophagic flux is a process that used to measure autophagic degradation activity (Xiaojuan *et al*, 2018).

Autophagy is not only an adaptive response to the stress, but it is also involved in a many of other physiological mechanisms, such as intracellular quality control, inhibition of cellular ageing, differentiation and development of the cells, apoptosis, and both innate and adaptive immunity (Yin *et al*, 2016). Therefore, weaken or deficiency in this pathway result in severe pathologies, like specific types of cancer, neurodegeneration, and muscular dystrophies (Jiang and Mizushima, 2014). As a result, it is very important to quantify autophagic process in specific, accurate, and reliable way to be able to assess the changes of autophagic flux process and eventually the blocked step when this pathway is defect (Orhon and Reggiori, 2017).

One of the main steps in the autophagy regulation is the binding of microtubule-associated protein 1 light chain 3 (LC3) with phosphatidylethanolamine and forming microtubule-associated protein 1 light chain 3 lipidated form (LC3-II). The Lipidated LC3 bind to the elongated phagophore after that the phagosome fuse with lysosomes. Then, LC3-II can be either delipidated and recycled or hydrolyze by lysosomal enzymes. This pathway is called LC3-associated phagocytosis (LAP). The gold standards for measuring autophagy activity are monitoring LC3-II conversion, LC3 distribution or its flux through the pathway of autophagy. Therefore, measuring the formation of cellular autophagosome puncta that contain LC3, and quantifying the ratio of LC3-II to LC3-I enable monitor autophagy flux activity (Kono *et al*, 2021).

Asare *et al* (2020) found that the materials contained in cigarette smoke weaken LC3-associated phagocytosis (LAP) pathway, leading to chronic inflammatory response which is initiate the lung injury and dysfunction of airway that is associated with chronic obstructive pulmonary disease COPD. While, many Previous studies have shown

expression of myd88 gene and the level of TNF- α . In particular, cigarette smoke acts as a double-edged sword that either exacerbates pathological immune responses or attenuates the normal defensive function of the immune system, possibly owing to the complexities and functional diversities of cigarette smoke components and individuals' medical condition. Nevertheless, smoking plays a harmful rather than beneficial role in either case. The results illustrated that changes in gene expression may have an important role in susceptibility of smokers to various diseases.

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