

## Chapter 2

# ACTIVATION OF INFLAMMATORY SIGNALING PATHWAYS IN METABOLIC SYNDROME: CHANGES IN ADIPOKINES AND CYTOKINES

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### INTRODUCTION

Metabolic syndrome (MetS) is constellation of metabolic abnormalities such as glucose intolerance, dyslipidemia, central obesity, hypertension, insulin resistance, hyperinsulinemia and microalbuminuria <sup>(1)</sup>. All these risk factors increase the risk of cardiovascular disease development <sup>(2)</sup>. Subjects with MetS have five fold greater risk of developing Type 2 Diabetes Mellitus (T2DM) and twice fold greater risk of developing cardiovascular diseases <sup>(3)</sup>. In addition; individuals diagnosed with MetS, have higher risk of polycystic ovary syndrome, non alcoholic fatty liver disease, gallstone, sleep disorders, asthma and some type of cancers such as breast, colorectal, pancreas cancers <sup>(4,5)</sup> and mortality from all causes <sup>(6)</sup>. Although the mechanism underlying MetS development is not fully understood, overweight/obesity, dietary patterns, genetic disposition physical inactivity <sup>(7)</sup>, race/ethnicity, hormone imbalances, inflammation and chronic low-grade

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inflammatory cytokine and ferritin levels at reference intervals is seen as an important step in protecting individuals against MetS development. Activation of inflammatory signaling pathways in metabolic syndrome causes changes in pro-inflammatory and anti-inflammatory cytokine levels, causing systemic inflammation and tissue damage. It is thought that studies on the relationship between inflammation and MetS and its components may help to develop new strategies to prevent and treat the MetS and related disorders.

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