ORIGINAL ARTICLE Iran J Allergy Asthma Immunol February 2020; 19(1):1-8. Doi: 10.18502/ijaai.v19i1.2409

Analysis of Methylation and Expression Profile of *Foxp3* Gene in Patients with Behçet's Syndrome

Jafar Farhadi^{1,2}, Mohammad Nouri³, Alireza Khabbazi², Nasser Samadi⁴, Zohreh Babaloo⁵, Mahdi Azad⁶, Somayeh Abolhasani⁷, Shahriar Alipour⁷, Golamreza Jadideslam^{1,2}, Sam Seydi Shirvani^{1,2}, and Ebrahim Sakhinia⁸

¹ Molecular Medicine, Faculty of Advanced Medical Sciences, Tabriz University of Medical Sciences, Tabriz, Iran ² Connective Tissue Diseases Research Center, Tabriz University of Medical Sciences, Tabriz, Iran

³ Department of Biochemistry, Faculty of Medicine, Tabriz University of Medical Sciences, Tabriz, Iran ⁴ Cancer Biochemistry, Cancer Biotechnology, Faculty of Medicine, Tabriz University of

Medical Sciences, Tabriz, Iran

⁵ Department of Immunology, Faculty of Medicine, Tabriz University of Medical Sciences, Tabriz, Iran ⁶ Department of Medical Laboratory Sciences, Faculty of Allied Medicine, Qazvin University of Medical Sciences, Qazvin, Iran

⁷ Department of Biochemistry, Faculty of Medicine, Urmia University of Medical Sciences, Urmia, Iran ⁸ Department of Genetic, Faculty of Medicine, Tabriz University of Medical Sciences, Tabriz, Iran

Received: 26 December 2018; Received in revised form: 12 February 2019; Accepted: 24 February 2019

ABSTRACT

Forkhead box P3 (Faxp3) gene is an important means in the Treg cells function, in both maintenances of immune tolerance and regulation of response. Epigenetic modifications of the faxp3 gene at its regulatory regions control the chromatin accessibility for the transcription factors and other transcriptional regulators in order to control Foxp3 expression. In addition, the methylation status of CpG islands within the Foxp3 promoter and regulatory elements regulate the expression of Foxp3. This study was performed to assess the role of the faxp3 gene in patients with Behçet's syndrome (BS).

Venous blood samples were collected from all participants and peripheral blood mononuclear cells (PBMC) were extracted through Ficoll-Hypaque method. Genomic DNA was randomly sheared by sonication and immunoprecipitated with a monoclonal antibody. The status methylation of the foxp3 gene was estimated in 108 blood samples of active BS patients and healthy individuals (controls); using methylation DNA immunoprecipitation (MeDIP) technique. Expression analysis was carried out; using Real-time PCR.

The expression of faxp3 gene in the patients' group (mean±SD: 1.79±1.12) was significantly lower than the healthy group (mean±SD: 2.73±1.33) (p<001). Also, the methylation levels of Foxp3 promoter showed that its level in patients (mean±SD: 2.3±1.16) was higher than the healthy group (mean±SD: 1.85±0.59). However, this increase was not statistically significant (p>0.05). Also, these results indicated that increasing the amount of methylation of the faxp3 gene by reducing its expression leads to an increase and intensifying of the disease.

The decrease in Foxp3 expression is possibly associated with hypermethylation of the gene, and it can be considered as a risk factor for BS. Future studies may be needed to identify the capability of specific DNA methylation alterations in this syndrome.

Keywords: Behçet's syndrome; DNA methylation; Foxp3; DIP

Corresponding Author: Ebrahim Sakhinia, PhD: Department of Genetics, Faculty of Medicine, Tabriz University of Medical Sciences, Tabriz, Iran. Tel: (+98 935) 602 2153, E-mail: esakhinia@yahoo.co.uk

Copyright© February 2020, Iran J Allergy Asthma Immunol. All rights reserved.

Published by Tehran University of Medical Sciences (http://ijaai.tums.ac.ir)