

# The Association of The High -Mobility Group Box 1 Gene and Its Product with Rheumatoid Arthritis in Basra Province - Iraq

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*Abstract*— Background: Rheumatoid arthritis (RA) is an autoimmune disease. High mobility group box 1 (HMGB1) is a nuclear protein which considered to be a biomarker in several active systemic autoimmune diseases such as RA. The study aimed to investigate HMGB1 gene and its product in RA patients in Basra.

Methods: The study included 26 patients and 10 healthy controls aged 21-53 yrs.*HMGB1* gene expression was detected using qRT-PCR and the serum levels of HMGB1 and Rheumatoid factors (RF)including (IgG and IgM) of study subjects were estimated by ELISA.

Results: The results showed that the mRNA expression levels of *HMGB1* was significantly higher in RA patients in compare with healthy control( $4.14\pm 3.89$  and 1, respectively). In contrast, the results showed no significant differences (p=0.72) between the mean of HMGB1 serum level in RA patients ( $3.55\pm 2.90$  ng/ml) and healthy controls ( $3.79\pm 1.30$ ng/ml) and*HMGB1* expression showed weak positive correlation to its serum levels. The level of IgG and IgM were significantly higher in RA patients in compare with healthy controls with a strong positive correlation. Moreover, there was weakly positive correlation between serum HMGB1 level and the RF-IgG(0.11) and no correlation between serum HMGB1 levels and RF-IgM(0.009) in RA patients.

Conclusion: Rheumatoid arthritis patients have elevated levels of *HMGB1* gene expression but their protein levels were normal due to the course of treatment. Furthermore, the high *HMGB1* gene expression level correlates positively with IgG levels.

Keywords— Rheumatoid arthritis, HMGB1, gene expression, qRT-PCR

## I. INTRODUCTION

Autoimmune diseases(AIDs) occur when the immune system attacks the body causing damage in different tissues and organs. AIDs includes many diseases but the most common one are rheumatic diseases like rheumatoid arthritis  $(RA)^1$ . Rheumatoid arthritis can be defined as a chronic inflammatory disease characterize by swelling and tenderness of the joints in addition to the destruction of cartilages and bones. The cause of this

destruction is the production of autoantibodies including rheumatoid factor (RF) (IgG and IgM) and anti-citrullinated autoantibody (anti-CCP)<sup>2</sup>.

The high -mobility group box 1 (HMGB1) was first identified by Goodwin and Colleagues in 1973 in Calf thymus<sup>3</sup>. It belongs to the high –mobility group (HMG) protein family which gain its name from its high electrophoretic mobility during electrophoresis in polyacrylamide gels <sup>4</sup>. It actsas a cytokine and it is released into the extracellular milieu. This protein is released passively in case of cellular necrosis and nearly all the nucleated cells can release it and by doing so they alert other cells to the damage happening <sup>5</sup>. HMGB1 has different functions including inflammation trigger, attracting cells, stem cell recruiting and activating dendritic cells as it sustains the proliferation of antigen specific T cell and help them to develop to Th1<sup>4</sup>.In acute and chronic inflammatory disease HMGB1 binds to soluble molecules and receptors such as RAGE, TLR2 and TLR4 leading to the activation of the cell carrying these receptors $^{6}$ .

The aim of our study is to evaluate the *HMGB1* gene expression and the level of HMGB1 protein in serum of RA patients in Basra and investigate its correlations between each other and with RF.

#### II. MATERIALS AND METHODS

## A. Study subjects

Adult Rheumatoid arthritis patients(n=26) from both sexes, age 21-53 yrs. are eligible for our study. All patients were visitors of Rheumatoid arthritis center located in AL-Basra Teaching hospital. The collection of samples starts in February till August of 2018. In addition, samples (n=10) were also collected from age matched healthy control (HC).

## B. Samples collection and processing

Five ml of venous blood were withdrawal from each studysubjects,1 ml were placed in EDTA tube for further molecular analysis while the remaining 4 ml were placed in a serum separation gel tube. Serum samples were separated by centrifugation at 3000 rpm for 20 min. then 3 aliquots for each sample were made and kept at -20  $^{\circ}$ C until further analysis.

## C. RNA Extraction

RNA was extracted from the study subjects blood samples using GENE ZOL<sup>™</sup> Tri RNA pure kit(Korea). The quality and concentration of the extracted RNA samples were measured by Nanodrop spectrophotometer (Quawell, USA).

#### D. Revers transcription of RNA

cDNA was synthesized from RNA samples, 100 ng of total RNA from each sample was used to synthesized cDNA using Bioneer two step Accupower® RT-PCR pre Mix (Korea) and according to the manufacturer protocols. **Note:** different volumes of total RNA were used depending on the concertation of the extracted RNA (high concentrations required small volumes or diluted volumes while low concentrations required RNA concentrating method using vacuumed centrifugation)

#### E. Measuring the expression of HMGB1

The *HMGB1* expression was measured using Quantitative Real time- polymerase chain reaction (qRT-PCR),20 ng of cDNA from each samples were used to

detect the expression of *HMGB1* and *GAPDH* (reference gene). The forward and the reverse primers used to show in Table 1. TaqMan was used to detect the gene expression and BYRT green was also used for the samples failed to be detected by TaqMan. The thermal cycler program used is shown in Table 2.The relative expression of the real time-PCR data was calculated according to the  $\Delta\Delta$ Ct method <sup>7</sup>.

## F. Serological study

The sera of all study subjects were tested for the level of Human High -mobility group box 1(HMGB1) protein ELISA Kit (Shanghai yehua Biological Technology Co., China).RF-IgG and IgM levels using Immunoenzymetric Assay by Rheumatoid Factor IgG and IgM kits (Demeditec /German). All kits were performed according to the manufacturer protocols.

#### G. Statistical analysis

The data underwent statistical analysis using a normality test at beginning followed by parametric (two sample T-test and confidence interval) and non-parametric test (Mann-Whitney U test)in MINTAB program. In addition, the Pearson correlation used to investigate the correlation between the study factors.

#### Table 1 Primers sequences of HMGB1 and GAPDH genes

Gene	Primers sequences (5'-3')	qRT-PCR chemistry	References
HMGB1	F: GCTGTGCAAAGGTTGAGAGC	TaqMan	http://bioinfo.ut.ee/primer30.4.0
	R: CTCGGGTACACAGGACACAC		
	Probe:HEXGATCGTCCCATCACAGTGTTG-BHQ-1		
GAPDH	F: CAACGAATTTGGCTACAGCA R: AGGGGTCTACATGGCAACTG	TaqMan	http://bioinfo.ut.ee/primer30.4.0
	Probe: FAM-CCACCAGCCCAGCAAGAGC-BHO-		
	Prope: FAM-CCACCAOCCACAAOAOC-BHQ- 1		
HMGB1	F: TAACTGAATAGGGGCGTGGTCT	BYRT green	8
	R: GAAAATGTGCT GGCTGTAGTGG		
GAPDH	F: CACCCACTCCTCCACCTTTG	BYRT green	9
	R: CCACCACCCTGTTGCTGTAG		

Table 2q RT- PCR Program for amplification	HMGB1 and GAPDH
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Step	Cycle	Temperature	Time
GoTaq® Hot Start Polymerase activation	1	95°C	5 min.
Denaturation	45	95°C	15 Sec.
Annealing and extension		60°C	1 min.

#### III. RESULTS

## A. Expression of HMGB1 in RA

qRT- PCR were used to measure *HMGB1* gene expression for both RA patients and HC using *GAPDH* as a reference gene. The current results showed that the mRNA expression level of *HMGB1* was significantly

increased in RA patients in comparison with healthy controls  $(4.14\pm 3.89 \text{ and } 1$ , respectively) as shown in Figure 1. For the BYRT green chemistry, the dye binding was specific to our target gene represented by one peak in the melting curve data as shown in Figure 2.

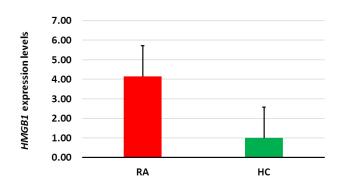


Figure 1 Relative Expression of HMGB1 Gene in RA patients and HC.

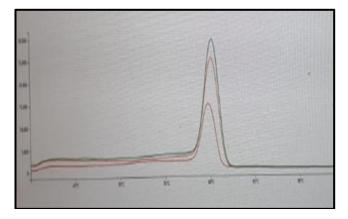


Figure 2 Melting curve for HMGB1 gene for three different samples showing the one Peak curve using BYRT green chemistry in quantitative RT-PCR

## B. Serum Human High mobility group B1 protein (HMGB-1) levels in RA

The results showed no significant differences (p = 0.36) between the mean of serum Human HMGB1 levels in RA patients ( $3.55 \pm 2.90$  ng/ml) and healthy controls ( $3.79\pm 1.30$  ng/ml) (Figure 3).

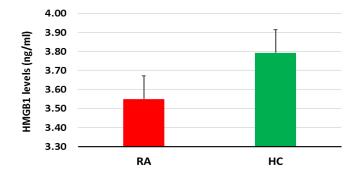


Figure 3 HMGB1 levels in RA patients and HC. No significant differences between RA and HC groups. The HMGB1 levels were measured by ELISA

## C. Correlation between HMGB1 expression and serum HMGB1 levels

The results of *HMGB1* expression showed weak positive correlation to its serum levels as the gene expression results showed high levels of expression in compare with the serum levels of HMGB1 measured by ELISA. The correlation coefficient was 0.14 and it was non-significant under p < 0.05 as shown in Figure 4.

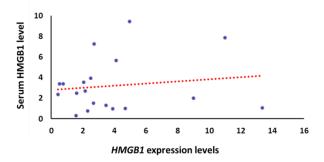
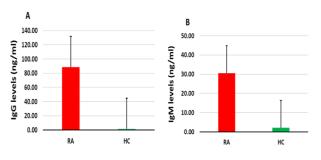


Figure 4 The correlation of HMGB1 expression with its serum levels The correlation coefficient was 0.14 and non-significant under p < 0.05.

## D. Rheumatoid factor (IgG level and IgM level) in RA

The level of IgG was significantly higher in RA patients in compare with healthy controls (p=0.028), the mean of IgG level was  $88.33 \pm 171.84$  ng/ml while the mean level of IgG in healthy controls was  $1.55 \pm 4.37$  ng/ml (Figure 5 A & B). Regarding the positive and negative cases of RA patients in terms of IgG levels, the results showed that 31.81 % of RA patients were positive for RF IgG leaving 68.18% of RA patients as negative cases, for healthy controls 100% were negative. The current study showed that level of IgM in RA patients in compare to healthy controls (p=0.10) and the means were  $30.55 \pm 77.24$  ng/ml and 2.12 ±3.32 ng/ml, respectively. Regarding the positive and negative cases of RA patients in terms of IgM levels. The results showed that 27.27% of RA patients were positive for IgM leaving 72.72% of RA patients as negative cases. For healthy controls 100% were IgM negative.



**Figure 5** RF (IgG, IgM) levels in RA patients and HC. **A.** IgG levels in RA patients and HC. **B.** IgM levels in RA patients and HC. The RF levels were measured by ELISA

From plotting the serum IgG values with the serum IgM levels in RA patients, the results showed strong positive correlation between the two factors. The R value was 0.65 (Figure 6)

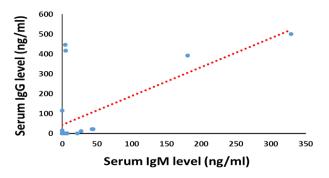
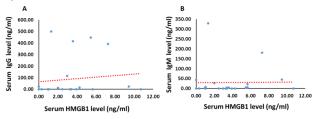


Figure 6 The correlation between IgG and IgM in RA. The R value was 0.65

## E. The correlation of serum levels of HMGB1 with RF in RA

The correlation between the serum levels of HMGB1 and RF (IgG and IgM) can be shown as follow:

From plotting the HMGB1level values with the IgG levels in RA patients, the results showed a weak positive correlation between the two factors. The R value was 0.11(Figure 7 A).while plotting the HMGB1 level values with the IgM levels in RA patients showed no correlation between the two factors. The R value was 0.01 (Figure 7B).



**Figure 7** The correlation between HMGB1 and other study parameters. **A.** The correlation between HMGB1 and RF-IgG levels in RA. The results showed a weak positive correlation. **B**. The correlation between HMGB1 and serum RF-IgM levels in RA. The results showed no correlation.

#### IV. DISCUSSION

Normally RA patients have high levels of HMGB1 in serum and in joints specially in the pannus regions. HMGB1 have functions involved in promoting proteolytic enzymes activity and the maturation of osteoclasts<sup>10, 11</sup>. Our results of HMGB1 expression in RA showed a significant elevation in the expression of the gene in compare to healthy controls as reported previously in many studies<sup>12, 13</sup>. Technically the gene expression data was reliable as the BYRT green dye showed specificity to the target genes as the one peak of melting curve showed. This elevation can be explained by the function of HMGB1 as a pro-inflammatory cytokine that is released from macrophages<sup>14-16</sup>. HMGB1 also can be released from neutrophils, monocytes and endothelial cells which are stimulated by inflammation. It can activate macrophage to produce TNF- $\alpha$ , IL-1 and other pro-inflammatory cytokines and it promotes the recruitment of inflammatory cells during RA which is a chronic inflammatory disease<sup>17</sup>. Moreover, the implication of HMGB1 in the

pathogenicity of arthritis is documented<sup>18, 19</sup>. It may contribute to the disease progress, as it is also promotes cell proliferation, modulation, migration and adhesive features. In addition, the extracellular HMGB1prolonged pro-inflammatory process<sup>20, 21</sup>.

Unlikely, the serum levels of HMGB1 in RA was normal which might be explained by the effect of antiinflammatory drugs that has been taken by the patients like methotrexate which reported to inhibit T- cells activation<sup>22</sup>. Similar results were reported for other AIDs such as medium vessel vasculitis (MVV), large vessel vasculitis (LVV), and Juvenile Idiopathic Arthritis(JIA) <sup>19</sup>in which the levels of HMGB1 gradually decreased to normal after treatment of the disease <sup>23</sup>.In addition, there are many factors that may affect the serum level of HMGB1 in AID such as age, sex, environmental and clinical factors<sup>19, 24</sup>.

Although the serum levels of HMGB1 in RA were normal but the correlation betweenHMGB1 gene and its protein in the serum showed a weak positive correlation which confirm the elevation of the protein but the interference of many factors (drug, age and sex) may affect the result of the protein level in the serum<sup>23, 25</sup>.

The RF level (IgG, IgM) in RA is high in compare to healthy controls <sup>2, 26</sup>, both antibodies showed a strong correlation in RA patients <sup>26</sup>. Despite this elevation in IgG and IgM some RA patients are negative to these antibodies as in many other study reported previous<sup>27</sup>.

The positive correlation between HMGB1 and IgG can be explained by the normal presence of IgG in chronic inflammation which is the case in RA <sup>28</sup>while IgM normally found in acute inflammation which explain that there is no correlation between HMGB1 and IgM<sup>29</sup>. Moreover, IgM has a relatively low affinity for modified self-components<sup>30</sup>.

In conclusion, HMGB1 gene expression is associated with RA patients in Basra. Despite the effector factors on its level in serum with confirmation of its correlation to IgG which is a characteristic feature of chronic inflammation such as in RA.

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