

Effects of mannuronic acid (M2000) on gene expression profile of signal transducer and activator of transcription proteins (STATs) in rheumatoid arthritis patients

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SUMMARY

Rheumatoid arthritis (RA), a form of inflammatory arthritis, is a chronic joint disease characterized by pain and inflammation that affects 0.5% to 1% of the population worldwide. The safety, efficacy, tolerability, and potency of β -D-mannuronic acid (M2000) as a novel NSAID with immunosuppressive property has been reported by several *in vitro* studies, experimental models and clinical trials phase I/II and III in ankylosing spondylitis and rheumatoid arthritis (RA) patients. This research is designed to study the therapeutic efficacy of oral administration of mannuronic acid in RA patients who had inadequate response to conventional drugs and to assess the effect of this drug on gene expression of the signal transducer and activator of transcription (STATs) protein (STAT1, STAT3, STAT4, and STAT6). The study has included 15 RA patients who had an insufficient response to the conventional therapy. The oral dose of mannuronic acid was 1000mg divided into two 500 mg doses per day for 3 months as an addition to conventional therapy. There were 15 healthy volunteer in the control group. Blood samples were collected from both groups, once from healthy controls and twice from RA patients before and after treatment by M2000. The peripheral blood mononuclear cells (PBMCs) were isolated to assess the gene expression level of STAT1, STAT3, STAT4, and STAT6 using the real-time PCR method. Results obtained in this study demonstrated a significant difference in the gene expression level of STAT1 between healthy controls and patients before treatment as well as a significant reduction in RA patients after treatment compared with the level before treatment. In addition, the gene expression level of STAT3 and STAT4 showed a significant reduction in RA patients after treatment compared to patients before treatment, while there was no significant difference between RA patients before treatment and the healthy control group for both molecules. On the other hand, there was no change in the gene expression level of STAT6 among all groups. The outcomes of this study confirmed that β -D-mannuronic acid (M2000) has the ability to control the levels of STAT1, STAT3 and STAT4 in RA patients, and might be beneficial in the management and therapy of RA.

Key words: Mannuronic acid M2000; STAT1; STAT3; STAT4; STAT6; arthritis; rheumatoid; gene expression.

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INTRODUCTION

Rheumatoid arthritis (RA) is a widespread common autoimmune disease associated with progressive disability, systemic complications, early death, and socioeconomic cost (1). RA is a persistent autoimmune inflammatory synovitis that affects about 1% of the population and

contributes to functional disability (2). The etiology of RA has not yet been identified. However, autoimmune mechanisms play a key role in the pathogenesis of the disease (3). Chronic inflammation in RA patients causes a synovial proliferation causing cartilage and bone absorption (4). RA is a persistent autoimmune inflammatory disease leading to progressive joint degeneration

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tion, disability, and elevated risk of cardiovascular complications, that represent the major cause of mortality among RA patients (5). The etiopathogenesis of RA is multifactorial and not fully understood, like in most autoimmune diseases. The autoimmune process leads to the dysregulation of cytokine synthesis, disturbances in the migration of immunocompetent cells and abnormal apoptosis. Up to date, the mechanisms through which inflammatory mediators, specifically cytokines, influence effector cells have not been fully studied. One of the transduction pathways that was found to have this influence is the Janus tyrosine kinase/signal transducer and activator of transcription (JAK/STAT) pathway. So far, seven homologous proteins of the STAT family have been identified in mammalian cells: STAT1, -2, -3, -4, -5A, -5B and -6. A single STAT type can be stimulated by many different ligands, but in general cytokines preferentially use a specific transduction pathway, e.g., INF γ usually selects STAT1 (6). When a cytokine is bound to a cell membrane receptor, the associated JAK tyrosine kinase is activated and the process of phosphorylation takes place. The molecules bind with subunits of cytoplasmic STAT proteins, which undergo dimerization. This complex is then translocated into the nucleus *via* importing. In the nucleus, STAT is released and binds with a DNA fragment responsible for the expression of a specific arrangement of genes (7). Previous studies have investigated the deregulation of the JAK/STAT pathway in RA. Elevated expression of the STAT1, STAT-3 and STAT-4 and proteins were found during active RA synovitis (6). Up-regulated levels of STAT3 mRNA in mononuclear cells from peripheral blood and synovial fluid, and elevated STAT1 expression in the synovial fluid have been observed in active RA (8). STAT3 has been found to be responsible for joint degeneration in RA (9). STAT1 controls cell growth proliferation, apoptosis, and functions of the immune system. Polymorphisms of STAT1 have been related to an increased risk of malignancy (10). In humans with autosomal-recessive

deficiency of STAT1, this deficiency leads to the progression of a primary immunodeficiency syndrome characterized by susceptibility to viral infections and bacterial pathogens (11). STAT1 has a role in inhibiting certain autoimmune disorders. Conversely, STAT1 mediates anti-proliferative and proapoptotic effects of IFNs, suggesting that STAT1 has also the potential for suppressing inflammation (12).

STAT3 is a pleiotropic transcription factor, which can play a role in the signal of different cytokines involving IL-6, IL-10, and other gp130 cytokines. Additionally, it could be included in immune and somatic cell abnormalities (12, 11). Recent research demonstrated that STAT3 plays an important role in inducing and maintaining a pro-carcinogenic inflammatory microenvironment. Therefore, STAT3 is believed to contribute to promoting oncogenesis (13, 14). In general, STAT3 has a complex role, as it can have different effects on various cells, depending on the cell type and activation status (15). Contrary to other STATs, that are expressed on a wide range of cell types, STAT4 is predominantly expressed by immune cells and the testis (16). STAT4 is activated by type I IFN in humans through interaction with STAT2 (17, 18). STAT4 is an important transcription factor for the biological function of several immune cells, such as macrophages, mast cells, natural killer cells, dendritic cells, T helper (Th) cells, regulatory T cells, follicular helper T cells, CD8+T cell and B cells (19-22). STAT4 can affect the activities of these cells that play a role in the pathogenesis of autoimmune diseases, such as systemic lupus erythematosus, RA, inflammatory bowel disease, multiple sclerosis, type 1 diabetes, systemic sclerosis, psoriasis and experimental autoimmunemyocarditis (19). By activating IL-12, STAT4 has an important role in the generation and proliferation of Th1 cells and is essential for the progression of Th17 cells (23, 24).

Up to 40% of patients with rheumatoid arthritis (RA) are insufficient responders (IR) to conventional disease-modifying antirheumatic drugs (DMARDs) or tumour necrosis factor α inhibitor (TNFi) biological

agents (25, 26). Patients who are intolerant and/or show an inadequate response to traditional DMARDs (DMARD-IR) are often treated with a biologic agent (27). For DMARD-IR patients, biologics are usually combined with traditional DMARDs, primarily MTX, but some biologics and also tofacitinib have shown to be effective as monotherapy as well (28-30). Adverse events (primarily gastrointestinal symptoms, respiratory symptoms, and hepatotoxicity) are the main reasons (>75%) for MTX withdrawal (31).

A systematic review reported two pivotal, international, double-blind randomized controlled trials, sponsored by the manufacturers, BEACON and BUILD, which both enrolled adult patients inadequately controlled on bDMARDs (TNF inhibitors) and, in the latter, patients were inadequately controlled on cDMARDs. Both studies had a 24-week double-blind treatment period in which baricitinib 2 mg and baricitinib 4 mg were compared with placebo. The primary outcome in each study was the proportion of patients achieving an American College of Rheumatology (ACR) improvement criteria of at least 20% (ACR20) response at 12 weeks. Key secondary outcomes that were accounted for multiplicity included health-related quality of life (HRQoL) on the Health Assessment Questionnaire-Disability Index (HAQ-DI), the Disease Activity Scale-28 and high-sensitivity C-reactive protein (DAS28-hs-CRP), and the Simplified Disease Activity Index (SDAI).

The primary outcome for both BUILD and BEACON was the proportion of patients achieving ACR20 at week 12. In both BEACON and BUILD (48.9% of baricitinib patients and 27.3% of placebo patients) and BUILD (65.9% versus 39.5%), more participants in the baricitinib group than in the placebo group achieved ACR20, and these differences were statistically significant between the groups. The proportion of patients achieving ACR20 at 24 weeks was higher with baricitinib than with placebo in BEACON (44.8% versus 27.3%) and BUILD (61.1% versus 42.1%). Interesting subgroup analyses were per-

formed on the primary outcome (ACR20 responses at week 12) based on the prior reason for failure on bDMARDs (lack of efficacy, adverse event, etc.) and for the number of previous bDMARDs in BEACON. Results for the subgroup which had a lack of efficacy were 49.1% for the baricitinib group versus 27.1% for the placebo group. In the *other* subgroup, there were only three patients in both groups. Therefore, it is difficult to understand why 50% responded in the baricitinib group and 100% in the placebo group. In both studies, baricitinib reduced (improved) the Health-Related Quality of Life (HAQ-DI), EQ-5D Health State Index/Self-Perceived Health score, DAS28-hs-CRP and MCID on the SDAI from baseline to week 12 in RA patients compared with placebo (32).

The β -D-mannuronic acid (M2000) (DE-102016113018.4), a novel NSAID with immunosuppressive property, is a safe agent with no toxicity for the GI tract and renal function (33). It has shown therapeutic benefits with the greatest tolerability, safety, and efficacy in various animal disease models, such as experimental autoimmune encephalomyelitis (EAE), adjuvant-induced arthritis (AIA), acute glomerulonephritis and nephrotic syndrome (34-37). The treatment with M2000 in combination with the conventional therapy demonstrated considerably greater efficacy along with a high safety profile compared to patients treated with the conventional therapy. Therefore, M2000 might be recommended as an appropriate choice in the remission of RA (38). The multinational phase III clinical trial showed that mannuronic acid is an effective and safe drug and is generally well-tolerated in patients with RA (39).

■ MATERIALS AND METHODS

Clinical characterization of patients and controls

15 rheumatoid arthritis patients with active disease were enrolled in this study. 80% were females and 20% were males. They were suffering from inadequate response to conventional drugs. The conventional therapy included DMARDs [Methotrexate

(MTX) 15-20 mg/week, Sulfasalazine (SSZ) 500-1000 mg/day and hydroxychloroquine (HCQ) 400 mg/day], corticosteroids (Prednisolone (PRD) 5-15 mg/day) and NSAIDs. As to the inclusion criteria, the RA patients were to use the drugs at least for 6 months before this study. The mean age was 52.33 ± 1.65 and the mean disease duration was 8.08 ± 1.60 years. The selected criteria for evaluating the immunological parameters of RA patients were based on the American College of Rheumatology recommendations, which considered DAS ≥ 3 in their clinical evaluation. The dose of mannuronic acid was 1000 mg per day divided by 2, then 500 mg in each capsule for three months, which was simultaneously administered with conventional drugs. Before starting our investigation, all the patients were informed and asked to sign an informed consent. Then, the patients received the follow-up appointments at baseline, 4 weeks and 12 weeks at the Department of Rheumatology in Loghman Hakim hospital Tehran, Iran, and the Division of Rheumatology Research, Rheumatism Center. Further follow-up was done by telephone in order to assess the adverse events of mannuronic acid. In addition, 15 individuals were enrolled as a healthy control group.

Ethical statement

The ethical approval with reference code number IR.TUMS.VCR.REC.1395.621-2016-09-14 was obtained from the Ethical Committee of Tehran University of Medical Sciences (TUMS) for this research study. A written and signed informed consent was obtained from all participants in the research. The research was conducted under the guidelines established by the American College of Rheumatology (ACR) and the Helsinki manifesto and its later amendments or comparable ethical standards.

β -D-mannuronic acid preparation and intake

The patented small molecule (DE-102016 113018.4) β -D-mannuronic acid (M2000) with molecular formula $C_6H_{10}O_7$ was

synthesized from alginic acid sodium salt (Sigma-Aldrich, St. Louis, MO, USA) based on a modified method of the acid hydrolysis tested (32). The purity of M2000 was approved by characterizing the hydrolytic products using Fourier transform infrared (FTIR) spectroscopy and carbon-13 nuclear magnetic resonance (^{13}C NMR) spectroscopy for asserting its molecular weight (194.139 g/mol) and exact/monoisotopic mass. The preclinical studies showed that M2000 is an anti-inflammatory agent with high biocompatibility and no toxicity on functions of kidneys and the GI tract (40-43).

Sample collection

Blood samples were collected once from 15 healthy volunteers and twice from 15 RA patients before and after treatment. Afterward, the peripheral blood mononuclear cells (PBMCs) were isolated using Ficoll-Paque (Amersham Pharmacia Biotech, Uppsala, Sweden) and stored at $-70^\circ C$. The entire amount of RNA was extracted from 2×10^6 - 5×10^6 cells using GeneAll[®] Hybrid-RTM kits (Qiagen, Valencia, CA, USA) according to manufacturer's guidelines and placed into 50 μ L of RNase-free water. The NanoDrop 2000 UV-Vis Spectrophotometer (Thermo Fisher Scientific, Waltham, MA, USA) was used to calculate the concentration of total RNA. It was then concentrated or diluted to a concentration of < 300 ng/ μ L for cDNA synthesis.

Quantitative real-time polymerase chain reaction

Reverse-transcribed random SolisBio Dye primers were used for real-time PCR based on cDNA Synthesis Kit protocol (Yekta E Takara Co., Ltd., Dalian, China). Quantitative real-time PCR was performed using 5X HOT FIREPol[®] EVAGreen[®]q PCR Mix Plus (ROX) with a specified primer (Table I) according to the available instructions. The analysis of STAT1, STAT3, STAT4, and STAT6 was fulfilled by ABI StepOnePlus[™] Real-Time PCR system (Applied Biosystems, Foster City, CA, USA). The relative quantities of selected genes mRNA were compared against one internal control,

Table I - The designed forward and reverse primers.

Gene Name	Forward primer	Reverse primer
STAT1	ACCTAACGTGCTGTGCGTAG	GAGACATCCTGCCACCTTGT
STA3	TCCTGAAGCTGACCCAGGTA	AGGTGAGGGACTCAAACCTGC
STAT4	CATCTCAACAATCCGAAGTGATTCA	GTCAGAGTTTATCCTGTCATTGAGCAG
STAT6	ACTCCAGGAGAGGTGTGAAAG	GAAGCAACTGGTGACGAGG
GAPDH	GAGAAGGCTGGGGCTCATTT	TAAGCAGTTGGTGGTGCAGG

GAPDH Mrna, which was calculated by considering the Δ CT method using an amplification plot (fluorescence signal vs cycle number). The difference (Δ CT) between the mean values in the duplicate samples of target genes and GAPDH mRNA was calculated. Afterward, the variation in ($\Delta\Delta$ CT) values of the gene expression of selected molecules among the groups was measured and reported as $2^{-\Delta\Delta$ CT.

Statistical analysis

All statistical analyses were carried out by SPSS software (24.0; IBM Corporation North Castle, Armonk, NY, USA). All data were expressed as means, standard deviations and $p < 0.05$ was considered statistically significant. The Kolmogorov-Smirnov test was used to check the normality of all data. The Analysis of Variance (ANOVA) test was used to compare the quantitative variables between the groups and the Newman-Keuls test to determine significant differences in the gene expression level between the untreated and treated groups. Statistical significance was classified as $*p < 0.05$, $**p < 0.01$ or $***p < 0.001$. A p-value of ≤ 0.05 was considered statistically significant.

RESULTS

Patient's response to treatment

After 3 months of mannuronic acid therapy, the medical reports showed an improvement in the clinical parameters of patients. At the end of the treatment, a significant decrease was observed in the level of Disease Activity Score for the 28-joint count (DAS28), tender joints count, swollen joints, patient's assessment and erythrocyte sedimentation rate (ESR), while there was

Table II - Clinical characterizations of patients before and after M2000 therapy.

Index	Before treatment	After treatment	P-value
DAS28	4.46±0.23	2.83±0.11	0.001
Tender joint count	4.00±0.56	1.00±0.27	0.001
Swollen joint count	2.33±0.64	0.50±0.19	0.011
Patient assessment of pain	66.67±3.95	40.00±4.76	0.005
ESR	22.33±3.83	14.08±2.65	0.016
CRP	41.7% (Positive)	33.3% (Positive)	1.000
RF	58.3% (Positive)	50% (Positive)	1.000

DAS28, Disease Activity Score of 28-joint, ESR, erythrocyte sedimentation rate; RF, rheumatoid factor; CRP, C-reactive protein.

no significant difference in the level of rheumatoid factor (RF) and C-reactive protein (CRP) (Table II).

The effects of β -D-mannuronic acid on STAT1 gene expression

Our findings demonstrated a significant difference in the gene expression level of STAT1 in the healthy control group compared to RA patients before treatment, as well as a significant difference between RA patients before and after a 12-week therapy with M2000 (Figure 1). Consequently, STAT1 as an inflammatory marker appeared significantly decreased following the treatment with M2000 in RA patients ($p < 0.01$).

The effects of β -D-mannuronic acid on STAT3 gene expression

In the gene expression level of STAT3, there was no significant difference between the healthy control group and RA patients before the M2000 therapy. Vice versa, there was a statistically significant difference in the gene expression level of STAT3 in the comparison between RA patients group be-

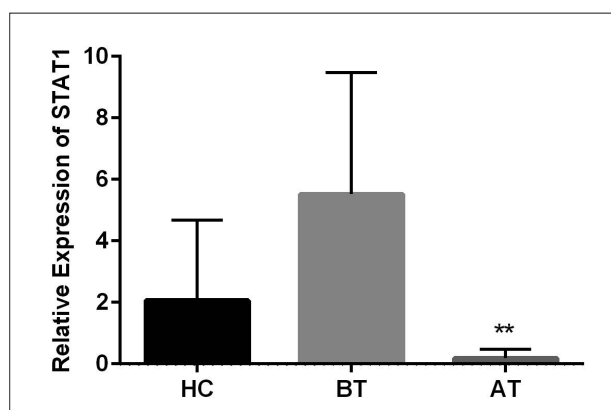


Figure 1 - Effects of β -D-mannuronic acid on STAT1 gene expression. HC, healthy control; BT, before treatment; AT, after treatment. The outcomes are shown as mean \pm SEM. p-value ≤ 0.05 was considered statistically significant. A significant reduction can be seen compared to before treatment: ** $p \leq 0.01$.

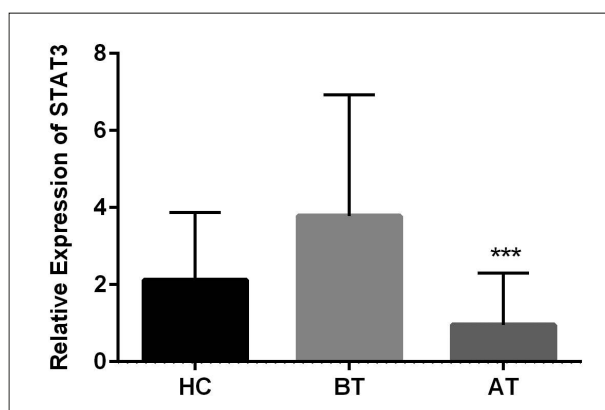


Figure 2 - Effects of β -D-mannuronic acid on STAT3 gene expression. HC, healthy control; BT, before treatment; AT, after treatment. The outcomes are shown as mean \pm SEM. p-value ≤ 0.05 was considered statistically significant. A significant reduction can be seen compared to before treatment: *** $p \leq 0.001$.

fore and after treatment with mannuronic acid for 12 weeks ($p \leq 0.001$) (Figure 2).

The effects of β -D-mannuronic acid on STAT4 gene expression

The gene expression level of STAT4 showed a statistically significant difference in the comparison between RA patients before and after treatment with M2000 for 3 months ($p \leq 0.00$). In addition, the clinical parameters reduced following the treat-

ment. On the other side the difference in the gene expression level of STAT4 between the healthy control group and the RA patients before treatment was not significant (Figure 3).

The effects of β -D-mannuronic acid on STAT6 gene expression

In contrast to previous groups of STATs, there was no significant difference in the gene expression level of STAT6 either in

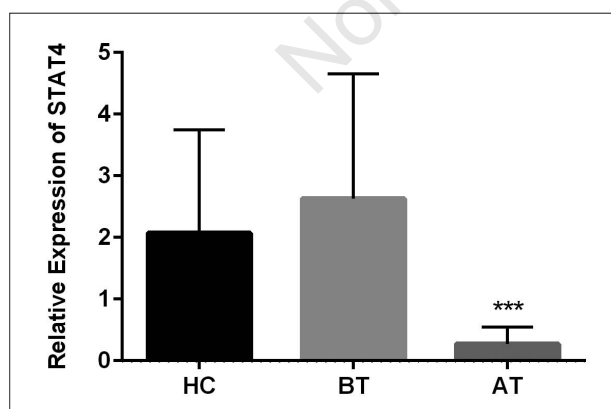


Figure 3 - Effects of β -D-mannuronic acid on STAT4 gene expression. HC, healthy control; BT, before treatment; AT, after treatment. The outcomes show mean \pm SEM. p-value ≤ 0.05 was considered statistically significant. A significant reduction can be seen compared to the value before treatment: *** $p \leq 0.001$.

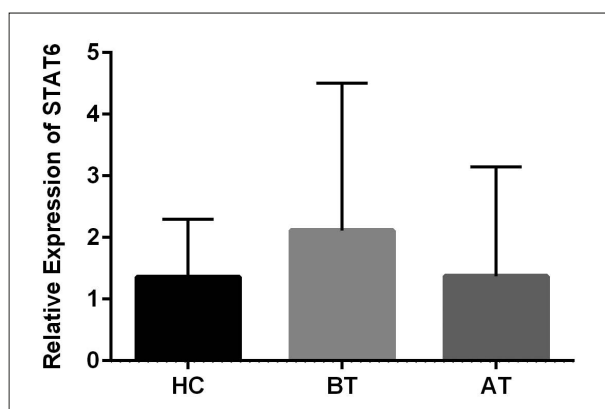


Figure 4 - Effects of β -D-mannuronic acid on STAT6 gene expression. HC, healthy control; BT, before treatment; AT, after treatment. The outcomes show mean \pm SEM. p-value ≤ 0.05 was considered statistically significant. No significant difference was identified in the gene expression level of STAT6 between all groups: $p \leq 0.093$.

healthy controls or in RA patients before and after the treatment with β -D-mannuronic acid ($p \leq 0.093$) (Figure 4).

■ DISCUSSION

Rheumatoid arthritis (RA) is a chronic autoimmune disorder that affects ~1-2% of the population worldwide. RA is characterized by inflammation, autoantibody production, cartilage and bone damage, and synovial hyperplasia. Inflammation promotes the synthesis of pro-inflammatory cytokines both systemically and in the joints, such as tumor necrosis factor- α and interleukin-6, which play a significant role in joint and other organ damage in this disease. Considering the role of the signal transducer and activator of transcription factors (STATs) in the signalling of these cytokines, these proteins may participate in the pathogenesis of RA. The expression and activity of STATs can contribute to the onset, progression, and virulence of RA. All STAT family members (STAT1, STAT2, STAT3, STAT4, STAT5a, STAT5b, and STAT6) have been related to autoimmune diseases, as highlighted in numerous research works (15). Despite several existing treatments and recent progress in disease therapy, remission rates and morbidity remain a critical concern for RA patients (44). Given the irreversible devastating damages caused by joint inflammation and the availability of effective disease-modifying drugs, it is widely recognized that earlier treatment is needed for more effective management of RA. There is also some evidence that early intervention can potentially be a curative treatment (45). Given the availability of various alternative biologic treatments and other novel disease-modifying antirheumatic drugs (DMARDs) for the treatment of patients with RA, it is increasingly challenging for clinicians to identify the optimal treatment. The biologic therapies are usually combined with traditional DMARDs, primarily methotrexate (MTX), but some biologics and tofacitinib have shown to be effective also as a monotherapy (46).

According to BEACON and BUILD, ba-

ricitinib is reported to improve clinical parameters in RA patients compared with placebo at 12 weeks, yet it was associated with serious adverse effects, such as upper respiratory tract infection, occurring in 9% of baricitinib patients and 5% of placebo patients in BEACON and in 6% of baricitinib and 8% of placebo patients in BUILD. Herpes zoster as a serious adverse event occurred in 1% of patients in each group in BEACON and 2% of baricitinib patients *versus* no placebo patients in BUILD. In addition, it also caused other notable events, including malignancies, thrombotic events, dyslipidemia, and elevations in hepatic enzymes. In addition, there were no clear differences between groups within BEACON and BUILD. There was a higher risk of elevated platelet counts with baricitinib *versus* placebo in BUILD (19% *versus* 5%); however, there was a much smaller difference between groups in BEACON (18% *versus* 14%). Low neutrophil counts were seen in 6% of baricitinib patients and 2% of placebo patients in BEACON and 8% of baricitinib patients *versus* 4% of placebo patients in BUILD. No patients had gastrointestinal perforation in either studies (32). Mannuronic acid as a novel and natural anti-inflammatory drug, which proved its safety, efficacy, tolerability, and potency in *in-vivo* and *in-vitro* tests and clinical trials in phase I/II in ankylosing spondylitis and in phase I/II and III in RA patients has been used in this study for 3 months in RA patients in order to evaluate its effect on STATs molecules. The results showed a significant response in the proposed gene expression as well as in clinical and para-clinical results. The previous clinical trial in RA patients evaluated different genes, but, in this study, we investigated the effect of this new drug on STATs which play an important role in the pathogenicity of this disease. In the synovia of RA patients and in experimental arthritis, STAT1 expression is increased and both STAT1 and STAT3 are in an activated state (47-49). STAT1 as an inflammatory factor was significantly different between the healthy control group and RA patients, even before the treatment, showing the importance of

this molecule in the pathogenicity of RA. In addition, the difference between patients before treatment and after treatment with mannuronic acid over 3 months was highly significant. Also, the gene expression of STAT1 was decreased following the treatment, thus proving that this drug is promising in controlling STAT1 transcription factors, which participate in regulating Th17 differentiation and controlling cell infiltration in inflamed joints (50). Not only did M2000 have an effect on STAT1, but it also had a significant effect on the gene expression level of STAT3 and STAT4 in RA patients after treatment compared to patients before treatment, as it reduced the gene expression of both molecules, which were elevated during the disease. These data indicated that this drug is effective in controlling both transcription factors in RA patients and suitable for the treatment of the disease. On the contrary, there was no significant difference in the gene expression levels of both molecules in the healthy control group compared to RA patients before treatment. This confirms again the efficacy of this drug on controlling both transcription factors, which participate in the pathogenicity in RA patients and reducing the level of gene expression of both molecules, which increased during the disease. On the other hand, this drug has not shown a significant effect on the gene expression level of STAT6 among the groups.

In conclusion our findings confirm the results of previous studies of mannuronic acid and its efficacy in different autoimmune diseases. This drug showed high efficacy in controlling the gene expression level of the transcription factors STAT1, STAT3 and STAT4, which have a significant role in the pathogenicity of RA. This drug which has shown its safety, efficacy, tolerability, and potency in different studies, has proven effective also in the treatment of RA patients and could be considered as a drug of choice in the near future.

Data availability

Data utilized to support the findings of this study are available from the corresponding author upon request.

Conflict of interest

The authors declare that they do not have any conflicts of interest.

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