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*Review*

# Evaluation of serum levels of IL-6, IL-10, and TNF-$\alpha$ in alopecia areata patients: A systematic review and meta-analysis

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## ABSTRACT

**Background**: The role of pro and anti-inflammatory cytokines in autoimmune and inflammatory disorders has always been discussed; several studies have assayed serum levels of Interleukin-6 (IL-6), Tumor Necrosis Factor-$\alpha$ (TNF-$\alpha$), and Interleukin-10 (IL-10) in alopecia areata (AA) patients. Determining the cytokine profile of AA patients will help us understand the role of the immune system in the pathogenesis of AA. Due to a lack of comprehensive studies in this regard, we have performed this meta-analysis to evaluate previously mentioned cytokines in AA patients. **Method**: We explored PubMed, Scopus, web of science, and ScienceDirect databases for research without a limited start date until May 2021. A number of 1098 studies were found in the initial database search and reference lists of relevant studies; 16 studies were finally included in the meta-analysis. Differences in the levels of IL-6, TNF-$\alpha$, and IL-10 in sera of controls and patients were pooled as standardized mean differences. A random-effects model was used in this study. Begg's and Egger's tests were carried out to investigate publication bias. **Results**: 16 studies were included. This study found significantly elevated levels of IL-6 in AA patients compared to healthy subjects (SMD = 1.57, 95% CI 0.17 to 2.97). The levels of TNF-$\alpha$ were also significantly higher in the serum of AA patients (SMD = 2.05, 95% CI 0.98 to 3.13). Though IL-10 levels were lower in serum of AA patients, this difference was not significant (SMD = -0.22, 95% CI -0.95 to 0.50). **Conclusion**: According to the crucial role of cytokines in autoimmunity, alternation in the serum levels of cytokines in AA patients was not unexpected; our study shows that cytokines might have an essential role in the pathogenesis of AA, though further studies are needed to clarify the exact role of cytokines in the emergence and persistence of AA.

**Key words:** Alopecia areata, Interleukin-10, Interleukin-6, Tumor Necrosis Factor-$\alpha$

## INTRODUCTION

Alopecia areata (AA) is an autoimmune, inflammatory, and nonscarring disease that leads to hair loss on the body or scalp. It has two main subgroups: Alopecia Totalis (AT) and Alopecia Universalis (AU). The former includes those patients with a total absence of terminal scalp hair, and the latter include those patients with complete loss of terminal scalp and body hair[1]. Few studies of prevalence and incidence have reported a 0.1 – 0.2% incidence with a lifetime risk of 1.7% equally in men and women. Furthermore, about 2% of new dermatology patients in the United States and the United Kingdom suffer from AA[1,2]. AA mostly shows itself through a sudden hair loss in well-demarcated sites. The area is commonly an oval or round patch of alopecia, and it can be numerous or isolated. Previous documents noted that though the most common complication of this disease is the involvement of the scalp hair, it may affect body hair such as underarm hair, beard, eyelashes, eyebrows, and pubic hair[2–5].

Circumstances such as infection and tissue injury can result in inflammation[6]. The inflammatory state can represent itself by elevated inflammatory cytokines such as IL-6, TNF$\alpha$, and CRP, and a decrease of i-inflammatory cytokines such as IL-10[7,8]. An inappropriate autoimmune response may stem from a chronic inflammatory condition[9]. It has been demonstrated that defects in cytokines and their signaling can lead to autoimmune diseases[10]. It is confirmed that cytokines play a crucial role in the CD4$^+$ T cell-mediated inflammatory response in AA[11].

IL-6 and TNF-$\alpha$ are two inflammatory cytokines that play an important role in different inflammatory disorders like rheumatoid arthritis and psoriasis[12,13]. Recent studies on the serum levels of these two inflammatory cytokines in AA patients have shown contradictory results[11,14,15]. Furthermore, recent studies on the serum levels of IL-10 in AA patients have shown paradoxical results, including both in-

creasing and decreasing[16,17].

Several studies have explored the links between AA and IL-6, IL-10 and AA and TNF-$\alpha$[11,16,18]. Due to inconsistent results and the lack of comprehensive studies, we performed a meta-analysis to determine the role of cytokines in AA and to provide more evidence for future related studies.

## METHODS

### Search method

PubMed, Scopus, Web of Science, and ScienceDirect were searched following the Preferred Reporting of Items for Systematic Reviews and *Meta-Analyses* (PRISMA) guidelines[19]. We used the following keywords: 'Alopecia areata' and 'Interleukin-10' or 'IL-10', 'Interleukin-6' or 'IL-6', 'TNF-$\alpha$' or 'tumor necrosis factor-alpha'. The reference lists of relevant studies were also reviewed. No restriction for the time was applied, and we searched databases from inception to May 2021.

### Exclusion and inclusion criteria

Only English articles were included in this study. The title and abstract of identified documents were reviewed to select relevant articles, and in case of doubt, full texts were reviewed. Duplicate studies and irrelevant studies such as review articles, case series, case reports, animal studies, genetic studies, and studies that did not report sufficient data were excluded.

### Data Extraction and Quality Assessment

We extracted the following information from the included studies: first author, publication year, country, disease type, measurement method, age of patients and healthy controls, and serum levels of IL-10, IL-6, or TNF-$\alpha$. If the mean and SD values were not mentioned in the text of manuscripts, we estimated them from given data such as median, range, and IQR.

The risk of bias was evaluated by NIH Quality Assessment Tools[20]. Quality assessment was performed by two authors.

### Statistical analysis

Random-effects meta-analysis was used to estimate standardized mean difference (SMD). The publication bias was assessed using Begg's and Egger's tests[21]. The heterogeneity of results between studies was checked by $I^2$ statistic[22], STATA/SE 11.0, and was utilized for statistical analysis (Stata-Corp, College Station, TX, USA).
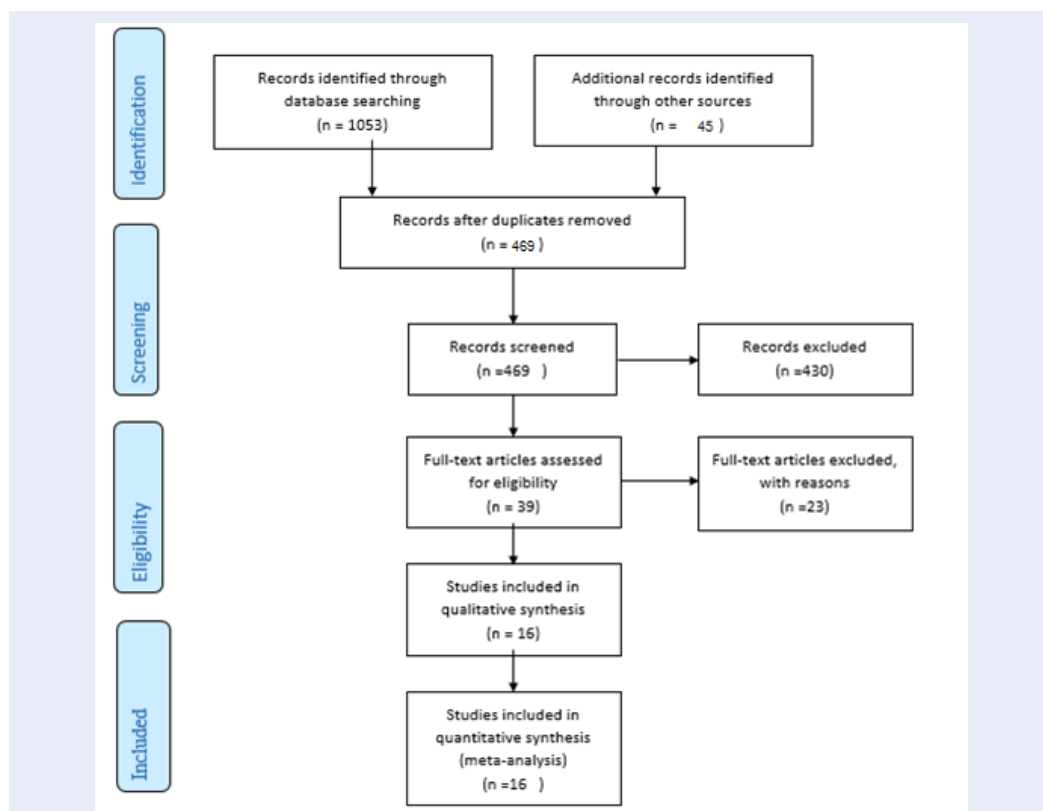


**Figure 1**: **The flow diagram regarding the flow of the number of records identified, included and excluded, and the reasons for exclusions through the different phases of a systematic review.**

**Table 1: Characteristics of included studies in the analysis of IL-6 serum levels**

| First author | Year | Country | Subgroups/Severity of disease | Number of cases/ healthy control | IL-6 levels in patients | IL-6 levels in healthy controls | IL-6 measurement | Age of cases/controls |
|---|---|---|---|---|---|---|---|---|
| Atwa et al.[23] | 2016 | Egypt | Single patch AA, Multiple AA, Ophiasis, AU, AT | 47/40 | $17.18 \pm 3.08$ | $4.59 \pm 1.66$ | ELISA | $22.68 \pm 8.62$ / $23.22 \pm 8.95$ |
| Bilgic et al.[11] | 2015 | Turkey | Mild, Moderate, Severe | 40/40 | $25.70 \pm 9.7$ | $11.80 \pm 9.2$ | ELISA | $28.7 \pm 7.9$ / $30.5 \pm 8.4$ |
| Ataseven et al.[14] | 2011 | Turkey | Not mentioned | 43/30 | $1.36 \pm 0.39$ | $1.44 \pm 0.56$ | ELISA | $23.42 \pm 11.41$ / $26.73 \pm 4.70$ |
| Tomaszewska et al.[24] | 2020 | Poland | Patchy, and Sever | 33/30 | $121.38 \pm 31.01$ | $75.26 \pm 21.15$ | ELISA | $18.64 \pm 8.56$ / $19.95 \pm 13.08$ |
| Barahmani et al.[18] | 2009 | USA | Transient AA, Persistent AA, AT or AU | 269/18 | AAP: 1.20* AAT: 1.50* AT / AU: 1.30* | 0.69* | Mutli-analyte proteome array | $39 \pm 18.8$ / $44 \pm 13.6$ |

**Abbreviations**: **AA**: Alopecia areata, **AU**: Alopecia universalis, **AT**: Alopecia totalis, **AAP**: persistent alopecia areata, **AAT**: transient alopecia areata, **IL-6**: Interleukin-6, *: median

**Table 2: Characteristics of included studies in the analysis of IL-10 serum levels**

| First author | Year | Country | Subgroups/Severity of disease | Number of cases/ healthy control | IL-10 levels in patients | IL-10 levels in healthy controls | IL-10 measurement | Age of cases/controls |
|---|---|---|---|---|---|---|---|---|
| Gautam et al.[16] | 2019 | India | Localized and Extensive AA | 40/40 | 3.144* | 1.050* | Sandwich ELISA | 29.4 ± 8.3 / 22.6 ± 3.8 |
| Loh et al.[25] | 2018 | Korea | Patchy, AT/AU, ADTA, Ophiasis | 55/15 | 4.54 ± 0.001 | 3.25 ± 0.071 | ELISA | 38.16 / 41.17 |
| Ma et al.[17] | 2017 | china | Severe AA in active or stable phase | 100/50 | | 21.20 | ELISA | Active phase: 41.2 ± 13.5, Stable phase: 40.9 ± 14.4/43.5 ± 16.1 |
| Ataseven et al.[14] | 2011 | Turkey | Not mentioned | 43/30 | 5.94 ± 1.17 | 6.32 ± 0.85 | ELISA | 23.42 ± 11.41 / 26.73 ± 4.70 |
| Tembhre et al.[26] | 2013 | India | Extensive AA,AT,AU | 51/45 | 9.47* | 10.47* | ELISA | 24.19 ± 6.69 / 26.42 ± 4.35 |
| Barahmani et al.[18] | 2009 | USA | Transient AA, Persistent AA, AT or AU | 269/18 | AAP: 0.85* AAT: 0.90* AT / AU: 0.85* | 0.69* | Mutli-analyte proteome array | 39 ± 18.8 / 44 ± 13.6 |

**Abbreviations**: **AA**: Alopecia areata, **AU**: Alopecia universalis, **AT**: Alopecia totalis, **ADTA**: acute diffuse and total alopecia, **AAP**: persistent alopecia areata, **AAT**: transient alopecia areata, **IL-10**: Interleukin-10, *: median

**Table 3: Characteristics of included studies in the analysis of TNF-$\alpha$ serum levels**

| First author | Year | Country | Subgroups/Severity of disease | Number of cases/ healthy control | TNF-alpha levels in patients | TNF –alpha levels in healthy controls | TNF –alpha measurement | Age of cases/controls |
|---|---|---|---|---|---|---|---|---|
| Kasumagic-Halilovic etal.[27] | 2011 | Bosnia and Herzegovina | LAA,AT/AU | 60/20 | 10.31 ± 1.20 | 9.59 ± 0.75 | ELISA | 35.6 / 32.6* |
| Abdel Halim et al.[28] | 2018 | Egypt | Multiple patches | 20/20 | 102.44 ± 16.16 | 39.03 ± 13.57 | Elisa | 28.4 ± 11.408 / 29.02 ± 13.57 |
| Atwa et al.[23] | 2016 | Egypt | Single patch, Multiple patches, Ophiasis ,AT ,AU | 47/40 | 19.94 ± 3.59 | 9.95 ± 2.42 | ELISA | 22.68 ± 8.62 / 23.22 ± 8.95 |
| Abd El-Raheem et al.[29] | 2020 | Egypt | Mild, Moderate, Severe | 75/75 | 8.8* | 1.4* | ELISA | 24.8 ± 18.6 / 26.3 ± 4.328 |
| Bilgic et al. [11] | 2015 | Turkey | Mild, Moderate, Severe | 40/40 | 20.60 ± 6.20 | 10.20 ± 7.50 | ELISA | 28.7 ± 7.9 / 30.5 ± 8.4 |
| Loh et al.[25] | 2018 | Korea | Patchy, AT /AU, ADTA), Ophiasis | 55/15 | 12.76 ± 0.003 | 3.33 ± 0.011 | ELISA | 38.16 / 41.17 |
| A. Alzolibani et al.[30] | 2016 | Saudi Arabia | Patchy persistent | 25/26 | 27.70 | 14.70 | Sandwich ELISA | 30.2 ± 8.43 / 32.3 ± 10.8 |
| Serarslan et al.[15] | 2020 | turkey | Patch pattern, Patch and ophiasis pattern, AT,AU | 36/34 | 113.68 ± 106.34 | 121.15 ± 55.31 | sandwich ELISA | 31.33 ± 9.47 / 30.5 ± 9.11 |
| Teraki et al. [31] | 1996 | Japan | Localized form | 7/7 | 8.30 ± 0.90 | 7.60 ± 0.20 | Radioimmunoassa kit | 24 / 21 |
| I. Omar et al. [32] | 2021 | Egypt | Patchy, AU, AT, Ophiasis | 72/75 | 5.3* | 3.9* | ELISA | Adults: 37.6 ± 12, Children: 11 ± 3.3/ Adults: 37.4 ± 13.7, Children: 12.9 ± 2.8 |
| Barahmani et al.[18] | 2009 | USA | Transient AA, Persistent AA, AT or AU | 269/18 | AAP: 3.58* AAT: 3.06* AT / AU: 2.97* | 2.46* | Mutli-analyte proteome array | 39 ± 18.8 / 44 ± 13.6 |

**Abbreviations**: **AA**: Alopecia areata, **AU**: Alopecia universalis, **AT**: Alopecia totalis, **ADTA**: acute diffuse and total alopecia, **AAP**: persistent alopecia areata, **AAT**: transient alopecia areata, **TNF-alpha**: Tumor necrosis factor-alpha, *: median

## RESULTS

A total of 1098 articles were found in the aforementioned databases. 629 duplicate papers were excluded. 453 studies were excluded after a review of the titles and abstracts. Finally, 16 studies were selected to be involved. The detailed search process is shown in **Figure 1**. Extracted data for IL-6, IL-10, and TNF-$\alpha$ are reported in **Tables 1 and 2** and **Table 3**, respectively. The levels of IL-10 in the serum of healthy subjects were higher than AA patients, but the difference was not significant (SMD = -0.22, 95% CI -0.95 to 0.50). Among six studies that report IL-10 levels in the sera of AA patients, one study shows reduced levels of IL-10 in AA patients compared to healthy controls[17], one study shows a raised level[16], and four studies show no significant difference[14,18,25,26]. There was a significant elevation of IL-6 in the serum of AA patients compared to healthy controls (SMD = 1.57, 95% CI 0.17 to 2.97).

Among 5 studies that describe IL-6 levels, three studies show elevated levels of IL-6 in AA patients[11,23,24], and two studies show no significant difference between AA patients and healthy subjects[14,18]. Significantly elevated levels of TNF-$\alpha$ in the serum of AA patients compared to healthy subjects were found (SMD = 2.05, 95% CI 0.98 to 3.13).

Among 11 studies that describe TNF-$\alpha$ levels in AA patients and healthy controls, eight studies describe higher levels of TNF-$\alpha$ in AA patients[11,23,25,27–30,32], and no significant difference was seen in three studies[15,18,31]. Forest plots of TNF-$\alpha$, IL-6, and IL-10 are shown in **Figures 2 and 3** and **Figure 4**, respectively. The p-values of Begg's tests (p = 0.014) and Egger's tests (p = 0.021) for IL-6 indicate evidence of publication bias.

## DISCUSSION

Alopecia areata is an autoimmune disorder that causes small patches of baldness on the scalp or/and body, and it is linked to some coexisting autoimmune disorders such as rheumatoid arthritis, vitiligo, and psoriasis[33,34]. The pathogenesis of AA is not completely understood. It seems that it is associated with a complex interplay between genetic susceptibility and immune system function[33,35]. The abnormal levels of pro-inflammatory cytokines (IL-6 and TNF-$\alpha$) and anti-inflammatory cytokine (IL-10) in AA were confirmed by previous documents[11,16]. The present meta-analysis discovers a significant link between AA and the inflammatory markers, with SMD for TNF-$\alpha$ (SMD = 2.05, 95% CI 0.98 to 3.13), followed by IL-6 (SMD = 1.57, 95% CI 0.17 to 2.97) and then IL-10 (SMD = -0.22, 95% CI -0.95 to 0.50).

The molecular mechanisms responsible for these alterations in immune function need to be explored more fully[36]. The objective of this meta-analysis was to collate data from published literature to evaluate the serum TNF-$\alpha$, IL-6, and IL-10 levels in patients with AA.

TNF-$\alpha$ is manufactured by a wide range of cells, including immune and non-immune cells, and is considered to be a pro-inflammatory cytokine that has an essential role in the pathogenesis of many inflammatory diseases such as arthritis, rheumatoid, and psoriasis[37–39]. Our meta-analysis shows increased serum levels of TNF-$\alpha$ in AA patients compared to healthy individuals, which was expected due to the nature of AA which seems to be a Th1 dominant autoimmune disease[40]. Hoffmann *et al.* showed that TNF-$\alpha$ could abrogate hair growth *in vitro*; it has been found that TNF-$\alpha$ can result in vacuolation of matrix cells in follicle bulbs and can diminish the matrix size[41,42]. Alzolibani *et al.* showed that mRNA expression of TNF-$\alpha$ is higher in PBMC of AA patients than healthy controls[30]. It has been found that lesioned skin biopsies of AA patients have elevated levels of TNF-$\alpha$ in comparison with non-lesioned biopsies or biopsies from healthy controls[43].

IL-6 is a pro-inflammatory cytokine that plays an important role in the pathogenesis of different autoimmune diseases or chronic inflammatory conditions[44]. This study showed that IL-6 levels are significantly raised in AA patients compared to healthy individuals. Transcript of IL-6 has been shown to have higher expression in balding dermal papilla cells than nonbalding dermal papilla cells and IL-6 can also inhibit elongation of the hair shaft in a dose-dependent manner[45]. IL-6 levels tend to be elevated in autoimmune diseases regardless of the dominant response; it is elevated in multiple sclerosis patients (Th1 dominant), SLE (TH2 dominant), and psoriasis (in which Th17 plays a crucial role in its pathogenesis). Therefore, elevated levels of IL-6 in AA patients are not unexpected[46–51].

IL-10 is an anti-inflammatory cytokine, and its dysregulation can increase the chance of autoimmunity[52]. Contrary to our expectations, pooled anti-inflammatory IL-10 was not significantly decreased in patients with AA. The literature on AA proposes that IL-10 imperfection might contribute to its pathogenesis[16]. It seems that disequilibrium in cytokine production, with a relative surplus of pro-inflammatory and Th1 types versus anti-inflammatory cytokines, could have a role in the permanence of alopecia lesions[53]. Freyschmidt-Paul *et al.* found that IL-10 deficient mice are more resistant to the induction of
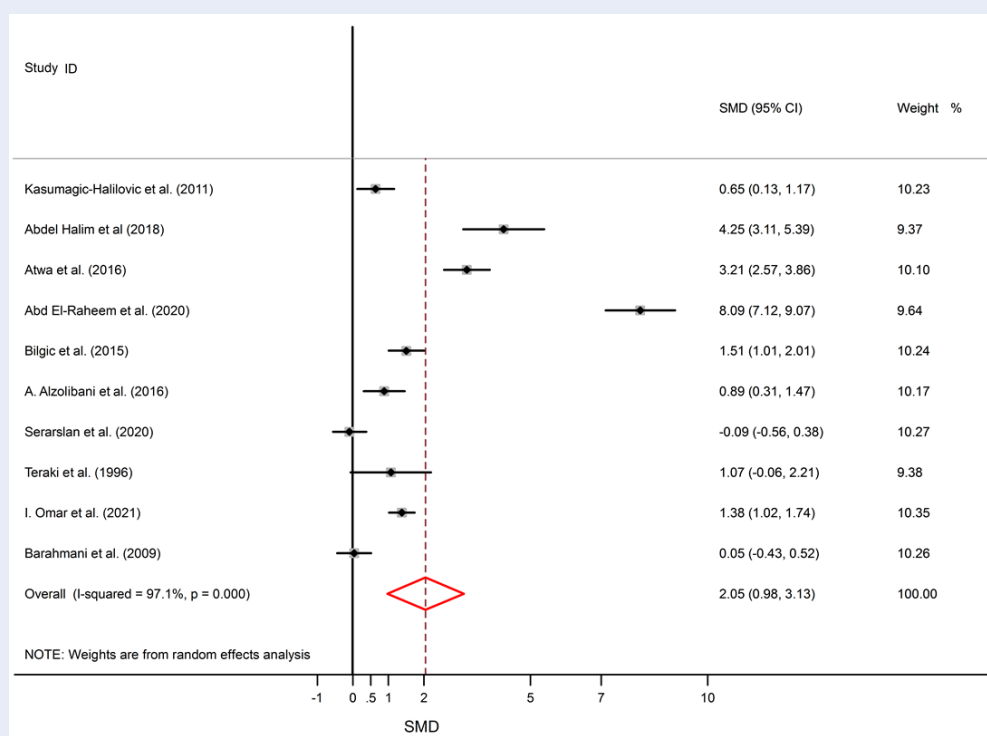
**Figure 2**: **Forest plot of serum TNF-$\alpha$ levels in AA patients and controls**. The square represents the point estimate of the effect for each study, and its size is proportional to the weight of this included study. The horizontal line through the square represents the 95% CI around the point estimate. The vertical line is the line of no effect. The diamond represents the pooled standardized mean difference and the width of the diamond represents the 95% CI around this estimate. https://doi.org/10.6084/m9.figshare.16910314.v1

AA[54]. Hoffman *et al.* showed increased mRNA levels of IL-10 in AA biopsies after treatment with Diphenylcyclopropenone, and IL-10 transcript levels in untreated biopsies of AA patients were nearly similar to healthy controls[55].

Altogether, these lines of evidence show that activation of the immune system is one of the biological processes associated with the pathogenesis of AA. Thus, the present study could be worthwhile in providing an evidence pool for future treatment suggestions.

There were a few limitations associated with our meta-analysis. First, insufficient data in some studies was a significant issue in our conducted meta-analysis. Second, due to the inconsistency of the classification methods, the different disease subtypes of AA could not be separately analyzed using the meta-analysis. Third, the limited number of studies may lead to less accurate results in meta-analysis studies. Fourth, publication bias could affect the results of this meta-analysis study.

## CONCLUSIONS

Altogether, these lines of evidence show that activation of the immune system is one of the biological processes which may be associated with the pathogenesis of AA. The present study could be worthwhile in providing an evidence pool for future treatment suggestions. Despite of our findings, we still need further researches with a stronger design to develop knowledge in this regard.

## ABBREVIATIONS

**AA**: Alopecia areata

**IL-6**: Interleukin-6

**IL-10**: Interleukin-10

**TNF-$\alpha$**: Tumor Necrosis Factor-$\alpha$
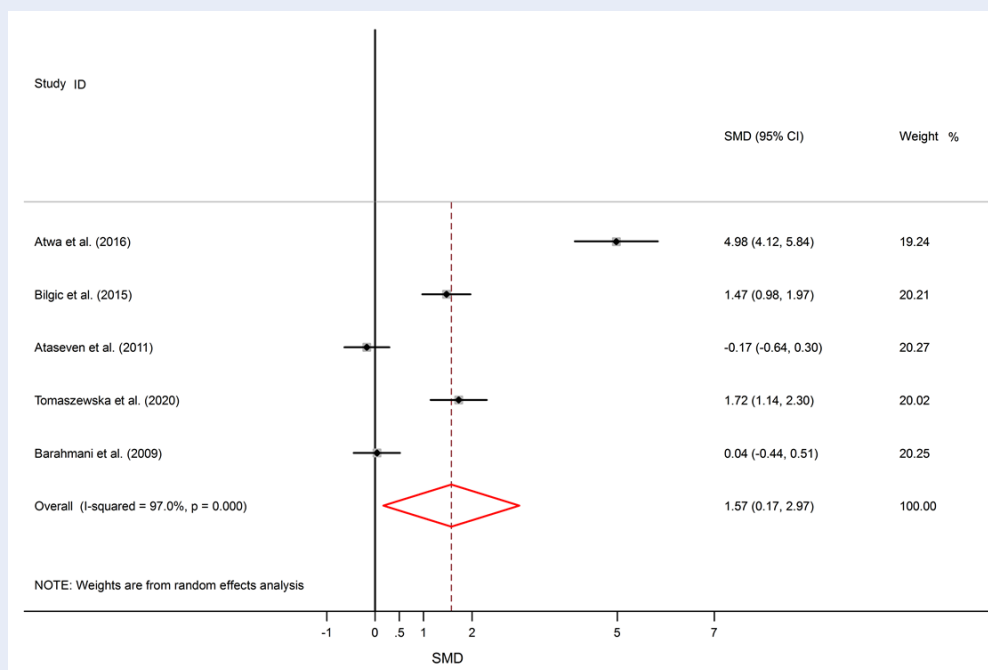
## ACKNOWLEDGMENTS

**Figure 3**: **Forest plots of serum IL-6 levels in AA patients and controls.** The square represents the point estimate of the effect for each study, and its size is proportional to the weight of this included study. The horizontal line through the square represents the 95% CI around the point estimate. The vertical line is the line of no effect. The diamond represents the pooled standardized mean difference and the width of the diamond represents the 95% CI around this estimate. https://doi.org/10.6084/m9.figshare.16910317.v1

## AUTHOR'S CONTRIBUTIONS

Authors equally contributed to this work. All authors read and approved the final manuscript.

## FUNDING

None.

## AVAILABILITY OF DATA AND MATERIALS

Data and materials used and/or analyzed during the current study are available from the corresponding author on reasonable request.

## ETHICS APPROVAL AND CONSENT TO PARTICIPATE

This study was conducted in accordance with the amended Declaration of Helsinki. The institutional review board approved the study, and all participants provided written informed consent.

## CONSENT FOR PUBLICATION

Not applicable.

## COMPETING INTERESTS

The authors declare that they have no competing interests.

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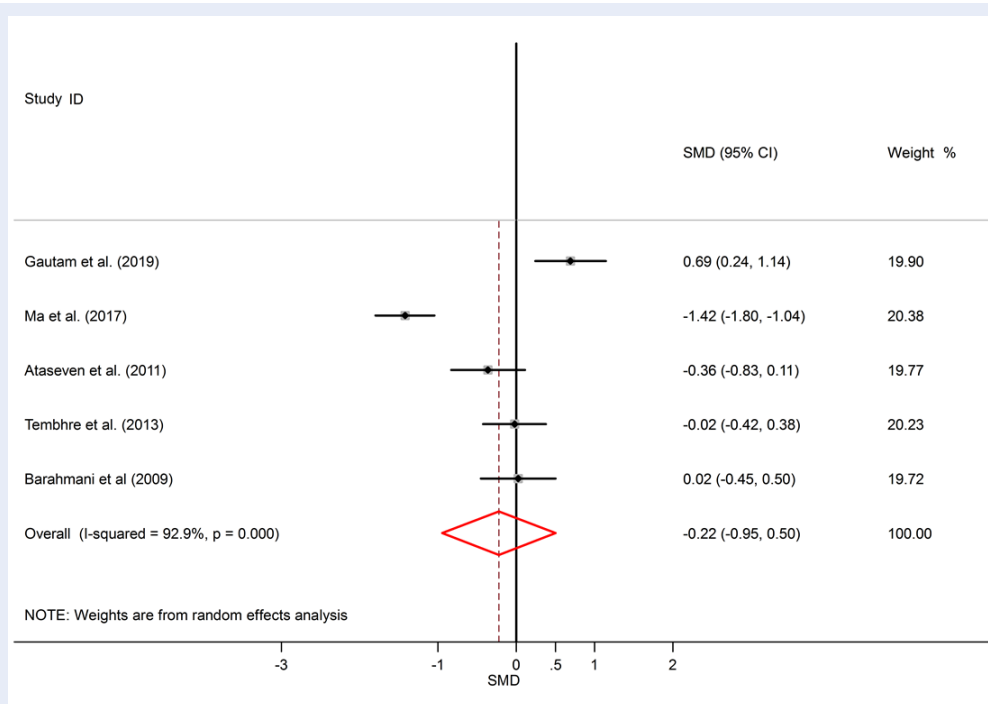
**Figure 4**: **Forest plots of serum IL-10 levels in AA patients and controls**. The square represents the point estimate of the effect for each study, and its size is proportional to the weight of this included study. The horizontal line through the square represents the 95% CI around the point estimate. The vertical line is the line of no effect. The diamond represents the pooled standardized mean difference and the width of the diamond represents the 95% CI around this estimate. https://doi.org/10.6084/m9.figshare.16910320.v1

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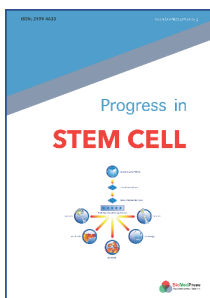**ISSN**: 2198-4093
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