The effect of disease-modifying anti-rheumatic drugs on skeletal muscle mass in rheumatoid arthritis patients: a systematic review with meta-analysis

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Abstract

Introduction: Rheumatoid arthritis (RA) is an autoimmune disease, characterized by chronic and systemic inflammation. Besides, it is known that RA patients may present several comorbidities, such as sarcopenia, a condition where patients present both muscle mass and muscle quality impairment. RA treatment is mostly pharmacological and consists in controlling systemic inflammation and disease activity. Despite that, the effect of pharmacological treatment on sarcopenia is not well characterized.

Objective: To summarize the effects of disease-modifying anti-rheumatic drugs (DMARDs) on skeletal muscle tissue in rheumatoid arthritis (RA) patients.

Methods: A systematic review of randomized clinical trials and observational studies was conducted using MEDLINE, Embase, Cochrane Library, and Web of Science. We selected studies with rheumatoid arthritis patients treated with disease-modifying anti-rheumatic drugs (DMARDs) that analyzed muscle mass parameters such as lean mass and appendicular lean mass. Methodological quality was assessed using the Newcastle-Ottawa Quality Assessment Scale. Standardized mean difference (SMD) and 95% confidence intervals (CI) were set. A meta-analysis of observational studies was performed using the R software, and we considered significant statistics when \( p < 0.05 \).

Results: Nine studies were included in this systematic review. In the meta-analysis, DMARD treatment had no positive difference (\( p = 0.60 \)) in lean mass. In the same way, in the appendicular lean mass parameter, our results showed that DMARDs did not have changes between baseline and post-treatment analysis (\( p = 0.93 \)).

Conclusion: There is no evidence of a significant effect of DMARD therapy, either synthetic or biological, on muscle mass. However, this association should be investigated with more studies.

Keywords: Systematic review, Rheumatoid arthritis, Sarcopenia, Muscle loss, Lean mass, Appendicular lean mass, Treatment, Drugs, DMARD

Introduction

Rheumatoid arthritis (RA) is a chronic, autoimmune disease characterized by systemic inflammation that affects mainly the joints [1]. In addition, RA leads to several comorbidities, such as cardiovascular disease and metabolic syndrome [2, 3]. Furthermore, RA
patients are often associated with changes in body composition [3, 4] such as reduced skeletal muscle mass [5], decreased muscle strength [6], and poor physical function [4, 5, 7, 8]. The alterations in RA patients regarding body composition can be a description of a sarcopenic patient that carries the risk of physical incapacity, low quality of life, and death [9, 10]. The prevalence of sarcopenia ranges from 25.9 to 43.3% between cohort studies, a wide variation due to the differences in sample, age, gender, race, and definitions and methods of diagnosing sarcopenia [9, 11–13].

Muscle impairment in RA and during sarcopenia is associated with several mechanisms triggered by inflammatory signaling [14]. Inflammatory mediators, such as tumor necrosis factor α (TNF-α) and interleukin 1β (IL-1β), are pointed out as triggers of catabolic effects in muscle tissue [14]. Thus, interleukin 6 (IL-6) has a role in driving catabolism in muscle mass and anabolism in fat mass [15, 16].

The available treatments for RA aim to attenuate disease activity by blocking inflammatory mediators and their signaling or inducing anti-inflammatory and regulatory pathways [17]. Disease-modifying anti-rheumatic drugs (DMARDs) significantly improve disease activity and prevent joint damage in RA by targeting the key inflammatory pathways [17, 18]. The classification of therapeutic drugs are as follows: conventional synthetic DMARDs (csDMARDs), which comprehend methotrexate, leflunomide, sulfasalazine, and hydroxychloroquine; biological DMARDs (bDMARDs); and targeted synthetic DMARDs (tsDMARDs) [19, 20].

bDMARDs include targeting monoclonal antibodies against TNF (infliximab, adalimumab, certolizumab, and golimumab) and IL-6 (tocilizumab and sarilumab), soluble receptor for TNF (etanercept), an inhibitor of T-cell co-stimulation (abatacept), and anti-CD20 B-cell depleting monoclonal antibody (rituximab) [21]. The tsDMARDs are inhibitors of the Janus tyrosine kinase family (JAK), which targets intracellular signaling of type I and II cytokines (tofacitinib, baricitinib, upadacitinib, and filgotinib). Thus, tsDMARD effects are T cell reduction and decreased leukocyte recruitment to joint, resulting in less synovial inflammation and prevented joint damage in RA patients [22, 23]. In a recent narrative review published by our group [8], we found that csDMARDs, tsDMARDs, and bDMARDs have no benefits on muscle mass when used to treat RA patients. However, tocilizumab, an IL-6 inhibitor, may improve muscle mass by increasing appendicular lean mass and total lean mass in RA patients [8]. Also, glucocorticoids (CG) that can control disease activity in RA have known negative effects on the skeletal muscle in RA patients [24, 25]. Targeting inflammatory cytokines seems to have a positive role in muscle wasting.

Data from many studies have shown that cytokine inhibition has been effective at preventing or treating muscle wasting. TNF-α blockade can partially revert muscle atrophy by suppression of the NF-κB pathway in several animal models and can prevent survival in aging mice [26–28]. Additionally, IL-6 have been independently implicated in some forms of muscle atrophy, and its deficiency attenuates atrophy in sepsis, diabetes mellitus, and Duchenne muscular dystrophy [16, 29–31].

Nevertheless, sDMARDs and bDMARDs can control disease activity by blocking inflammatory signaling, but their effect on skeletal muscle tissue in RA patients remains unclear. Thus, this systematic review aims to summarize the current evidence on the effect of pharmacological treatment on skeletal muscle tissue in RA patients.

Materials and methods
We conducted this systematic review in accordance with the PRISMA [32] guidelines after registering the protocol with the PROSPERO platform (CRD42021279386).

PICOS/PECOS format
This systematic review with meta-analysis was based on a focused question described in a PICO/PECO format [33]. We established the following: Patient/Problem/Population= Rheumatoid arthritis patients, Intervention/Exposure= Chronic treatment with biological and synthetic DMARD and glucocorticoids, Comparison= Baseline and post treatment, Outcomes= Muscle mass parameters, such as muscle mass, fat-free mass, appendicular lean mass and lean mass; and Study= Randomized clinical trials and Observational studies.

Data sources
The electronic databases used were Cochrane Library, PubMed, Embase, and Web of Science (DATA). We used a comprehensive search strategy tailored to each database. We contacted the authors, when necessary, for more information on the statistical methodology of the articles chosen as a reference. However, in some cases, we have not received any feedback.

Search terms
Keywords and medical subject headings (MeSH) for the following terms: “Rheumatoid arthritis,” “Antirheumatic agents,” “Methotrexate,” “Leflunomide,” “Sulfonamides,” “Hydroxychloroquine,” “Glucocorticoids,” “Tumor necrosis factor,” “Interleukin-6,” “Janus Kinases,” “Muscle,” “Skeletal,” “Body composition,” “Cachexia,” “Sarcopenia,” and related terms were selected. The term OR was used for Union of MeSH terms and “entry terms,” and the
The results obtained for lean mass and appendicular lean mass. After the authors' agreement, nine studies were included in this review. The baseline mean and after-treatment mean were extracted and converted and included authors' names, date of publication, journal of publication, number of participants in the study, the age group of the population, type of population, type of treatment, duration of treatment, treatment posology, and results obtained for lean mass and appendicular lean mass. After the authors' agreement, nine studies were included in this review. The baseline mean and after-treatment mean were extracted and converted and the delta of the mean (difference of final mean and baseline mean) for meta-analysis. In one study [34], we estimated the baseline mean from graph bars with the ImageJ software.

**Methodological quality assessment**

Methodological quality was assessed by the Newcastle-Ottawa Scale for cohort studies or for randomized clinical trials [35–37] by two independent reviewers (Santos, LP, and Portes, JKS). In these scales, each study was judged by questions about groups of criteria: selection of cohort, comparability of the study, and ascertainment of the outcomes for cohort studies and selection, comparability, and exposure for randomized clinical trials. For each item, in the selection, outcome, and exposure groups, a maximum of one star can be assigned, and for the comparability group, a maximum of two stars can be assigned. So, the maximum possible score was 9 stars. Based on the scale, studies with scores of 3 or 4 in the selection, 1 or 2 scores in comparability, and 2 or 3 in outcome or exposure were classified as good-quality studies. On the other hand, studies with 2 stars in the selection, 1 or 2 stars in comparability, and 2 or 3 stars in outcome and exposure were classified as fair-quality studies. Finally, studies with scores of 0 or 1 in the selection, 0 scores in comparability, or 0 or 1 score in outcome or exposure were classified as poor-quality studies.

**Risk of bias assessment**

The risk of bias in the randomized clinical trials was assessed using the Risk of Bias Tool 2.0 (RoB2) from Cochrane to randomized clinical trials [38]. The evaluators examined the randomization process, deviations from intended interventions, missing outcome data, measurement of the outcome, and selection of the reported results. Thus, the studies were classified into low, moderate, or high risk of bias.

**Statistical analysis**

The meta-analysis was conducted using the mean change and SD change from each study. All outcome measures were continuous variables. A meta-analysis, representing the effects of interventions, was performed: the random-effects model with the mean difference (MD) MD was performed when studies reported outcomes using the same assessment scale or assessment instrument.

The 95% confidence intervals (CI) were used, and the heterogeneity of the studies included in the meta-analysis was assessed using the inconsistency test ($I^2$). We considered low, moderate, and high inconsistency in the approximated values of 25%, 50%, and 75%, respectively [39, 40]. The software used for statistical analysis was RevMan (Review Manager 5.4.1, The Cochrane...
and we considered it significant statistically when $p < 0.05$.

**Results**

**Search strategy**

We identified 1123 possible studies (134 duplicate publications) based on our search items. First, the title and abstract of the 1244 studies were screened. After this process, 32 articles were included for the full-text screening. Finally, after the full-text reading, we included nine articles: Engvall et al. [41] and Marcora et al. [42] as randomized clinical trials and Al Khayyad et al. [43], Vial et al. [44], Tournadre et al. [45], Toussirot et al. [46], Ferraz-Amaro et al. [47], Metsios et al. [48], and Chikugo et al. [34] as observational studies. The search and inclusion/exclusion criteria are described in Fig. 1.

**Characteristics of the studies**

Of the nine studies included, four of them were performed in France [42, 44–46], one of them in the UK [48], one in Spain [47], one in Japan [34], one in Sweden [41], and one in Italy [43]. Studies included were published between 2007 and 2020. Only one study was performed using female patients [34] while the other four were performed with male and female patients [45–48]. Included papers reported sample size ranged from 8 to 146 subjects, patients’ age means from 50 to 61 years. Studies also showed baseline DAS-28 ranged from 3.0 to 6.1 [34, 45–48]. Characteristics of the included studies are described in Tables 1 and 2.

**Characteristics of treatments**

Among the nine papers included, the treatments used in the studies were tocilizumab [45, 46], anti-TNF [47, 48],...
| First author name | Country   | Gender | Year | Sample size | Age      | Treatment        | Dose | Measure | Lean mass (kg) | App. lean mass (kg) | Method of BC assay | Pre-DAS-28          | Post-DAS-28       |
|--------------------|-----------|--------|------|-------------|----------|-------------------|------|---------|---------------|---------------------|-------------------|-------------------|-------------------|
| Tournadre [23]     | France    | M/F    | 2017 | 21          | 57.8 ± 10.5 | Tocilizumab 12 months | NI   | Lean mass | Baseline: 42.1 (± 11.1) | Final: 43.2 (± 11.3) | DEXA              | 4.94 ± 1.25 | 28 ± 1.5 |
| Toussirot [22]     | France    | M/F    | 2020 | 107         | 56.6 ± 13.5 | Tocilizumab 12 months | 8 mg/kg (monthly) | Lean mass | Baseline: 40.76 (± 8.4) | Final: 42.11 (± 8.9) | DEXA              | 4.93 ± 1.3  | 23 ± 1.3 |
| Ferraz-Amaro [24]  | Spain     | M/F    | 2011 | 16          | 50.8 ± 14.6 | Anti-TNF 12 months | Varied | Lean mass | Baseline: 53.7 (NI) | Final: 50.5 (NI)    | BIA               | 5.58 ± 0.87 | 2.89 ± 1.37 |
| Metsios [25]       | United Kingdom | M/F | 2007 | 20          | 61.1 ± 6.8  | Anti-TNF 3 months | NI   | Lean mass | Baseline: 50.9 (± 12.7) | Final: 51.1 (± 12.5) | BIA               | 5.66 ± 0.7  | 3.59 ± 0.7 |
| Chikugo [26]       | Japan     | F      | 2018 | 4           | 55.3 ± 19.5 | Tofacitnib 6 months | NI   | App. lean mass | –              | –                   | BIA               | 5.1 ± 0.8  | NI     |
| Al Khayyat [27]    | Italy     | F      | 2021 | 20          | 65 ± 12.9   | Rituximab 18 months | Eight infusions of 500 mg ~1 g | Lean mass | Baseline: 39.94 ± 8.74 | Final: 38.64 ± 8.19 | DEXA              | 4.21 ± 1.1  | NI     |
| Vial [28]          | France    | Male/Female | 2021 | 83          | 58.5 ± 10.8 | Biologic DMARD (TNFi and non-TNFi) | NI   | Lean mass | TNFi Baseline: 49.6 ± 10.8 | Final: 50.7 ± 11.3 Non TNFi Baseline: 47.7 ± 11.0 | DEXA              | 4.21 ± 1.1  | NI     |

*BIA* bioimpedance, *DEXA* dual-energy X-ray absorptiometry, *NI* not informed, *TNF* tumor necrosis factor, *TNFi* TNF inhibitor, *BC* body composition, *M* male, *F* female
Table 2 Characteristics of the clinical trials included in the systematic review with meta-analysis

| First author name | Country | Gender | Year | Sample size | Treatment | Mean age | Dose | Measure | Method of BC assay | Pre-DAS-28 | Post-DAS-28 |
|--------------------|---------|--------|------|-------------|-----------|----------|------|---------|-------------------|------------|-------------|
| Marcora [29]       | France  | M/F    | 2006 | 24          | Group 1: etanercept Group 2: MTX | NI       | Group 1: 50 mg/week Group 2: 7.5–15 mg/week | Lean mass | DEXA             | Group 1: 6.1 ± 0.7 Group 2: 5.8 ± 1.1 | Group 1: t3.2 ± 1.5 Group 2: 3.1 ± 1.5 |
| Engvall [30]       | Sweden  | M/F    | 2010 | 40          | Group 1: MTX + SSZ + HCQ Group 2: infliximab + MTX | Group 1: 59 Group 2: 560 | Group 1: 20 mg/week MTX + 2000 mg/day SSZ + 400 mg/day HCQ Group 2: 20 mg/week MTX + 3 mg/kg infliximab (weeks 0, 2, 6 and every 8 weeks) | Lean mass App. lean mass | DEXA             | Group 1: 4.3 Group 2: 4.8 | NI          |

BIA bioimpedance, DEXA dual-energy X-ray absorptiometry, NI not informed, TNF tumor necrosis factor, BC body composition, M male, F female, MTX methotrexate, SSZ sulfasalazine, HCQ hydroxychloroquine
JAKi [34], rituximab [43], bDMARDs [44], etanercept [42], methotrexate [41, 42], sulfasalazine [41], and hydroxychloroquine [41].

Methods of assessment of the muscle mass and treatment effect
Three of nine studies (33%) used bioimpedance as a measurement method [34, 47, 48], while the other six (66%) used dual-energy X-ray absorptiometry (DEXA) [41–46]. Despite being different methods of assessing muscle mass, studies have shown that these methods have good validity and agreement [49, 50]. Toussirot et al. used lean mass, and the proposed treatment showed significant improvement in this parameter (3.3%) [46]. Tournadre et al. analyzed parameters lean mass and appendicular lean mass showing significant benefits in both parameters after one year of treatment with tocilizumab (2.6% in lean mass and 5.6% in appendicular lean mass) [45]. Ferraz-Amaro et al. (5.9%) and Metsios et al. (0.39%) used lean mass as a parameter, but both showed no significant improvement after anti-TNF treatment [47, 48]. Chikugo et al. used appendicular lean mass as a parameter and showed no significant changes in this parameter after JAKi treatment (0.49%) [34]. Al Khayyat et al. [43] used both lean mass and appendicular lean mass as parameters and showed a decrease (3.3%) in lean mass and an increase (10%) in appendicular lean mass. Vial et al. [44] used lean mass as a parameter and showed an improvement of lean mass in the TNFi group (2.2%) and a decrease (0.7%) of lean mass in the non-TNFi group.

Methodological quality of the studies
In the methodological quality assessment of the nine studies, eight studies [34, 41–45, 47, 48] were classified as good-quality studies, and one was classified as a poor-quality study [46]. Data were described in Table 3.

Risk of bias of studies
In the risk of bias analysis, two of the two studies [41, 42] were classified with a high risk of bias. Data was described in Table 4.

Meta-analysis of lean mass
Four out of seven observational studies performed lean mass measures [45–48]. About this outcome, we performed two different methods in our meta-analysis: a general analysis comparing the four studies, and a subgroup analysis comparing types of treatment. Two studies used tocilizumab as treatment, and the other two used anti-TNF therapy. Despite the lack of significant difference, in the general analysis, five [44–46, 48] of eight groups analyzed have shown a positive delta of lean mass, and three [43, 44, 47] groups have shown a negative delta. In general analysis, the treatment with DMARD was not able to increase lean mass in patients (mean = 0.47; 95% CI [−0.92 to 1.87]; \(I^2, 0\% p = 0.91\) (Fig. 2). In the subgroup analysis, tocilizumab treatment (mean = 1.32; 95%

### Table 3 Methodological quality of the studies

| Author            | Year | Cohort selection | Comparability | Outcome ascertainment | Total score | Quality        |
|-------------------|------|------------------|---------------|-----------------------|-------------|----------------|
| Chikugo et al.    | 2018 | -                | ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★         | 7           | Good quality  |
| Ferraz-Amaro et al. | 2011 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 8           | Good quality  |
| Metsios et al.    | 2007 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 7           | Good quality  |
| Tournadre et al.  | 2017 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 6           | Good quality  |
| Toussirot et al.  | 2020 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 5           | Poor quality  |
| Al Khayyat et al. | 2021 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 6           | Good quality  |
| Vial et al.       | 2021 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 8           | Good quality  |
| Engvall et al.    | 2010 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 7           | Good quality  |
| Marcara et al.    | 2006 | ★ ★ ★ ★ ★ ★ ★    | ★ ★ ★ ★ ★ ★ ★ | ★ ★ ★ ★ ★ ★ ★ ★★ | 6           | Good quality  |

### Table 4 Risk of bias analysis

| Study ID | Randomization process | Deviations from intended interventions | Missing outcome data | Measurement of the outcome | Selection of the reported result | Overall |
|----------|-----------------------|----------------------------------------|----------------------|---------------------------|---------------------------------|---------|
| Engvall et al. | High               | Some concerns                           | Low                  | Low                       | Low                            | High    |
| Marcara et al.   | High               | Some concerns                           | Low                  | Low                       | Low                            | High    |
Fig. 2 In general analysis, the treatment with DMARD was not able to increase lean mass in patients.
CI [−0.87 to 3.52]; $I^2$, 0%; $p = 0.95$) and TNFi treatment (mean = 0.21; 95% CI [−2.63 to 3.04]; $I^2$, 0%; $p = 0.47$) had positive mean (Fig. 3). In rituximab (mean = −1.30) and bDMARD nTNFi (mean = −0.30) treatment, the mean was negative.

**Meta-analysis of appendicular lean mass**

Regarding appendicular lean mass, three studies have measured this outcome. Still, one of these studies has performed a trial with two groups of treatment: one group treated with tofacitinib, and one group treated with other bDMARDs [34]. Regardless of the increase of mean appendicular lean mass, treatment with DMARD showed no significant change in appendicular lean mass delta (mean = 1.11, 95% CI [−0.58 to 2.79]; $I^2$, 0%; $p = 0.91$) (Fig. 4).

**Discussion**

As a result of this systematic review with meta-analysis, we found that DMARD treatment did not appear to induce significant muscle mass changes in RA patients. Still, regarding lean mass measurement, we described in subgroup analysis that anti-IL-6 and anti-TNF treatments were more related to the gain of lean mass than other DMARD therapies. Besides, considering the slight mass gain in both lean mass and appendicular lean mass and the small number of studies, we cannot exclude the possibility of a beneficial effect, particularly in anti-IL6 and anti-TNF therapy. This systematic review with meta-analysis is the first to verify the effect of DMARD treatment and its subclasses in muscle mass parameters.

Considering the muscle mass loss present in RA sarcopenia [10], DMARD treatment not only prevented this but also showed a trend of a slight gain of muscle mass in analyzed parameters. Dao et al. (2021) [51] investigated the associations between RA treatment and sarcopenia prevalence. Inherently, the authors showed that RA patients on csDMARD treatment had a lower prevalence of sarcopenia compared to RA patients on bDMARD. tsDMARD treatment had no association with sarcopenia. As we also saw in our review, Dao et al. emphasized the small number of papers in the literature and pointed out that it could be the reason for the lack of associations.

In our review, studies showed that IL-6 inhibition tended to be related to slight lean mass gain. IL-6 can bind the membrane IL-6 (IL-6R) and induce intracellular signals [52]. Still, IL-6 can also bind to soluble receptors (sIL-6R) creating a complex able to stimulate cells that do not have the membrane receptor [53]. Due to these mechanisms, the IL-6 effect in cells can be dualistic, being either inflammatory or anti-inflammatory [16]. Indeed, in an acute exercise setting, IL-6 secreted by muscle cells can drive muscle growth signaling, muscular regeneration, and activation of muscle stem cells [54]. On the other hand, chronic expression of IL-6 by inflammatory and immune cells is related to the induction of muscle atrophy and protein degradation [55, 56]. These effects occur by IL-6R binding leading to activation of the JAK/STAT complex [57] and signaling the increased expression of catabolic genes, such as muscle RING-finger protein-1 (MURF1), ubiquitin-proteasome subunits, caspases and cathepsins [14]. Thus, we consider that anti-IL6 therapy could have a positive effect on muscle mass in conditions of chronic inflammation based on its influence on important routes of inflammatory signaling and on its role as a locally secreted myokine [58]. Differently from other pro-inflammatory cytokines, which are mostly secreted by inflammatory cells and their action is generally systemic, IL-6 is secreted by muscle cells for paracrine communication leading to potent local signaling [59].

**Table 1**

| Study         | Mean   | MRAW   | 95% CI          | Weight |
|---------------|--------|--------|-----------------|--------|
| Tournadre 2017|        |        |                 |        |
| Chikug (t) 2018|        |        |                 |        |
| Chikug (d) 2018|        |        |                 |        |
| Al Khayyat 2021|        |        |                 |        |
| Tournadre 2017| 0.00   | [−5.45; 5.55] | 9.4%            |        |
| Chikug (t) 2018| 0.10   | [−4.43; 4.63] | 13.9%           |        |
| Chikug (d) 2018| 1.63   | [−0.74; 4.00] | 50.9%           |        |
| Al Khayyat 2021| 1.11   | [−0.58; 2.79] | 100.0%          |        |

Forest plot of the effect of DMARD treatment on appendicular lean mass mean (n=3 studies). Chikug (t): tofacitinib treated group; Chikug (d): other DMARD treated group. $I^2$: Heterogeneity of studies; MRAW: untransformed means by R software; SD: standard deviation; SMD: standardized mean difference; 95% CI: 95% confidence interval; IV: inverse variance; Random: random effects model.

**Fig. 3** Positive mean of tocilizumab treatment and TNFi treatment.
Fig. 4 Regardless of the increase of the mean appendicular lean mass, treatment with DMARD showed no significant change in appendicular lean mass delta.
TNF-α is another key factor in muscle impairment in RA [60]. TNF-α inhibition therapy also seemed to have a positive effect on lean mass in AR patients. At a molecular level, TNF-α is the main responsible for the NFκB activation pathway [61, 62], a transcript factor known to drive the subsequent expression of inflammatory mechanisms [63]. With the meta-analysis results, we speculate that despite its approved effect against RA disease activity, blocking systemic inflammation, anti-TNF treatment tended to have a local effect to block TNF downstream in the muscle being able to prevent AR muscle loss [64]. Interestingly, in both randomized clinical trials mentioned in our review, Marcora et al. and Engvall et al. did not present, in their results, significant change in both lean mass and appendicular lean mass, when patients were treated with anti-TNF drug [41, 42].

JAKi treatment, a more recent approach, has been demonstrated to be effective against RA inflammation [65]. The JAK/STAT pathway is known for acting together with cytokine receptors carrying the intracellular signal through the phosphorylation of STATs [57, 65, 66]. For example, JAK/STATs are attached to IL-6 membrane receptors and are responsible for activating the transcription of inflammatory genes [68]. In our review, we showed that JAKi treatment did not present a significant effect on appendicular lean mass. Still, its effect was similar to DMARD treatment performed in the same study [34]. We believe that JAKi analysis was limited by the lack of studies and the study sample size.

In this review, we used Newcastle Ottawa to describe the quality of each study included in our systematic review with meta-analysis. The majority of studies were identified with good quality. Finally, this systematic review with meta-analysis has some limitations. First, there were a small number of studies included. Furthermore, the studies included were performed by enrolling both male and female patients, and it is known that men have higher muscle mass than women.

We conclude that DMARDs have no effect on muscle mass parameters in rheumatoid arthritis patients. Indeed, we showed that DMARD treatment was not able to have a positive effect both in lean mass (total lean mass including trunk) and appendicular lean mass (lean mass of arms and legs only), results that coincide with clinical trials available in the literature. However, this review could be a path to better understanding the treatment of RA muscle loss, being the first to systematically analyze the literature about it. We believe that the limitations found in our review, such as the small number of studies and sample size, may have been relevant for not having found differences in our analyses. Emphasizing this is important to drive and induce researchers to develop investigations about it. In addition, the enlightenment of how DMARDs act in muscle mass is important for the formulation of treatment protocols that can treat not only autoimmune and inflammatory diseases but also muscle-wasting conditions such as sarcopenia and cachexia. Finally, by summarizing and qualifying the data about the relationship between DMARDs and muscle mass in RA, this systematic review is crucial to enlighten the evidence presented in the literature.

Conclusion
We conclude that this review was the first to summarize the data about the relationship between DMARDs and muscle mass. In addition, we have that DMARD treatment has no positive effect on rheumatoid arthritis muscle mass loss. With this review, we contribute to enlightenment in DMARD treatment in rheumatoid arthritis once it does not have any approved pharmacological therapy for comorbidities such as muscle loss.

Abbreviations
RA: Rheumatoid arthritis; MD: Mean difference; SMD: Standardized mean difference; CI: Confidence interval; DMARD: Disease-modifying anti-rheumatic drug; bDMARD: Biological disease-modifying anti-rheumatic drug; csDMARD: Conventional synthetic disease-modifying anti-rheumatic drug; TNF-α: Tumor necrosis factor-alpha; IL-6: Interleukin 6; sIL-6R: Soluble interleukin 6 receptor; IL-1β: Interleukin 1 beta; JAK: Janus kinase; IL-2: Interleukin 2; IL-7: Interleukin 7; DEXA: Dual-energy X-ray absorptiometry; BIA: Bioimpedance; Murf1: Muscle Ring-finger protein 1; TGF-β: Transforming growth factor-beta; JAK/STAT: Janus kinase/signal transducer and activator of transcription; MTX: Methotrexate.

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Authors’ contributions
T.H. participated in the design of the study, carried out the initial selection of the studies, participated in the data analysis, drafted the manuscript, performed the meta-analysis, and prepared all the figures. L.P. participated in the design of the study, carried out a full-length selection of the studies, and performed the methodological quality and risk of bias analysis. B.J. participated in the design of the study, carried out the initial selection of the studies, and participated in the data analysis. J.P. participated in the design of the study, carried out a full-length selection of the studies, and performed the methodological quality and risk of bias analysis. R.C. participated in the design of the study, helped in the selection of the studies, helped in the methodological quality and risk of bias analysis, helped in the meta-analysis, and helped in the preparation of the figures. R.M. participated in the design of the study, helped in the manuscript draft, and helped in the preparation of figures. The author(s) read and approved the final manuscript.

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Availability of data and materials
The authors declare that all data supporting the findings of this study are available within the article and its supplementary information files.
**Declarations**

**Ethics approval and consent to participate**

Where applicable, I confirm that all human and animal studies have been approved by the appropriate ethics committee and have therefore been performed in accordance with the ethical standards laid down in the 1964 Declaration of Helsinki and its later amendments. Where applicable, I confirm that all persons gave their informed consent prior to their inclusion in the study. Details that might disclose the identity of the subjects under study have been omitted.

**Consent for publication**

All authors give their consent to the publication of this study.

**Competing interests**

The authors declare that they have no competing interests.

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