

# Comparative efficacy and safety of bimekizumab in axial spondyloarthritis: a systematic literature review and network meta-analysis

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## Abstract

**Objectives:** To compare the efficacy and safety of bimekizumab 160mg every 4 weeks, a selective inhibitor of interleukin-17F and 17A, with biologic/targeted synthetic disease-modifying anti-rheumatic drugs (b/tsDMARDs) in non-radiographic axial spondyloarthritis (nr-axSpA) and ankylosing spondylitis (AS).

**Methods:** A systematic literature review identified randomised controlled trials until January 2023 for inclusion in Bayesian network meta-analyses (NMAs), including three b/tsDMARDs exposure networks: predominantly-naïve, naïve, and experienced. Outcomes were Assessment of SpondyloArthritis international Society (ASAS)20, ASAS40, and ASAS partial remission (PR) response rates at 12–16 weeks. A safety NMA investigated discontinuations due to any reason and serious adverse events at 12–16 weeks.

**Results:** The NMA included 36 trials. The predominantly-naïve network provided the most comprehensive results. In the predominantly-naïve nr-axSpA analysis, bimekizumab had significantly higher ASAS20 response rates vs secukinumab 150mg (with loading dose [LD]/without LD), and comparable response rates vs other active comparators. In the predominantly-naïve AS analysis, bimekizumab had significantly higher ASAS40 response rates vs secukinumab 150mg (without LD), significantly higher ASAS-PR response rates vs secukinumab 150mg (with LD), and comparable response rates vs other active comparators. Bimekizumab demonstrated similar safety to other b/tsDMARDs.

**Conclusion:** Across ASAS outcomes, bimekizumab was comparable with most b/tsDMARDs, including ixekizumab, TNF inhibitors and upadacitinib, and achieved higher response rates vs secukinumab for some ASAS outcomes in predominantly b/tsDMARD-naïve nr-axSpA and AS patients at 12–16 weeks. In a pooled axSpA network, bimekizumab demonstrated comparable safety vs other b/tsDMARDs.

## Keywords

Axial spondyloarthritis, Systematic literature review, Network meta-analysis, b/tsDMARDs, nr-axSpA, r-axSpA

## Key messages

- Bimekizumab achieves higher response rates versus secukinumab for some ASAS outcomes in nr-axSpA and AS.
- Bimekizumab is associated with similar response rates versus other b/tsDMARDs across ASAS outcomes.
- Bimekizumab demonstrates similar safety and tolerability to other b/tsDMARDs.

## Introduction

Axial spondyloarthritis (axSpA) is a chronic inflammatory disease that predominantly affects the axial skeleton (sacroiliac joints and spine) (1, 2). AxSpA comprises patients with evident radiographic damage to the sacroiliac joints (ankylosing spondylitis [AS], also known as radiographic axSpA [r-axSpA]) and those without definitive radiographic sacroiliitis (non-radiographic axSpA [nr-axSpA]) (3, 4). Age of disease onset is typically mid-twenties (5), with an estimated 10–40% of nr-axSpA patients progressing to AS over 2–10 years (6).

Historically, nr-axSpA emerged as a subclassification of axSpA; however, axSpA is now widely considered a single disease spectrum, encompassing both nr-axSpA and AS (7). AS and nr-axSpA share a similar clinical presentation and disease burden (8–10); both are associated with chronic back pain, fatigue, and morning stiffness, affecting mobility and the ability to perform daily activities (11, 12). Many patients with axSpA also have peripheral musculoskeletal manifestations, with peripheral arthritis and enthesitis most common (affecting an estimated 28–30% and 29–35% of patients, respectively) (13). Some patients also present with extra-musculoskeletal manifestations, including

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3 acute anterior uveitis, psoriasis, and inflammatory bowel disease (12, 14-16). As such, axSpA has a  
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5 considerable impact on quality-of-life (17-20).  
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8 The initial pharmacological treatment for axSpA is non-steroidal anti-inflammatory drugs (NSAIDs).  
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10 For patients with active disease and an inadequate response, intolerance, or contraindication to  
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12 NSAIDs, available therapies include biologic disease-modifying anti-rheumatic drugs (bDMARDs),  
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14 comprising tumour necrosis factor (TNF) inhibitors, interleukin (IL)-17A inhibitors, and targeted  
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16 synthetic disease-modifying anti-rheumatic drugs (tsDMARDs) like the recently approved Janus  
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18 kinase (JAK) inhibitors (21, 22).  
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22 Despite available treatments, many patients do not achieve a sufficient treatment response or partial  
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24 remission, and some lose their clinical response to treatment over time (23, 24). Furthermore, clinical  
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26 response to second-line bDMARDs is lower than in bDMARD-naïve patients (25); hence, there is a  
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28 considerable need for treatment options achieving deep and sustained responses via a novel  
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30 mechanism of action (23, 24).  
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33 Bimekizumab is a humanised monoclonal immunoglobulin G1 antibody that recently received  
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35 marketing authorization in EU and UK, and that selectively inhibits interleukin (IL)-17F in addition to  
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37 IL-17A. Interleukin-17A and IL-17F are pro-inflammatory cytokines and key mediators of  
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39 inflammation and new bone formation which leads to structural damage in axSpA (26-28). Unlike IL-  
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41 17A-specific inhibitors, bimekizumab enables neutralisation of IL-17F/F in addition to IL-17A/A and  
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43 IL-17A/F. Preclinical data demonstrate that dual blockade of IL-17A and IL-17F is required for  
44  
45 optimal inhibition of downstream inflammatory and tissue remodelling responses (30). In the Phase  
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47 III trial programme for axSpA, bimekizumab resulted in significant and rapid improvements in  
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49 efficacy outcomes vs placebo (BE MOBILE 1 [NCT03928704] and BE MOBILE 2 [NCT03928743])  
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51 (31).  
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55 The aim of this analysis was to establish the comparative efficacy, safety, and tolerability of  
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57 subcutaneous (SC) bimekizumab 160 mg every 4 weeks (Q4W) versus b/tsDMARDs in axSpA using  
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59 a systematic literature review (SLR) and Bayesian network meta-analyses (NMA). The current  
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3 analysis provides an up-to-date synthesis of available evidence, including the BE MOBILE studies  
4 (31), which were published since the completion of previous SLRs/NMAs in axSpA. Although TNF  
5 inhibitors were included in the analysis, the relative efficacy of these is already well-established (32,  
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7 33), so this NMA focuses on comparisons between recently approved IL-17A, IL-17A/F, and JAK  
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9 inhibitors.  
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## 16 **Methods**

### 17 **Systematic literature review**

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19 A clinical SLR was initiated in May 2012 and updated eight times, most recently on 10<sup>th</sup> January  
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21 2023, to identify randomised controlled trial (RCT) evidence assessing bimekizumab and relevant  
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23 b/tsDMARDs for the treatment of adult patients with AS or nr-axSpA with an inadequate response to,  
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25 intolerance of, or contraindication to NSAID therapy (see Supplementary Data S1 for dates of all SLR  
26  
27 updates, available at *Rheumatology* online). Studies were required to report outcome measurements  
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29 after a minimum of 12 weeks of follow-up, but before switch/cross-over or early escape (34, 35).  
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31 Eligible interventions comprised IL-17A inhibitors, IL-17A/F inhibitors (i.e. bimekizumab), TNF  
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33 inhibitors, and JAK inhibitors. Eligible comparators comprised any of the aforementioned  
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35 interventions, conventional DMARDs, NSAIDs, or placebo. The pre-specified population,  
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37 intervention, comparator, outcomes, and study design (PICOS) elements used to assess study  
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39 eligibility are presented in Supplementary Table S1, available at *Rheumatology* online.  
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45 The SLR was performed in accordance with best practice guidelines from the Cochrane  
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47 Collaboration, Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA), and  
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49 the Centre for Reviews and Dissemination (CRD) (36-38). The Ovid platform was used to search  
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51 Embase, MEDLINE, the Cochrane Central Register of Controlled Trials (CENTRAL), and the  
52  
53 Cochrane Database of Systematic Reviews (CDSR) on 10<sup>th</sup> January 2023. Electronic database search  
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55 strings were developed for Embase, then translated for the other databases to account for differences  
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57 in syntax and subject headings (Supplementary Table S2–S4, available at *Rheumatology* online).  
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59 Title/abstract screening and full-text screening were both performed by two independent reviewers.  
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3 Any conflicts regarding eligibility were resolved through discussion, and where necessary, arbitration  
4 was provided by a third reviewer. Hand-searching of conference proceedings, HTA submissions,  
5 clinical trial registries, and the reference lists of relevant SLRs/NMAs were used as supplementary  
6 measures to ensure all relevant studies were captured (Supplementary Data S2, available at  
7 *Rheumatology* online). Relevant unpublished clinical study reports for bimekizumab were also  
8 eligible for inclusion. Data extraction and risk of bias assessment were performed by one reviewer,  
9 with all datapoints and risk of bias judgments checked by a second independent reviewer. Risk of bias  
10 was assessed using the CRD 7-item checklist for RCTs, currently recommended by the National  
11 Institute for Health and Care Excellence (NICE) (39).

### 22 23 **NMA feasibility assessment**

24 Additional eligibility criteria (Supplementary Table S5, available at *Rheumatology* online) were  
25 applied to identify studies suitable for inclusion in the NMA. Licensed b/tsDMARDs were the  
26 comparators of primary interest. Reasons for exclusion from the NMA included atypical axSpA  
27 classification criteria, biosimilar studies, and treatments with limited licensing or for which the  
28 development programme has been terminated. While secukinumab can be increased from 150 mg to  
29 300 mg in clinical practice, the 300 mg dose could not be included in the NMA due to insufficient  
30 trial data on the approved SC 300 mg dose (40). Although the MEASURE 3 trial reports data for  
31 secukinumab 300 mg SC Q4W at week 16, loading was by intravenous (IV) infusion, which is  
32 currently not approved.

### 43 44 **Network meta-analysis**

45 For efficacy outcomes, separate Bayesian NMAs were performed for patients with nr-axSpA and AS.  
46 Analyses were performed for three subpopulations, defined by patients' prior b/tsDMARD exposure:

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52 • **Predominantly (>50%) b/tsDMARD-naïve network:** Studies where >50% of the enrolled  
53 patients were b/tsDMARD-naïve or where it can be assumed that >50% of patients were  
54 b/tsDMARD-naïve.
- 55  
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57 • **100% b/tsDMARD-naïve network:** Studies where either 100% of the enrolled patients were  
58 b/tsDMARD-naïve or where studies reported separate data for a b/tsDMARD-naïve subgroup.  
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- **100% b/tsDMARD-experienced network:** Studies where either 100% of the enrolled patients were b/tsDMARD-experienced or where studies reported separate data for a b/tsDMARD-experienced subgroup.

Based on the timepoint at which the included studies reported primary and secondary efficacy results, the timepoint used for the NMA was 12–16 weeks; for inclusion in the NMA, studies had to report outcomes after a minimum of 12 weeks of follow-up (41), but before switch/cross-over or early escape. ASAS-defined improvement criteria (used in clinical trials) of interest for the efficacy NMA were ASAS20, ASAS40, and ASAS-PR (Table 1).

Two tolerability and safety outcomes at week 12-16 were also analysed: discontinuation due to any reason, and serious adverse events (SAEs). These analyses were conducted in a combined nr-axSpA and AS population irrespective of previous TNF exposure, due to the small number of patients experiencing events, and because patient characteristics and dose exposure of treatments were similar across the two indications.

Bimekizumab was compared to both placebo and active comparators. Doses for bDMARD comparators included in the NMA were:

- IL-17A inhibitors: ixekizumab 80 mg Q4W SC, secukinumab 150 mg Q4W SC (some patients may first receive initial loading doses (LDs) of secukinumab 150 mg SC at Weeks 0, 1, 2, 3 and 4 followed by Q4W thereafter; herein referred to as ‘with LD’ or ‘without LD’, respectively).
- IL-17A/F inhibitors: bimekizumab 160 mg Q4W.
- TNF inhibitors: adalimumab 40 mg twice-weekly (Q2W) SC, certolizumab pegol 200 mg Q2W or 400 mg Q4W SC, etanercept 25 mg BIW or 50 mg QW SC, golimumab 50 mg Q4W SC or 2mg/kg Q8W IV, infliximab 5 mg IV Q6W. TNF inhibitors were pooled as a single treatment class because the relative efficacy of TNF inhibitors in axSpA is already well-

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3 established (42, 43). Moreover, HTA publications, such as NICE TA383, conclude that TNF  
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5 inhibitors should be considered as a single class with broadly similar effects (44).  
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8 Doses for tsDMARD comparators were:  
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11 • JAK inhibitors: upadacitinib 15 mg once-daily (QD) oral, tofacitinib 5 mg twice-daily (BID)  
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13 oral (AS only).  
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## 16 17 18 **Statistical analysis**

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20 A Bayesian framework was chosen for the NMAs as Bayesian analysis is a standard approach that has  
21  
22 been extensively used by researchers due to the fact that, as described in the NICE Decision Support  
23  
24 Unit (DSU) guidance, “simulation from a Bayesian posterior distribution supplies both statistical  
25  
26 estimation and inference, and a platform for probabilistic decision making under uncertainty” (45).  
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29 The NMAs were conducted using standard methods for clinical data synthesis in WinBUGs using  
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31 validated model code for binomial outcomes, using a binomial model with logit link, available from  
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33 the NICE DSU (45-49, 51, 52, 54, 55). The WinBUGs models were run for a minimum burn-in of  
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35 10,000 iterations to maximise convergence. Subsequently, three chains of at least 1,000 samples  
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37 (3,000 simulations) were drawn from the posterior distributions. Both random and fixed effect, and  
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39 unadjusted and placebo-adjusted models were fitted to the data, with the mean residual deviance and  
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41 the deviance information criteria (DIC) used to estimate how well the predicted values fitted the  
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43 observed dataset. An alternative form of the DIC was also estimated (total residual deviance +  
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45 posterior variance) as for some analyses, the number of effective parameters was estimated to be  
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47 negative (56). However, this parameter did not always differentiate between models and other factors,  
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49 such as poor convergence or unrealistically large credible intervals (CrIs), would eliminate a  
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51 particular model. Fixed-effect (nr-axSpA; combined axSpA) and fixed effect placebo-adjusted (AS)  
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53 models offered preferred model fit. Results are expressed as odds ratios (ORs). Significant differences  
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55 between treatments were based on the 95% CrI for the OR crossing 1. Surface under the cumulative  
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57 ranking curve (SUCRA) values were calculated for each treatment, with higher values representing  
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3 higher ranked treatments (57). Note that SUCRA values should not be considered in isolation but  
4 interpreted alongside OR point estimates and CrIs (57).  
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## 10 **Results**

### 11 **Systematic literature review**

12 Overall, 341 publications reporting on 65 unique trials were included in the SLR (Figure 1 and  
13 Supplementary Figures S1 and S2, available at *Rheumatology* online); the feasibility assessment  
14 determined that 36 trials were suitable for inclusion in this NMA, comprising 10 in nr-axSpA, and 27  
15 in AS (note one reported separate data for both populations [RAPID-axSpA (10)]). Baseline patient  
16 and disease characteristics of the included studies are provided in Supplementary Tables S6–S9,  
17 available at *Rheumatology* online. Reasons for exclusion of the 29 remaining trials are provided in  
18 Supplementary Table S10, available at *Rheumatology* online. Trials were broadly similar in terms of  
19 their main baseline characteristics (where reported). Enrolment of patients with nr-axSpA was based  
20 on meeting the 2009 ASAS axSpA classification criteria (58), with the presence of objective signs of  
21 inflammation defined by bone marrow edema on magnetic resonance imaging (MRI) and/or elevated  
22 C-reactive protein (CRP). For most AS trials (24 of 27) enrolment of patients was based on the 1984  
23 modified New York (mNY) criteria (59). However, the entry criteria for three studies (COAST-V,  
24 COAST-W, and Xue 2022) restricted patients to those meeting the more recent ASAS classification  
25 criteria (58), which uses the imaging criterion of the mNY criteria for radiographic axSpA plus  
26 additional criteria (>1 feature of axSpA). Agreement between the mNY and ASAS r-axSpA criteria is  
27 reported to be very high (4) and these populations are likely to have considerable clinical overlap,  
28 despite differences in classification. In BE MOBILE 2, patients were enrolled with mNY criteria, and  
29 met ASAS criteria for the classification of r-axSpA. It was therefore assumed that the trial populations  
30 were sufficiently similar to be directly compared in the NMA.  
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54 The risk of bias for individual trials included in the NMA is provided in Supplementary Table S11,  
55 available at *Rheumatology* online. Overall, the included RCTs had a low risk of bias, with some  
56 elements of the assessment ranking unclear due to missing reporting. One area of weakness was that a  
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3 small number of baseline characteristics differed across the randomised treatment groups within five  
4 AS trials and one nr-axSpA trial (60–65). However, no studies were deemed unsuitable for inclusion  
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6 in the NMA based on concerns regarding risk of bias.  
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10 The network diagram for the predominantly b/tsDMARD-naïve networks (in nr-axSpA and AS) are  
11 presented in Figure 2. The 100% b/tsDMARD-naïve, experienced (for efficacy) and combined axSpA  
12 population (tolerability and safety) networks are provided in Supplementary Figure S3–S5, available  
13  
14 at *Rheumatology* online.  
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## 20 **Network meta-analysis**

### 21 ***Predominantly (>50%) b/tsDMARD-naïve network***

22 The predominantly b/tsDMARD-naïve network provided the most comprehensive results across  
23 outcomes and comparators comprising all 10 nr-axSpA studies and 25 out of 27 AS studies. Across  
24 the network, approximately 90% of patients were b/tsDMARD-naïve (67%–100% naïve in nr-axSpA  
25 and 61%–100% naïve in AS).  
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### 33 ***ASAS20***

34 Ten nr-axSpA studies and 24 AS studies reported ASAS20 (Supplementary Table S12 and S13,  
35 available at *Rheumatology* online). In nr-axSpA, bimekizumab was associated with significantly  
36 higher ASAS20 versus secukinumab 150 mg Q4W without LD (OR 2.18, 95% CrI: 1.13, 4.24) and  
37 secukinumab 150 mg Q4W with LD (OR 2.31, 95% CrI: 1.20, 4.53). Bimekizumab was comparable  
38 with all other active comparators; no further significant differences between bimekizumab and active  
39 comparators were observed. In AS, bimekizumab was associated with similar ASAS20 response rates  
40 compared with other active treatments in AS (Figure 3). Predicted probabilities for ASAS20 response  
41 by treatment are presented in Figure 4 and relative risk (RR) estimates for bimekizumab compared  
42 with other b/tsDMARDs are presented in Supplementary Table S14, available at *Rheumatology*  
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60 online.

**ASAS40**

Ten nr-axSpA studies and 22 AS studies reported ASAS40 (Supplementary Table S12 and S13). In nr-axSpA, bimekizumab was associated with similar (95% CrI crosses 1.0) ASAS40 response rates compared with other active treatments. In AS, ASAS40 response rate was significantly higher with bimekizumab in AS versus secukinumab 150 mg Q4W without LD (OR 1.60, 95%CrI: 1.01, 2.60). No other significant differences between bimekizumab and active comparators were observed (Figure 3). Predicted probabilities for ASAS40 response by treatment are presented in Figure 4 and RR estimates for bimekizumab compared with other b/tsDMARDs are presented in Supplementary Table S14.

**ASAS-PR**

Nine nr-axSpA studies and 16 AS studies reported ASAS-PR (Supplementary Table S12 and S13). In nr-axSpA, bimekizumab was associated with similar ASAS-PR response rates compared with other active treatments. In AS, bimekizumab was associated with significantly higher ASAS-PR compared with secukinumab 150 mg with LD (OR 1.65, 95% CrI: 1.08, 2.51) and similar ASAS-PR response rates compared with the other active treatments in AS (Figure 3). Predicted probabilities for ASAS-PR response by treatment are presented in Figure 4 and RR estimates for bimekizumab compared with other b/tsDMARDs are presented in Supplementary Table S14.

No other significant differences between bimekizumab and active comparators were observed in the predominantly b/tsDMARD-naïve network. League tables of pairwise comparisons for all treatments in the predominantly b/tsDMARD-naïve network are presented in Supplementary Tables S15–S20, available at *Rheumatology* online.

**100% b/tsDMARD-naïve network**

Results of the 100% b/tsDMARD-naïve network were broadly consistent with results of the predominantly b/tsDMARD-naïve network; no significant differences between bimekizumab and active comparators were observed for any outcome (Supplementary Table S21, available at *Rheumatology* online).

### ***100% b/tsDMARD-experienced network***

Two AS studies enrolled 100% b/tsDMARD-experienced patients (COAST-W and SELECT-AXIS 2 [Study 1]) and a further seven AS studies reported data for the subgroup of b/tsDMARD-experienced patients enrolled in the trial (Supplementary Table S13). These nine AS studies were included in the 100% b/tsDMARD-experienced analysis for ASAS20 and ASAS40. Seven of these studies also reported ASAS-PR for this subgroup; however, the analysis did not converge as there were too few patients in the subgroup and zero events in some of the placebo-control arms. No significant differences between bimekizumab and active comparators were observed for any outcome (Table 2). A b/tsDMARD-experienced network was not feasible in nr-axSpA as too few studies reported data for b/tsDMARD-experienced patients.

### ***Combined axSpA population safety network***

The combined axSpA population network for discontinuation due to any reason and for SAEs included 25 and 24 studies, respectively (Supplementary Table S22, available at *Rheumatology* online). Bimekizumab demonstrated comparable discontinuations due to any reason and comparable SAEs, relative to all active treatments in the network (Supplementary Table S23 and S24, available at *Rheumatology* online).

## **Discussion**

This SLR and NMA provides an up-to-date synthesis of available evidence to determine the relative efficacy, tolerability, and safety of bimekizumab compared with b/tsDMARDs in adult patients with nr-axSpA or AS with an inadequate response to, intolerance of, or contraindication to NSAID therapy. Separate analyses were undertaken across three ASAS efficacy outcomes in the following patient subpopulations: predominantly (>50%) b/tsDMARD-naïve, 100% b/tsDMARD-naïve, and 100% b/tsDMARD-experienced. Tolerability and safety analyses were conducted in a combined axSpA population.

Prior to the approval of IL-17A and JAK inhibitors for axSpA, treatment options were relatively limited, with TNF inhibitors being the only available targeted therapies (21, 22). Today, a wider range

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3 of treatments are available. The comparative efficacy of available b/tsDMARDs is of great interest to  
4 patients, clinicians, and payers alike; it is important to understand the relative efficacy of available  
5 therapies to determine best practices. Several placebo-controlled RCTs demonstrate the efficacy of  
6 recently available therapies, but head-to-head trials are lacking, and so NMA can assist in assessing  
7 their comparative effectiveness (66). Whilst previous NMAs in axSpA evaluated the relative efficacy  
8 of some therapies (42, 67-75), this is the first analysis that incorporates upadacitinib and  
9 bimekizumab.

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19 The predominantly bDMARD-naïve network provided the most complete set of results across  
20 outcomes and comparators. In nr-axSpA, bimekizumab was associated with significantly higher  
21 ASAS20 response rates vs secukinumab 150 mg (in both the with and without LD comparisons), and  
22 in AS, ASAS40 and ASAS-PR response rates were significantly higher with bimekizumab vs  
23 secukinumab 150 mg without LD and secukinumab 150 mg with LD, respectively. Besides these  
24 comparisons, no significant differences were found; bimekizumab was associated with similar  
25 response rates compared with all other b/tsDMARDs.

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35 Inclusion of recent Phase III trials of bimekizumab (BE MOBILE 1 and BE MOBILE 2 (76, 77))  
36 enabled analysis of 100% b/tsDMARD-naïve and experienced subpopulations. Broadly, conclusions  
37 of the 100% b/tsDMARD-naïve network were consistent with those of the predominantly  
38 b/tsDMARD-naïve network, so the latter can be used as a proxy for the former (78). Importantly, the  
39 predominantly-naïve network used a more robust dataset (intention-to-treat/full analysis set) than an  
40 analysis using subgroup data, allowing more comparators to be included in the networks. In the 100%  
41 b/tsDMARD-experienced network, analyses were possible in AS for ASAS20 and ASAS40, and no  
42 significant differences between bimekizumab and active comparators were observed. These  
43 b/tsDMARD-experienced analyses are a novel addition to the literature, albeit with low trial and  
44 patient numbers, and should be interpreted with caution. Additionally, the safety and tolerability of  
45 bimekizumab at week 12-16 was comparable to all active comparators in the combined axSpA  
46 population.

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3 To the best of our knowledge, this research represents the most recent and comprehensive SLR/NMA  
4 for axSpA, building upon previous meta-analyses in b/tsDMARD-naïve patients (44, 69, 73-77). A  
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6 2018 NMA published by Deodhar et al compared the efficacy of TNF, IL-17, and JAK inhibitors  
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8 (tofacitinib only) in AS . The authors concluded that tofacitinib, golimumab IV, and infliximab had  
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10 the highest SUCRA values for efficacy; however, differences in efficacy were not significantly  
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12 different and analyses were based on one small Phase II study for tofacitinib . The current analysis  
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14 includes Phase III trials for bimekizumab (58) and JAK inhibitors (78-80) which were not available at  
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16 the time of the previous NMA, as well as data for nr-axSpA. The present analysis also includes new  
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18 outcomes relative to the Deodhar study, including ASAS40, ASAS-PR, and safety/tolerability  
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20 outcomes. For ASAS20, the only outcome included in both studies, results in the predominantly-naïve  
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22 network were consistent with those previously published, with bimekizumab now featuring amongst  
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24 the most efficacious treatments.  
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### 29 ***Limitations***

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31 Whilst some baseline characteristics differed between studies, no studies were deemed unsuitable for  
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33 inclusion in the NMA based on differences in baseline characteristics, and the overall risk of bias was  
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35 low. However, for 100% b/tsDMARD-naïve and experienced analyses, some evidence comes from  
36  
37 trial subgroup data for which baseline characteristics are not available; this introduces uncertainty  
38  
39 regarding how balanced the study arms are. Results are based on fixed-effect (nr-axSpA; combined  
40  
41 axSpA) and fixed-effect placebo-adjusted (AS) models, which may underestimate uncertainty in the  
42  
43 treatment effects. However, these results estimated some wide 95% CrIs for the relative treatment  
44  
45 effects, even using the fixed-effect models, and thus additional clinical study data would be beneficial  
46  
47 to reduce the uncertainty in the findings. Any additional study data would also help to enable a more  
48  
49 rigorous assessment of between-study heterogeneity and placebo-effects adjustment. There is a  
50  
51 paucity of RCT data for b/tsDMARD-experienced patients, and this network was not feasible in nr-  
52  
53 axSpA. Nine studies reported data for this subgroup subpopulation in AS, leading to a network of  
54  
55 eight treatments (including placebo). For the b/tsDMARD-experienced network, the comparisons  
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57 against bimekizumab were based on subgroup data from BE MOBILE 2; however, trial randomisation  
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3 was not designed to enrol a sufficient number of bDMARD-experienced patients to detect a difference  
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5 between bimekizumab and placebo in this subgroup . The placebo arm of BE MOBILE 2, that  
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7 connects to the rest of the network, contained just 17 patients, and so the analysis is subject to  
8  
9 uncertainty. The safety analyses are also associated with increased uncertainty due to the small  
10  
11 number of events. Further limitations of the NMA include different ages of included studies (2002  
12  
13 [infliximab] to 2022 [ixekizumab] (81, 82)), short-term efficacy analyses (12–16 weeks), and the lack  
14  
15 of published data on 300 mg secukinumab, which prevented inclusion of this higher secukinumab  
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17 dose in the NMA.  
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## 23 **Conclusion**

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25 Across ASAS outcomes, bimekizumab demonstrated comparable efficacy with most b/tsDMARDs,  
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27 including ixekizumab, TNF inhibitors and upadacitinib, and achieved higher response rates compared  
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29 with secukinumab at 12–16 weeks for ASAS20 in predominantly b/tsDMARD-naïve nr-axSpA  
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31 patients, and for ASAS40 and ASAS-PR in predominantly b/tsDMARD-naïve AS patients. In a  
32  
33 pooled axSpA network, bimekizumab demonstrated comparable safety compared with other  
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35 b/tsDMARDs. Overall, the present analyses provide evidence for bimekizumab being an efficacious  
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37 option in the management of both b/tsDMARD-naïve and experienced patients across the axSpA  
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39 spectrum, with similar safety and tolerability to existing treatments.  
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3 **Conflict of Interest:** D. Pritchett and M. Orme are employees of Source Health Economics and  
4 ICERA Consulting Ltd, respectively, the consultancy companies that conducted the systematic  
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6 Willems are employees of UCB Pharma. D. Willems and V. Taieb are shareholders in UCB Pharma.  
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33 are available from the corresponding author on reasonable request.  
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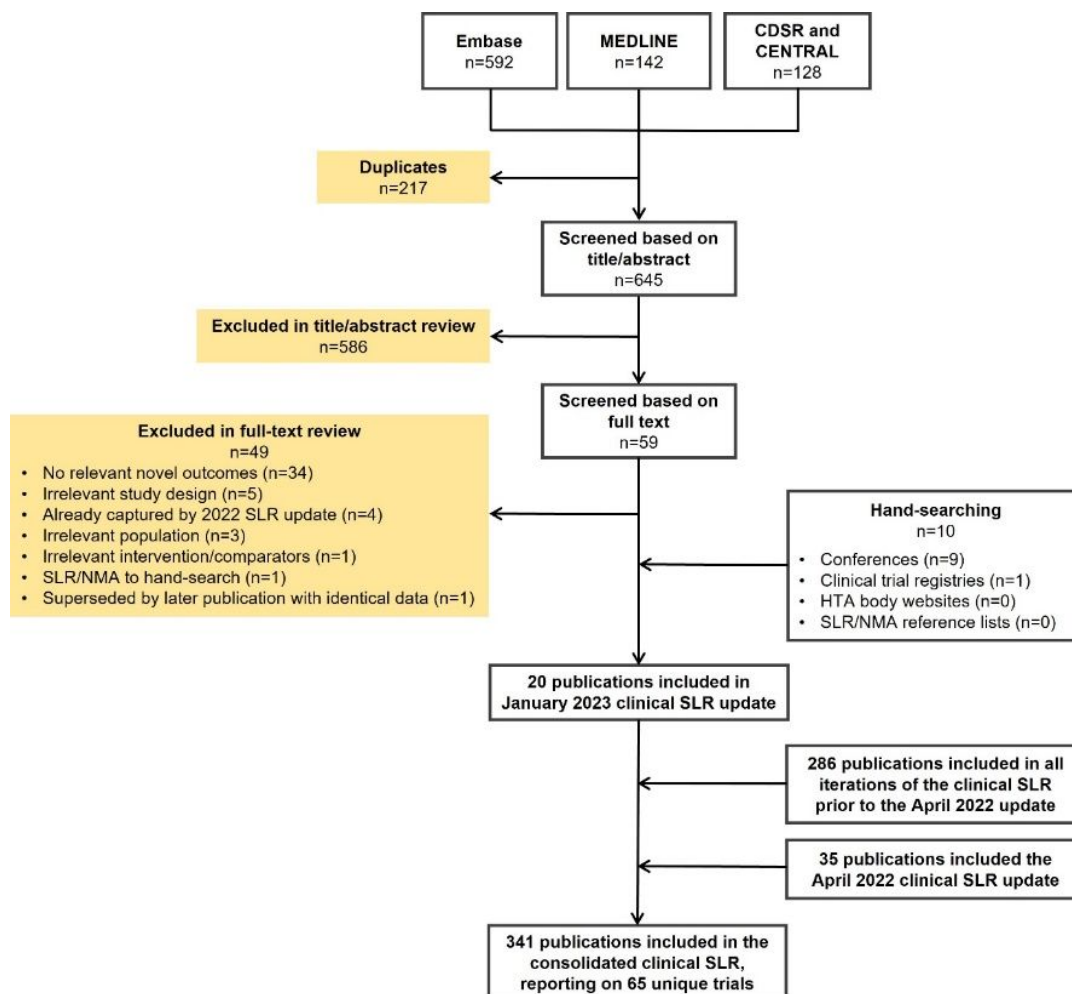
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## Tables and figures

**Table 1: Outcomes included in the NMA**

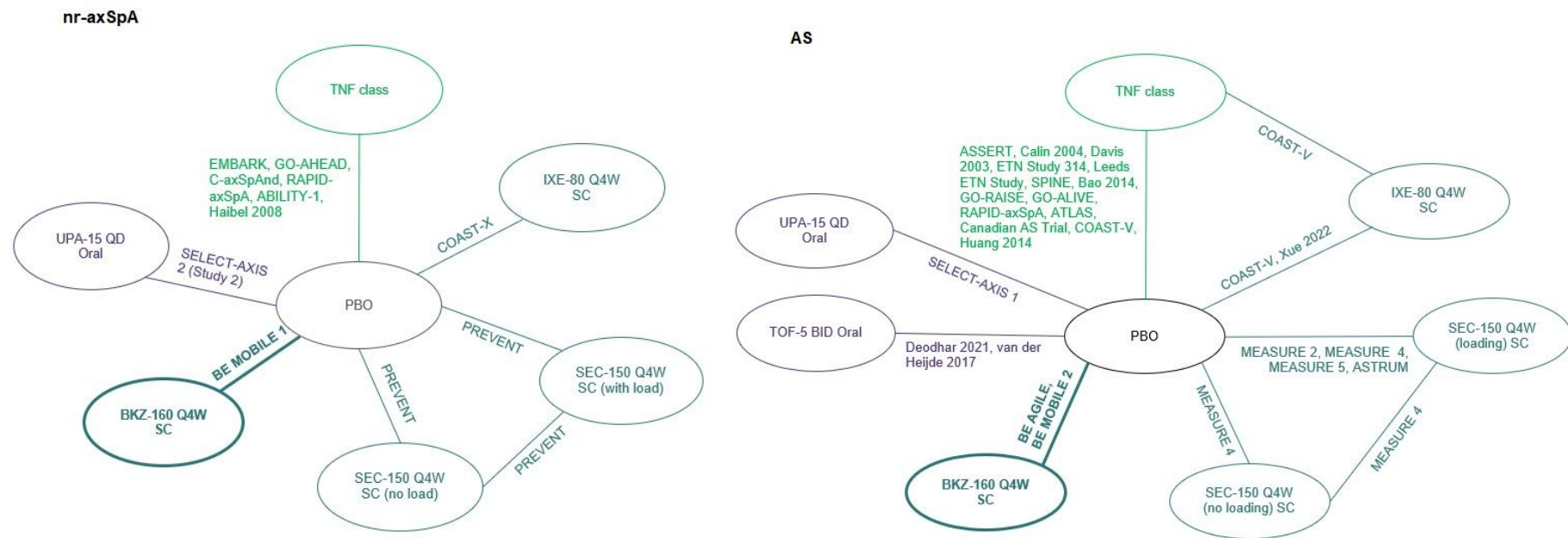
Outcome	Definition
ASAS20 and ASAS40	<p>ASAS20 and ASAS40 represent a relative improvement from baseline of <math>\geq 20\%</math> and <math>\geq 40\%</math>, respectively, together with a predefined absolute improvement from baseline in three or more of the following domains:</p> <ul style="list-style-type: none"> <li>• Patient global assessment</li> <li>• Spinal pain (BASDAI question 2)</li> <li>• Function (BASFI)</li> <li>• Inflammation (mean of BASDAI questions 5 and 6 on morning stiffness)</li> </ul> <p>ASAS20 and ASAS40 are benchmarks for measuring symptomatic improvement in patient disease status in clinical trials, and are the primary outcomes of interest in many comparator trials in the NMA .</p>
ASAS-PR	<p>ASAS-PR is defined as <math>&lt; 2</math> on a scale of 0–10 in each of the four ASAS domains :</p> <ul style="list-style-type: none"> <li>• Patient global assessment</li> <li>• Spinal pain (BASDAI question 2)</li> <li>• Function (BASFI)</li> <li>• Inflammation (mean of BASDAI questions 5 and 6 on morning stiffness)</li> </ul> <p>ASAS-PR corresponds to a state of partial remission/very low levels of disease activity and is also useful for cross-trial comparisons (86, 87). It is one of the most stringent clinical endpoints for axSpA; despite currently available treatments, many patients do not currently achieve a sufficient treatment response or very low levels of disease activity (23, 24, 83).</p>

Abbreviations: ASAS20, Assessment of Spondyloarthritis international Society-improvement of  $\geq 20\%$ ; ASAS40, Assessment of Spondyloarthritis international Society-improvement of  $\geq 40\%$ ; ASAS-PR, Assessment of Spondyloarthritis international Society partial remission; axSpA, axial spondyloarthritis; BASDAI, Bath Ankylosing Spondylitis Disease Activity index; BASFI, Bath Ankylosing Spondylitis Functional Index; NMA, network meta-analysis.

**Figure 1: PRISMA flow diagram of SLR study selection**

Abbreviations: CDSR, Cochrane Database of Systematic Reviews; CENTRAL, Cochrane Central Register of Controlled Trials; HTA, health technology assessment; NMA, network meta-analysis; PRISMA, Preferred Reporting Items for Systematic Reviews and Meta-Analyses; SLR, systematic literature review.

Figure 2: Network diagrams in nr-axSpA and AS (predominantly b/tsDMARD-naïve network)<sup>a, b, c</sup>



a Presented network diagrams are all studies that are included in one or more of the predominantly b/tsDMARD-naïve analysis; b See Supplementary Table S12 and S13 for list of studies that report ASAS20, ASAS40 or ASAS-PR outcome; c Network diagrams for 100% b/tsDMARD-naïve and experienced networks are provided in Supplementary Figure S3–S5. Abbreviations: AS, ankylosing spondylitis; ASAS20, Assessment of Spondyloarthritis international Society-improvement of  $\geq 20\%$ ; ASAS40, Assessment of Spondyloarthritis international Society-improvement of  $\geq 40\%$ ; ASAS-PR, Assessment of Spondyloarthritis international Society partial remission; BID, twice-daily; BKZ, bimekizumab; b/tsDMARD, biologic/targeted synthetic disease-modifying anti-rheumatic drugs; IXE, ixekizumab; nr-axSpA, non-radiographic axial spondyloarthritis; PBO, placebo; Q4W, every 4 weeks; QD, once-daily; SC, subcutaneous; SEC, secukinumab; TNF, tumour necrosis factor; TOF, tofacitinib; UPA, upadacitinib.



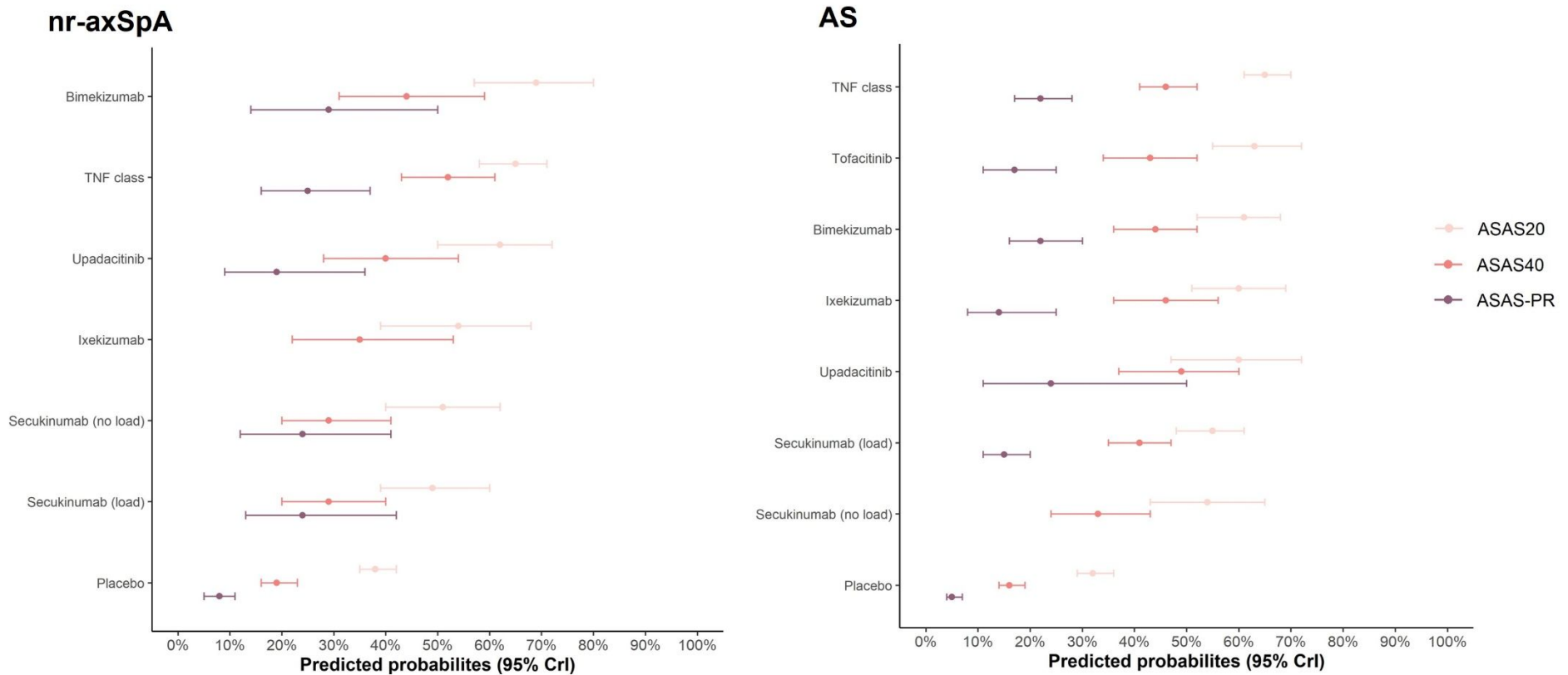
**Figure 3: ASAS20, ASAS40, and ASAS-PR outcomes in predominantly b/tsDMARD-naïve patients with nr-axSpA and AS<sup>a</sup>**



<sup>a</sup> Results expressed as ORs; higher ORs indicate better outcomes for bimekizumab. Bold denotes significance based on 95% CrI.

Abbreviations: ASAS, Assessment in Spondyloarthritis International Society; AS, ankylosing spondylitis; b/tsDMARD, biologic/targeted synthetic disease-modifying anti-rheumatic drugs; CrI, credible interval; nr-axSpA, non-radiographic axial spondyloarthritis; OR, odds ratio; PR, partial remission; SUCRA, surface under the cumulative ranking curve; TNF, tumour necrosis factor.

**Figure 4: Predicted probabilities of response in nr-axSpA and AS (predominantly b/tsDMARD-naïve network), ranked by ASAS20**



Abbreviations: AS, ankylosing spondylitis; ASAS, Assessment in Spondyloarthritis International Society; b/tsDMARD, biologic/targeted synthetic disease-modifying anti-rheumatic drugs; CrI, credible interval; IV, intravenous; nr-axSpA, non-radiographic axial spondyloarthritis; PR, partial remission; SC, subcutaneous; TNF, tumour necrosis factor.

**Table 2: NMA results in 100% b/tsDMARD-experienced network in AS only, bimekizumab vs comparator <sup>a, b, c</sup>**

Treatment	ASAS20		ASAS40		ASAS-PR	
	OR (95% CrI)	SUCRA	OR (95% CrI)	SUCRA	OR (95% CrI)	SUCRA
Bimekizumab	Reference	66%	Reference	65%	Analysis did not converge - too few b/tsDMARD-experienced patients/zero events in control arm	
TNF inhibitor class <sup>d</sup>	1.30 (0.11, 6.50)	52%	0.70 (0.17, 2.35)	83%		
Upadacitinib	0.82 (0.28, 2.95)	81%	0.85 (0.40, 1.78)	78%		
Tofacitinib	1.85 (0.07, 11.39)	35%	1.71 (0.50, 5.21)	35%		
Ixekizumab	1.50 (0.35, 4.75)	40%	1.94 (0.83, 4.43)	28%		
Secukinumab (load)	1.13 (0.36, 3.93)	58%	0.86 (0.39, 1.87)	77%		
Secukinumab (no load)	1.05 (0.28, 4.51)	62%	1.87 (0.65, 5.30)	31%		
Placebo	2.91 (0.93, 8.51)	6%	<b>3.46</b> <b>(1.67, 6.88)</b>	2%		

<sup>a</sup> Results expressed as ORs; higher ORs indicate better outcomes for bimekizumab. Bold denotes significance based on 95% CrI; <sup>b</sup> Network not feasible for nr-axSpA; <sup>c</sup> NMA based on 3 chains of 1,000 simulations; <sup>d</sup> Includes certolizumab pegol only.

Abbreviations: AS, ankylosing spondylitis; ASAS20, Assessment of Spondyloarthritis international Society-improvement of ≥20%; ASAS40, Assessment of Spondyloarthritis international Society-improvement of ≥40%; ASAS-PR, Assessment of Spondyloarthritis international Society partial remission; b/tsDMARD, biologic/targeted synthetic disease-modifying anti-rheumatic drugs; CrI, credible interval; NMA, network meta-analysis; OR, odds ratio; SUCRA, surface under the cumulative ranking curve; TNF, tumour necrosis factor.