

# Assessing Comorbid Inflammatory Arthritis Conditions and Swollen to Tender Joint Count Ratios in a Real-World Systemic Lupus Erythematosus Cohort

Cristi Cavanaugh, MHS, Greg Donadio, MS, Kathryn Starzyk, ScM, Michael Behling, MPH, Gary Curhan, ScD, MD, Rich Gliklich, MD | OM1, Inc, Boston, MA, USA



## Background

- Joint swelling and tenderness are common in patients with systemic lupus erythematosus (SLE) and other inflammatory arthritis conditions such as rheumatoid arthritis (RA), psoriatic arthritis (PsA), and ankylosing spondylitis (AS).
- Swollen to tender joint count ratio (STR) is an index originally used in RA which assesses severity of disease activity based on 28 joint counts with a higher score indicating greater likelihood of treatment response [1].
- Given the prevalence of joint involvement in SLE, STR has potential application in this patient population.

## Objective

To assess the overlap of SLE and other inflammatory arthritis conditions in a real-world cohort and characterize patients based on joint involvement, as defined by STR.

## Methods

- The OM1 SLE Registry (OM1, Boston, MA) follows more than 37,000 SLE patients longitudinally with deep clinical data, including laboratory, patient-reported and disease activity information, and linked administrative claims, starting from 2013.
- Patients  $\geq 16$  years of age with swollen and tender joint counts based on 28 joints on the same encounter were included. Patient cohorts were defined as SLE+ (RA, PsA, or AS) vs. SLE alone.
- STRs were calculated by inserting 1 if the denominator was 0 [2]. Patients were categorized by first available STR as having low (STR  $< 0.5$ ), moderate ( $0.5 \leq \text{STR} \leq 1.0$ ), or high (STR  $> 1.0$ ) disease activity [3].
- Clinical characteristics were summarized by disease activity group and STR group. Definitions of SLE treatments were based on 2019 EULAR recommendations [4].

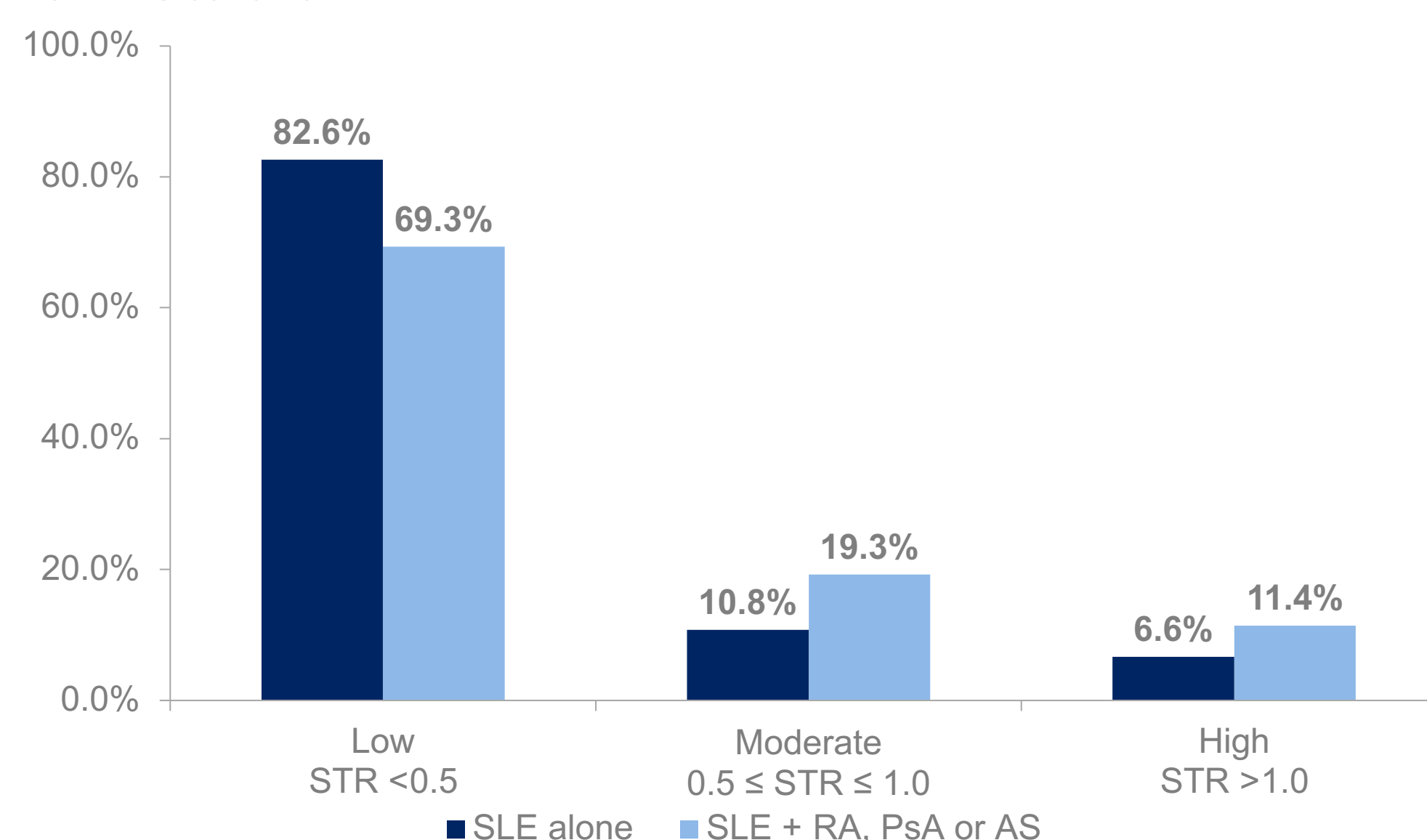
## Results

- The study included 9,919 patients with at least one STR available in the OM1 SLE Registry. Of the 17.0% of patients in the SLE+ cohort, 94.8% had RA, 6.7% had PsA, and 1.6% had AS. STR by cohort is displayed in **Figure 1**.
- Baseline characteristics by STR group for the full cohort are described in **Table 1**. When stratified by SLE cohort, SLE+ patients were older at 57.1 years (SD:13.1) compared to 51.1 (SD: 14.9) for SLE alone.
- Comorbid osteoarthritis was more common in the SLE+ cohort (46.3% vs. 26.0%).
- For SLE+, immunosuppressant use increased with higher STR from 44.5% (low STR) to 54.2% (high STR). For SLE alone, immunosuppressant use was lower overall (26.5%) with similar use among STR groups (**Figure 2**).
- Select disease-modifying antirheumatic drug (DMARD) use was higher among SLE+ patients (33.9% vs. 3.9%) with highest use in the moderate STR group (39.2%; **Figure 3**).
- Use of biologics (belimumab or rituximab) was higher among SLE+ (13.7% vs. 7.5%) with highest use in the low STR group (14.3%; **Figure 3**).

## Conclusions

- Higher STR and differences in treatment patterns were observed in the SLE+ cohort
- Additional research is needed to assess the relationship between STR and treatment response in SLE and relevant subphenotypes

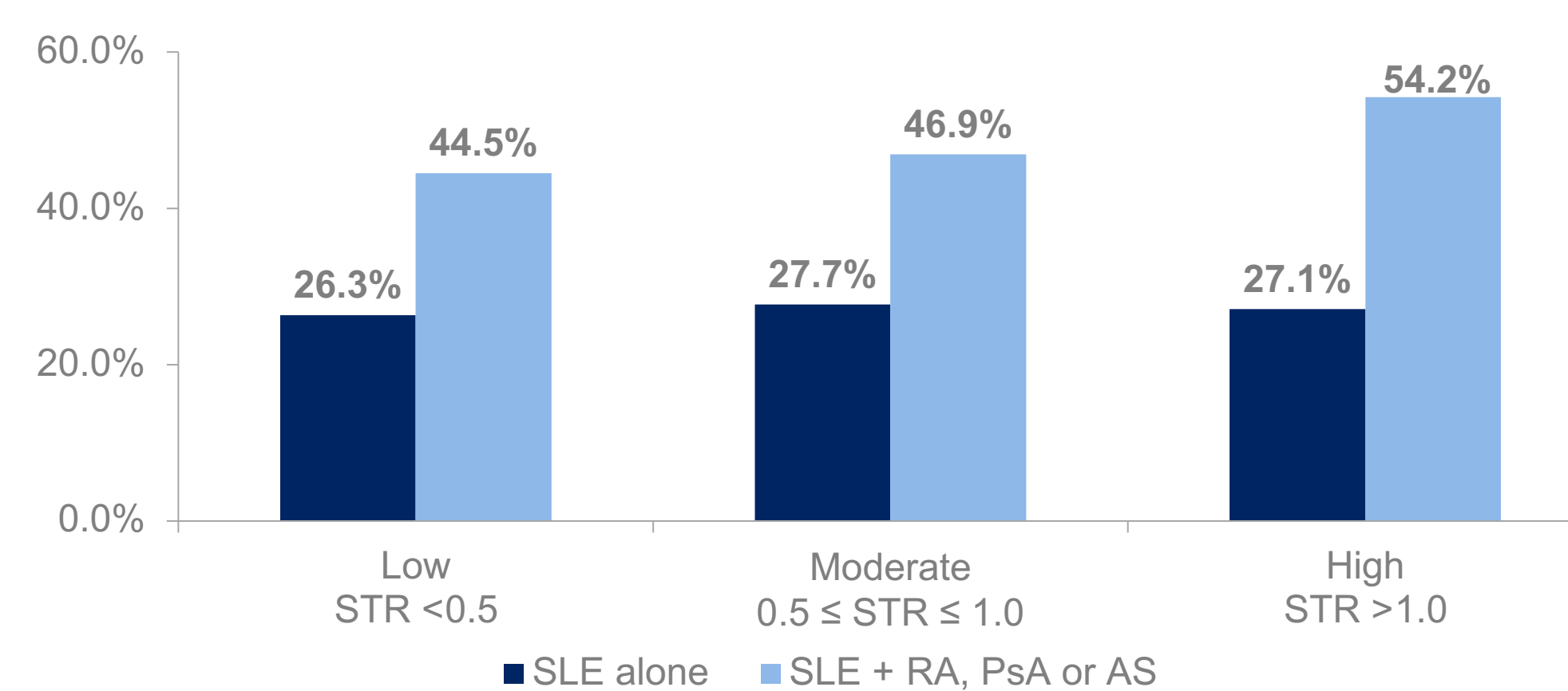
**Figure 1.** STR among patients with SLE alone or a comorbid inflammatory arthritis condition



**Table 1.** Baseline characteristics by STR group

	Low STR $\leq 0.5$ (N=7,970)	Moderate 0.5 $\leq$ STR $\leq 1.0$ (N=1,211)	High STR $> 1.0$ (N=738)
Age, mean (SD)	51.8 (14.9)	53.9 (14.3)	53.5 (14.9)
Female, n (%)	7,357 (92.3%)	1,103 (91.1%)	680 (92.1%)
<b>Race, n (%)</b>			
Black	1,546 (24.7%)	261 (28.3%)	128 (23.8%)
White	4,526 (72.2%)	636 (69.0%)	385 (71.7%)
Other	199 (3.2%)	25 (2.7%)	24 (4.5%)
Unknown	1,699	289	201
<b>Charlson Comorbidity Index, n (%)</b>			
0-1	4,693 (58.9%)	726 (60.0%)	446 (60.4%)
2-3	2328 (29.2%)	335 (27.7%)	206 (27.9%)
4-5	662 (8.3%)	87 (7.2%)	55 (7.5%)
$\geq 6$	287 (3.6%)	63 (5.2%)	31 (4.2%)

**Figure 2.** Prior immunosuppressant use by SLE cohort and STR group



**Figure 3.** Prior use of DMARDs and biologics by SLE cohort and STR group

