

A phase 2a trial of brepocitinib for cicatricial alopecia



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Background: Cicatricial alopecias are chronic, progressive scarring hair-loss conditions. Molecular dysregulation is not fully understood, hindering treatment development. Th1/IFN γ signaling and Janus kinase dysregulation has shown involvement, providing rationale for this phase 2a trial with Tyrosine kinase 2/Janus kinase 1 inhibitor brepocitinib.

Methods: Randomized, placebo-controlled phase 2a trial spanning 52 weeks. Adults (≥ 18 years of age) with lichen planopilaris, frontal fibrosing alopecia, or central centrifugal cicatricial alopecia diagnosis were randomized 3:1 to brepocitinib 45 mg daily or placebo for 24 weeks, after which all patients received brepocitinib for another 24 weeks, with a safety follow up 4 weeks later. Lesional scalp biopsies were collected at baseline, week 24, and week 48. Coprimary endpoints were changes in lesional expression of C-C motif chemokine ligand (CCL5), changes in lesional expression of fibrosis-related markers, and safety at week 24.

Results: Patients receiving brepocitinib showed significant downregulation in CCL5 expression at week 24 ($P = .004$). Enrichment analysis of a subset of fibrosis markers showed trending upregulation in placebo patients ($P < .1$). Brepocitinib was well tolerated and improved clinical severity scores.

Limitations: Single-dose regimen, small placebo group.

Conclusion: Brepocitinib significantly reduces CCL5 expression and was well tolerated at week 24, meeting coprimary endpoints. Brepocitinib reduces inflammatory biomarker expression and improves clinical severity, while maintaining favorable safety profile. (J Am Acad Dermatol 2025;92:427-34.)

Key words: central centrifugal cicatricial alopecia; cicatricial alopecia; fibrosis; frontal fibrosing alopecia; IFN γ ; JAK inhibitor; lichen planopilaris; phase 2a trial; scarring alopecia; systemic treatment; Th1.

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INTRODUCTION

Cicatricial alopecias (CA) are chronic, progressive hair loss disorders, encompassing frontal fibrosing alopecia (FFA), lichen planopilaris (LPP), and central centrifugal cicatricial alopecia (CCCA). CA is increasing in prevalence, representing about 7% of alopecia cases seen in specialized clinics.¹ FFA and LPP, while clinically distinct, share overlapping histopathological findings.² FFA primarily affects postmenopausal women causing frontal hairline and eyebrow loss.³ LPP affects all ages, causing inflammation and perifollicular hyperkeratosis, triggering scalp pruritus and dysesthesia.⁴⁻⁶ CCCA occurs in Afro-American women, causing scalp vertex shedding with associated tenderness and scale.^{7,8} Patients are often misdiagnosed and treated ineffectively, contributing to tremendous psychological burden.^{9,10} Commonly used treatments, such as topical/intralesional steroids and hydroxychloroquine, are used off-label and not well tolerated.¹¹⁻¹³ Currently, there are no Food and Drug Administration-approved treatments for CA, highlighting an unmet need.¹⁴

Histopathologic findings in CA have demonstrated underlying lymphocytic inflammation around the hair follicle, leading to perifollicular fibrosis; however, research on molecular dysregulation has been limited.¹⁵⁻¹⁷ Transcriptomic data across all 3 conditions has shown upregulation of fibrosis-related pathways.¹⁸ The pathologic fibrotic remodeling is likely due to upregulation of inflammatory cytokines, namely T helper (Th) 1/interferon (IFN) γ , and fibrosis-related markers.^{19,20} Our group has shown strong Th1 immune marker upregulation and pathway-level activation of Janus kinase (JAK) signal in FFA.²¹ Fibrosis-related genes also correlated with JAK/signal transducers and activators of transcription (STAT) signaling in lesional skin.²² JAK antagonism targets Th1/IFN γ cytokines by blocking STAT activation,²³ an effective approach in several other Th1-driven conditions, including alopecia areata.²⁴ Previous single patient reports and retrospective studies have shown promising outcomes for CA treatment with JAK inhibition, yet this research

was limited in scale and lacked molecular evidence.²⁵⁻²⁸ These findings provide the rationale behind this study, the first placebo-controlled clinical trial of a systemic treatment for CA.

Herein, we present the results from the first double-blind placebo-controlled single-center study evaluating the efficacy of brepocitinib, a tyrosine kinase (TYK) 2/JAK1 inhibitor, in the treatment of CA. Our mechanistic results measure changes in inflammatory and fibrosis biomarkers in lesional scalp, and our clinical results evaluate safety and clinical severity score changes.

CAPSULE SUMMARY

- Cicatricial alopecias are chronic scarring hair-loss conditions lacking effective treatments. Th1/IFN γ and Janus kinase dysregulation, which are involved in pathogenesis, may be targeted by brepocitinib.
- Brepocitinib is well tolerated with therapeutic potential for cicatricial alopecias, supported by improvement in clinical severity and expression of inflammatory pathways, including Th1/IFN γ and Janus kinase/signal transducers and activators of transcription.

METHODS

Study design and oversight

This single-center phase 2a, prospective, randomized, double-blind, placebo-controlled study (NCT05076006) evaluated the mechanistic and clinical efficacy and safety of brepocitinib for CA. The trial encompassed a 24-week double-blind treatment period and 24-week open-label period, with a safety follow up approximately 4 weeks after treatment cessation. Adults (≥ 18 years old) with clinically confirmed diagnosis of CCCA, LPP, or FFA and disease duration of 6 months to 7 years were eligible.

Patients with CCCA and LPP/FFA were randomized at baseline in a 3:1 ratio to receive either oral brepocitinib monotherapy 45 mg daily or placebo for 24 weeks. Medications related to CA were washed out before baseline (see Supplementary Appendix, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>). Investigators, personnel, and patients were blinded to trial group assignments. After the week 24 visit, all patients received brepocitinib 45 mg daily.

Study protocols were conducted in accordance with ethical principles of the Declaration of Helsinki. For additional study description and eligibility criteria see Supplementary Materials, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>.

Abbreviations used:

AE:	adverse event
CA:	cicatricial alopecias
CCCA:	central centrifugal cicatricial alopecia
CCL:	C-C motif chemokine ligand
CHLG:	Change in Hair Loss Grade
DLQI:	dermatology life quality index
FFA:	frontal fibrosing alopecia
FFASI:	Frontal Fibrosing Alopecia Severity Index
IFN:	interferon
JAK:	Janus kinase
LPP:	lichen planopilaris
LPPAI:	Lichen Planopilaris Activity Index
STAT:	signal transducers and activators of transcription
Th:	T helper
TYK:	tyrosine kinase

Efficacy and safety endpoints

The coprimary endpoints included mechanistic and safety related outcomes. Mechanistic endpoints included changes from baseline in expression of C-C motif chemokine ligand (CCL) 5, a surrogate for IFN γ activity, and biomarkers of fibrosis in lesional scalp at week 24. The primary safety endpoint was measured by incidence and severity of treatment-emergent adverse events (AEs) and clinically significant changes in vital signs and laboratory parameters, which were assessed every 4 weeks through week 52.

The secondary endpoint of this study was change from baseline at weeks 24 and 48 in lesional scalp gene expression of key inflammatory and fibrosis-related markers (Supplementary Tables I-III, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>). Additional secondary clinical endpoints included changes from baseline in corresponding CA clinical assessment scores: Lichen Planopilaris Activity Index (LPPAI), Frontal Fibrosing Alopecia Severity Index (FFASI) and Change in Hair Loss Grade (CHLG).²⁹⁻³¹

Biomarker and clinical evaluations

Lesional scalp biopsies (4.5 mm punch) were collected at baseline, week 24, and week 48 and analyzed by Taqman low-density array quantitative reverse transcription polymerase chain reaction (RT-PCR) to assess gene expression levels.

CA clinical assessments, eyebrow and eyelash scores, and patient reported outcomes, including dermatology life quality index (DLQI) and visual analog scale pruritus, were collected through week 48. Clinical photos of scalp and eyebrows were taken

at baseline and every 8 weeks thereafter until the end of the study.

The analysis for the mechanistic coprimary endpoints was performed with the use of analysis of covariance model to assess at week 24 the significance of change versus baseline in CCL5 and fibrosis related markers (eg RT-PCR expression levels, within, and between treatment groups). We fitted a linear model including treatment (brepocitinib/placebo) and tissue (nonlesional/lesional) as fixed effects, and baseline CCL5 expression as covariate.

Gene set variation analysis was performed for a subset of key genes associated with fibrosis, previously shown to be upregulated in CA^{21,22} (CTGF, SNAI2, PAI1, COL3A, COL1A) in order to calculate a Z-score, which reflects the aggregate expression level change in a predefined set of fibrosis genes.

We performed subgroup analysis after including the CA form (CCCA, LPP, and FFA) as an additional covariate.

The safety coprimary endpoint was evaluated using descriptive statistics. AE rates, severity, and relatedness were compared between treatment groups using Fisher's exact test.

Additional details of the statistical analyses are provided in the Supplementary Materials, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>.

RESULTS

Patient characteristics

Patients were enrolled from June 9, 2021 until study completion on September 14, 2023. Fifty patients (FFA/LPP $n = 26$; CCCA $n = 24$) were randomly assigned to brepocitinib 45 mg daily or placebo. Following study enrollment closure, one FFA/LPP patient was excluded due to failure to meet inclusion criteria.

Thirty-seven patients received brepocitinib and 12 received placebo. The mean (\pm SD) age of patients receiving brepocitinib was 54 (\pm 14) years and 59 (\pm 12) years in placebo.

Across groups, 39% of patients were white, 55% were black, and 6% were other. A majority of patients were female, and a predominant proportion of CCCA patients were black, reflecting reported demographic distributions. Baseline demographic characteristics were similar across groups (Supplementary Tables IV-VII, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

A majority of patients [brepocitinib 45 mg daily 28/37 (76%); placebo 10/12 (83%)] completed 24 weeks

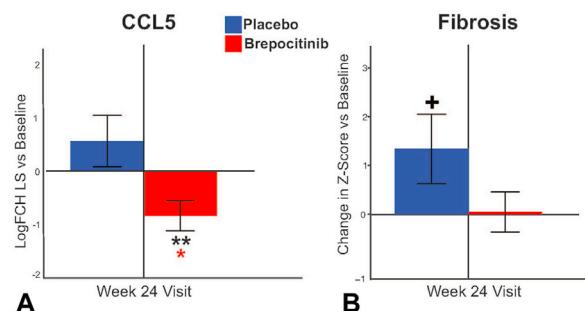


Fig 1. Primary mechanistic endpoint outcome. **A**, Change from baseline in gene expression of CCL5 in lesional scalp at week 24 based on TLDA/RT-PCR. Y-axis shows log2-fold changes in normalized RT-PCR expression and are presented as means \pm SEMs. **B**, Change from baseline in gene expression of fibrosis-gene set variation analysis in lesional scalp at week 24 based on TLDA/RT-PCR. Y-axis shows differences in Z scores vs baseline in normalized RT-PCR expression and are presented as means \pm SEMs. -dCt values are analyzed by mixed-effects model with time, treatment, and tissue interaction as a fixed effect and a random effect for each patient. *Black stars* indicate significance versus baseline; *red stars* indicate significance versus placebo. ** $P < .01$, * $P < .05$, + $P < .1$. *CCL*, C-C motif chemokine ligand; *LogFCH*, log2-fold change; *LS*, lesional; *SEM*, standard error mean.

of the double-blind treatment period; 38 patients continued into the open-label phase, of which 20 FFA/LPP and 8 CCCA patients completed the trial (Supplementary Fig 1, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Primary mechanistic endpoint outcome

CCL5 expression was significantly decreased from baseline in patients treated with brepocitinib at week 24 ($P = .004$). Change in CCL5 in the brepocitinib group was significantly different from placebo at week 24 ($P = .01$). No significant downregulation relative to baseline was observed in fibrosis-related markers in the brepocitinib group; however, enrichment analysis of a subset of fibrosis markers included in the panel showed trending upregulation at week 24 with placebo, while no worsening was observed in the brepocitinib group ($P < .10$; Fig 1).

Primary safety outcome

During the double-blind treatment period, AE were reported in 29 patients who received brepocitinib and 5 patients who received placebo ($P = .03$). The majority of AE were mild-to-moderate in severity, with low rates of trial discontinuation due to AE ($n = 2$). Among the most common AE were acne, COVID-19 infection, anemia, and elevation in serum creatinine level. Two patients receiving brepocitinib experienced serious AE (anemia and

pneumonia with gastroenteritis), leading to drug discontinuation. During the open-label period, 2 patients discontinued due to anemia and one patient discontinued due to elevated creatine phosphokinase. No significant difference in AE rates was found between the placebo cross over group and the brepocitinib group (Supplementary Tables VIII and IX, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Rates of mild, moderate, severe, and life-threatening AE were not significantly different between placebo and brepocitinib groups in either study phase. No cardiovascular events, thrombotic events, malignancies, or deaths in any treatment group were reported through week 52.

Key secondary endpoint outcomes

Brepocitinib significantly reduced lesional scalp expression of key inflammatory markers belonging to Th1/IFN γ (CXCL9, CXCL10), general inflammation (MMP12), T cell activation/Th1 (GZMB, IL2RA), and Th2 (CCL13, CCL18, CCL26) pathways at week 24 compared to baseline. Expression of Treg marker, FOXP3, and JAK/STAT marker, JAK3, was also significantly reduced at week 24 and remained so through week 48, compared to baseline (all $P < .05$). Th1 marker, IFN γ , and Th2 marker, IL4, were significantly downregulated at week 48 ($P < .05$; Supplementary Figs 2 and 3, Table X, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Compared to placebo, expression of markers of general inflammation (MMP12), T cell activation/Th1 (GZMB, IL2RA, IL15RB), Th1/IFN γ (IFN γ , CCL5, CXCL10), Th2 (CCL13, CCL18), Treg (FOXP3), and JAK/STAT (JAK3) were significantly lower with brepocitinib at week 24 (all $P < .05$; Supplementary Figs 2 and 3, Table X, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Evaluation of biomarker expression within disease subgroups (FFA/LPP and CCCA) showed that in the FFA/LPP group at week 24, brepocitinib significantly downregulated Th1/IFN γ (IFN γ , CCL5, CXCL9, CXCL10), general inflammation (MMP12), T cell activation (GZMB, IL2RA), and Th2 (IL4, CCL13, CCL18, CCL26) markers. FOXP3 and JAK3 were also significantly reduced (all $P < .05$). Expression of a majority of these biomarkers was also significantly lower with brepocitinib as compared to placebo. These markers remained significantly downregulated at week 48 in the FFA/LPP group (all $P < .05$; Supplementary Figs 4 to 6, Table XI, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

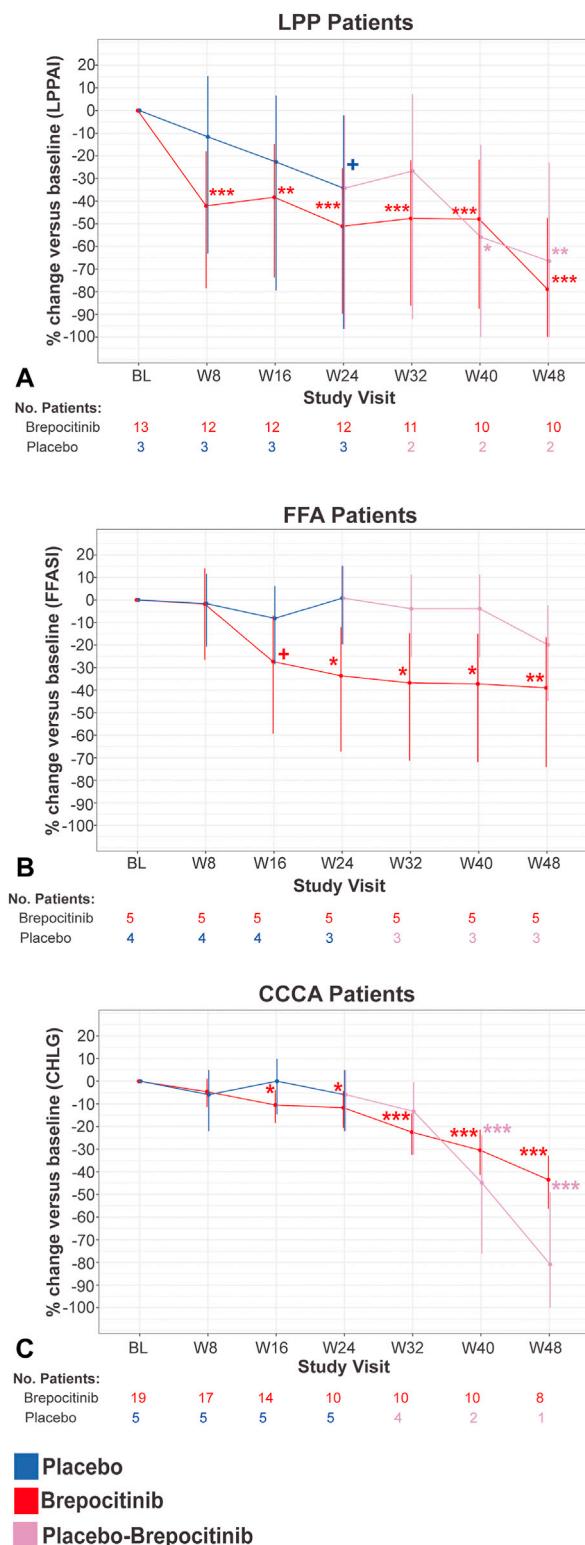


Fig 2. Secondary clinical efficacy endpoint. Mean percent change from baseline in (A) lichen planopilaris activity index (LPPAI) in LPP patients (B) frontal fibrosing alopecia severity index (FFASI) in FFA patients (C) central hair loss grade (CHLG) in CCCA patients. Red, blue, and pink represent brepocitinib, placebo, and brepocitinib after

Among CCCA patients receiving brepocitinib, significant downregulation of markers belonging to general inflammation (MMP12), Th1/IFN γ (CXCL10), Th2 (CCL13, CCL18, CCL26), and Th17/22 (IL23R, S100A7) was observed at week 24 as compared to baseline (all $P < .05$). No significant downregulation of fibrosis-related markers was observed in brepocitinib patients; however, trending upregulation of markers of fibrosis was observed in CCCA placebo group (Supplementary Figs 4 to 6, Table XI, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Key secondary clinical efficacy outcomes

Brepocitinib significantly decreased clinical severity scores in each CA subtype through week 48. At week 24, LPP patients receiving brepocitinib showed significant mean percent change in LPPAI of -51.0% (90% CI, -79.5 to -30.2) compared to baseline ($P < .001$). Further improvement in the brepocitinib group was observed through week 48 with mean percent change of -79.2% (90% CI, -100 to -53.6) compared to baseline ($P < .001$). LPP patients receiving placebo did not achieve a statistically significant change from baseline in LPPAI at week 24; however, placebo patients who crossed over into brepocitinib in the open-label phase showed significant mean percent change of -67.5% by week 48 compared to baseline (90% CI, -100 to -30.7) ($P = .002$; Fig 2, A, Supplementary Table XII, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

FFA patients receiving brepocitinib showed significant mean percent change in FFASI score by week 24 of -33.6% (90% CI, -67.2 to -12.1) compared to baseline ($P = .02$), with further improvement at week 48 and mean percent change of -39.0% (90% CI, -74.2 to -16.5) compared to baseline ($P < .01$). At week 24, FFA patients receiving placebo showed mean numerical increase in FFASI compared to baseline [0.9% (90% CI, -19.4 , 15.1)]; however, among FFA patients who crossed over into brepocitinib, numerical reduction was observed with mean

placebo (open-label), respectively. Error bars represent 90% confidence intervals. *** $P < .001$, ** $P < .01$, * $P < .05$, + $P < .1$. Red stars represent change from baseline in the brepocitinib arm. Blue stars represent change from baseline in the placebo arm. Pink stars represent change from baseline in the open-label phase of the placebo arm. Number of patients at each time point is denoted under the respective study week visit. CCCA, Central centrifugal cicatricial alopecia; FFA, frontal fibrosing alopecia; LPP, lichen planopilaris.

percent change of -19.6% (90% CI, -44.6 to -2.1) at week 48 from baseline (Fig 2, B, Supplementary Table XIII, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

CCCA patients receiving brepocitinib showed significant improvement in CHLG at week 16, with further improvement observed at weeks 24 and 48 with percent change in CHLG of -11.9% (90% CI, -20.8 to -4.4 ; $P < .05$) and -43.5% (90% CI, -56.2 to -33.0 ; $P < .001$) compared to baseline, respectively. CCCA patients receiving placebo showed minimal change in CHLG at week 24; however, following cross over to brepocitinib, at week 48, one patient maintained treatment and showed significant mean percent change in CHLG of -81.6% (90% CI, -100 to -50.8) compared to baseline ($P < .001$; Fig 2, C, Supplementary Table XIV, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>). Representative patient photos are shown in Supplementary Fig 7, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>. Eyebrow and eyelash scores remained stable in the drug group by week 48 (Supplementary Table XV, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Among patients receiving brepocitinib, DLQI was significantly reduced from baseline starting at week 8 ($P = .004$), with mean percent change at week 24 of -17.6% (90% CI, -24.7 , to -11.6 , $P < .001$). No improvement in DLQI was observed in the placebo group at week 24 (Supplementary Fig 8, Table XVI, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>). Patients' subjective measure of itch was also significantly reduced at week 24 in the brepocitinib group with mean percent change of -31.6% (90% CI, -57.8 , to -12.3 ; $P = .02$; Supplementary Fig 8, Table XVII, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>). When comparing the change in pruritus at weeks 24 and 48 across CA disease groups, a significantly greater reduction occurred in the LPP group compared to the CCCA and FFA groups ($P < .01$; Supplementary Table XVIII, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Post hoc analysis showed that patients with shorter disease duration (less than the study cohort median of 5 years) had significantly greater clinical score percent improvement than patients with long disease duration (>5 years) at weeks 24 and 48 (all $P < .05$; Supplementary Fig 9, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

Correlation analysis

Spearman correlations between change from baseline in clinical severity measures and change from

baseline in biomarkers in lesional scalp biopsy were conducted (Supplementary Table XIX, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>). Notably, week 24 change in FFASI positively correlated with CXCL10 ($r = 0.83$, $P = .01$). Total scalp margins, the assessment of total scalp hair loss within the FFASI scoring system, positively correlated with CCL5 ($r = 0.74$; $P = .04$). Change in pruritus, a subcategory of the LPPAI scoring system, positively correlated with changes in CCL5, CXCL9, JAK3, and CTGF ($r > 0.60$, $P < .05$ for all). Change in CHLG also positively correlated with change in IL23 subunit, IL23p19 ($r = 0.62$, $P = .02$; Supplementary Table XIX, available via Mendeley at <https://data.mendeley.com/datasets/nsjbw2d8r2/1>).

DISCUSSION

Herein we present the first randomized placebo-controlled study of a systemic dual TYK2/JAK1 inhibitor and assessment of biomarker changes in CA tissue following treatment. Brepocitinib successfully downregulated CCL5 expression after 24 weeks of treatment, meeting a coprimary endpoint. Existing literature points to a Th1-based cytotoxic T cell response and IFN γ activation in the pathogenesis of CA, and thus the downregulation of CCL5, an IFN γ -associated marker, is highly encouraging.^{21,22} As a TYK2/JAK1 dual inhibitor, brepocitinib targets STAT activation of Th1/IFN γ and γ c cytokines.³² Thus, it allows for biomarker downregulation beyond the Th1 axis, including modulation of markers associated with T-cell activation, JAK/STAT signaling, Th2, and Treg.

Previous studies suggest that certain proinflammatory signals, particularly JAK/STAT, are associated with fibrotic changes.^{21,22,33-36} Inflammatory changes likely precede fibrotic changes in CA. Thus, inhibition of inflammatory pathways may prevent downstream fibrosis. Expression of fibrotic markers remained stable in the brepocitinib group but worsened in the placebo group, suggesting that inhibiting inflammation may prevent fibrotic progression. Comparison of clinical improvement based on median disease duration showed greater improvement with shorter disease duration in the brepocitinib group. Earlier disease may be dominated by inflammatory dysregulation, underscoring the importance of prompt intervention, as seen in alopecia areata.^{37,38}

Brepocitinib was well tolerated through week 52, with a safety profile comparable to that of previous trials, meeting another of its coprimary endpoints.²⁴

Brepocitinib groups across all CA subtypes met the secondary endpoint of significant reduction in clinical severity scores by week 24, with continued improvement through week 48. Moreover,

molecular analysis correlated with clinical improvement, bridging mechanistic and clinical endpoints.

There were some limitations of the present study. The scoring schema of the CA subtypes is inherently limited by inter-rater variability and subjectivity. Future investigations are warranted to develop more reliable scoring systems. The study is also limited by a single-dose regimen, 3:1 design, and small placebo group, limiting significant comparisons between brepocitinib and placebo.

This study demonstrates that brepocitinib effectively decreases inflammatory biomarker expression and improves clinical severity measures at 24 and 48 weeks while also maintaining a favorable safety profile through 52 week follow up period. Larger scale studies utilizing brepocitinib for the treatment of CA are warranted to move towards formal approval.

Conflicts of interest

Dr Meariman is a consultant for Abbvie. Dr Ungar is an employee of Mount Sinai and has received research funds (grants paid to the institution) from: Incyte, Rapt Therapeutics, and Pfizer. He is also a consultant for Arcutis Biotherapeutics, Bristol Myers Squib, Castle Biosciences, Fresenius Kabi, Galderma, Janssen, Lilly, Pfizer, Primus Pharmaceuticals, Sanofi, and UCB. Dr Guttman-Yassky is an employee of Mount Sinai and has received research funds (grants paid to the institution) from: Abbvie, Celgene, Eli Lilly, Janssen, Medimmune/Astra Zeneca, Novartis, Pfizer, Regeneron, Vitae, Glenmark, Galderma, Asana, Innovaderm, Dermira, UCB. They are also a consultant for Sanofi Aventis, Regeneron, Stiefel/GlaxoSmithKline, MedImmune, Celgene, Anacor, AnaptysBio, Dermira, Galderma, Glenmark, Novartis, Pfizer, Vitae, Leo Pharma, Abbvie, Eli Lilly, Kyowa, Mitsubishi Tanabe, Asana Biosciences, and Promius. Drs Oemar, Mahling, and Peeva are Pfizer employees and own stock and stock option. Authors David, Shokrian, Hawkins, Sikand, Singer, Estrada, Bose, and Pulsinelli, and Drs Duca, Dubin, Andrews, and Da Rosa have no conflicts of interest to declare.

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