

Alopecia Areata and Atopic Dermatitis: Common Mechanisms and Emerging Therapeutics

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Abstract: Alopecia areata (AA) and atopic dermatitis (AD) have historically been classified as immunologically distinct—AA as a T helper 1 (T_h1) driven autoimmune condition and AD as a T helper 2 (T_h2) polarized inflammatory disease. However, emerging evidence demonstrates substantial immunological overlap, with both conditions exhibiting multi-axis cytokine dysregulation converging on Janus kinase-signal transducer and activator of transcription (JAK-STAT) dependent signaling and regulatory T cell insufficiency/dysfunction. Epidemiologically, these conditions frequently co-occur: up to 40% of AA patients have atopic comorbidities. Both diseases share genetic susceptibility loci, including interleukin 13 (IL-13) and human leukocyte antigen (HLA) associations, further supporting convergent pathogenic mechanisms. Key shared biomarkers include elevated interleukin 4 (IL-4), IL-13, and IgE, alongside dysregulated interferon-gamma and interleukin 15 signaling through the JAK-STAT pathway. These mechanistic insights have catalyzed the development of therapeutics for both conditions. JAK inhibitors—including baricitinib, ritlecitinib, upadacitinib, and abrocitinib—have demonstrated efficacy in AA and/or AD by interrupting cytokine signaling central to both diseases. Dupilumab, an interleukin -4 receptor alpha antagonist FDA-approved for AD, has demonstrated benefit in a specific subset of AA patients with concurrent atopic disease; these findings highlight pathophysiologic heterogeneity within the disease and support the need for endotype-stratified treatment strategies. Investigational therapies targeting the OX40/OX40 ligand costimulatory pathway, interleukin 2 receptor agonists (repegaldesleukin), interleukin 7 receptor antagonists (bempikibart), and tri-specific cytokine inhibitors (EQ101) represent the next generation of agents that may treat both conditions by restoring immune homeostasis rather than broadly suppressing immunity. In this narrative review, we synthesize current understanding of the shared immunopathogenesis of AA and AD and highlight approved and investigational therapeutics with dual-disease relevance, pointing toward an era of mechanistically targeted, potentially disease-modifying interventions.

Keywords: alopecia areata, atopic dermatitis, JAK inhibitors, dupilumab, Th2

Introduction

Alopecia areata (AA) and atopic dermatitis (AD) are common inflammatory skin disorders historically viewed as immunologically distinct, with AA being a cell-mediated autoimmune hair loss condition driven by T helper type 1 (T_h1) inflammation,^{1,2} and AD dominated by T helper type 2 (T_h2)-type pathways,^{3,4} a dichotomy that has increasingly been challenged as advances in immune-mediated inflammatory disease research reveal shared pathogenic mechanisms across conditions once considered unrelated.⁵ Clinically, up to 15–20% of AA patients may have concurrent AD, and AA has been linked to other atopic conditions like asthma and allergic rhinitis, with up to 40% demonstrating atopic disease overall.^{6–8} The primary trigger for AA is loss of immune privilege (IP), leading to an autoimmune attack on the hair follicle, driven by CD8⁺ T cells that target the follicular bulb. AD pathogenesis is multifactorial, with type 2 inflammation now recognized as a central driver that both arises from and perpetuates epidermal barrier dysfunction.^{9,10} Both diseases exhibit multi-axis cytokine dysregulation with many shared features of both T_h1 and T_h2 dysregulation,

converging on Janus kinase-signal transducer and activator of transcription (JAK-STAT) dependent cytokine signaling and regulatory T cell (Treg) insufficiency/dysfunction as shared pathogenic denominators.¹¹

The management of patients with comorbid AA and AD presents a particular clinical challenge: comorbid AD has been identified as a poor prognostic factor for AA treatment response and a predictor of relapse risk, and therapies effective for one condition may have limited or uncertain efficacy for the other — IL-13 inhibitor monotherapy, for example, is approved for AD but has not shown efficacy in AA thus far, while topical therapies that are central to AD management are largely ineffective in AA.

Notably, immunomodulators like Janus kinase (JAK) inhibitors and biologics such as dupilumab have transformed treatment of both AA and AD, and offer further therapeutic promise across both diseases by intervening on shared cytokine signaling pathways.^{12,13} In this narrative review, we discuss the immunological background of AA and AD with a focus on shared immune axes, highlight key biomarkers of disease activity, and review approved and investigative therapeutics that may successfully treat both diseases.

Epidemiology

The global prevalence of AA is approximately 2%,¹⁴ with roughly 5–10% of AA cases being alopecia totalis (AT) or alopecia universalis (AU), complete scalp and body hair loss, respectively.^{14,15} Across races, a cross-sectional study found a significantly higher odds of AA in Black patients compared with White patients and a lower odds of AA in Asian patients compared with White patients.¹⁶ Individuals aged 25–40 have the highest likelihood of developing AA.¹⁷ The incidence appears similar between males and females.¹⁸ AA is also the third most common dermatologic condition in children, and the annual number of new pediatric cases continues to rise.¹⁹ Although the underlying pathogenesis of AA has yet to be fully elucidated, studies have identified genetic susceptibility loci centered around innate and adaptive immunity, including human leukocyte antigen (HLA-DR), cytotoxic T-lymphocyte-associated protein 4 (CTLA4), interleukin-2 (IL-2)/IL-21, and interleukin-2 receptor alpha (IL-2RA).^{20,21} Of note, one genome wide association study identified interleukin-13 (IL-13), a central cytokine in AA pathogenesis, as one of the most strongly associated genes.²⁰ Family history of AA is a risk factor, as AA tends to present in multiple members within the same family and presents more often among siblings than among parents and offspring.¹⁸ Other risk factors for AA include having an autoimmune or atopic background such as atopic dermatitis, psoriasis, or inflammatory bowel disease.⁶

AD is a chronic inflammatory skin disease that can present at any age with recurrent eczematous lesions and intense itching. The global prevalence rate over one year of AD is approximately 2.6%, approximately 2.0% in adults and 4.0% in children, and the lifetime prevalence rate is up to 10% globally in adults, and up to 20% globally in children.^{22–24} AD is slightly more common in females (2.8%) than in males (2.4%).²² Although it may develop throughout life, between 60% and 73% of affected children present with AD within the first one to two years of age.^{22,25}

AD and AA are epidemiologically linked, with multiple studies demonstrating bidirectional increase in risk; providing support for AD as the most common comorbidity in AA patients.^{26,27} In several meta-analyses, AA patients were found to have an increased risk of AD, and conversely, AD patients were found to have a significantly increased risk of AA.^{26,28} A Danish population-based study reported this association to have as high an adjusted odds ratio as 26.31 for AD among patients with AA.²⁷ This genetic link is further supported by multiple Mendelian randomization studies looking at AD and AA, strengthening a potential underlying shared etiology in some patients.^{29,30} Clinically, the significance of this association extends beyond epidemiology, as comorbid AD has been identified as a poor prognostic factor for treatment response and a predictor of relapse risk in AA across several studies.^{6,31,32}

Biomarkers of Alopecia Areata

AA is believed to arise from the collapse of immune privilege (IP) in the hair follicle.³³ Under normal conditions, the anagen hair follicle maintains a distinct state of IP, characterized by minimal expression of major histocompatibility complex (MHC) proteins in its inferior segment, reduced levels of stress-induced ligands, local production of anti-inflammatory cytokines/hormones (eg. transforming growth factor-beta (TGF β), interleukin-10 (IL-10), alpha-melanocyte-stimulating hormone (α -MSH)), and a relative scarcity of immune cell infiltrates.^{34,35} For as yet unclear reasons, disruption of this protective environment leads to upregulated expression of self-antigenic peptides presented by

MHC molecules along the follicular epithelium, thus facilitating infiltration of self-reactive T cells and a dense immune cell infiltrate which histopathologically appears as a distinctive “bee-swarm pattern,” marked by dense lymphocytic infiltrates surrounding the bulbar region of anagen hair follicles.³⁴

In AA, peribulbar immune cell infiltrates are predominantly composed of CD4⁺ and CD8⁺ T cells, accompanied by increased numbers of natural killer (NK) cells, Langerhans cells, dendritic cells, and mast cells.³⁶ Among these, CD8⁺ T cells closely associated with the hair follicle were found to express NKG2D, an activating receptor typically associated with the NK cell lineage.³⁷ Notably, CD8⁺ NKG2D⁺ T cells are sufficient to induce disease in a cell-transfer murine model of AA, highlighting a potential pathogenic role.¹² Furthermore, activated CD8⁺ T cells produce high levels of pro-inflammatory cytokines, such as Interferon-gamma (IFN- γ), which skew the hair follicle microenvironment towards a proinflammatory state.³⁸

The primary T_h1 cytokine IFN- γ is hypothesized to be a key mediator of IP collapse within the hair follicle. Its ability to disrupt IP has been demonstrated ex-vivo: anagen-stage hair follicles collected from healthy scalps cultured in the presence of IFN- γ resulted in increased expression of MHC I molecules.³⁵ IL-2 is another key cytokine released by CD4⁺ cells which signals through JAK1 and JAK3 receptors, driving gene transcription programs that promote cellular expansion, survival, and maintenance.³⁹ In AA patients, elevated IL-2 expression is detected within the deep dermis, where the follicular bulb resides, and is significantly increased compared to levels of the upper dermis.³⁹ Furthermore, active lesions of AA patients also show higher expression of IL-2 than inactive areas.³⁹ IL-15, structurally related to IL-2 and essential for the survival of CD8⁺ memory T cells, further contributes to disease pathogenesis by supporting the expansion of T cells and NK cells and enhancing CD8⁺ T-cell production of granzymes, IFN- γ , and TNF- α .⁴⁰ Circulating IL-15 levels correlate with disease severity,⁴⁰ and both patients with AA and murine models exhibit increased IL-15 and IL-15R α expression in the outer root sheath of the hair follicle compared with controls.¹²

Beyond IFN- γ , IL-2, and IL-15, T_h2-associated cytokines, particularly IL-4 and IL-13, have more recently been implicated in AA. Genetic susceptibility involving IL-13 was identified in a large genome-wide study that included patients with and without atopy, while IL-4 polymorphisms were similarly associated with increased AA risk.^{11,20} Transcriptomic profiling of AA scalp tissues revealed IL-13 as one of the most highly upregulated cytokines, with significant increases in lesional versus nonlesional scalp, and serum IL-13 levels were also elevated in a cohort of 51 AA patients compared with controls.¹¹ Alongside T_h2 activation, elevations in T_h1-related and cardiovascular-associated proteins (MMP9, AGER, IL1RL1/ST2/IL33R) were detected in both scalp and serum, with the scalp exhibiting a greater degree of immune dysregulation than serum.⁴¹ Additional evidence for T_h2 involvement comes from elevated circulating levels of IL-4, IL-13 and IgE in AA patients.^{41–43} Furthermore, increased IL-13⁺/cutaneous lymphocyte-associated antigen (CLA)⁺ skin-homing CD4⁺ T and CD8⁺ T cells have been identified in the peripheral blood of AA patients, with IL-13⁺CD4⁺CLA⁻ T cell frequencies correlating with disease severity and IFN- γ ⁺CD8⁺ T-cell frequencies correlating with disease duration.⁴⁴

Biomarkers of Atopic Dermatitis

Immune dysregulation is central to the pathogenesis of AD, driving skin barrier disruption and the clinical manifestations of eczematous lesions and itch. AD has long been characterized as a T_h2-polarized disease, with elevated expression of IL-13 and IL-4, and numerous additional T_h2-associated markers.⁴⁵ Increasing evidence, however, supports its heterogeneity, with distinct endotypes demonstrating variable contributions from T_h1, T_h17, and T_h22 pathways. For example, Asian patients exhibit greater T_h17 activation, Black patients display fewer FLG loss-of-function mutations, and pediatric patients show higher expression of TSLP receptor, among other immunologic differences.⁴⁶ Several T_h2-related biomarkers, including TARC/CCL17, periostin, and serum IgE, correlate with disease activity and therapeutic response.²³

Epidermal barrier dysfunction is another hallmark of AD,⁴⁷ characterized by dysregulated expression of key structural proteins such as filaggrin (FLG), transglutaminases, keratins, and intercellular structural proteins. FLG, a major structural protein in the stratum corneum, is critical for maintaining epidermal integrity; FLG mutations are associated with an increased risk of severe AD with earlier onset, longer persistence, and skin infection.^{48,49} Immune activation may also influence FLG gene expression, as IL-4 and IL-13 downregulate FLG and contribute to impaired barrier function.⁵⁰ Critically, IL-4/JAK signaling-mediated suppression of antimicrobial peptides has direct clinical consequences in AD,

which is associated with cutaneous infections including *Staphylococcus aureus* colonization and eczema herpeticum, with no corresponding increased risk in AA. Beyond suppressing antimicrobial peptides (AMP), these cytokines promote allergic inflammation and chemokine production, underscoring their central roles in AD pathophysiology.⁵⁰ The clinical success of targeted therapies such as dupilumab (IL-4R α blockade) and the IL-13 inhibitors tralokinumab and lebrikizumab further highlights the pathogenic relevance of these cytokines.^{3,51–53}

Interleukin-22 (IL-22) is another proinflammatory cytokine implicated in AD due to its critical role in wound healing and immune stimulation. Produced primarily by T_h22 with additional production from T_h17 cells, IL-22 plays a central role in stimulating the proliferation, migration, and differentiation of the cells involved in tissue repair.⁵⁴ IL-22 can directly stimulate the production of the extracellular matrix components in fibroblasts via binding to the IL-22 receptor and triggering downstream JAK/STAT dimerization.⁵⁵ In keratinocytes, IL-22 induces hyperproliferation and decreases FLG expression, contributing to epidermal thickening and barrier dysfunction.⁵⁴ IL-22 may also modulate pruritus, as gastrin-related peptide (GRP), a mediator of non-histaminergic itch, is increased in AD and correlates with itch severity, with IL-22 upregulating GRP receptor expression in keratinocytes.⁵⁶ Although anti-IL-22 therapies have not shown broad clinical efficacy, additional analysis suggested that inhibiting IL-22 in the subset of patients with high expression of IL-22 may have clinical utility.⁵⁷ In this context, therapeutically engaging the IL-22 receptor itself may represent a more promising strategy, as emerging data from a Phase 2b clinical trial (NCT05923099) on temtokibart, a monoclonal antibody directed against the interleukin-22 receptor subunit alpha-1 (IL-22RA1), indicate encouraging early signals of efficacy.^{58–60}

CCL17, also known as thymus and activation-regulated chemokine (TARC), is a chemokine produced by dendritic cells, endothelial cells, keratinocytes, and fibroblasts that acts as a chemoattractant for CCR4⁺ T_h2 lymphocytes, playing a central role in the pathogenesis of AD by promoting T_h2 cell migration into inflamed skin.⁶¹ CCL17/TARC is considered one of the most robust and specific biomarkers for monitoring AD activity and treatment response, and is used clinically in several countries to guide management.^{45,62} Its specificity extends beyond general atopic diseases, with elevated CCL17/TARC levels are more strongly associated with AD than with asthma or allergic rhinitis.⁶³ CCL17/TARC is markedly overexpressed in lesional AD skin and elevated in the serum, correlating closely with disease activity and severity as measured by clinical scores such as SCORing Atopic Dermatitis (SCORAD).^{45,62} Elevated CCL17/TARC can be detected even before clinical onset of AD, supporting a role in early disease pathogenesis and as a predictive biomarker.⁶⁴ Other T_h2 chemokines, including CCL18, CCL22, and CCL26, are likewise elevated in AD and contribute to the self-perpetuating cycle of T_h2-driven inflammation.⁴⁵

Interleukin-31 (IL-31), a key pruritogenic cytokine produced primarily by T_h2 cells, acts directly on sensory neurons and keratinocytes to induce itch, impair barrier function, and drive inflammation.^{65–67} Elevated IL-31 levels correlate strongly with disease severity and propagate the itch-scratch cycle, worsening skin lesions and quality of life.⁶⁵ Nemolizumab, a humanized monoclonal antibody targeting the IL-31 receptor α (IL-31R α), has demonstrated rapid and sustained reduction in pruritus, improvement in skin signs, and enhanced quality of life in patients with moderate-to-severe AD inadequately controlled by topical therapies.⁶⁷

Thymic stromal lymphopoietin (TSLP) is highly expressed in the epidermis of patients with AD, with significantly elevated serum levels in both adults and children.⁶⁸ Its production is induced by environmental triggers such as allergens, microbes, diesel exhaust, cigarette smoke, and chemical irritants.⁶⁹ Notably, in a Korean birth cohort, skin tape stripping revealed increased TSLP expression in the skin of 2 month-old infants who later developed AD at 24 months, suggesting a pathogenic role early in disease development.⁷⁰ TSLP overexpression is specific to AD lesions, with no significant elevations detected in nonlesional AD skin or in lesions from nickel-induced allergic contact dermatitis.⁷¹

Shared Pathways

Having outlined the distinct biomarker landscapes of AA and AD, collectively, current biomarker and genetic data suggest that AA and AD, though clinically distinct, share several key pathogenic pathways, with JAK-STAT dependent cytokine dysregulation and impaired Treg function emerging as central convergent mechanisms (Figure 1). These common mechanisms help to explain both their epidemiologic association and their emerging overlap in therapeutic strategies.

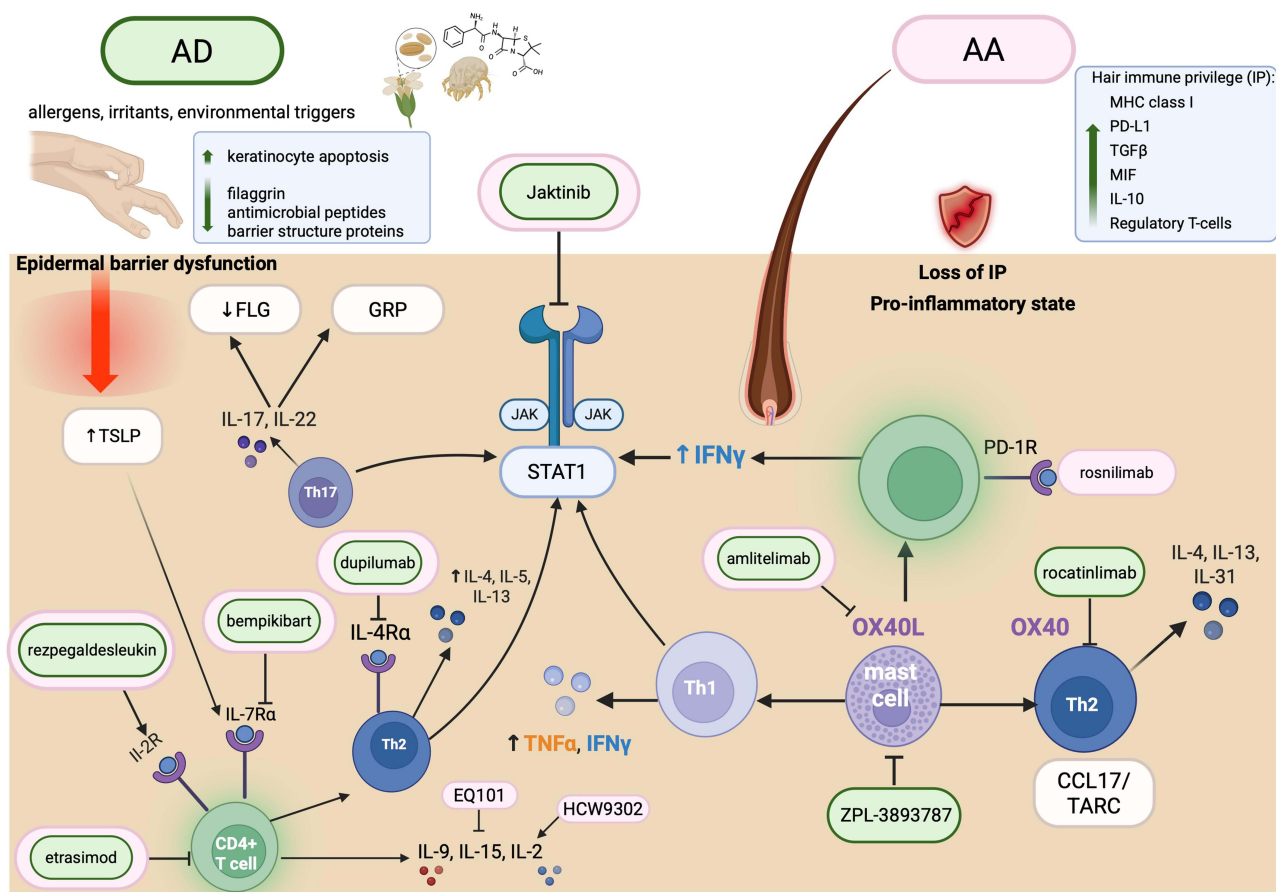


Figure 1 Convergent and divergent immunopathogenic pathways in atopic dermatitis (AD) and alopecia areata (AA), with therapeutic targets. AD (left) is characterized by a bidirectional interplay between type 2 inflammation and epidermal barrier dysfunction: T_H2 cytokines (IL-4, IL-13) suppress epidermal differentiation proteins — including filaggrin (FLG) and loricrin — and disrupt stratum corneum lipid composition, while barrier impairment in turn facilitates allergen penetration and keratinocyte-derived TSLP release, amplifying the inflammatory cycle. AA (right) arises from collapse of hair follicle immune privilege (IP), characterized by loss of local immunosuppressive mediators (TGFβ, IL-10, PD-L1, MIF) and $CD8^+$ T cell-mediated autoimmune attack on the follicular bulb. Both diseases converge on shared JAK-STAT signaling, T_H1/T_H2 cytokine dysregulation (IFN-γ, IL-4, IL-5, IL-13), mast cell activation, and OX40/OX40L costimulatory signaling. Therapeutic agents are color-coded by primary indication: green (AD) and pink (AA), shown with their respective molecular targets. Created in BioRender.

Abbreviations: FLG, filaggrin; GRP, gastrin-releasing peptide; IP, immune privilege; TSLP, thymic stromal lymphopoietin; CCL17/TARC, thymus and activation-regulated chemokine.

IFN-γ has been studied as a biomarker of disease severity in AA and disease chronicity in AD. IFN-γ and its downstream chemokines, particularly CXCL9 and CXCL10, are essential mediators of inflammatory amplification.⁷² In AA, IFN-γ is produced primarily by $CD8^+$ T cells and other lymphocyte subsets, contributing to the collapse of hair follicle IP, upregulation of MHC class I, and recruitment of additional inflammatory cells, which collectively drive follicular damage and hair loss.^{38,73} Elevated serum IFN-γ levels and increased IFN-γ gene expression in lesional tissue correlate with disease severity, disease activity, the extent and chronicity of hair loss, and the degree of $CD8^+$ T-cell perifollicular infiltration.^{74,75} Animal models further support its pathogenic contributions, as exogenous IFN-γ administration can induce AA-like lesions.⁷⁶ In AD, IFN-γ sustains T_H1 - and T_H2 -cell activation, induces keratinocyte apoptosis, and impairs the epidermal barrier, thereby increasing transepidermal water loss.⁷² Chronic AD is characterized by progressive enrichment of IFN-γ driven T_H1 pathways, paralleling the expansion of multiple immune pathways.^{77–79}

Although TNF-α is not considered a primary driver of AA or AD pathogenesis, it plays an important homeostatic role in skin immunity, helping to maintain the balance between pro- and anti-inflammatory signals that preserves both hair follicle immune privilege and epidermal barrier integrity. Reports of TNF-α inhibitor-induced AA highlight this role,⁸⁰ leading Perez et al to propose that TNF-α may contribute to the maintenance of hair follicle IP.⁸¹ Similarly, AD has been observed in patients receiving TNF-α inhibitors for inflammatory bowel disease.⁸² One proposed mechanism is that TNF-

α blockade may shift the immunologic environment toward T_H2 upregulation, with personal or familial atopic history predisposing individuals to anti-TNF- α -induced AD.⁸²

IL-2 is another T_H1 cytokine responsible for T-cell proliferation, including Treg survival and expansion, which positions it as a key homeostatic regulator across diverse inflammatory conditions.⁸³ As discussed above, in AA, IL-2 promotes $CD8^+NKG2D^+$ T cell-mediated hair follicle destruction.¹² In AD, impaired IL-2 production contributes to Treg dysfunction and loss of immune tolerance.⁸³ This dual role as a pathogenic effector in AA versus a Treg sustainer in AD depends on dose and receptor context: low doses of IL-2 preferentially expand Tregs, whereas higher doses of IL-2 promotes effector T cell activation.⁸⁴

The T_H2 axis, composed of $CD4^+$ T cells producing IL-4, IL-5, and IL-13, drives type 2 immunity, including eosinophil activation, IgE production, and barrier dysfunction.⁸⁵ This pathway represents a shared immunologic mechanism in both AA and AD (Figure 1). Both diseases feature expansion of circulating T_H2 and T_H22 cells, and reduced regulatory T cells, with disease severity correlating with the degree of $T_H1/T_H2/T_H17$ activation and keratin dysregulation.⁸⁵ In AA, suppression of T_H2 biomarkers with dupilumab correlates with clinical improvement and restoration of hair keratin expression, further supporting a pathogenic role of the T_H2 axis in AA.⁸⁶ Even less specific JAK pathway inhibitors such as ritlecitinib (JAK3/TEC) and brepocitinib (JAK1/TYK2) demonstrate that modulation of T_H2 responses is strongly associated with improvements in keratin expression and clinical outcomes.⁸⁷ In comorbid AA and AD, immune signatures differ based on atopic status. Kageyama et al demonstrated that atopic and non-atopic AA patients exhibit distinct transcriptomic profiles.⁵ Non-atopic AA is characterized by a more pronounced T_H1 -skewed immune activation, with higher ratios of $CD4^+IFN-\gamma^+$ cells to $IL-4^+$ or $IL-13^+CD4^+$ cells.⁵ In contrast, patients with coincident AD exhibit dense infiltration of both $CXCR3^+$ (T_H1 -associated) and $CCR4^+$ (T_H2 -associated) cells around the hair bulb.⁵ Furthermore, IgE levels are associated with dupilumab response in both atopic and non-atopic AA, suggesting that IgE is not exclusive to atopic disease.⁵ Dupilumab therapy significantly reduces $CCR4^+$ cells (T_H2) cell infiltration in atopic AA, whereas $CXCR3^+$ cell numbers (T_H1) remain unchanged. These findings underscore the immunologic heterogeneity of atopic and non-atopic AA and suggest that type 2 inflammation plays a particularly important role in AA associated with extrinsic AD, whereas non-atopic AA has more prominently type 1 inflammation.⁵

While IL-15 plays a well-characterized pathogenic role in AA — supporting $CD8^+$ T cell expansion and correlating with disease severity — its contribution to AD pathogenesis remains poorly understood, representing a point of mechanistic divergence between the two diseases and raising uncertainty about whether IL-15-targeting strategies will have therapeutic relevance in AD.

The OX40 costimulatory pathway further contributes to T-cell-mediated inflammation and the development of tissue resident memory (TRM) T cells. OX40 is an inducible receptor predominantly expressed on activated T cells and is upregulated by cytokines such as IL-1, IL-2, IL-4, and TNF- α .⁸⁸ Engagement of OX40 by OX40L promotes T-cell proliferation and survival, TRM cell generation, and proinflammatory cytokine production.⁸⁸ Persistent antigen exposure and prolonged expression of IFN- γ , IL-4, IL-17, and IL-22 sustain OX40 signaling, perpetuating a self-amplifying inflammatory loop.⁸⁹ In AD, $OX40^+$ T cells and $OX40L^+$ dendritic cells are enriched in lesional skin compared with nonlesional or psoriatic skin.⁸⁹

In AA, mast cells in the perifollicular region also exhibit increased OX40L expression.³⁶ As described above, $OX40L^+$ mast cells activate $CD8^+$ T cells through the OX40-OX40L interaction, enhancing IFN- γ production and promoting T_H2 cell maturation, while downregulating signals required for Treg induction.³⁶ These findings suggest that $OX40L^+$ mast cells undermine Treg-mediated immune tolerance and thereby contribute directly to AA pathogenesis.⁹⁰ Transcriptomic studies show robust upregulation of T_H1 and T_H2 -associated products in AA scalp, with OX40 signaling playing a particularly important role in AA comorbid with atopy.⁹¹ Both lesional and nonlesional AA tissue exhibit increased perifollicular infiltration by $OX40^+$ and $OX40L^+$ cells, which strongly correlate with hair-keratin downregulation.⁹¹ Notably, current hair loss episode duration correlates positively with $OX40^+$ infiltrates in atopic AA, but negatively in non-atopic AA, suggesting distinct immunologic trajectories.⁹¹ Finally, cutaneous OX40/OX40L upregulation is mirrored by elevations in circulating $OX40^+$ and $OX40L^+$ leukocytes, and OX40 is highly expressed on non-skin-homing Tregs in the context of increased circulating $OX40L^+$ antigen presenting cells, further suggesting that Treg dysfunction in AA may originate systemically.⁹¹

Early mechanistic studies established that IFN- γ and interleukin-15, key cytokines in AA, signal through JAKs, and clinical case series demonstrated hair regrowth with oral JAK inhibitors in recalcitrant disease.² More than 50 cytokines signal via the JAK – signal transducer and activator of transcription (STAT) pathway, wherein cytokines binding to transmembrane receptors trigger JAK dimerization, which then activate STATs to translocate to the nucleus and regulate transcription of target genes.^{92,93} The JAK–STAT pathway plays a central role in immune dysregulation in AD, contributing to exaggerated T_H2 polarization, eosinophil activation, B-cell maturation, and suppression of Tregs.⁹⁴ Additionally, IL-4–mediated activation of JAK–STAT signaling promotes AD pathogenesis by inducing epidermal chemokines, pro-inflammatory cytokines, and pro-angiogenic factors, while simultaneously suppressing antimicrobial peptide expression and key mediators of skin barrier integrity.⁹⁴

Despite these convergent inflammatory axes, the site of immune activation in AA and AD imposes important therapeutic constraints that limit the interchangeability of treatment strategies. In AA, the principal target of immune attack is the hair follicle bulb, an anatomically deep structure residing in the mid-to-lower dermis or sometimes even subcutis, rendering topical therapies largely ineffective due to presumed inadequate drug penetration, an observation consistent with failed trials of topical JAK inhibitors in AA.^{95–97} By contrast, AD pathogenesis involves a bidirectional interplay between type 2 inflammation and epidermal barrier dysfunction,^{51,98} positioning the skin surface as a site of immune-driven pathology and a pharmacologically accessible target. Thus, barrier-directed therapies remain central to AD management but have little therapeutic rationale in AA. Furthermore, the regulatory T-cell landscape diverges between the two diseases. AA patients exhibit significantly depleted Tregs compared to both controls and AD patients, and this Treg deficiency correlates with activated CD8+ memory T cells,⁴⁴ suggesting a fundamental collapse of peripheral immune tolerance coupled with cytotoxic effector activation — in contrast to AD, which is instead driven by CD4+ T_H2 cells with compensatory Treg expansion.

Traditional Therapies for Alopecia Areata

Traditional treatments for AA include intralesional and topical corticosteroids, systemic steroids, contact immunotherapy, and traditional immunosuppressants, although these approaches are limited by variable efficacy and side effects.⁹⁹ Intralesional corticosteroids remain the first-line treatment for adults with one to two small patches of hair loss, but their use is associated with pitting scars and limited effectiveness in more extensive disease.⁹⁹ Systemic corticosteroids, such as prednisolone (with dosing regimens commonly in the range of 0.5 mg/kg/day tapering over 6–12 weeks),¹⁰⁰ may be used in severe AA, but their utility is tempered by suboptimal durability of response and considerable adverse effects. Risks include weight gain, worsening diabetes or hypertension, and suppression of the hypothalamic–pituitary–adrenal axis, along with high relapse rates.¹⁰⁰ Other traditional modalities include contact immunotherapy which stimulates an allergic contact dermatitis reaction using agents such as diphencyclopropenone (DPCP) and squaric acid dibutylester (SADBE) to promote hair regrowth.¹⁰¹ Adverse effects include urticaria, regional lymphadenopathy. This strategy is generally contraindicated in patients with acute, rapidly progressive AA.¹⁰¹ For more severe forms of AA, systemic immunosuppressants such as methotrexate and cyclosporine have been used, although their efficacy is inferior to that of newer targeted therapies and they carry risks of renal, hepatic and other toxicities.^{102,103} Topical and low dose oral minoxidil are also commonly used, typically as adjunctive therapy in combination with immune-modulating approaches.¹⁰⁴ Limited discussion has centered around topical phosphodiesterase-4 inhibitors such as apremilast, which decrease proinflammatory cytokines and have shown moderate efficacy in smaller studies. However, evidence supporting their use in AA remains insufficient.⁹⁹

Traditional Therapies for Atopic Dermatitis

In patients with mild AD, treatment focuses on reducing exposure to aggravating factors and minimizing flare frequency. For mild-to-moderate AD, topical corticosteroids have long been the cornerstone of therapy, but the emergence of non-steroidal topical therapies has expanded treatment options. These include topical calcineurin inhibitors (pimecrolimus and tacrolimus), topical phosphodiesterase inhibitors (crisaborole and roflumilast), topical JAK inhibitors (ruxolitinib), and topical aryl hydrocarbon receptor modulators (tapinarof). Phototherapy with narrowband UVB is another therapeutic option,¹⁰⁵ while historically, systemic immunosuppressants (eg. methotrexate, cyclosporine, azathioprine, mycophenolate

mofetil) were used for moderate-to-severe AD,^{106,107} though the therapeutic landscape has shifted substantially with the development of targeted biologics and small-molecule inhibitors.

JAK Inhibitors

The clinical efficacy of JAK inhibitors across both AA and AD provides perhaps the strongest validation of JAK-STAT signaling as a key shared pathogenic axis underlying both conditions. JAK inhibitors are an established class of drugs with significant therapeutic overlap for both AA and AD. JAKs are a family of non-receptor tyrosine kinases that transmit signals from the cell membrane to the cell nucleus via STAT proteins.⁹³ Binding of cytokines to extracellular receptor domains induces JAK activation, making these kinases central regulators of cytokine-induced signal transport.⁹³ Unlike biologics which are often subcutaneous or IV injections, most JAK inhibitors are administered orally.⁹³ While delgocitinib 2% cream was recently approved for moderate to severe chronic hand eczema (CHE),¹⁰⁸ large clinical trials of delgocitinib ointment,⁹⁵ ruxolitinib 1.5% cream,⁹⁶ and tofacitinib 2% ointment⁹⁷ in AA failed to demonstrate marked clinical efficacy. All JAK inhibitors carry a black box warning for increased risk of major adverse cardiovascular events (MACE), venous thromboembolism, blood clots, severe infection, malignancy, and overall mortality as a result of the ORAL Surveillance trial.¹⁰⁹ This randomized post-authorization safety study evaluated the side effects and safety profile of tofacitinib in rheumatoid arthritis patients ≥ 50 years old with at least one additional cardiovascular risk factor.¹⁰⁹ It is worth noting that AA and AD patients treated with JAK inhibitors tend to be younger, often lack the cardiovascular comorbidities required for ORAL Surveillance enrollment, and are frequently not on concurrent methotrexate — factors that substantially limit direct extrapolation of these safety signals to dermatologic practice. When compared to dupilumab in AD patients, JAK inhibitors were associated with a greater risk of skin infections, herpes simplex, herpes zoster, anemia, neutropenia, thrombocytopenia, hyperlipidemia, and acneiform eruptions.¹¹⁰ However, risk of severe adverse events such as malignancy, VTE, or major adverse cardiovascular events were comparable between JAK inhibitor and dupilumab groups.¹¹⁰ Accordingly, the American Academy of Dermatology recommends reserving JAK inhibitors for patients who have failed or are not candidates for other systemic therapies and advises caution in patients with cardiovascular risk factors, history of malignancy, or chronic/recurrent infections.¹¹¹

JAK inhibitors differ in their selectivity profiles. JAK1-selective agents (upadacitinib, abrocitinib) and JAK3-selective agents (ritlicetinib) primarily suppress inflammatory cytokine signaling, while additional JAK2 blockade may contribute to impaired hematopoiesis, providing a mechanistic basis for anemia which has been reported with pan-JAK inhibitors (tofacitinib, baricitinib).^{112,113} Importantly, however, JAK selectivity is relative rather than absolute: at therapeutic doses, “selective” inhibitors retain meaningful cross-JAK activity, and both clinical efficacy and adverse events likely reflect suppression of multiple JAK-dependent signaling pathways rather than blockade of a single pathway.¹¹⁴ This may partly explain why both JAK1-selective and less-selective agents demonstrate efficacy across AA and AD despite their nominally distinct cytokine targets.

For moderate-to-severe AD, the American Academy of Dermatology recommends the JAK1-selective inhibitors upadacitinib and abrocitinib as second-line systemic therapy for patients who have failed or are not candidates for other systemic agents, including biologics such as dupilumab.¹¹⁵ In the JADE COMPARE trial, a Phase 3 head to head comparing 100 or 200mg abrocitinib to standard dupilumab dosing and placebo in adults with moderate to severe AD, abrocitinib 200 mg once daily demonstrated significantly higher response rates than placebo at week 12: 70.3% of patients achieved EASI-75 compared with 27.1% on placebo, and 48.4% reached an investigator global response (IGA) 0/1 response versus ~4.0% with placebo.¹¹⁶ Neither abrocitinib dose differed significantly from dupilumab for most secondary outcomes at later time points.¹¹⁶ The 200 mg dose showed superior itch relief at week 2 compared with dupilumab.¹¹⁶ A different head to head trial comparing upadacitinib 30mg to dupilumab (HEADS UP) showed that 72.4% of upadacitinib treated patients achieved the week 16 primary endpoint of EASI-75 compared to 62.6% of the dupilumab treated cohort.¹¹⁷ All ranked secondary end points also demonstrated the superiority of upadacitinib vs dupilumab at that time point.¹¹⁷

Focusing on the use of JAK inhibitors in AA, meta-analyses of randomized trials confirm that they are associated with significantly greater odds of achieving $\geq 50\%$ and $\geq 90\%$ improvement in SALT scores compared to placebo, with a favorable safety profile and no increase in severe adverse events over short-term follow-up.^{118,119} Real-world cohort

studies corroborate these findings, showing substantial regrowth in most patients over 6–12 months, although relapse is common after discontinuation.¹²⁰ In particular, the first FDA-approved treatment for AA was baricitinib, a JAK1/JAK2 inhibitor approved in 2022 for adult patients with severe AA.² Ritlecitinib, a JAK3/TEC inhibitor, and deuruxolitinib, JAK1/JAK2-selective, were approved in 2023 and 2024, respectively, with ritlecitinib currently being the only FDA approved JAK inhibitor for patients ≥ 12 years.¹²¹ In the pivotal phase 3 BRAVE-AA1 and BRAVE-AA2 trials, baricitinib (2 mg and 4 mg daily) was evaluated in adults with a Severity of Alopecia Tool (SALT) score ≥ 50 ($\geq 50\%$ scalp hair loss) and a current AA episode lasting > 6 months to < 8 years without spontaneous improvement. The primary outcome was a SALT ≤ 20 at week 36. At week 36, the estimated proportion of patients achieving a SALT ≤ 20 in BRAVE-AA1 was 38.8% with baricitinib 4 mg, 22.8% with 2 mg, and 6.2% with placebo; corresponding rates in BRAVE-AA2 were 35.9%, 19.4%, and 3.3%, respectively. In BRAVE-AA1, baricitinib 4 mg demonstrated a 32.6% improvement over placebo (95% CI, 25.6–39.5), while the 2-mg dose showed a 16.6% improvement (95% CI, 9.5–23.8), with both doses significantly outperforming placebo ($p < 0.001$ for each comparison).¹³ Findings from the BRAVE-AA1 randomized withdrawal study highlight the chronicity of AA: after 52 weeks of baricitinib, approximately 80% of patients who discontinued treatment relapsed by week 152, compared to only 7% of those who continued therapy.¹³ Relapse typically began to manifest around 8 weeks after stopping the drug.¹³ Expert consensus currently recommends considering discontinuation of systemic therapy only after complete and stable regrowth has been maintained for at least 6 months, or when disease activity is sufficiently controlled to allow transition to topical therapy. However, for most patients with severe AA, long-term maintenance therapy is required to sustain regrowth.¹³ A new development in JAK inhibition for both AA (NCT05051761)¹²² and AD (NCT05526222)¹²³ is represented by the compound jaktinib (also known as gecacitinib), a broad JAKi.

Dupilumab

Treatment of moderate-to-severe AD was revolutionized with the approval of dupilumab, the first targeted therapy shown to be effective for AD in 2017.³ Dupilumab is a fully human monoclonal antibody that targets the IL-4R α subunit, thereby inhibiting signaling of both IL-4 and IL-13, which are central drivers of T_H2 inflammation, implicated in AD and other allergic diseases.^{3,4,124,125} By blocking the shared IL-4R α subunit, dupilumab disrupts the downstream inflammatory cascade, reducing T_H2-associated cytokines, chemokines, and markers of epidermal hyperplasia, while promoting restoration of skin barrier function.¹²⁵

Robust evidence from phase 3 randomized controlled trials demonstrates that dupilumab significantly improves clinical signs, symptoms, and quality of life in adults with moderate-to-severe AD inadequately controlled by topical therapies.³ In the SOLO 1 (NCT02277743) and SOLO 2 (NCT02277769) trials, 16 weeks of dupilumab (300 mg subcutaneously weekly or every other week) resulted in a significantly higher proportion of patients achieving clear or almost clear skin and at least 75% improvement in Eczema Area and Severity Index (EASI-75) compared to placebo, with additional reduction in pruritus (an improvement of at least 4 points in patient-reported pruritus score by week 16) and improvement in mental health outcomes (a reduction of at least 4 points in Dermatology Quality of Life Index surveys).^{3,4} Long-term safety of dupilumab is favorable with FDA approval as young as age 6 months.¹²⁶ Dupilumab was also effective in treating anti-TNF- α -induced AD, suggesting a potential role for T_H2 inhibition in patients with AD induced during anti-TNF- α therapies.¹²⁷ Molecular studies confirm that dupilumab progressively normalizes lesional skin transcriptomes and suppresses systemic type 2 biomarkers,⁵¹ a finding that provides a biological rationale for its potential utility in AA. Indeed, initial case series showed that AD patients with AA treated with dupilumab not only experienced improvement in their skin, but also significant hair regrowth.¹²⁸ This paved the way for current clinical trials investigating the potential benefit of dupilumab in treating AA in both adult and pediatric patients (NCT03359356).¹²⁹ Thus far, AA patients with and without concomitant AD have shown improvement in SALT scores in response to dupilumab.¹²⁹ Interestingly, responders were more likely to have personal or familial atopy and/or high serum total IgE levels compared to non-responders. A possible explanation to these observations may be that increased levels of serum IgE are indicative of T_H2-associated inflammation in the hair follicle, which may respond better to T_H2-targeting approaches.¹²⁹

Dupilumab paved the way for more biologics in AD targeting the type 2 inflammatory axis that may also show some potential in treating AA. Two anti-IL-13 antibodies, tralokinumab and lebrikizumab, have shown efficacy in AD and

were recently FDA approved in June 2021 and September 2024 for tralokinumab and lebrikizumab, respectively.^{52,53,130} In phase 3 clinical trials, 16–22% of patients on tralokinumab achieved an IGA of 0–1 (indicating clear or almost clear skin) compared to 7–10% in placebo,¹³¹ while 33–43% of patients on lebrikizumab achieved an IGA of 0–1 compared to placebo rates of 10–12%.¹³² Although a few reports have been published describing cases of AA successfully treated with tralokinumab,^{133,134} a Phase 2 randomized, double-blind, placebo-controlled pilot study (NCT02684097) on adults affected by moderate to severe AA did not show significant clinical efficacy.¹³⁵ While the underlying mechanistic explanation for the failure of tralokinumab in AA remains unclear, one possibility is that IL-13 blockade alone is insufficient in AA, and that dupilumab's broader dual IL-4/IL-13 blockade, combined with patient enrichment for atopic background and elevated IgE, may represent necessary conditions for type 2–targeted efficacy in AA. Alternatively, the dose that demonstrated efficacy in AD (300 mg q2weeks) may be insufficient to adequately suppress the more complex inflammatory milieu of AA.

Investigative Treatments

Building on the approved therapies discussed above, and given the overlapping immunologic mechanisms underlying AD and AA pathogenesis, the therapeutic landscape for agents capable of treating both conditions is highly promising. Anti-OX40/OX40L agents are among the most advanced investigational systemic therapies for moderate-to-severe AD (Table 1).^{136,137} A phase 2b trial of amlitelimab (anti-OX40L) demonstrated significant reductions in Eczema Area and Severity Index (EASI) scores, with durable clinical responses persisting for months after treatment cessation, suggesting potential disease-modifying effects,¹³⁶ and a recent phase 3 trial (COAST 1) met all primary (IGA of 0–1)

Table 1 Investigational Therapies for Atopic Dermatitis

Drug	Mechanism	Phase	Trial (NCT)	Age	Primary Endpoint	Efficacy Results	Safety
Rocatinlimab ^a ;	Anti-OX40 mAb	3	ROCKET-IGNITE (NCT05398445); ROCKET-HORIZON (NCT05651711)	≥12 years	vIGA-AD 0/1 (≥2-point reduction) and EASI-75 at week 24	Met: IGNITE EASI-75 42.3% vs 12.8% (PBO); vIGA 0/1 23.6% vs 8.7%. HORIZON EASI-75 32.8% vs 13.7%; vIGA 0/1 19.3% vs 6.6% (P<0.001)	Pyrexia, chills, headache most common; GI ulceration <1%
Amlitelimab	Anti-OX40L mAb	3	COAST 1 (NCT06130566)	≥12 years	vIGA-AD 0/1 (≥2-grade improvement) and EASI change at week 24	Met: vIGA-AD 0/1 21.1% (Q4W), 22.5% (Q12W) vs 9.2% (PBO), P<0.01; EASI-75 35.9% (Q4W), 39.1% (Q12W) vs 19.1% (PBO), P<0.001	Well tolerated; ISRs 2.2% vs 0.7% (mild); pyrexia 1.1% vs 0.7%
Rezpegaldesleukin ^b	IL-2 pathway agonist (Treg selective)	2b	REZOLVE-AD (NCT06136741)	≥18 years	Mean % improvement in EASI at week 16	Met: EASI-75 up to 62% (extended dosing to week 24); vIGA 0/1 28–38%	TEAEs 80% vs 58% (PBO); ISRs, pyrexia, nasopharyngitis; no conjunctivitis or increased infections

(Continued)

Table 1 (Continued).

Drug	Mechanism	Phase	Trial (NCT)	Age	Primary Endpoint	Efficacy Results	Safety
Etrasimod	Oral S1P1,4,5 receptor modulator	2b	ADVISE (NCT04556734)	18–70 years	% change in EASI from baseline at week 12	Not met: –57.2% (2 mg) vs –48.4% (PBO), P=0.18; vIGA 0/1 29.8% vs 13.0%, P=0.045	TEAEs 59.6% (2 mg) vs 47.8% (PBO); nausea, constipation, UTI, back pain, dizziness; no SAEs
Jaktinib	Broad-spectrum JAK inhibitor (JAK1/ JAK2/ JAK3/ TYK2)	2/3	NCT04539639 (Phase 2); NCT05526222 (Phase 3)	≥ 18 years	Proportion achieving EASI-50 at week 12	Met (50 mg): 80.9% (50 mg BID) vs 54.5% (PBO), P=0.016	All AEs Grade 1–2
ZPL-3893787	Histamine H4 receptor antagonist	2a	NCT02424253	18–65 years	% EASI change from baseline at week 8	Met: –50% (drug) vs –27% (PBO); placebo-adjusted EASI reduction 5.1, P=0.01	Well tolerated; 2 discontinuations (headache, somnolence); no SAEs
Bempikibart	IL-7R α antagonist	2a	SIGNAL-AD (NCT05509023)	≥ 18 years	% change in EASI from baseline at week 14	Not met: 74% improvement (drug) vs 76% (PBO)	No SAEs or Grade \geq 3 treatment-related AEs

Notes: ^aMonotherapy in IGNITE/HORIZON; TCS/TCI permitted in SHUTTLE. ^bBiologic/JAK inhibitor-naïve patients.

Abbreviations: AEs, adverse events; BID, twice daily; EASI, Eczema Area and Severity Index; EASI-50/75, 50%/75% improvement in EASI from baseline; GI, gastrointestinal; IL, interleukin; ISRs, injection site reactions; JAK, Janus kinase; mAb, monoclonal antibody; NCT, National Clinical Trial identifier; OX40/OX40L, tumor necrosis factor receptor superfamily member 4/ligand; PBO, placebo; Q4W/Q12W, every 4 weeks/every 12 weeks; S1P, sphingosine-1-phosphate; SAEs, serious adverse events; TCI, topical calcineurin inhibitors; TCS, topical corticosteroids; TEAEs, treatment-emergent adverse events; Treg, regulatory T cell; UTI, urinary tract infection; vIGA-AD, validated Investigator's Global Assessment for Atopic Dermatitis.

and key secondary endpoints (at least a 4 point reduction in patient reported pruritus score).¹³⁸ Phase 3 trials for rocatinlimab (anti-OX40) in moderate-to-severe AD (ROCKET, ROCKET HORIZON) also recently showed statistically significant improvement in IGA and EASI-75 at 24 weeks compared to placebo.^{137,139} Since it has been established that blocking OX40 signaling may dampen autoreactive T-cell expansion, reduce inflammatory cytokine production, and restore immune tolerance at the hair follicle,¹⁴⁰ amltelimab is also currently in phase 2 trials for severe AA (NCT06444451) as the concept of tempering pathogenic T cell responses could also apply to AA.¹⁴¹

As discussed above, the role of IL-2 in the immunopathogenesis of both AA and AD makes this cytokine an appealing therapeutic target, supporting the development of novel IL-2–modulating agents for both conditions (Table 2). The Treg-selective IL-2 receptor agonist repegaldesleukin is currently the most extensively studied molecule within this pharmacologic class. While current biologics like dupilumab focus on neutralizing the effects of IL-4 and IL-13, repegaldesleukin is an agonist that stimulates the expansion and activity of regulatory Tregs. Repegaldesleukin demonstrated a favorable safety and tolerability profile, with consistent pharmacokinetics across dosing cohorts (AD, psoriasis).⁸³ In the AD cohort, patients receiving the higher dose achieved a mean 83% improvement in EASI score after 12 weeks.⁸³ Notably, EASI scores were sustained for up to 36 weeks post-treatment discontinuation in a majority of week 12 responders (71% and 80%, respectively).⁸³ A phase 2b trial for this drug in AD (REZOLVE-AD) was recently completed with significant reductions in EASI scores after 16 weeks of therapy.¹⁴²

Table 2 Investigational Therapies for Alopecia Areata

Drug	Mechanism	Phase	Trial (NCT)	Age	Primary Endpoint	Efficacy Results	Safety
Dupilumab	Anti-IL-4R α mAb (blocks IL-4/IL-13)	2a/2	NCT03359356 (adult); NCT05866562 (pediatric)	≥ 18 years; 6–17 years	Change from baseline in SALT at week 24	Prevented worsening vs PBO (SALT +2.2 vs -6.5, $P < 0.05$); week 48: SALT30 32.5%, SALT50 22.5%, SALT75 15%	No new safety signals; consistent with known profile
Rezpegaldesleukin ^a	IL-2 pathway agonist (Treg selective)	2b	REZOLVE-AA (NCT06340360)	18–70 years (F); 18–60 years (M)	Mean % reduction in SALT at week 36	Dose-dependent effect; SALT ≤ 30 , SALT ≤ 20 , SALT ≤ 10 responses observed (topline Dec 2025)	Consistent with prior trials; details pending
Tralokinumab	Anti-IL-13 mAb	2	NCT02684097	18–75 years	Change in gene expression (Th2, Th22, Th1, Th17) at week 24	Recruiting/ongoing	Recruiting/ongoing
Amlitelimab	Anti-OX40L mAb	2/3	NCT06444451	≥ 18 years	Change from baseline in SALT at week 36	Recruiting/ongoing	Recruiting/ongoing
EQ101	Trispecific inhibitor targeting common IL-2/IL-9/IL-15 subunit	2	NCT05589610	18–60 years	% meeting SALT ≤ 20 at week 24	20% achieved SALT ≤ 20 at week 24; 29% of moderate-severe patients achieved SALT ≤ 20	No SAEs; 98.9% AEs Grade 1–2; 2 Grade 3 (lymphocytopenia, fatigue)
Bempikibart	IL-7R α antagonist (blocks IL-7/TSLP)	2a	SIGNAL-AA (NCT06018428)	18–75 years	Mean relative % change in SALT at week 24	16% SALT reduction vs 2% (PBO) at week 24; SALT ≤ 20 responses through week 36	Well tolerated; no SAEs or Grade ≥ 3 treatment-related AEs
HCW9302	IL-2 fusion protein (Treg selective)	1	NCT07049328	18–70 years (F); 18–60 years (M)	Safety (Phase 1)	Recruiting/ongoing	Recruiting/ongoing
Etrasimod ^b	Oral SIPI _{4,5} receptor modulator	2	NCT04556734	18–70 years	% change in SALT from baseline at week 24	Not met: -21.4% (3 mg), -13.8% (2 mg), +0.35% (PBO)	TEAEs 67.7–80.0%; consistent with known profile

(Continued)

Table 2 (Continued).

Drug	Mechanism	Phase	Trial (NCT)	Age	Primary Endpoint	Efficacy Results	Safety
Rosnilimab	PD-1 agonist	2a	AZURE (NCT05205070)	18–75 years	Change from baseline in SALT at week 24	Not met: No patients achieved SALT ≤ 20 ; development discontinued	Generally well tolerated; no dose-limiting toxicities
Jaktinib	Broad-spectrum JAK inhibitor (JAK1/ JAK2/ JAK3/ TYK2)	3	NCT05051761	18–65 years	% meeting SALT ≤ 20 at week 24	Recruiting/ongoing	Recruiting/ongoing

Notes: ^aJAK inhibitor/biologic-naïve patients. ^bJAK inhibitor-naïve patients.

Abbreviations: AEs, adverse events; F, female; IgE, immunoglobulin E; IL, interleukin; JAK, Janus kinase; M, male; mAb, monoclonal antibody; NCT, National Clinical Trial identifier; OX40L, tumor necrosis factor receptor superfamily member 4 ligand; PBO, placebo; PD-1, programmed cell death protein 1; SIP, sphingosine-1-phosphate; SAEs, serious adverse events; SALT, Severity of Alopecia Tool; SALT $\leq 20/30$, SALT score of 20/30 or less; SALT30/50/75, 30%/50%/75% improvement in SALT from baseline; TEAEs, treatment-emergent adverse events; Th1/Th2/Th17/Th22, T helper cell subtypes 1/2/17/22; Treg, regulatory T cell; TSLP, thymic stromal lymphopoietin.

Rezpegaldesleukin is also currently being studied in a phase 2b trial (NCT06340360) for severe to very severe AA in patients aged 12 years and older.¹⁴³ After 36 weeks of therapy, patients on rezpegaldesleukin achieved a mean reduction in SALT of 28.2% - 30.3% compared to 11.2% in placebo, providing promising results for future phase 3 trials.¹⁴⁴ A novel IL-2 fusion protein (HCW9302) has entered Phase 1 trial for moderate-to-severe AA as of August 2025 (NCT07049328).¹⁴⁵ This molecule selectively engages IL-2 receptors predominantly expressed on regulatory T cells, functioning similarly to rezpegaldesleukin to enhance Treg-mediated immune regulation. Notably, IL-2 has been incorporated as one of the targets in a tri-specific antibody currently under investigation for AA, reflecting how the simultaneous blockade of multiple immunologic pathways is emerging as a new therapeutic frontier. EQ101 is a first-in-class tri-specific inhibitor for AA that targets the common receptor subunit shared by IL-2, IL-9, and IL-15, thereby selectively blunting those cytokines' activity without broad JAK inhibition.¹⁴⁶ This strategy aims to more precisely inhibit pathogenic T cell signals in AA, particularly IL-15-driven cytotoxic T cells, with IL-2 and IL-9 contributing to T cell and mast cell activation. Preclinical data showed EQ101 reversed alopecia in a mouse model more effectively than ruxolitinib, and phase 2 data has demonstrated favorable safety, tolerability, and efficacy in adults.¹⁴⁶ However, the efficacy of EQ101 for AD has yet to be examined and its specific targets may not address the pathogenic mechanisms underlying AD.

Another compound of interest is bempikibart (ADX-194), a monoclonal antagonist of the IL-7 receptor α -chain. IL-7 expression is critical for T cell survival and expansion, and blockade of IL-7R α suppresses inflammatory T cell responses while relatively sparing Treg populations.¹⁴⁷ In AA, IL-7R α inhibition halted disease progression and even reversed early-stage disease in murine models, underscoring the relevance of IL-7 signaling in AA pathogenesis.¹⁴⁷ Although IL-7 is not considered a primary driver of AD, IL-7 and TSLP share structural similarities and both signal through the IL-7R α subunit.¹⁴⁸ This overlap provides a mechanistic rationale suggesting that IL-7R α -mediated pathways may influence immune dysregulation in AD as well. Thus, IL-7 signaling may have pathogenic relevance in both AA and AD by modulating the balance between inflammatory T-cell activity and regulatory control. For these reasons, bempikibart has advanced into Phase 2 development for both AA and AD. In AA, early efficacy was observed in the Phase 2a SIGNAL-AA trial (NCT06018428), where adults treated over 24 weeks, with follow-up through 36 weeks, registered mean SALT reductions of 16% at Weeks 24 versus 2% with placebo, and 9% of treated patients achieved SALT ≤ 20 compared with 0% in the placebo group.¹⁴⁹ In AD, the two-part Phase 2a SIGNAL-AD study (NCT05509023) evaluated subcutaneous dosing every two weeks in adults with moderate-to-severe AD. In Part A (n=15), mean EASI improvements at Week 14

were 58% with 2 mg/kg, 84% with 3 mg/kg, and 72% pooled, versus 38% for placebo. However, in Part B (n=106), bempikibart 200 mg Q2W produced a 74% mean EASI improvement at Week 14, essentially identical to placebo (76%), and thus the primary endpoint was not met. This failure may be due to inadequate study design given the high placebo response rate in Part B. Post-hoc analysis suggested some efficacy in the bempikibart compared to control group, but further studies will be needed to prove efficacy or lack thereof. The drug was well-tolerated with no serious or Grade \geq 3 treatment-related adverse events. As a result, current development is now primarily focused on AA rather than AD.¹⁵⁰

Antihistamines and histamine receptor antagonists may also play a dual role in AA/ AD. The histamine H4 receptor on eosinophils and T cells mediates itch and inflammation. A drug candidate, ZPL-3893787, an H4 antagonist, reached Phase 2 in AD and showed a reduction in eczema severity and itch compared to placebo.¹⁵¹ There is some clinical evidence for using antihistamines in alopecia areata as an adjunctive therapy, consistent with the concept of an allergic or atopic immune component in AA.^{152–154}

Finally, several early (Phase 1 and Phase 2) investigational agents with novel mechanisms of action are also being explored. Although these molecules have yet to establish clinical efficacy, they may help refine future therapeutic targets. Among them, etrasimod, a sphingosine 1-phosphate (S1P) receptor modulator that retains lymphocyte subsets within skin-draining lymph nodes to limit migration into inflamed tissue,¹⁵⁵ showed a statistically significant decrease in IGA (29.8%) compared to placebo (13%) in a 12-week phase 2b trial for AD.¹⁵⁶ On the other hand, it failed to meet primary and secondary endpoints at week 24 in a Phase 2, randomized, double-blind, placebo-controlled trial evaluating the efficacy and safety of etrasimod in adults with moderate-to-severe AA (NCT04556734).¹⁵⁷ While any mechanistic explanations remain speculative given lack of published data, this failure may reflect the inability of S1P-mediated lymphocyte retention in lymph nodes to adequately address the established tissue-resident memory T cell infiltrate in AA, which by definition is not recirculating and therefore not susceptible to this mechanism of action. Similarly, rosnilimab, a programmed death (PD) 1 agonist that was examined recently in phase 2 studies for AA, failed to meet primary or secondary endpoints in both groups.¹⁵⁸ Rosnilimab's failure may reflect downregulation or dysfunction of PD-1 on chronically activated perifollicular CD8+ T cells in established AA, limiting the efficacy of checkpoint-based suppression in this context, or may alternatively reflect insufficient trial power. Nevertheless, its mechanism of action as an immune checkpoint agonist, potentially broadly decreasing T cell overactivity, is a novel approach to temper the T-cell mediated immune response seen in AA and AD.

Conclusion

Once regarded as immunologically distinct disorders, AD and AA are now understood to share convergent inflammatory pathways — centrally, JAK-STAT dependent cytokine dysregulation and Treg insufficiency/dysfunction — that together explain both their epidemiologic association and their emerging therapeutic overlap. There are several important limitations of the current evidence base. Much of the data supporting shared mechanisms derives from transcriptomic data. Agents effective in one disease may not adequately address the full pathogenic architecture of the other. Critically, the shared mechanism framework appears most robust for the atopic-associated AA endotype, wherein T_H2 dysregulation plays a prominent pathogenic role, and less so for non-atopic AA, which is characterized by predominantly T_H1-skewed inflammation. As the field advances, endotype-stratified approaches, selecting therapies based on atopic background, serum IgE, and transcriptomic immune signatures, represent the most promising path toward personalized, potentially disease-modifying interventions for both AA and AD. Continued advances in our understanding of hair follicle immune privilege, T-cell dysregulation, and type 1 and type 2 immune circuits are rapidly reshaping the treatment landscape. Emerging shared therapeutic strategies, ranging from JAK inhibitors to OX40/OX40L blockade and IL-2–modulating compounds, point toward a future in which interventions are not merely immunosuppressive but restore immune homeostasis and potentially modify disease course.¹⁵⁹

Data Sharing Policy

Data sharing is not applicable to this article as no data were created or analyzed in this study.

Author Contributions

A.T.: Conceptualization, Data curation, Writing – original draft, Writing – review & editing, Visualization. G.R.: Data curation, Writing – original draft. F.M.M.: Conceptualization, Validation, Writing – review & editing. J.L.: Data curation, Writing – original draft. E.D.D.: Conceptualization, Validation, Writing – review & editing, Visualization. B.U.: Conceptualization, Validation, Writing – review & editing, Visualization, Supervision, Project administration.

All authors gave final approval of the version to be published; have agreed on the journal to which the article has been submitted; and agree to be accountable for all aspects of the work.

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