

Mechanistic insights into Ritlecitinib-mediated immunomodulation in Alopecia Areata

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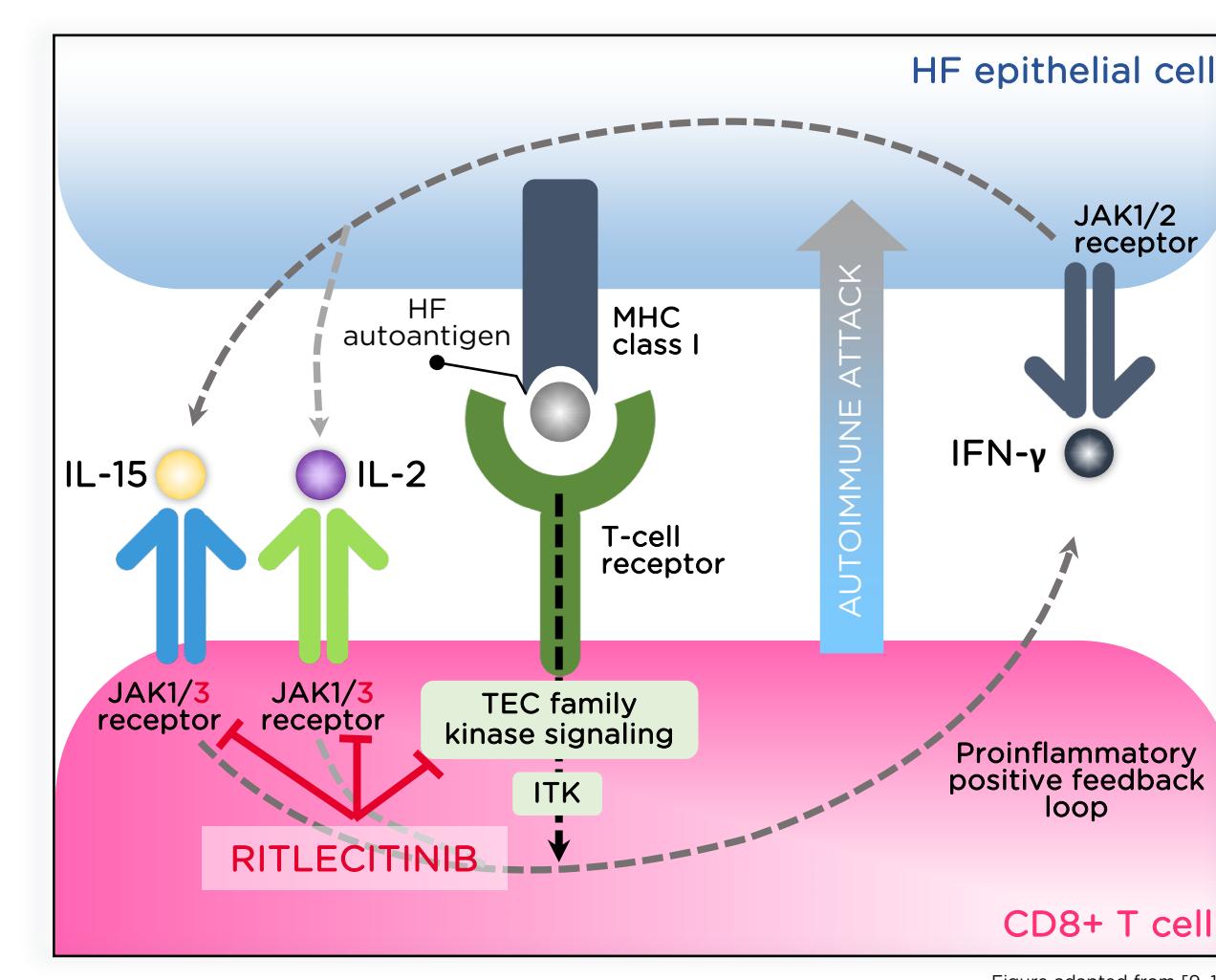
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Background

Alopecia areata (AA) is an autoimmune-mediated inflammatory hair loss disorder in which IFN γ , secreted by CD8+ T-cells, exacerbates peri- and intrafollicular inflammation and induces hair follicle (HF) immune privilege (IP) collapse, characterized by increased expression of MHC class I and II, downregulation of IP guardians and increased expression of pro-inflammatory mediators [1-3]. T-cell activation is regulated by T-cell receptor and IL-2 receptor stimulation and subsequent downstream induction of JAK3 and TEC family kinases (Figure on the right) [4]. The clinical relevance of JAK3/TEC family kinases signalling in AA has been demonstrated by the approval of the JAK3/TEC family kinases inhibitor ritlecitinib for the treatment of severe AA [5].

Aim of the Study

Here we analysed how inhibition of JAK3/TEC family kinases signalling by Ritlecitinib affects alopecia areata pathogenesis in human HFs induced with an alopecia areata-like phenotype by activation of JAK3/TEC family kinases signalling, and in alopecia areata lesional skin *ex vivo*.

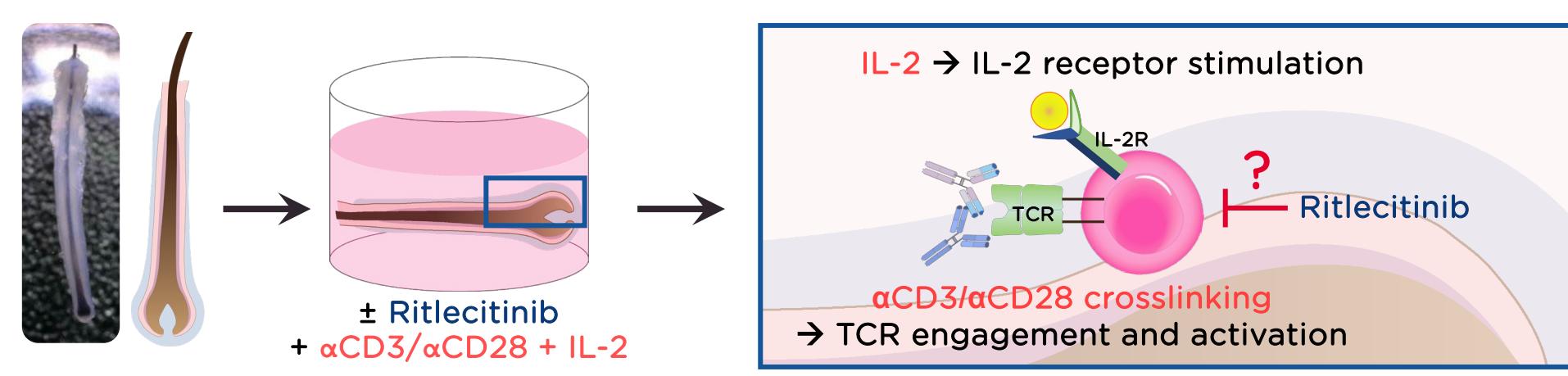


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Engagement of the T-cell (TCR) and IL-2 (IL2R) receptors induce a positive proinflammatory feedback loop via activation of the interleukin-1-inducible T-cell kinase (ITK) and the JAK3/TEC receptor complex [6]. This activation enhances the inflammatory milieu in AA receptor. Furthermore, TCR and IL-2R activation regulate receptor, activation, differentiation and maintenance of CD8+ cells. Cytotoxic CD8+ cells contribute to the IP collapse observed in AA-affected hair follicles. The JAK3/TEC inhibitor Ritlecitinib is approved for the treatment of severe AA [5-8].

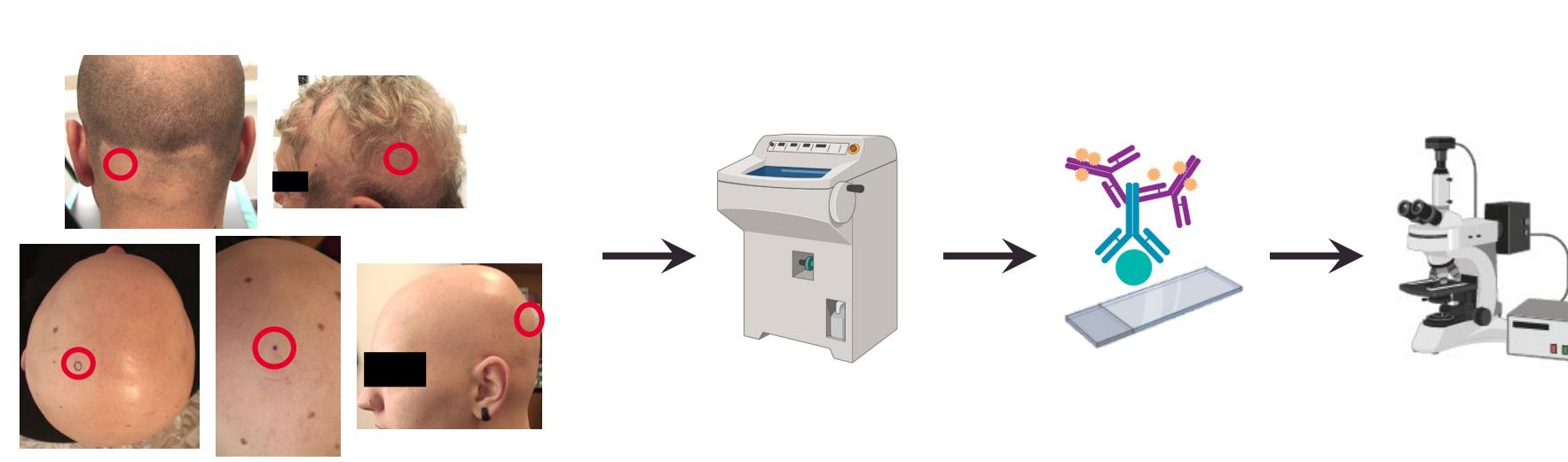
Material & Methods

Experimentally induced AA-like phenotype *ex vivo*

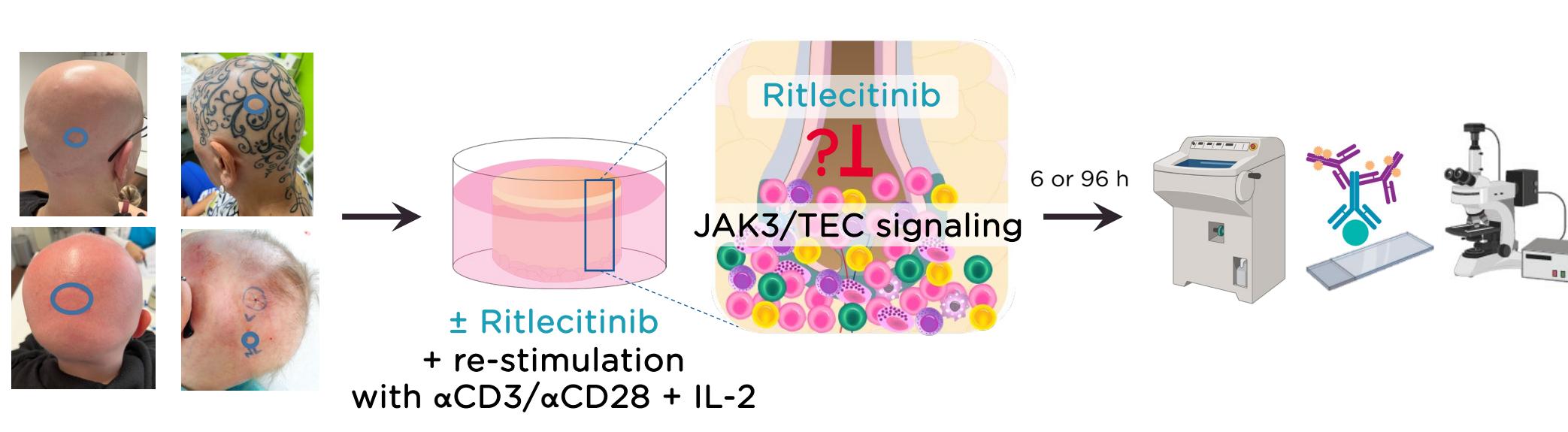


Microdissected, full-length, healthy human hair follicles (HFs) were treated with vehicle control or anti-CD3 and anti-CD28 antibodies (α CD3/ α CD28/IL-2) for 5-8 days *ex vivo* to induce T-cell activation on resident perifollicular T cells via TCR (TEC signalling activation and IL-2 receptor stimulation (JAK3 signalling activation), respectively. 2 μ M Ritlecitinib was added one day prior to the cytokines. Afterwards, CD3+ T-cell numbers and proliferation, immune privilege status and the release of pro-inflammatory and cytotoxic mediators were assessed.

Lesional AA scalp skin *in situ*



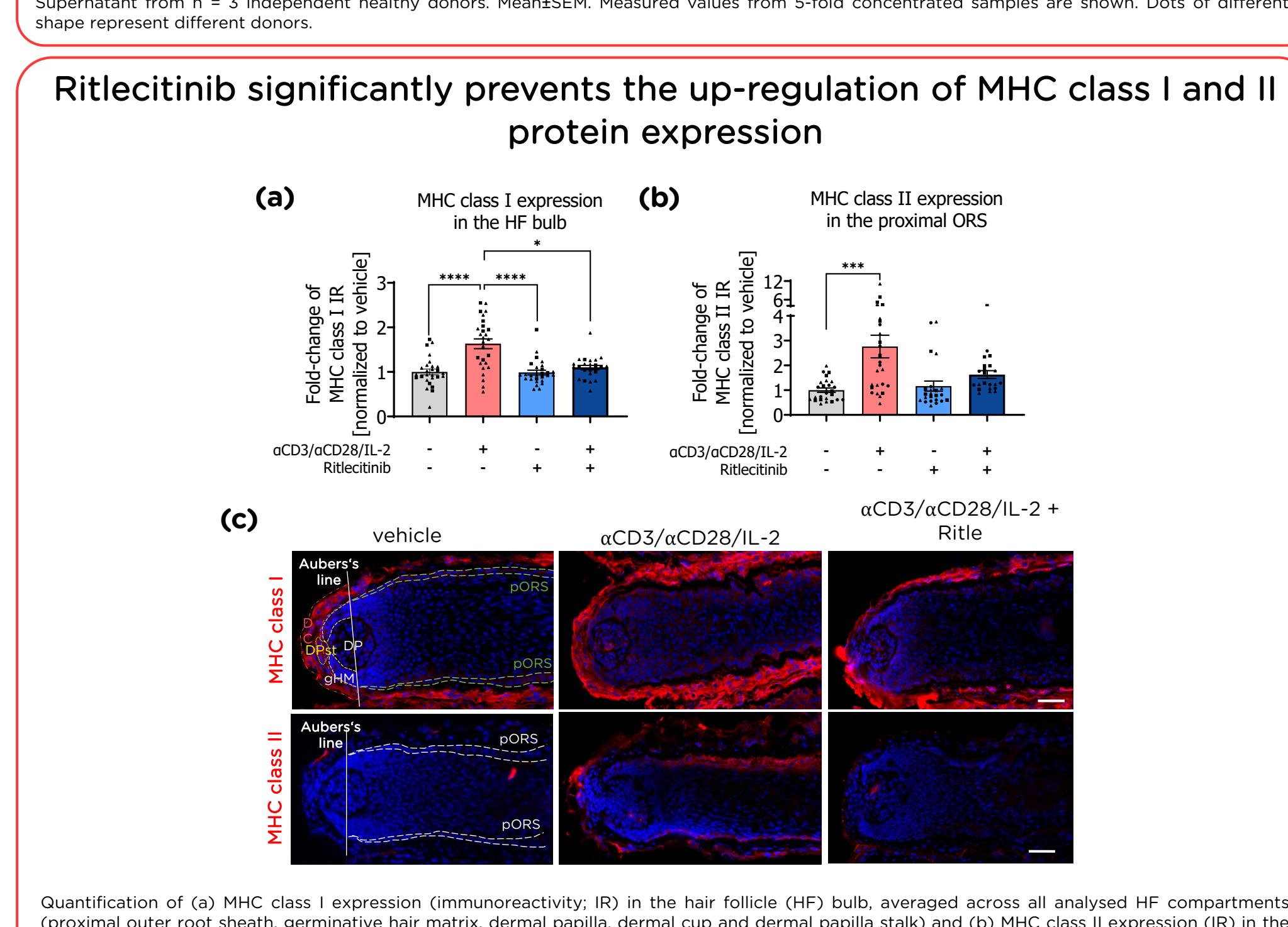
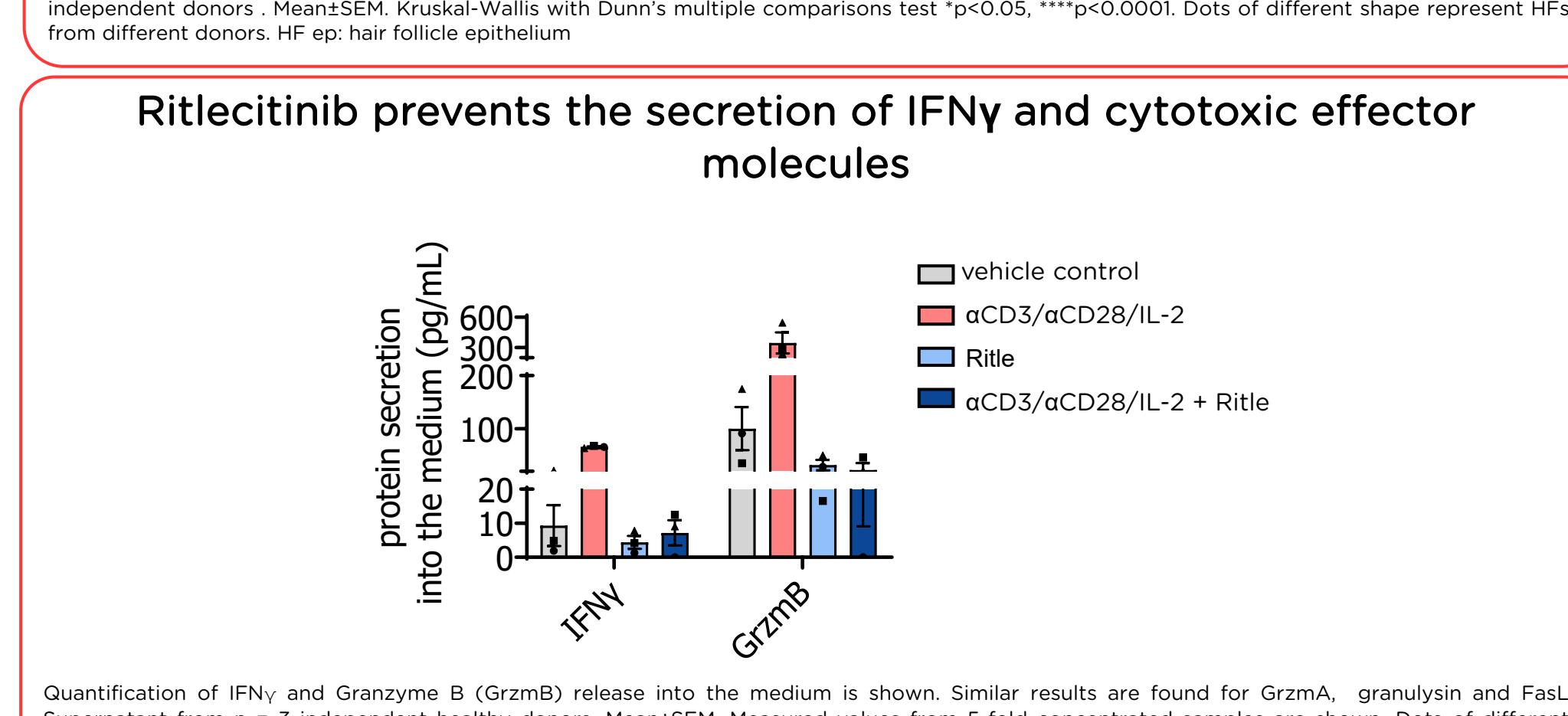
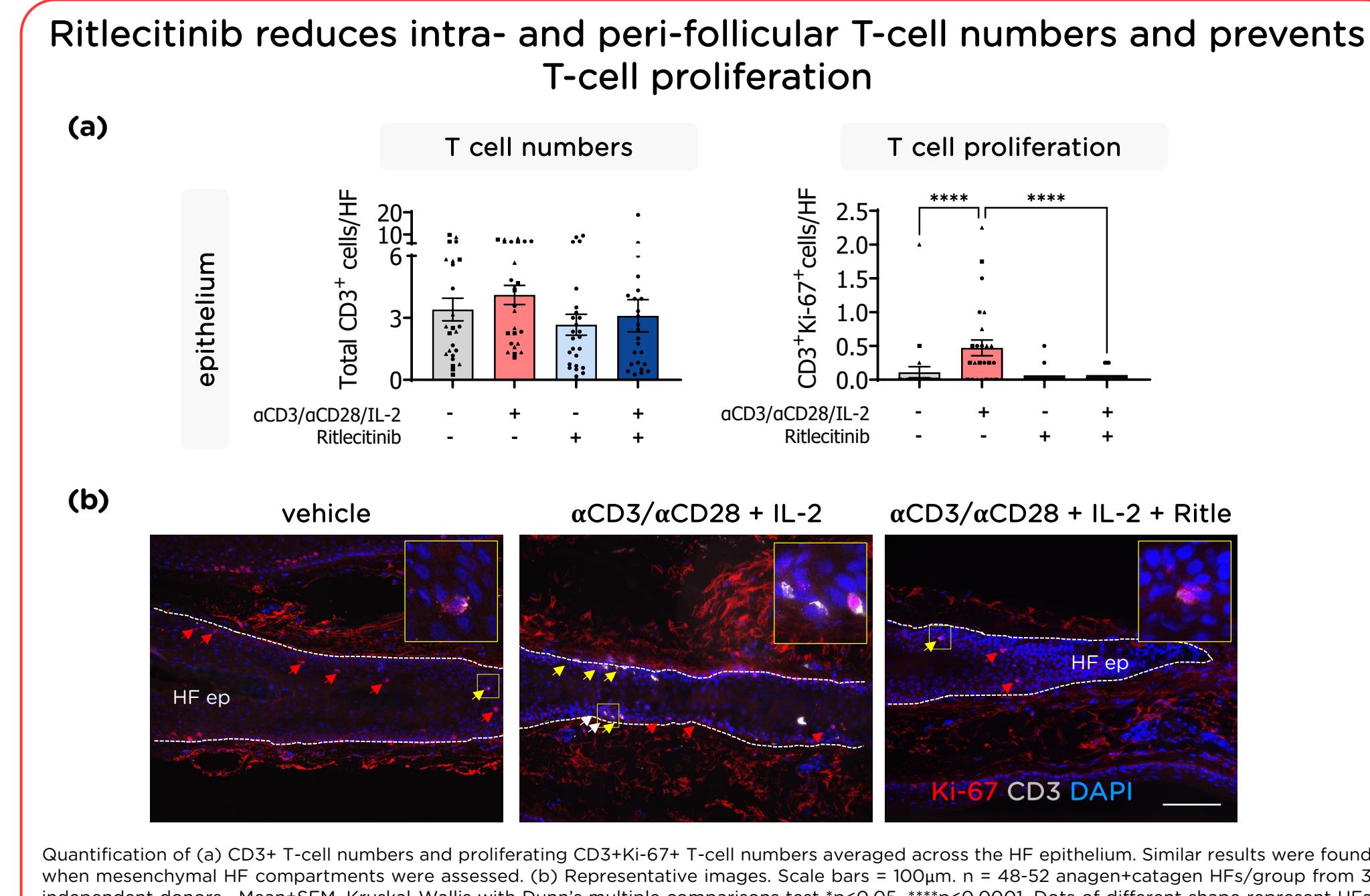
Lesional AA scalp skin *ex vivo* + Ritlecitinib treatment



Chronic lesional AA scalp skin was obtained from 4 patients. Scalp skin was re-stimulated with α CD3/ α CD28 + IL-2 and treated with vehicle control or Ritlecitinib. After 6 or 96 hours of culture, samples were embedded for cryosectioning and quantitative immunofluorescence analysis was performed on activity markers for JAK3 and TEC signalling.

Results

Experimentally induced AA-like phenotype *ex vivo*



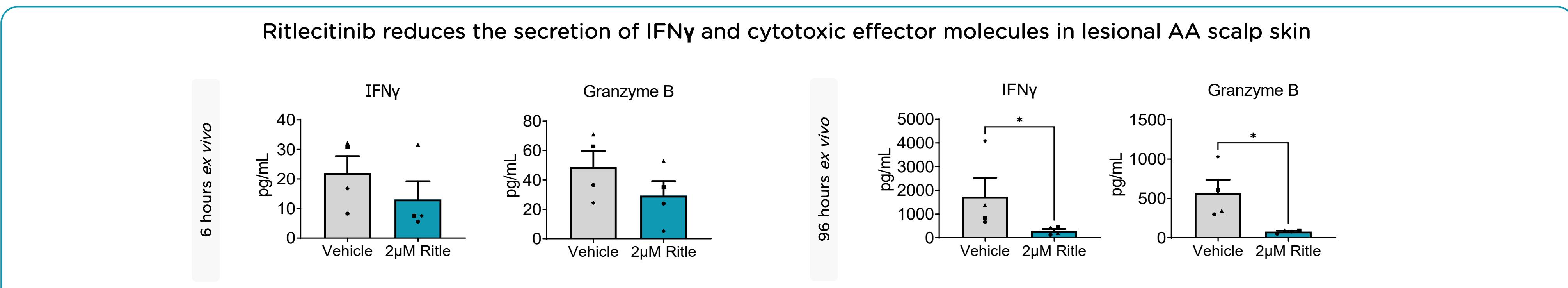
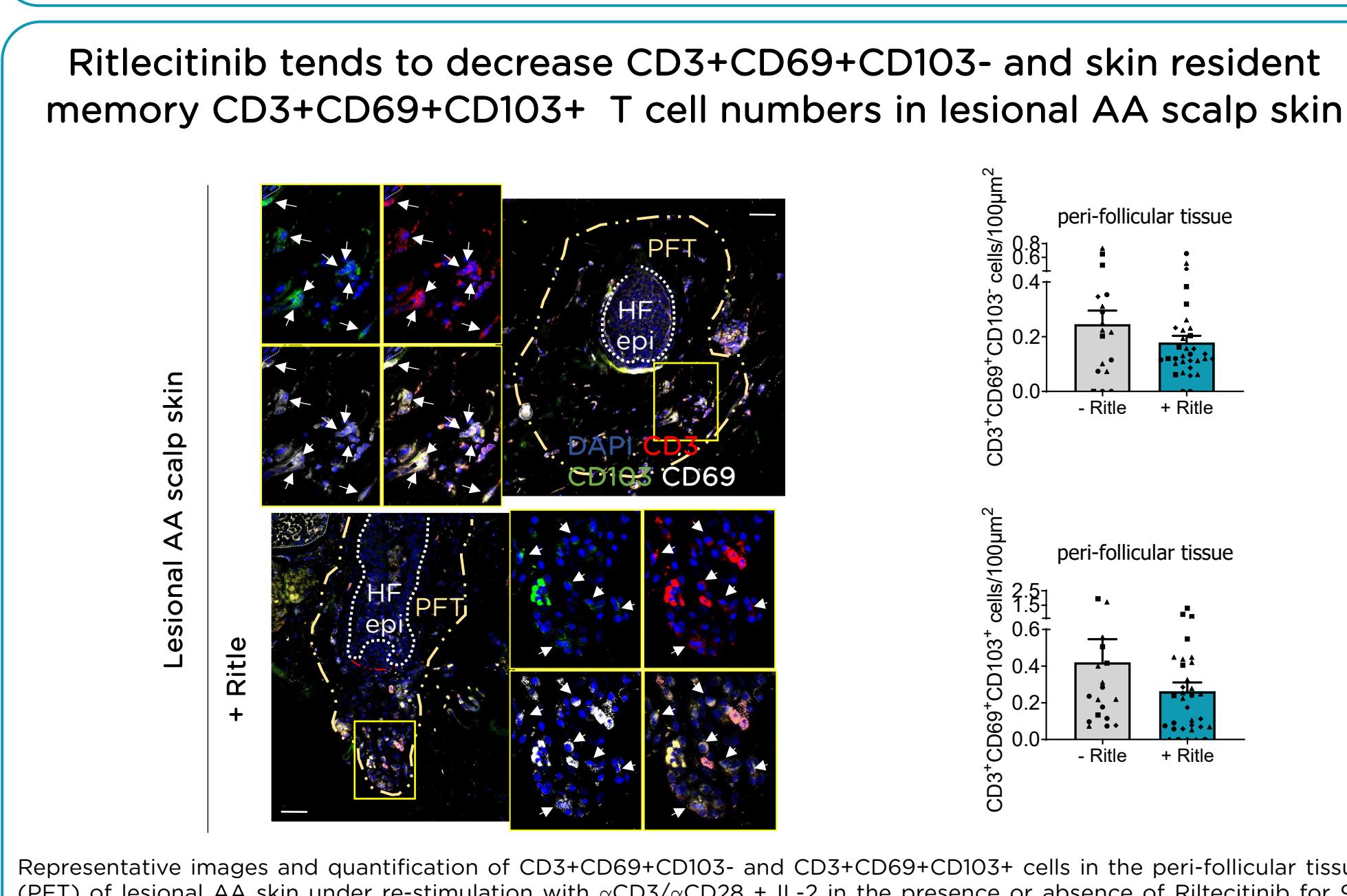
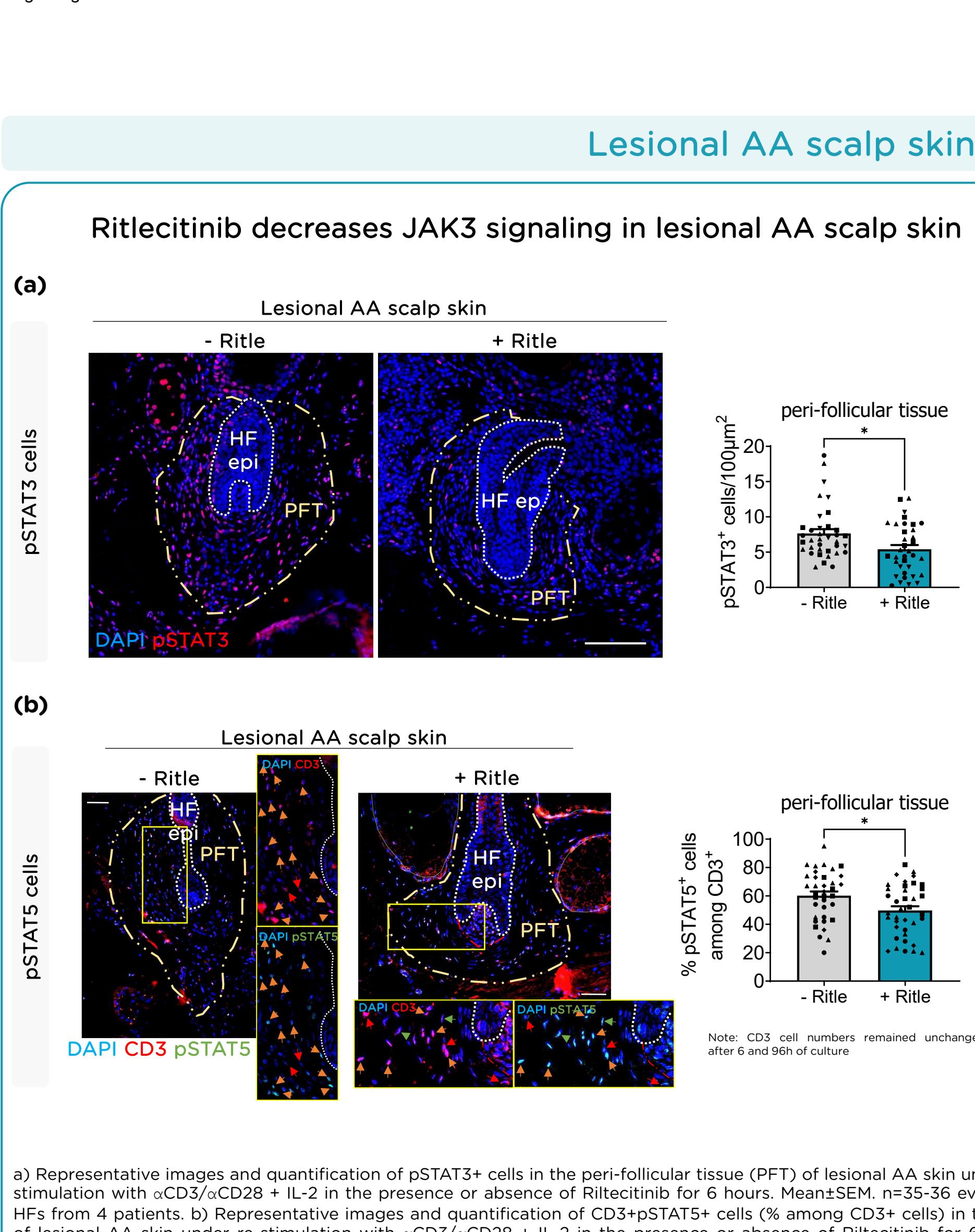
Lesional AA scalp skin *in situ*

JAK3/TEC signaling is active in chronic lesional AA scalp skin

Cells	Healthy biopsies		Typical chronic AA		Chronic AA with active inflammation	
	HF epi	+ surrounding tissue	HF epi	CTS + Pb inf	HF epi	CTS + Pb inf
CD3+ pSTAT5+	-	++	-+	+	++	+++
pSTAT3+	-	+	-+	++	+	+++
pSTAT6+	+	-+	++	-+	++	+
IRF4+	+	-+	+	-+ / +	++	-+ / +
CD8+ GranzB+	-	+	-+	+	-	++

+ very few positive cells + few positive cells ++ positive cells +++ many positive cells +/+ donor dependent

Lesional AA scalp skin *ex vivo* + Ritlecitinib treatment



Quantification of IFN γ and Granzyme B (GrzB) release into the medium from lesional AA scalp skin under re-stimulation with α CD3/ α CD28/IL-2 in the presence of vehicle control or Ritlecitinib for 6 and 96 hours. Similar results are found for IL-17A, IL-6, IL-10, TNFa, FasL, GranzinB and GrzB both after 6 and 96 hours *ex vivo*. Supernatant from n = 4 patients. Mean \pm SEM. Dots of different shape represent patients.

Our data highlight the clinical relevance of targeting JAK3/TEC kinases pathways for the treatment of AA.

References: [1] Bertolini M. et al. *Exp Dermatol*. 2020. [2] Passeron T. et al. *Front Immunol*. 2023. [3] Gilhar A. et al. *N Engl J Med*. 2012. [4] Marchingo J. M. et al. *Science* 2014. [5] King B. et al. *Lancet*. 2022. [6] Nayar R. et al. *PNAS*. 2012. [7] Shah K. et al. *Target Ther*. 2021. [8] Kapnick S. M. et al. *J Immunol*. 2017. [9] Rébéri C. et al. *JAKSTAT*. 2013. [10] Shepers H. et al. *JAKSTAT*. 2012. [11] Karpathiotis G. et al. *Pathol Res Pract*. 2021. [12] Dahabreh D. et al. 2023. [13] Divito S. J. *Kupper TS*. 2014