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Hidradenitis suppurativa: the “bright side” of autoinflammation and hidden diseases

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Hidradenitis suppurativa (HS) is a chronic-relapsing, debilitating inflammatory disease primarily affecting the pilosebaceous unit.¹ It is clinically characterized by recurrent, painful, deep-seated nodules, abscesses and sinus tracts ending in hypertrophic scarring that typically involve apocrine gland-rich areas of the body, notably axillary and inguinoperineal regions.¹ HS is estimated to affect 1% of the population,¹ but, in our opinion, the incidence and prevalence of the disease has significantly increased in the last few years. The pathophysiology of HS is the result of a complex interplay between genetic and environmental factors cross-talking with both innate and adaptive immunity dysfunction, as well-described in the review article by Napolitano et al.,² providing an update on HS pathogenesis. Heterozygous mutations in the γ -secretase genes – presenilin enhancer 2 (PSENEN), presenilin (PSEN1) and nicastrin – were the first reported genetic changes in HS.^{3,4} The above mutations cause inactivation of Notch signaling which is responsible for an altered homeostasis of hair follicle and apocrine gland leading to the production of the so-called damage-associated molecular pattern (DAMP) molecules. These molecules induce an abnormal activation of the inflammasome, a molecular platform triggering the inflammatory process in HS as in the classic monogenic autoinflammatory diseases like familial Mediterranean fever.⁵ The important autoinflammatory component in the pathogenesis of the disease is supported also by the upregulation of interleukin(IL)-1 β ^{6,7}, which is a pivotal cytokine in autoinflammation.⁸ On the other hand, some studies found γ -secretase mutations only in a minority of HS cases,^{9,10} suggesting γ -secretase mutation alone is not sufficient to produce the HS phenotype.¹¹ Interestingly, our group reported mutations involving a number of autoinflammatory genes in the recently described syndromic variant of HS known as PASH (pyoderma gangrenosum, acne, suppurative hidradenitis),^{12,13} giving rise to considering HS a polygenic autoinflammatory condition in which innate immunity dysfunction plays a key role. From an immunological point of view, IL-17, cytokine merging innate and adaptive immunity, has also been reported overexpressed in the lesional skin of HS.¹⁴ Of note, with respect to IL-1 β and IL-17 expression, HS resembles PASH¹⁵ as well as two prototypic neutrophilic dermatoses, pyoderma gangrenosum and Sweet's syndrome,¹⁶ making justified, to include HS in the spectrum of neutrophilic dermatoses,¹⁷ based also on the high number of skin infiltrating neutrophils especially in later stages of the disease.⁷ The interesting article by Pescitelli et al.¹⁸ supports the recent view on HS as a systemic disease linked to several comorbidities, with which HS shares genetic factors, environmental triggers and inflammatory pathways. Reports of elevated circulating levels of tumour necrosis factor (TNF)- α ¹⁹ in HS are in line with systemic inflammatory activation. Obesity and metabolic syndrome are the most common associated conditions in HS patients¹⁸ but inflammatory bowel diseases, particularly Crohn's disease,²⁰ and spondyloarthritis²¹ also occur more frequently in these patients than in the general population. Thus, clinical intervention for HS must include consideration of these comorbidities and complications with a multidisciplinary approach that is outlined also in the article by Veraldi et al.²² Finally, the paper by Lacarrubba et al.²³ reviews and discusses the important contributory role of imaging techniques, particularly ultrasonography.²⁴ Ultrasonography may reveal features not appreciable at clinical examination, notably fistulous tracts, allowing a more accurate staging, treatment planning and monitoring response to therapy in this debilitating disease.

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