Correspondence

Generalized eruptive keratoacanthoma of the Grzybowski type: some considerations on treatment and pathogenesis

Dear Editor,

A 64-year-old male patient presented with a 2-month history of a generalized mucocutaneous eruption characterized by suspected keratoacanthomas (KAs), umbilicated papules, and small follicular papules, which mainly involved the face, upper limbs, and trunk; the oral mucosa was affected as well (Fig. 1a–c).

Histopathologic analysis of several lesions revealed the typical features of KA (Fig. 2a and b). Both the absence of similar cases in the patient’s family and the late onset of these skin tumors made a familial form of KA unlikely. A diagnosis of generalized eruptive keratoacanthoma (GEKA) was made. No concomitant head and neck/oropharyngeal carcinoma was present.

Due to the intense itching, the progressive onset of new lesions involving more and more extensive mucocutaneous areas, and the negative impact on his quality of life, we decided to treat the patient with acitretin 20 mg/day and topical tretinoin 0.05% cream once a day. Marked improvement was observed at the end of the second month of combined treatment (Fig. 1a–c).

GEKA, first described by Grzybowski, is a very uncommon variant of KA with unknown etiology that involves widespread areas of the skin. The course of the disease is chronic, and its response to therapy is usually poor or partial. There are several studies focused on the relationship between human papillomaviruses (HPVs) and KA, especially in the solitary form. On the other hand, only a few studies have addressed the prevalence of viral sequences in multiple KAs. In particular, Forslund et al. and Stockfelt et al. detected the presence of cutaneous HPV DNA by the use of PCR in 20-50% of specimens obtained from keratoacanthomas patients. Strumia et al. found HPV-16 with a high prevalence in KAs and normal skin from immunocompetent individuals. On the contrary, Haas et al. failed to detect HPV-positive biopsy specimens in a patient with a 9-year history of generalized eruptive KA of the Grzybowski type.

In order to assess possible involvement of viral infections in the pathogenesis of this GEKA, we analyzed a skin biopsy of a retroauricular lesion by nested PCR for the presence of HPV together with specific human herpesviruses (HHVs) that have a possible association with skin lesions: the α-herpesviruses (herpes simplex virus types 1 and 2 [HSV-1, HSV-2] and varicella-zoster virus [VZV]), the human β-herpesvirus type 6 (HHV-6), and the Kaposi’s sarcoma-associated γ-herpesvirus (HHV-8).

Isolation of DNA was performed as previously described. We avoided specimen contamination by keeping the DNA extraction area strictly separated from the PCR mixture preparation and sample handling. PCR analysis was performed in triplicate for each set of primers. We used positive (viral genomic DNA) and negative (PCR mixture without DNA template) controls. The sensitivity of the PCR reactions was determined by amplification of known amounts of target sequences (data not shown). To test the absence of major PCR inhibitors, the same amount of DNA used for virus-specific PCR analysis was amplified for human beta actin gene. Nested PCRs for VZV, HHV-6, HHV-8, HSV-1, and HSV-2 were performed as previously described. Because of the low amount of DNA that was available from the skin biopsy, it was necessary to select the HPV genus to be analyzed. HPV genera beta, gamma, mu, and nu are considered to be correlated with benign cutaneous lesions, and only occasionally have they been found in skin cancers. With cutaneous β-HPVs that are ubiquitous in the general population. On the contrary, HPV-6/11, usually associated with mucosa, were frequently found in benign, premalignant, and malignant cutaneous lesions. Since the significance of the presence of HPV 6/11/16 in skin lesions remains unknown, we selected to evaluate these specific HPV types. HPV DNA presence was determined using a nested PCR as previously described. HPV-16 DNA presence was determined using a nested PCR specific for HPV-16-E6 region that we set up within the original amplicon published by Ferre and Garduno and a nested PCR with degenerate primers already described by de Roda Husman et al.

Our data revealed only the presence of HPV-16 DNA among the viruses analyzed (Fig. 2), while HHVs were not present in the biopsy, even using high sensitive nested PCRs. The presence of HPV-16 DNA harbored in GEKA lesions does not allow any conclusive consideration on the etiological role of HPV infection and the role of HPV in keratoacanthomas remains thus elusive. However, its potential involvement in the pathogenesis of this disorder supports the retinoids as the first therapeutic option. Retinoids could be effective in GEKA treatment not only because of their antitumor properties but also by altering keratinization and inhibiting HPV replication and assembly. In keeping with this, an inverse relation has been observed between concentration of retinoids and HPV DNA within infected epithelial cells. We recommend the evaluation of the role of HPV infection in GEKA as meritorious of further investigation.
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Conflict of interest: The authors declare that they have no conflict of interest.

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