CASE REPORT



An Unusual Gingival (Peripheral) Tumor with Features of Keratoameloblastoma with Cytologic Atypia or Possible Malignant Transformation Exhibiting *ARID1A* Mutation

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Abstract

Background Keratoameloblastoma is a poorly characterized and rarely reported odontogenic neoplasm that can exhibit overlapping histopathologic features with conventional ameloblastoma and keratocystic odontogenic tumor (KCOT), with an ambiguous relationship to the so-called solid KCOT.

Methods A peripheral maxillary tumor causing bone saucerization in a 54-year-old male is described and investigated with immunohistochemistry and Next-Generation Sequencing (NGS).

Results Microscopically, the tumor comprised of a predominantly plexiform proliferation of odontogenic epithelium with central keratinization and evidence of surface origin. Peripheral cells exhibited nuclear palisading with variable reverse polarization, while stellate reticulum-like areas were observed internally. A few follicles and a few foci in the lining of cystic spaces revealed increased cellularity with cells exhibiting small but conspicuous nucleoli, focal nuclear hyperchromatism, and a few mitoses mostly seen in the peripheral outer cell layer. Nuclear staining for ki-67 was increased in those areas when compared with the other cystic, follicular, and plexiform areas. These features were interpreted as cytologic atypia suggesting also the possibility of a malignant process. Immunohistochemically, the tumor was positive for CK19 and negative for BRAF VE1, calretinin, and CD56. Ber-Ep4 was only focally positive. By sequencing, an *ARID1A* c.6527_6538delAG frameshift mutation (VAF: 5.8%), classified as likely oncogenic, and an FBXW7 c.1627 A > G missense mutation (VAF: 8.0%), classified as a variant of uncertain significance, were detected. Two mutations, probably germline (VAF~50%), were recorded for *RNF43* and *FBXW7*. No pathogenic variants were identified in *PTCH1, BRAF, NRAS, HRAS, KRAS, FGFR2*, or *SMO* genes. **Conclusion** The significance of an *ARID1A* variant in keratoameloblastoma is uncertain since this variant has not been reported in ameloblastoma or KCOT, to date. Alternatively, it may characterize malignant transformation in the present case since *ARID1A* mutations have been encountered in various cancers. Sequencing of additional cases is necessary to determine whether this may represent a recurrent genomic event.

 $\label{eq:constraint} \begin{array}{l} \mbox{Keratoameloblastoma} \cdot \mbox{Meloblastoma} \cdot \mbox{Keratocystic odontogenic tumor} \cdot \mbox{Odontogenic keratocyst} \cdot \mbox{Carcinoma cuniculatum} \\ \end{array}$

Introduction

Keratoameloblastoma (KA) is a rare odontogenic tumor that is thought to present with overlapping clinicopathologic features with the so-called solid keratocystic odontogenic tumor/odontogenic keratocyst (KCOT/OKC). The distinction between the two has been, at least to some,

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problematic or even impossible while others suggest that it is unnecessary to separate the two entities. There is also a thought that cases of KA represent ameloblastomatous transformation of KCOT/OKC. All the above postulates are discussed in a recent review paper by Liam Robinson et al. [1], where the clinicopathologic characteristics of apparent cases of KA are presented. These authors shared their personal experience of seven examples of KA and add them to the nineteen already published since the first report of the entity is under the name KA by Altini et al. [2]. One should note here that the first reported case is apparently by Ralph Spaulding in 1956 as correctly pointed by Professor Fumio Ide et al. [3].

Besides delineating the controversies in the KA and sKCOT/OKC, the significance of the work by Robinson et al. [1] lies on the fact that it provides evidence in support of KA as a variant of ameloblastoma (AMEL), apart from sKCOT/OKC and another rare variant of AMEL known as papilliferous keratoameloblastoma (PKA). Robinson et al. [1] presented as defining histomorphologic characteristics (a) the presence of variably solid and cystic epithelial follicular proliferations, (b) focal stellate reticulum-like central areas, (c) subnuclear vacuolization in the basal cell/ameloblastic layer, (d) overt and often pronounced lamellated-type central parakeratinization, and (e) occasionally, central areas of calcification [1]. Of interest was also the capricious immunohistochemical negativity of KA for calretinin [1], an immunohistochemical stain expected to be positive in ameloblastomas [1, 4]. There is, however, no reference on the evaluation for $BRAF^{V600E}$ [Mendez et al.] or CD56 immunohistochemical reactivity. These antibodies have shown invariable positive staining for conventional and cystic ameloblastomas.

Progress in the characterization of molecular events of KCOT/OKC and AMEL has disclosed *PTCH1* alterations nearly universally occurring in the former [5] and MAPK pathway alterations, in the absence of *PTCH1* alterations, in almost 90% of cases in the latter [6]. *BRAF*^{V600E} mutant ameloblastoma localizes to the mandible while maxillary ameloblastoma tends to be *BRAF*-wild type and enriched for *SMO* mutations [6, 7].

The mutually exclusive mutational profiles of KCOT/ OKC and ameloblastoma raise the possibility that molecular characterization of KA may inform appropriate classification and treatment, given the more aggressive biologic behavior of ameloblastoma relative to KCOT. In fact, Zhang et al. [8] have tried to separate the two entities at the molecular level by investigating 9 cases diagnosed either as KA or sKCOT/OKC for *PTCH1* and *BRAF* mutations. In their cohort there was lack of *BRAF* mutations and presence of *PTCH1* missense mutation in 1 out of the 9 cases.

The vast majority of KA are, as expected, central tumors. A peripheral example KA has been published [9]. Herein, we report of an unusual peripheral tumor with ostensible features of KA and present some interesting, in our opinion, histopathologic, immunohistochemical, and targeted sequencing findings.

Case Report

Clinical Findings

A 54-year-old male presented with a "tender" swelling in the area of the right maxillary premolars causing saucerization of the bone. A periapical radiograph revealed bone loss characterized by radiolucency with foci of subtle opacities (Fig. 1A). A mucosal flap was raised, and a conservative excision was performed.

Histopathology

At gross examination, the specimen comprised a soft tissue mass with calcified areas measuring $1.5 \times 1.3 \times 1.0$ cm. Upon decalcification, the histologic findings were those of an epithelial mostly cystic tumor of apparent odontogenic origin characterized by interconnecting or plexiform cystic spaces and scattered small follicles both featuring areas of palisading in the peripheral basal cell layer with occasional reverse polarization, infrequent subnuclear vacuolization, and central areas of mostly lamellated keratinization (Fig. 1B–D). The cystic lining was a few layers in thickness akin to KCOT/OKC. A few follicles featured foci of early keratinization in form of keratin within pseudoductal structures (Fig. 1E). Also, in some follicles and in foci of the lining of cystic spaces, cells revealed small but conspicuous nucleoli (Fig. 1F), infrequent nuclear hyperchromatism, and a few mitoses mostly seen in the peripheral outer or basal cell layer. Basal cell hyperplasia was also occasionally noted. The stroma contained abundant keratin, deriving from burst cysts, which was calcified in many areas and accompanied by an intense plasmacytic inflammatory response.

Because the lesion was present throughout the submitted tissue and involved the surgical borders, a wider local excision was recommended which took place less than 3 months of the initial procedure. The surgical specimen measured $2.0 \times 1.4 \times 0.6$ cm and included surface mucosa. The origin of the tumor from the surface epithelium was confirmed (Fig. 2A). Essentially similar histopathologic patterns of growth, i.e., plexiform, follicular, and areas were evident as were a few foci of calcifications. A heavy plasmacytic infiltrate was also noted. However, some plexiform areas and follicles featured foci of increased cellularity characterized by sheets of basophilic cells, with inconspicuous cytoplasm and round or oblong nuclei occasionally overlapping nuclei and small nucleoli. Also, foci of nuclear hyperchromatism and few scattered mitoses, the latter mostly in the periphery of the cellular areas, were observed (Fig. 2B, C).

Immunohistochemical study of both surgical tissue materials was undertaken. Cytokeratin 19 (A53-B/



Fig. 1 A Radiolucent area in the right side of maxilla in the area of the premolars causing saucerization of the alveolar bone. Small radiopacities corresponded to calcified keratin. **B** Low-power microphotograph depicting interconnecting or plexiform cystic spaces of various sizes and scattered small follicles (H&E, original magnification \times 40). **C** Plexiform pattern featuring a peripheral layer of frequently low columnar cells exhibiting reverse polarization. Internally, a stellate reticulum-like pattern was appreciated with abundant parakeratin (asterisk) is present. Also focally in the connective tissue, a few dystrophic calcifications (arrow) can be appreciated. (H&E, original

magnification \times 200). **D** Tadpole-like follicle with areas of reverse polarization of elongated basal cells with spongiosis, occasional subnuclear vacuolization, and central area of lamellated parakeratinization. Internally, the pattern may be interpreted as either condensed stellate-like reticulum or the spinous layer of keratocystic odontogenic tumor (odontogenic keratocyst), which explains the difficulty encountered. (H&E, original magnification \times 400). **E** Small follicle revealing pseudoductal foci of early keratinization (arrow). (H&E, original magnification \times 400). **F** Small nests featuring cells with small but conspicuous nucleoli (black arrow) and a few mitoses in the periphery (red arrow) (H&E, original magnification \times 600)



Fig. 2 A Neoplastic plexiform proliferation of cords with features of ameloblastoma and area of keratinization (H&E, original magnification \times 40). *Inset* reveals palisading in the basal cell layer and stellate reticulum-like internal areas (H&E, original magnification \times 200). **B** Plexiform pattern with areas of increased cellularity in the periphery. Also, small follicles are present. Pacinian-like keratin present in cystic areas (H&E, original magnification \times 100). **C** High-power microphotograph of the periphery exhibiting nuclear hyperchromatism, cells with small conspicuous nucleoli, and mitotic activity primarily in the periphery. (H&E, original magnification \times 600).

D CK19 decorated mostly suprabasal cells, less often basal cells, and keratin. (CK19-ABC, original magnification \times 100). **E** BeR-EP4 decorated the cytoplasmic membranes selectively and especially in extensions of the surface epithelium (Ber-EP4-ABC, original magnification \times 200). **F** Variations in ki67 nuclear staining between areas of increased cellularity and cystic area (ki67-ABC, original magnification \times 100) *Inset* reveals another area where ki67 positivity was increased in a cellular area but only present in the basal cell/peripheral cell layer in the cystic areas (ki67-ABC, original magnification \times 200)

A2.26, predilute, Cell Marque/Ventana) plasmalemmal staining was observed (Fig. 2D). Ber-EP4 (BerEP4, predilute, Cell Marque/Ventana) highlighted only focally the cytoplasm of the surface epithelium (Fig. 2E) exhibiting ameloblastomatous characteristics. There was no staining for calretinin (polyclonal, 1:100, Invitrogen), BRAF V600E (BRAFV600E; VE1, predilute, Ventana) and CD56 (MRQ-42, predilute, Cell Marque/Ventana). Immunohistochemical evaluation with Ki67 (30-9, predilute, Ventana) (Fig. 2F) revealed nuclear staining mostly in the basal and lower suprabasal cells of the surface epithelium in the areas of keratoameloblastomatous transformation, the basal cell layer lining of some keratin-filled cysts. In some nests, only rare positive cells were noted. However, increase in the number of nuclei stained was observed in the areas of increased cellularity where cytologic atypia was noted. The histopathologic features were interpreted as peripheral keratoameloblastoma with foci of at least cytologic atypia. The possibility of malignant transformation could not be excluded. Also, the surgical material of the second surgery, representing additional enucleation, was received in pieces and evaluation of tumor-free borders was impossible.

After the second surgery, repeated calls for patient reevaluation were unsuccessful and the patient remains lost to follow-up 25 months post second surgery.

Targeted Next-Generation Sequencing (NGS)

DNA extracted from FFPE tissue was subjected to multiplex PCR enrichment and next-generation sequencing at University Hospitals Cleveland Medical Center (Cleveland, OH) using the Oncomine Comprehensive Assay v3 (DNA) and Oncomine Focus Assay (RNA) on the ion Torrent S5 Prime platform (ThermoFisher Scientific, Waltham, MA), in which 150 cancer-associated genes were interrogated for SNVs, CNVs, indels, and 23 driver genes were analyzed for gene fusions. Neoplastic tissue accounted for about 10% of sample tissue due to extensive inflammatory infiltrate. Sequencing identified an *ARID1A* c.6527_6538delAG frameshift mutation (VAF: 5.8%) classified as likely oncogenic and an FBXW7 c.1627 A > G missense mutation

(VAF: 8.0%) classified as a variant of uncertain significance, in addition to two other variants of uncertain significance which were detected at a VAF of roughly 50% (suspected germline) (Table 1). Paired normal tissue was not available for testing.

Discussion

KA was last included in the WHO classification in 1992, where it was considered a variant of ameloblastoma with extensive keratinization. However, it has since been omitted on account of its poorly defined diagnostic criteria and rarity [10]. Occasional examples, including cases with prominent papillary architecture, termed PKA, continue to be published, but an improved understanding of this unusual variant of ameloblastoma remains desirable. KA shows nearly equal male-to-female ratio and approximately 2:1 mandibular predilection, somewhat less dominant than conventional AMEL [1]. KA can occur at a wide age range, and has been treated variably by resection or enucleation, with recurrences noted following enucleation [11].

Only a peripheral example has appeared in the literature which was treated, apparently, with conservative enucleation since the mass was "well-encapsulated" [9]. Follow-up information was not provided. Histopathologically, the provided figures reveal similar pattern of growth with the present case. However, there are mostly low to medium power and of suboptical quality when it comes to the detailed depiction of the cytologic neoplastic characteristics. However, there is no reference of cytologic atypia as described herein.

Reports prior to the study by Robinson et al. [1] state that KA can present substantial histopathologic overlap with the rarely reported sKCOT/OKC with some authors postulating ameloblastomatous transformation in sKCOT/OKC [3, 8, 12]. Whitt et al. [13] proposed classifying KA into four different subtypes (a) papilliferous, (b) simple, (c) simple with OKC-like features, and (d) complex. Still, tumors reported as KA and sKCOT/OKC represent diagnostic challenges, despite the fact that this terminology implies apparent relationships to well-known entities. Histopathologically, nuclear palisading and hyperchromatism of peripheral/

Table 1 Summary of findings detected by targeted DNA/RNA panel

Call	Gene	Transcript	Amino acid	Codon	Effect	VAF
Likely oncogenic	ARID1A	NM_006015.4	p.Gln2176fs	c.6527_6528delAG	Frameshift	5.769231
Uncertain significance	SLX4	NM_032444.2	p.Lys458Glu	c.1372 A>G	Missense	49.77489 Probably germline
Uncertain significance	FBXW7	NM_033632.3	p.Arg543Gly	c.1627 A>G	Missense	7.959479
Uncertain significance	RNF43	NM_017763.5	p.Gly336Ser	c.1006 G>A	Missense	48.78049 Probably germline

basal cells is seen in both conventional KCOT/OKC and conventional ameloblastoma, and parakeratosis is not unique to KCOT as acanthomatous change with keratinization can be seen in conventional ameloblastoma [1].

There is only one molecular study which tried to investigate gene aberrations in sKCOT/KA. Zhang et al. [8] reported 9 cases without subclassifying each case as KA or sKCOT, on account of this substantial histopathologic overlap. None of the 9 cases harbored a *BRAF*^{V600E} mutation, in good agreement with the findings in the present case. Two cases with fresh tissue were subjected to *PTCH1* sequencing, one of which harbored a missense mutation. These findings led the authors to conclude that solid KCOT and KAB should not necessarily be segregated into two separate entities but should be best classified altogether as solid KCOT; however, this opinion should be substantiated by stronger data. A study utilizing next-generation sequencing to identify the molecular characteristics of KA is necessary.

In the present tumor, sequencing failed to identify any pathogenic variants in *BRAF*, *NRAS*, *HRAS*, *KRAS*, *FGFR2*, or *SMO*, as seen in ameloblastoma, or any pathogenic variant in *PTCH1*, as seen in KCOT. Additionally, immunohistochemistry for BRAF VE1 calretinin and CD56 was negative, the latter being also positive in cases of ameloblastoma and occasionally in KCOT/OKC [14].

Interestingly, targeted NGS revealed an *ARID1A* alteration. *ARID1A* (AT-rich interactive domain-containing protein 1 A) encodes a large subunit of the SWI/SNF chromatin remodeling complexes. This gene family plays an important role in maintaining chromosome accessibility to tissue-specific enhancers and localizing SWI/SNF complexes to them [15]. SWI/SNF complexes are mutated in approximately 20% of human cancers, representing the most frequently mutated epigenetic regulator in cancer, and are among the most commonly mutated tumor suppressors in cancer [16, 17]. *ARID1A* mutations have been identified in endometrioid and clear cell ovarian carcinoma, gastric adenocarcinoma, urothelial carcinoma, and hepatocellular carcinoma, among several other tumor types [16].

The presence of an *ARID1A* alteration in the present case is of unclear clinical significance, as *ARID1A* alterations have not been reported in ameloblastoma or KCOT, to date, and SWI/SNF complexes are not known to modulate MAPK pathway signaling. However, *ARID1A* functions as an epigenetic regulator in dental root patterning and development, and can regulate sonic-hedgehog (SHH) signaling in this context [18–20]. More work is needed to determine if *ARID1A* alterations play any role in or modulate SHH signaling in odontogenic neoplasia. *ARID1A* mutations have been identified in 45% of basosquamous carcinomas, as a secondary event following *PTCH1* alterations, and may confer a squamous phenotype in these tumors genetically related to basal cell carcinoma [21]. In the present case, there was no evidence of *PTCH1* alterations. Thus, sequencing of examples of KA are necessary to determine if *ARID1A* inactivation is a recurrent genomic event or if it occurs secondarily to a more commonly occurring mutation, conferring plasticity in differentiation.

Of interest in the current case are the (a) increased cellularity in the periphery of a few follicles with the presence of nuclear hyperchromatism and few scattered mitoses, (b) the presence of keratin in the stroma, the apparent result of burst cystic areas with frequent calcification of the keratin as it interacted with the stroma, (c) intense plasmacytic infiltrate, (d) negative immunostaining for calretinin, BRAF V600E and CD56, and (e) increased ki67 nuclear reaction in areas of cytologic atypia. This may suggest either a possible malignant transformation (lowgrade peripheral ameloblastic carcinoma with KA features?) or proliferation of cells to form new neoplastic follicles and nests. One should take into consideration the relatively short time between the first and the second surgery and a possible impact this may have had in the proliferative capacity of the neoplastic cells. As mentioned, the first specimen did not reveal the atypia noted in the second. The presence of occasionally calcified keratin in the stroma has been reported in examples of KA [1] and explains the radiopacities seen radiographically. The heavy plasmacytic infiltrate is apparently an inflammatory response to the ruptured cysts.

The pattern of growth in the present case was reminiscent to squamous cell carcinoma cuniculatum (CC) an entity that was discussed in the differential diagnosis of KA by Robinson et al. [1]. We are of the opinion that the histomorphologic features herein are overwhelmingly of an ameloblastomatous tumor and we do not know of any reports in the literature of oral CC exhibiting such distinct phenotype.

In conclusion, we presented a 54-year-old male with a gingival tumor fulfilling diagnostic criteria of KA with, however, foci of cytologic atypia or possible malignant transformation. The tumor was found to harbor an ARID1A mutation on targeted sequencing. No alterations in MAPK pathway signaling genes or in PTCH1 were identified. ARID1A mutations have not been reported in AMEL or KCOT/OKC to date, rendering interpretation of this pathogenic variant challenging. Sequencing of additional cases of KA will be necessary to determine if KA harbors a unique mutational profile relative to AMEL or KCOT/OKC or if additional, more commonly implicated genes are found to be consistently mutated. Additionally, sequencing of future cases of sKCOT will be necessary to establish if PTCH1 inactivation is a recurrent finding in that tumor, and to ultimately confirm if KA and sKCOT/OKC are separate entities.

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Data Availability The data generated during the study that include clinical notes, clinical photographs, pertinent radiographs, hematoxylin and eosin preparations, immunohistochemical stains, and Next-Generation Sequencing data are available on request from the corresponding author (IGK).

Code Availability Non-applicable.

Declarations

Competing Interests The authors declare no conflict of interest.

Ethical Approval Immunohistochemistry and new generation sequencing (NGS) were performed for investigative and diagnostic purposes and that part of the study was exempt from the Institutional Review Board. The NGS part of the study was carried out at the University Hospitals Cleveland Medical Center, Cleveland OH. The NGS part of the study was approved by the University of Minnesota Institutional Review Board (ID: 00012651). The University of Minnesota reviewed the study and determined that the proposed activity is not research involving human subjects as defined by DHHS and FDA regulations. To arrive to this determination, the University of Minnesota IRB used "WORKSHEET: Human Research (HRP-310)." Based on that, ongoing IRB review and approval for this activity is not required.

Consent to Participate The histopathologic, immunohistochemical, and NGS parts of the study were exempt from the institutional review board (University of Minnesota IRB: ID: 00012651).

Consent for Publication The patient's information, excluding age and gender, cannot be identified. Please note that the patient was lost to follow-up. Repeated attempts to communicate with the patient were unsuccessful. The NGS part was performed for diagnostic/investigative purposes.

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