HEAD AND NECK



Adenoid cystic carcinoma of the salivary glands: a pilot study of potential therapeutic targets and characterization of the immunological tumor environment and angiogenesis

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Abstract

Background Adenoid cystic carcinoma (ACC) is a rare type of cancer commonly occurring in salivary glands. It is characterized by slow but infiltrative growth, nerve infiltration and overall poor prognosis, with late recurrence and distant metastasis. The treatment of ACC is still limited to surgery and/or (adjuvant) radiotherapy. Till now no promising systemic therapy option exists. However, various studies deliver promising results after treatment with anti-angiogenetic agents, such as anti-EGFR-antibody Cetuximab or Tyrosinkinase inhibitor Lenvatinib.

Methods By using of immunohistological methods we analyzed and compared the macrophage and lymphocyte populations, vascularization, and PD-L1-status in 12 ACC of the salivary glands.

Results All cases showed a significant elevation of macrophages with M2 polarization and a higher vascularization in ACC compared to normal salivary gland tissue. The CD4/CD8 quotient was heterogenous. ACC does not show relevant PD-L1 expression.

Conclusions The predominant M2 polarization of macrophages in ACC could be responsible for elevated vascularization, as already been proved in other cancer types, that M2 macrophages promote angiogenesis.

Keywords Adenoid cystic carcinoma · Macrophage polarization · Neovascularization · Salivary gland cancer

Introduction

Salivary gland malignancies are overall rare and almost one fifth of them are adenoid cystic carcinomas (ACC) [1]. Based on its growing pattern, ACC is divided in three

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Andreas Mamilos Andreas.Mamilos@ukr.de histological types, namely, cribriform, tubular and solid. In total the cribriform type has the best and solid type the worst prognosis [2]. An infiltration of nerves is also a typical feature of ACC and is correlated with a poorer outcome of the disease [3].

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Former studies have shown an elevated expression of epidermal growth factor receptor (EGFR) in ACC [4, 5]. This suggests a high tendency to neovascularization, so that ACC might be susceptible to anti-angiogenetic agents, such as Cetuximab or Lenvatinib. On the other hand ACC show a weak to no programmed death ligand-1 (PD-L1) expression [6], a decisive biomarker for the use of Pembrolizumab in head and neck squamous cell carcinoma, which makes the benefit from immune checkpoint inhibitors questionable. ACC are low immunogenic tumors with low infiltration rates through CD8-positive lymphocytes [6], which is a crucial part of the anticancer immune response in the tumor microenvironment that can affect the outcome [7-9]. Furthermore, the CD4/CD8 ratio is in some cancer types, like the triple negative breast cancer, a useful prognostic tool [10], but can also be used as indicator to monitor the course of a therapy with immune checkpoint inhibitors [11]. However, its relevance in ACC is not yet analyzed.

Angiogenesis is a critical part of tumor growth and progression in general [12] and previous reports indirectly suggest elevated angiogenesis in ACC [13, 14], whereas a similar immmunohistological study did not show any significant difference between intratumoral and peritumoral vascularization [15]. Macrophages are a heterogenous group of cells that polarise to a M1 or M2 phenotype and are able to retain their plasticity and transform according to environmental signals [16–18]. M2-polarized macrophages, which have been linked to angiogenesis and cancer growth in pancreas [19], are also strongly represented in ACC [20, 21] and perhaps, at least partially, responsible for enhanced angiogenesis in this cancer. Till Dato no study to our knowledge compared both aspects.

In the era of personalized medicine, where monoclonal antibodies revolutionised oncology, the treatment of ACC does not appear to profit from these developments, as its curative therapy is still limited to surgery with/or radiation [22] and the therapy outcome is still not satisfactory. Up to now, no systemic treatment seems to be able to achieve satisfactory results [23–25].

In our study we used immunohistochemical methods to investigate the immunological microenvironment including vascularization of tumor and normal salivary gland tissue to examine if there are any abnormalities, that could be used as basis for translational development of new therapy regimens for ACC.

Materials and methods

Patients

Twelve Patients (8 female, 4 male) with mean age at time of diagnosis of 59.5 years, treated at the University Hospital of Regensburg, Germany were retrospectively included in this study (Table 1). The diagnosis of ACC was reviewed and confirmed. Tissue from the pathological routine diagnostics from these 12 primary ACC of the salivary glands was used for examination. After achieving a positive ethics votum (22-2753-104) an in situ immunohistochemical characterization of the immunological tumor environment was performed. For the in-situ characterization, standard routine diagnostic procedures and antibodies were applied. First, a new hematoxylin–eosin slide was prepared (Fig. 1).

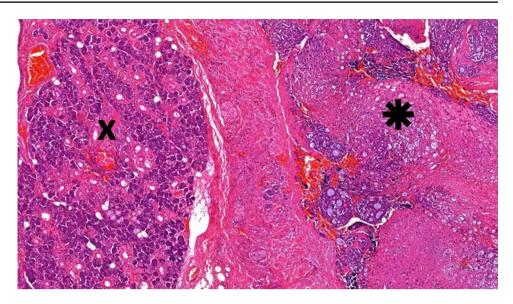
The tumor and residual salivary gland tissue were identified. In three tissue samples no normal salivary gland tissue was available for examination. Then, immunohistochemical reactions for PD-L1 (Dako Anti-Human PD-L1 Clone 22C3), CD4 (Ventana anti-CD4 Rabbit Monoclonal Primary Antibody Clone SP35) and CD8 (Ventana anti-CD8 Rabbit Monoclonal Primary Antibody, Clone SP57) for lymphocytes, CD68 (Dako Anti-Human CD68, Clone

Age	Localisation	UICC 8th Ed	TNM 8thEd	Pn	CPS	IC (%)	TPS (%)
46	Right submandibular gland	IVA	pT4a,cN0,cM0	Pn1	<1	<1	0
39	Right parotid gland	III	pT3,cN0,cM0	Pn1	0	0	0
87	Left sublingual gland	IVA	pT4a,cN0,cM0	Pn1	3	2	0
45	Left submandibular gland	II	pT2,pN0,cM0	Pn1	0	0	0
39	Hard palate	Ι	pT1,cN0,cM0	Pn0	0	0	0
69	Right parotid gland	IVA	pT4a,pN0,cM0	Pn1	0	0	0
45	Left parotid gland	Ι	pT1,pN0,cM0	Pn0	0	0	0
72	Left parotid gland	IVA	pT3,pN2b,cM0	Pn1	0	0	0
70	Right parotid gland	IVA	pT3,pN2b,cM0	Pn1	0	0	0
36	Left submandibular gland	Ι	pT1,pN0,cM0	Pn1	0	0	0
88	Left parotid gland	IVA	pT4a,cN0,cM0	Pn1	0	0	0
73, 67	Left parotid gland	Ι	pT1,pN0,cM0	Pn0	0	0	0
	46 39 87 45 39 69 45 72 70 36 88	 46 Right submandibular gland 39 Right parotid gland 87 Left sublingual gland 45 Left submandibular gland 45 Left parotid gland 469 Right parotid gland 45 Left parotid gland 45 Left parotid gland 46 Left parotid gland 47 Left parotid gland 48 Left parotid gland 	46Right submandibular glandIVA39Right parotid glandIII87Left sublingual glandIVA45Left submandibular glandII39Hard palateI69Right parotid glandIVA45Left parotid glandI72Left parotid glandIVA70Right parotid glandIVA36Left submandibular glandI88Left parotid glandIVA	46Right submandibular glandIVApT4a,cN0,cM039Right parotid glandIIIpT3,cN0,cM087Left sublingual glandIVApT4a,cN0,cM045Left submandibular glandIIpT2,pN0,cM039Hard palateIpT1,cN0,cM069Right parotid glandIVApT4a,pN0,cM045Left parotid glandIpT1,pN0,cM072Left parotid glandIVApT3,pN2b,cM070Right parotid glandIVApT3,pN2b,cM036Left submandibular glandIpT1,pN0,cM088Left parotid glandIVApT4a,cN0,cM0	46Right submandibular glandIVApT4a,cN0,cM0Pn139Right parotid glandIIIpT3,cN0,cM0Pn187Left sublingual glandIVApT4a,cN0,cM0Pn145Left submandibular glandIIpT2,pN0,cM0Pn139Hard palateIpT1,cN0,cM0Pn069Right parotid glandIVApT4a,pN0,cM0Pn145Left parotid glandIpT1,pN0,cM0Pn072Left parotid glandIVApT3,pN2b,cM0Pn170Right parotid glandIVApT3,pN2b,cM0Pn136Left submandibular glandIpT1,pN0,cM0Pn188Left parotid glandIVApT4a,cN0,cM0Pn1	46Right submandibular glandIVApT4a,cN0,cM0Pn1<139Right parotid glandIIIpT3,cN0,cM0Pn1087Left sublingual glandIVApT4a,cN0,cM0Pn1345Left submandibular glandIIpT2,pN0,cM0Pn1039Hard palateIpT1,cN0,cM0Pn0069Right parotid glandIVApT4a,pN0,cM0Pn1045Left parotid glandIpT1,pN0,cM0Pn0072Left parotid glandIVApT3,pN2b,cM0Pn1070Right parotid glandIVApT3,pN2b,cM0Pn1036Left submandibular glandIpT1,pN0,cM0Pn1088Left parotid glandIVApT4a,cN0,cM0Pn10	46Right submandibular glandIVApT4a,cN0,cM0Pn1<1<139Right parotid glandIIIpT3,cN0,cM0Pn10087Left sublingual glandIVApT4a,cN0,cM0Pn13245Left submandibular glandIIpT2,pN0,cM0Pn10039Hard palateIpT1,cN0,cM0Pn10069Right parotid glandIVApT4a,pN0,cM0Pn10045Left parotid glandIpT1,pN0,cM0Pn10072Left parotid glandIVApT3,pN2b,cM0Pn10070Right parotid glandIVApT3,pN2b,cM0Pn10036Left submandibular glandIpT1,pN0,cM0Pn10088Left parotid glandIVApT4a,cN0,cM0Pn100

Pn Perineural-invasion, UICC Union for International Cancer Control

Table 1Patientcharacterization, tumorlocation and stadium, as PD-L1Expression

Fig. 1 Overview of an evaluated case with salivary gland (X) and adenoid cystic carcinoma (★). H.E.-Stain



KP1) for macrophages and CD163 (NovocastraTM Lyophilized Mouse Monoclonal Antibody CD163, Clone 10D6) for M2-macrophages were performed. In addition, a CD31 (Dako Monoclonal Mouse Anti-Human CD31, Clone JC70) reaction was used to identify the vessel endothelial cells. All immunohistochemical stains were performed on tissue sections (2–4 μ m thickness), prepared from-formalin-fixed (4% neutral buffered formalin) paraffin-embedded tissue blocks. Immunohistochemical staining was performed using a Roche Ventana Benchmark Ultra automated slide stainer (Ventana Medical Systems, Roche, France).

All slides were scanned (3DHISTECH Ltd.Pannoramic slide scanner 250) and evaluated using a virtual microscopy software (3DHISTECH Ltd. Case Viewer Ver.2.2) (Figs. 2, 3).

For PD-L1 the three common diagnostic scores [tumor proportion score (TPS), Immune Cell Score (IC) and combined positive Score (CPS)] were assessed according the guidelines by an experienced consultant pathologist [26].

To quantify the lymphocytes, ten high power fields (HPF) of the invasion tumor front were examined and the positive cells were counted. The averages and then the quotient of CD4 to CD8 lymphocytes were calculated.

The CD68 reaction was used to visualize all macrophages. To evaluate the quantity of the macrophages ten HPF of the invasion tumor front and normal salivary gland tissue were examined and the positive cells were counted. Then, the mean value (Confidence interval 95%, CI 95%) of macrophages within the tumor front and the salivary gland tissue was calculated. The macrophages were then further characterized concerning their subpopulations. M2-macrophages were detected using an anti-CD163-antibody. Utilising the same, previously described procedure for the quantification of CD68-positive macrophages, the M2-macrophages were also quantified. Using the numbers of CD68-positive-macrophages and CD163-positive-M2-macrophages the number of M1-macrophages per HPF was calculated.

For the quantification of blood vessels, the endothelial cells were marked using a CD31-reaction. The vessel count in cross sections was performed within the tumour and normal salivary gland tissue, examining the same ten HPF, respectively, followed by calculation of the mean values like previously described.

Statistical analysis

All acquired results were documented in Microsoft Excel 16.58 (Microsoft Corporation, Redmond, WA, U.S.A.). The calculations and graphs were also produced with GraphPad Prism version 9.3.1 for MacOS (GraphPad Software, San Diego, California USA). To investigate if there are significant differences between M1 and M2 macrophages, but also vessels and CD4 + and CD8 + lymphocytes in tumor and healthy salivary gland tissue, we applied the Wilcoxon signed ranged test. A *p* value ≤ 0.05 in two tailed tests was considered as statistically significant.

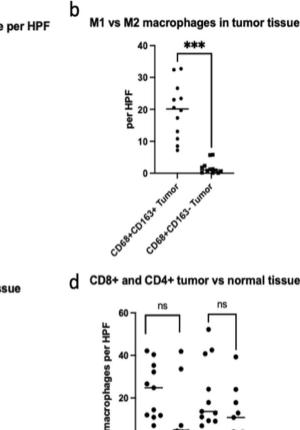
Results

The majority of macrophages in ACC shows an M2 polarization

In eight samples statistically significant more macrophages were observed in the tumor front in comparison with normal salivary gland tissue (Fig. 2a). In one sample nearly the same number of macrophages was found within the salivary gland tissue and the tumor, whereby this sample also showed signs of sialadenitis. For the three remaining samples a

Fig. 2 Comparison of the vascularization and immune cells in ACC and normal salivary gland tissue. Dots representing the mean number of counted cells or vessels per high power field (HPF). a CD68+CD163macrophages (M1) versus CD68+CD163+macrophages (M2) only in tumor tissue. **b** CD68 + macrophages, c CD31 + vessels and d CD8+and CD4+lymphocytes in tumor and normal salivary gland tissue. p < 0.05; **p<0.005; ***p<0.001

а CD68+ tumor vs. normal tissue per HPF 50 40 per HPF 30 20 10 n С Vessels tumor vs normal tissue 10 per HPF 2 CD31*Tumor CD31* Normal Used



comparison of macrophages within the tumor and salivary gland was not possible due to lack of tumor free salivary gland tissue. Despite this, the macrophages of the tumor were calculated in all cases and further characterized in subpopulations of M1 and M2 macrophages. Interestingly, the majority of macrophages was polarized to an M2-phenotype (p=0.0005), whereas only a small number of macrophages represented the M1-phenotype (Figs. 2b and 3).

Elevated intratumoral vascularization compared to normal salivary gland tissue

In the analyzed tissues, in which both tumor and normal salivary gland tissue were available, we observed a much higher vascularization in the tumor compared to normal tissue. Intratumoral vessels were in average elevated by 2.89 times in the tumor tissue compared to local salivary gland tissue (p = 0.0039) (Figs. 2c and 4).

Heterogeneous lymphocytes infiltration in ACC

Cove Worms These

CD8* Tumor CD8* Normal Tiss

CDA* TUMO

Besides three cases, where no normal salivary gland tissue was available and another one in which no more tissue was available for examination, the intratumoral CD4 positive lymphocytes where elevated from 1.7 to 9.9 times compared to normal tissue, but the presence of CD4-positive lymphocytes was not in the same range. In six cases the range of intratumoral CD8-positive lymphocytes was elevated up to 17 times. In general though, comparing the number of lymphocytes in the tumor and normal tissue, revealed no statistically significant difference (Fig. 2d). After evaluating and comparing the quotient of CD4-to-CD8 intratumoral lymphocytes, we observed heterogenous results. In four out of eleven cases more CD4 lymphocytes were found and in seven cases the CD8 lymphocytes compared to CD4 lymphocytes dominated.

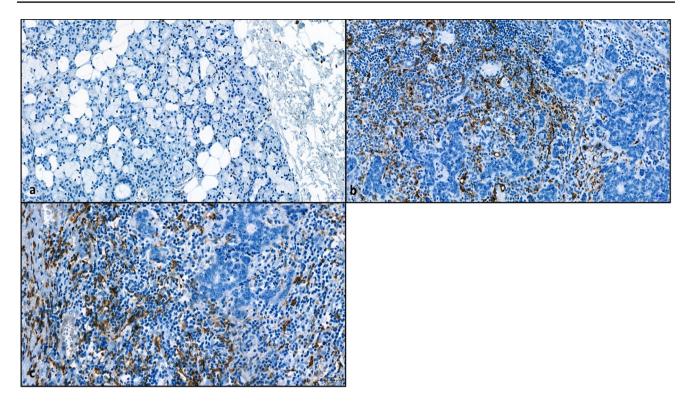


Fig. 3 Immunohistochemical comparison of CD68 + Macrophages in salivary gland **a** ACC, **b** and CD163 + Macrophages in ACC (**c**). Immunohistochemical reaction for CD68 and CD163 in brown

PD-1 status in ACC

In terms of PD-L1 status no relevant expression was observed. More specifically all cases showed a TPS of 0. The vast majority of the samples (10 out of 12) also showed an IC and CPS of 0. In two cases we observed slightly positive IC and CPS scores with a maximum of 2% and 3%, respectively (Table 1). This shows that the tumor cells show no immuno-histochemical expression of PD-L1. In addition, the immunal cells show no or minimal positivity for the immunohistochemical reaction for PD-L1.

Discussion

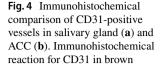
In the present study we clearly demonstrated an elevated number of macrophages in the tumor environment in comparison with normal salivary gland tissue but even more interesting is the finding of the almost complete polarization of macrophages in CD163-positive–M2-macrophages. It is well-known that the polarization of macrophages affects in various ways the behavior of tumors [27, 28]. M2 macrophages promote tumor aggression and progress [19, 29] in contrary to M1-macrophages which have an antitumoral effect [28]. They are also known for their role in the tumor angiogenesis [19, 30]. This is also consistent with our findings, showing significantly more vessels within the tumor tissue compared to normal salivary gland tissue (p=0.0039), as well as significantly more M2 macrophages in comparison with M1 macrophages.

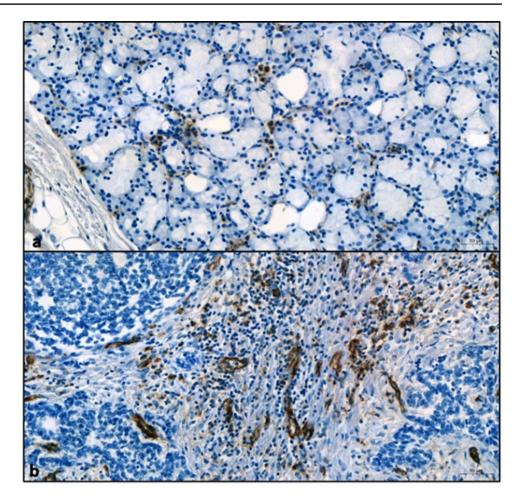
Recently Yang et al. showed in their experiments, that in ACC the mechanism of CCL2/CCR2 axis in the interactions between tumor cells and tumor-associated macrophages during the progression of salivary ACC, promotes a polarization to M2 tumor-associated macrophages [20], which explains the significantly higher polarization (p = 0.0005) of macrophages in the M2-subpopulation found in our study.

Adenoid cystic carcinomas are rare tumors and represent a challenge for clinicians. Confirming already performed studies, ACC are negative or extremely low positive for PD-L1 (all three scores), also suggesting a failure of immune check point inhibitor therapy.

The CD4/CD8-ratio of tumor infiltrating lymphocytes possesses in some malignancies—like gastric cancer [31] or as previously mentioned in the triple negative breast carcinoma [10]—a predictive value. In our study the ACC shows great heterogeneity in this matter and of course a larger collective is necessary to proof its clinical value.

These results, especially concerning the elevated macrophage infiltrate of the tumor front, their obvious polarization in M2 Macrophages and the also significantly elevated neovascularization, could help to develop new





strategies of treating ACC. Potential inhibition of the angiogenesis, using well-known anti-angiogenetic agents may bring an additional effect regarding the cure of ACC. Our results support various studies and reports involving Cetuximab in the treatment of ACC showing promising results [32–36]. In addition, the tyrosine kinase inhibitor (TKI) Lenvatinib achieved good results in the treatment of patients with recurrent or metastasized ACC [37, 38], as well as other TKI-like sorafenib [39, 40] and axitinib [41]. However, there is a lack of prospective randomized controlled phase III studies that confirm the effectiveness of such therapies, mainly because of the rarity and slow growth of these tumors. Furthermore, controlling the population of M2-macrophages in tumor is already being investigated as a potential cancer treatment with promising results, but this concept is still far from finding its way to routine daily practice [42]. Nevertheless, targeting tumor-associated macrophages might be a breakthrough in the treatment of ACC, especially in patients with metastatic or advanced disease, which lack options of effective therapy. As soon as such substances are available, they should be investigated in animal models and also phase II and III studies to find their way in the clinical practice if they prove to be effective. It is, therefore, necessary that large cancer centers cooperate and perform together such studies to collect the essential number of patients, so that highly qualitative and representative results can lead to novel therapies for ACC. ACC is a complicated deadly systemic disease so a wider therapeutical approach, targeting various features of the tumor itself or its environment is in our opinion the key to treat this tumor entity or at least control its course.

In the study of Li et al. showed recently that macrophage migration inhibitory factor was overexpressed in ACC. MIF might be a risk factor for ACC patients [43] and seems also to promote the perineural invasion in ACC patients [44]. Until now there are no clear data, how this factor affects the polarization of macrophages. This is a key point for examining in further studies.

Next step is to examine the general effect of M2-macrophages in ACC using molecular pathology techniques, where protein and RNA-based studies shall be performed. We are aware that the small number of examined cases poses as a limitation of our study. However, all cases showed statistically consistent results, so that the power of the results is credible and confirmed. It is, therefore, necessary to perform larger studies to confirm our results and taking in consideration the rarity of ACC this would be easier with the cooperation of other large cancer centers.

Conclusion

The fact that the ACC invasion front is being dominated by a large number of macrophages with M2 phenotype and also the highly elevated angiogenesis in the tumor tissue maybe the clue to treat a stubborn until today disease. Larger studies that require cooperation of many cancer centers are urgently required and we are willing to be a part of such national or international initiative.

Author contributions Conceptualization: IM, CB, AM. Methodology: IM, AM. Validation: IM, JK, CB and AM. Formal analysis: IM, AM. Investigation: IM, AM. Resources: TE, JK, PB, CB, CB. Data curation: IM, TE, AM. Writing—original draft preparation: IM, AM. Writing—review and editing, all authors. Visualization: IM, AM. Supervision: JK, CB, AM. Project administration: AM, CB. Funding acquisition, CB and CB. All authors contributed to the article and approved the submitted version.

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Data availability The data sets presented in this article are not freely available because of patient confidentiality and participant privacy terms.

Declarations

Conflict of interest The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

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