

Neuroendocrine Markers Insulinoma-Associated Protein 1, Chromogranin, Synaptophysin, and CD56 Show Rare Positivity in Adenocarcinoma Ex-Goblet Cell Carcinoids

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Abstract

Background: Adenocarcinoma ex-goblet cell carcinoid (AdexGCC) was considered a neuroendocrine adenocarcinoma, despite majority of tumor cells being negative for conventional neuroendocrine markers such as chromogranin and synaptophysin. Recently, insulinoma-associated protein 1 (INSM1) has been identified as a novel neuroendocrine marker that is more sensitive than chromogranin, synaptophysin, and CD56 in pulmonary neuroendocrine tumors.

Methods: We studied this marker in conjunction with chromogranin, synaptophysin, and CD56 in 36 appendiceal AdexGCCs (21 primaries, 15 metastatic).

Results: Primary AdexGCCs showed staining for INSM1, chromogranin, synaptophysin, and CD56 in 13/21 (62%), 18/21 (86%), 18/21 (86%), and 9/19 (47%) cases, respectively. However, the mean proportion of tumor cells stained for INSM1, chromogranin, synaptophysin, and CD56 was only 8.0% (median 1%, range 0-70%), 15.7% (median 2%, range 0-70%), 19.9% (median 5%, range 0-90%), and 5.6% (median 0%, range 0-50%), respectively. Metastatic AdexGCCs showed staining for INSM1, chromogranin, synaptophysin, and CD56 in 8/15 (53%), 11/15 (73%), 12/15 (80%), and 3/14 (21%) cases. The mean proportion of tumor cells stained for INSM1, chromogranin, synaptophysin, and CD56 in metastatic tumors was 1% (median 1%, range 0-3%), 12% (median 1%, range 0-85%), 17% (median 5%, range 0-85%), and 2% (median 0%, range 0-20%), respectively.

Conclusions: Primary and metastatic AdexGCCs showed no difference in INSM1, chromogranin, synaptophysin, or CD56 staining. INSM1 exhibits low expression in AdexGCCs and is expressed by a

lower proportion of tumor cells compared to chromogranin and synaptophysin.

Keywords: Adenocarcinoma ex-goblet cell carcinoid; Neuroendocrine; Insulinoma-associated protein 1; Chromogranin; Synaptophysin

Introduction

Appendiceal goblet cell carcinoid (GCC) and adenocarcinoma ex-goblet cell carcinoid (AdexGCC) have previously been described as two entities in a spectrum of neoplasms showing both glandular/mucinous and neuroendocrine differentiation [1]. Conventional neuroendocrine markers, such as chromogranin and synaptophysin, however, are often negative or focally positive in these tumors [2, 3]. Hristov et al previously demonstrated that chromogranin and synaptophysin were only expressed in 37% and 30% of AdexGCCs, respectively [2]. This is in keeping with ultrastructural studies in GCCs that showed the presence of abundant mucinous vacuoles, in comparison with the extremely difficult to locate, occasional membrane-bound, electron-dense granules [3]. In contrast, conventional carcinoids are often diffusely positive for chromogranin and synaptophysin, with abundant electron-dense granules on electron microscopy [4].

Molecular and proteomics profiling of neuroendocrine neoplasms have identified several proteins that play a pivotal role in neuroendocrine differentiation. Amongst them, one of the best characterized is insulinoma-associated protein 1 (INSM1), a zinc-finger transcription factor initially isolated from pancreatic insulinomas [5]. INSM1 is the upstream transcription factor for chromogranin [6]. As laboratory evidence began to accumulate, INSM1 has gained attention in clinical medicine as well and has been recently evaluated as a diagnostic marker of neuroendocrine differentiation in human tumor tissue samples [7, 8]. Rosenbaum et al was the first to demonstrate INSM1 to be 92.7% sensitive and 95.8% specific in showing neuroendocrine or neuroepithelial differentiation [7]. Rooper et al further established INSM1 as a pan-neuroendocrine marker more sensitive than chromogranin, synaptophysin, and CD56 combined in pulmonary neuroendocrine neoplasms [8]. However, no studies have looked at INSM1 expression in GCCs or AdexGCCs. Studies to date have mainly focused on pure

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Table 1. Primary Adenocarcinoma Ex-Goblet Cell Carcinoid Results (N = 21)

	0	1+	2+	3+	4+
CG	3	14	2	2	0
SYN	3	13	0	3	2
INSM1	8	11	1	1	0
CD56	10	8	1	0	0

CG: chromogranin; SYN: synaptophysin. Scoring criteria: 0 (< 1% cells staining), 1+ (1-25%), 2+ (26-50%), 3+ (51-75%), and 4+ (76-100%).

neuroendocrine neoplasms, neglecting tumors with mixed phenotype. The purpose of this study was to investigate this novel marker in AdexGCCs, an entity thought to show mixed glandular/mucinous and neuroendocrine differentiation.

Materials and Methods

Case selection

The study was approved by the Institutional Review Board of Washington University School of Medicine. The surgical pathology archives from 1990 to 2017 of the Washington University Pathology and Immunology Department were searched for appendiceal AdexGCCs (primary and metastatic). Archived hematoxylin and eosin slides and immunohistochemistry stains were retrieved and reviewed in consensus by three pathologists (CY, LZ, and DC) to confirm the diagnosis. To ensure consistency in classification, only cases featuring both the diagnostic crypt-like architecture [1] and diffuse strong SATB2 positivity, as shown in our previous studies [9, 10] were included. In total, 36 appendiceal AdexGCCs (21 primaries, 15 metastases) were enrolled in our study.

Immunohistochemistry

Whole tissue paraffin blocks containing adenocarcinoma component for each case was selected. Sections were cut at 5 μ m thickness, deparaffinized and subjected to antigen retrieval. Immunohistochemistry was performed on all cases with monoclonal antibodies to INSM1 (Santa Cruz Biotechnology, Clone A8, 1:200 dilution), chromogranin (Ventana, clone LK2H10, prediluted), synaptophysin (Cell Marque, Clone MRQ-40, prediluted), and CD56 (Ventana, clone MRQ-42, prediluted) on a Ventana Benchmark-XT automated stainer using the Ventana ultraView DAB detection kit. Positive control (lung carcinoid) and negative control (incubation with secondary antibody only) were included for each run of immunohistochemistry stains. The tests were performed with full ethical compliance as a human study.

Scoring and statistical analysis

Only nuclear staining for INSM1, cytoplasmic staining for

Table 2. Metastatic Adenocarcinoma Ex-Goblet Cell Carcinoid Results (N = 15)

	0	1+	2+	3+	4+
CG	4	9	0	1	1
SYN	3	9	1	1	1
INSM1	7	8	0	0	0
CD56	11	3	0	0	0

CG: chromogranin; SYN: synaptophysin. Scoring criteria: 0 (< 1% cells staining), 1+ (1-25%), 2+ (26-50%), 3+ (51-75%), and 4+ (76-100%).

both chromogranin and synaptophysin, and membranous staining for CD56 were considered as positive. The percentage of tumor cells stained was visually estimated and semi-quantitatively scored as 0 (< 1% cells staining), 1+ (1-25%), 2+ (26-50%), 3+ (51-75%), and 4+ (76-100%). Chi-square test was used to compare the staining patterns, and Mann-Whitney U test to compare the mean percentage of tumor cells stained. A P-value of less than 0.05 was considered statistically significant.

Results

Primary adenocarcinoma ex-goblet cell carcinoids

Primary AdexGCCs showed staining for INSM1, chromogranin, synaptophysin, and CD56 in 13/21 (62%, 1+ in 11, 2+ in one, 3+ in one), 18/21 (86%, 1+ in 14, 2+ in two, 3+ in two), 18/21 (86%, 1+ in 13, 3+ in three, 4+ in two), and 9/19 (47%, 1+ in eight, 2+ in one) cases (Table 1), respectively (INSM1 vs. chromogranin, $P = 0.079$; INSM1 vs. synaptophysin, $P = 0.079$; INSM1 vs. CD56, $P = 0.356$; chromogranin vs. synaptophysin, $P = 1.000$; chromogranin vs. CD56, $P = 0.010$; synaptophysin vs. CD56, $P = 0.010$). The mean number of tumor cells stained for INSM1, chromogranin, synaptophysin, and CD56 was 8.0% (median 1%, range 0-70%), 15.7% (median 2%, range 0-70%), 19.9% (median 5%, range 0-90%), and 5.6% (median 0%, range 0-50%), respectively (INSM1 vs. chromogranin, $P = 0.096$; INSM1 vs. synaptophysin, $P = 0.021$; INSM1 vs. CD56, $P = 0.433$; chromogranin vs. synaptophysin, $P = 0.368$; chromogranin vs. CD56, $P = 0.030$; synaptophysin vs. CD56, $P = 0.009$). High versus low grade areas showed no difference in staining.

Metastatic adenocarcinoma ex-goblet cell carcinoids

Metastatic AdexGCCs showed staining for INSM1, chromogranin, synaptophysin, and CD56 in 8/15 (53%, 1+ in eight), 11/15 (73%, 1+ in nine, 3+ in one, 4+ in one), 12/15 (80%, 1+ in nine, 2+ in one, 3+ in one, 4+ in one), and 3/14 (21%, 1+ in three) cases (Table 2), respectively (INSM1 vs. chromogranin, $P = 0.026$; INSM1 vs. synaptophysin, $P = 0.121$; INSM1 vs. CD56, $P = 0.077$; chromogranin vs. synaptophysin, $P = 0.666$; chromogranin vs. CD56, $P = 0.005$; synaptophysin vs. CD56, $P =$

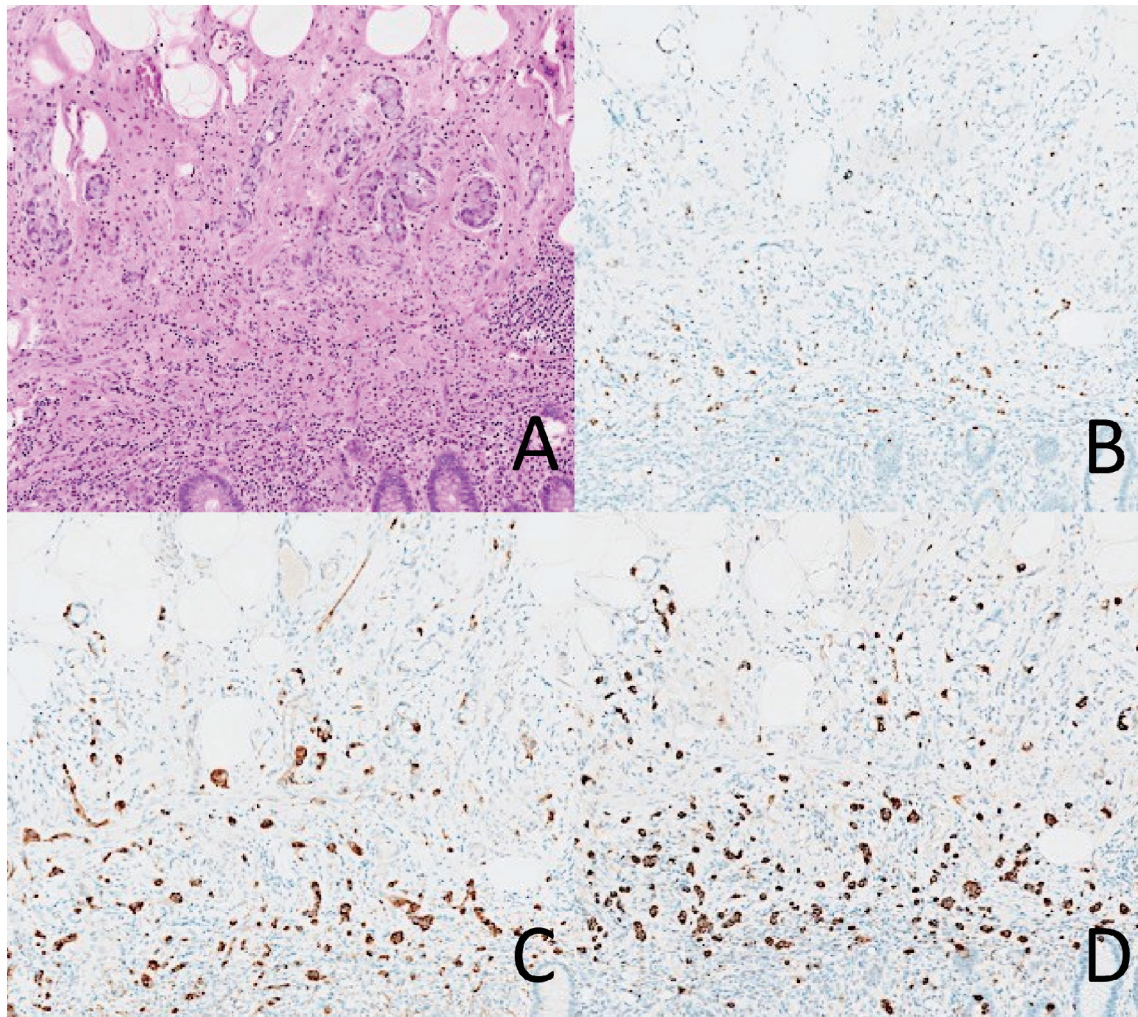


Figure 2. A case of adenocarcinoma ex-goblet cell carcinoid with positive staining for INSM1, chromogranin, and synaptophysin in the goblet cell component. There is diffuse staining of all three markers in the tumor cells. (A) Hematoxylin and eosin. (B) INSM1. (C) Chromogranin. (D) Synaptophysin.

ity of cases with less than 5% of cells positive for these neuroendocrine markers. Neuroendocrine phenotype, if present, is definitely not predominant, and would explain why neuroendocrine therapy regimens are not recommended for the treatment of AdexGCCs. As shown in our study, only five of the primary AdexGCCs show diffuse (> 50%) staining for any neuroendocrine marker, and only two of the metastatic AdexGCCs show diffuse (> 50%) staining for any neuroendocrine marker (Fig. 2). Most cases show either very focal positivity (Fig. 3) or are completely devoid of any neuroendocrine marker expression (Fig. 4). While there are still cases that show greater than 50% tumor cells staining for either chromogranin or synaptophysin, both these markers are less specific in tumors with abundant mucin given their non-nuclear staining pattern. We have also observed chromogranin and synaptophysin to be positive in approximately 11% and 30% of tumor cells in gastric signet ring cell carcinomas, respectively (unpublished data).

Recent molecular studies have shown these tumors to be different genetically from conventional carcinoids and colo-

rectal adenocarcinomas. Mutational landscape studies on AdexGCCs showed tumor cell mutation which were not commonly seen in colorectal adenocarcinomas such as *USP9X*, *NOTCH1*, *CTNNA1*, *CTNNB1*, *TRRAP*, *ARID1A*, *ARID2*, *CDH1*, *RHPN2*, *MLL2*, *RHOA*, *SOX9*, and *KDM6A* [14-16]. The mutations usually associated with colorectal adenocarcinomas, such as *TP53*, *KRAS*, and *APC*, were only identified in rare cases of AdexGCCs [16]. Neuroendocrine tumors did not show driver mutations in one study, but this could be due to the small number of cases enrolled [14]. Johncilla et al [15] demonstrated that there is overlap in mutational landscape in GCC and AdexGCC, further evidence supporting that GCC and AdexGCC are within spectrums of the same entity, rather than two distinct histological types. Overall, the mutational landscape of GCCs and AdexGCCs are different from both conventional neuroendocrine tumors and colorectal adenocarcinomas, suggesting that they are a separate entity.

Prognostic studies also show that these tumors are distinct from neuroendocrine tumors and actually share more simi-

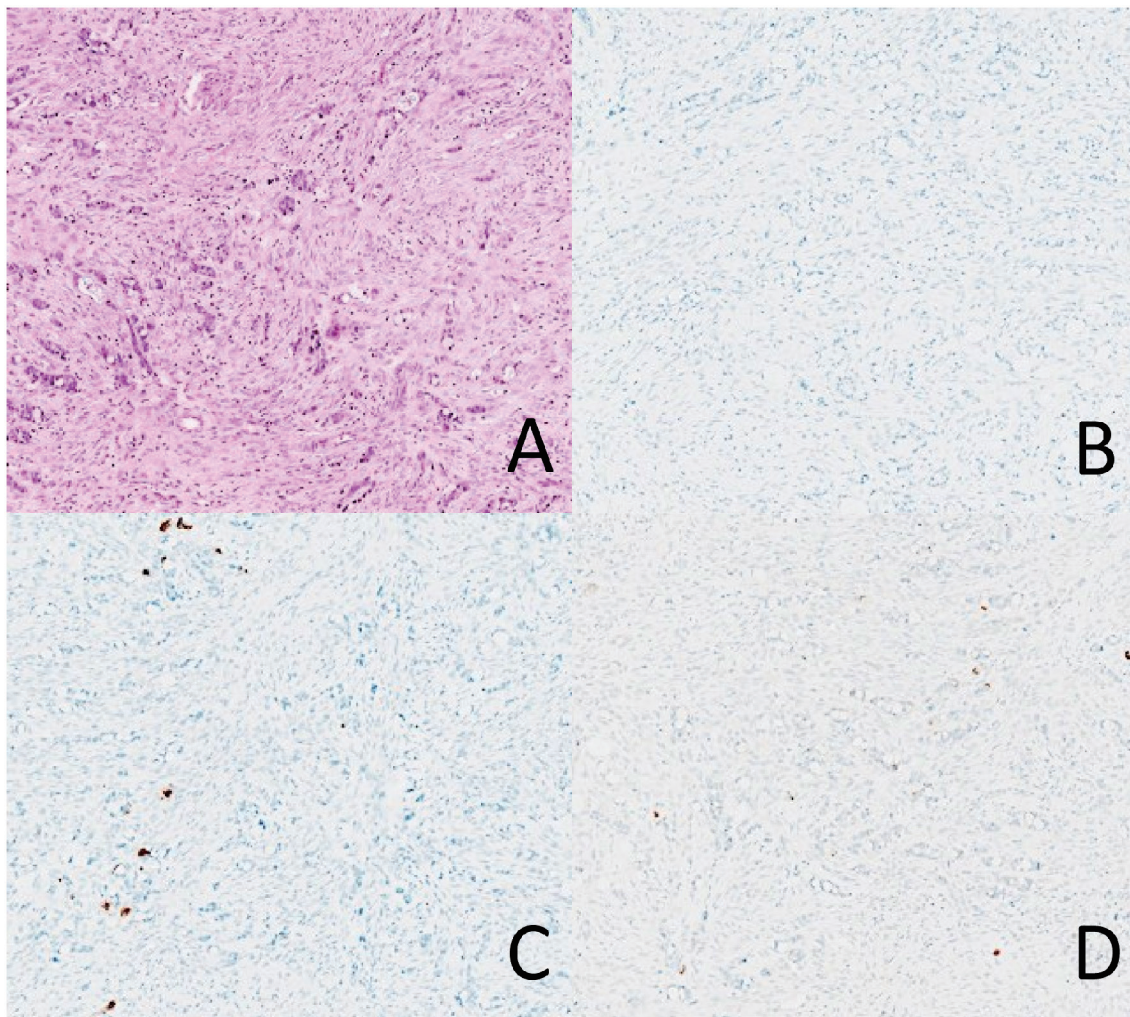


Figure 3. A case of adenocarcinoma ex-goblet cell carcinoid with weak positive staining for chromogranin and synaptophysin, but negative for INSM1. Note that chromogranin and synaptophysin are only expressed in a small proportion of tumor cells. (A) Hematoxylin and eosin. (B) INSM1. (C) Chromogranin. (D) Synaptophysin.

larities with colorectal adenocarcinomas. Most recently, Yozu et al [17] showed using their model of grading and staging AdexGCCs like other tubal gut adenocarcinomas, they can better separate patients into prognostic groups. Current surgical and oncologic approaches would favor them to be classified like an adenocarcinoma since mainstream treatment includes right hemicolectomy. The American Joint Committee on Cancer/Tumor-Node-Metastasis classification staging for AdexGCC of the appendix is similar to staging of primary adenocarcinoma of the appendix [18].

It had been shown in biological studies that cells in GCCs and AdexGCCs produce lysozyme and have a secretory component, which is more similar to intestinal crypts and not consistent with neuroendocrine cells [19]. This is in keeping with recent findings that supports the fact that GCCs and AdexGCCs have been hypothesized to origin from the crypt cells in the intestine, through differentiation through the MATH1 pathway rather than the NOTCH pathway [20, 21], which can be seen in the secretory crypt cells. The same cells that are thought to dif-

ferentiate through this pathway are neuroendocrine and Paneth cells. This might be one of the reasons that neuroendocrine markers such as synaptophysin and chromogranin are not proportionally, and only focally, expressed in these tumors. The hypothesis is that Paneth cells are thought to be differentiated through GFII/SOX9 pathway, while goblet cells are differentiated through GFII/KLF4 pathway, and finally neuroendocrine cells are differentiated with neurogenin 3 pathway. All three pathways are downstream of the MATH1 pathway. It is likely because of the close proximity in terms of the location in the tubal gut mucosa of these three cell types, and the fact that they share a common MATH1 pathway, that these three cell types can be seen intermixed in GCCs and occasionally AdexGCCs. Our study highlights the fact that AdexGCCs show either negative or limited expression of the four neuroendocrine markers studied, and in conjunction of additional evidence in molecular and prognostic studies, support separating AdexGCC from carcinoids or neuroendocrine tumor.

While our analysis is the first to comprehensively evalu-

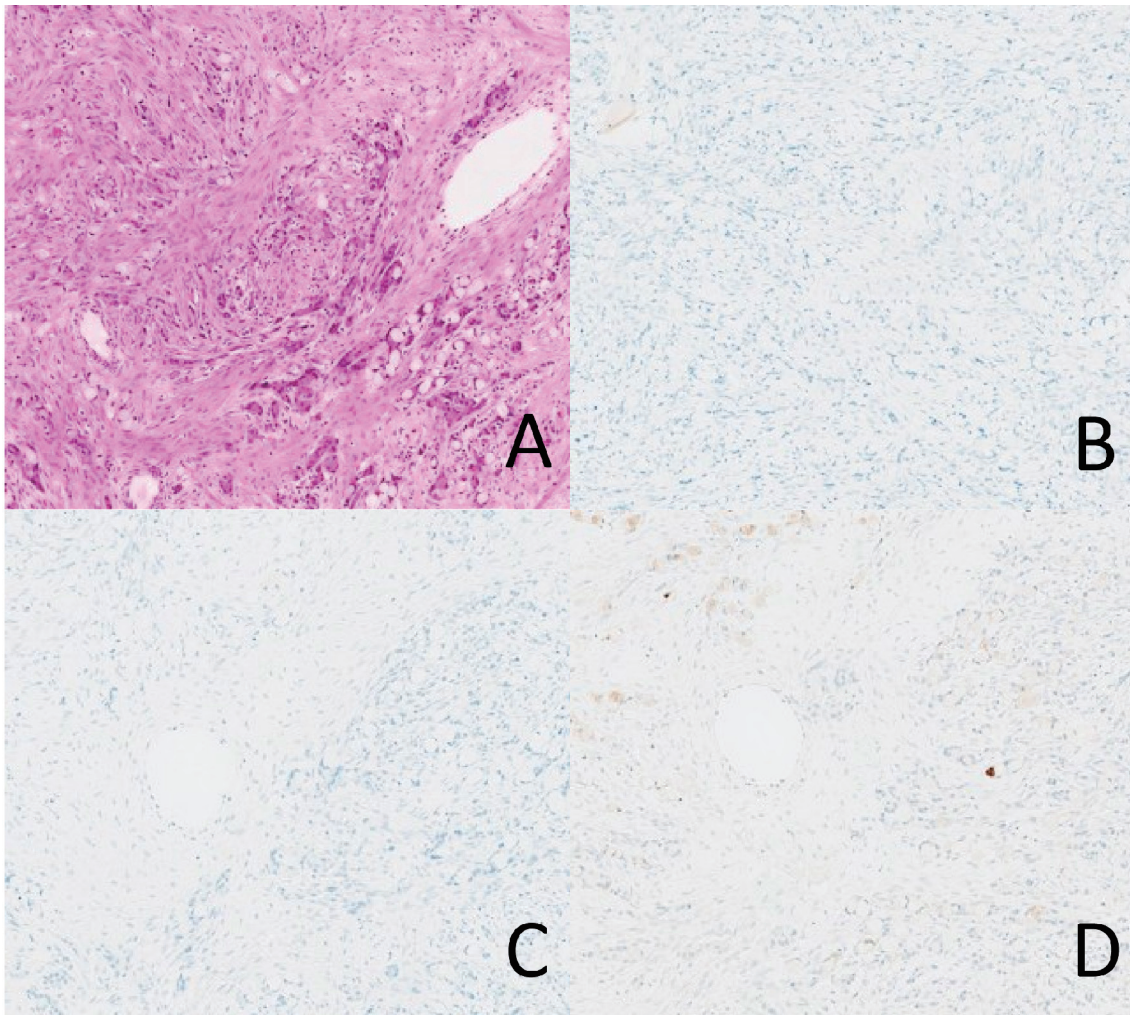


Figure 4. A case of adenocarcinoma ex-goblet cell carcinoid negative for INSM1, chromogranin, and synaptophysin. None of the tumor cells are staining with any of the markers. (A) Hematoxylin and eosin. (B) INSM1. (C) Chromogranin. (D) Synaptophysin.

ate novel neuroendocrine marker INSM1 in AdexGCCs along with three conventional neuroendocrine markers chromogranin, synaptophysin, and CD56, there are several limitations that merit discussion. First, our study was a retrospective immunohistochemistry study, and was limited by the inherent bias in studies of this type. Second, the number of cases included in our study is fairly small, due to the relatively low prevalence of these tumors. Thirdly, all the cases came from the same institution, thus a patient population selection bias could not have been avoided. Finally, follow-up information was not available for most of the cases, so the relevance of neuroendocrine marker expression and prognosis cannot be elucidated. Further prospective multi-institutional studies with more clinical intervention data would be needed to address the above issues.

Conclusions

We investigate novel neuroendocrine marker INSM1 in

AdexGCCs and found that it is often negative or shows only focal positivity in limited tumor cells. Our study highlights that INSM1, along with conventional neuroendocrine markers, such as chromogranin, synaptophysin, and CD56 shows a different staining pattern when compared with carcinoids or true neuroendocrine carcinomas. Along with emerging clinicopathologic and molecular evidence, AdexGCCs should be distinguished from a neuroendocrine tumor.

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Conflict of Interest

None of the authors have any relationships with, or financial interest in, any commercial companies pertaining to this article

Informed Consent

Informed consent was waived by the Human Research Protection Office of Washington University in St. Louis due to the retrospective nature of this study.

Author Contributions

All authors have contributed to the study. CY designed the study, collected the data, and drafted the manuscript; LZ and IG contributed to the study design and critically reviewed the manuscript; DC designed the study, oversaw the project, and critically reviewed the manuscript.

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