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Prognostic impact of the loss of E-cadherin and *de novo* expression of N-cadherin at
the invasive front of primary and recurrent oral squamous cell carcinoma

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- Aus der Sektion Medizin -



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1. Abbreviations

CAFs	Cancer-Associated Fibroblasts
CBCT	Cone-beam computed tomography
CCI	Charlson Comorbidity Index
CI	Confidence interval
CT	Computed tomography
CTC	Circulating tumor cells
EBV	Epstein-Barr virus
ECE	Extracapsular extension
ECAD	E-cadherin
ECM	Extracellular matrix
EGFR	Epidermal growth factor receptor
EMT	Epithelial–mesenchymal transition
EMT-TFs	EMT-promoting transcription factors
FISH	Fluorescence in-situ hybridization
GBM	Glioblastoma multiforme
GEKID	Gesellschaft der epidemiologischen Krebsregister in Deutschland
HIF-1	Hypoxia-inducible factor-1
HNSCC	Head and neck squamous cell carcinoma
HPV	Human papillomavirus
HR	Hazard Ratio
IHC	Immunohistochemistry
miR-134	microRNA
MET	Mesenchymal–epithelial transition
MDSCs	Myeloid-derived suppressor cells
MRI	Magnetic resonance imaging
MAPK/ERK	mitogen-activated protein kinases/ extracellular signal-regulated kinases
mTOR	Mammalian target of rapamycin
NCAD	N-cadherin
OPL	Oral premalignant lesion
OSCC	Oral Squamous Cell Carcinoma
PET-CT	Positron emission tomography–computed tomography
PD-1	Programmed cell death protein 1
PDCD7	Programmed Cell Death 7
pr DFS	post-recurrence disease-free survival
pr OS	post-recurrence survival

pr OCSS	post-recurrence oral cancer-specific survival
PTEN	Phosphatase and tensin homolog
PKM2	pyruvate kinase M2
ROI	Region of interest
ROS	Reactive oxygen species
TAMS	Tumor-associated macrophages
TGF	Transforming growth factor
TMA	Tissue microarray
US	Ultrasound
RT	Radiotherapy
RCT	Radio-Chemotherapy
RKI	Robert Koch-Institut
WHO	World Health Organization
Wnt	Wingless-INT

2. Introduction

Oral cancer encompasses malignancies of the floor of the mouth, tongue, gum, lip, palate, and gingiva and is one of the most common kinds of squamous cell carcinoma of the head and neck. Oral squamous cell carcinoma accounts for 91% of all instances of oral cancer ¹.

Head and neck squamous cell carcinomas (HNSCCs) are the most common malignancies of the head and neck, arising from the mucosal epithelium of the oral cavity, larynx and pharynx. The risk of HNSCC varies by nation and area, and it has been associated with exposure to tobacco-derived carcinogens, excessive alcohol use, or both. Previous infections with the carcinogenic strains of human papillomavirus (HPV), like HPV-16 and HPV-18, increase the risk of oropharyngeal tumors ²⁻⁴.

Successful vaccination campaigns might prevent HPV-positive HNSCC across the globe since the two most prevalent carcinogenic HPVs, HPV-16 and HPV-18, are covered by FDA-approved HPV vaccines. Smoking is still a significant risk factor for oral cavity and laryngeal HNSCCs, now known as HPV-negative HNSCCs. However, a comprehensive physical examination is still the best method for early OSCC identification since there hasn't yet been a reliable screening procedure. Even though only a small number of oral pre-malignant lesions (OPLs) that show up as leukoplakia (white patches) or erythroplakia (red patches) turn into invasive cancer, most people with advanced stage HNSCC do not have pre-cancerous lesions. Generally speaking, OSCC is often treated with surgical resection followed by adjuvant radiation (RT) or radiochemotherapy (RCT), depending on the stage of the malignancy ⁵.

2.1 Incidence and prevalence

With 890,000 new cases and 450,000 mortalities in 2018, HNSCC is the sixth most frequent malignancy worldwide ⁵. The global cancer observatory predicts that the incidence of HNSCC would climb by 30% (or 1.08 million new cases per year) by 2030 ⁶.

⁷. The high prevalence of HNSCC in Southeast Asia and Australia is linked to the intake of specific carcinogen-containing substances, whereas in the United States and Western Europe is linked to growing rates of oropharyngeal infection with HPV ⁸⁻¹⁰. In general, men are two to four times more likely than women to develop HNSCC. The median age of HNSCC patients without a viral association is 66 years, but the median ages of patients with HPV-associated oropharyngeal cancer and Epstein-Barr virus (EBV)-associated nasopharyngeal cancer are 53 years and 50 years, respectively ^{11, 12}.

Each year, approximately 10,000 individuals in Germany develop malignant tumors of the oral cavity and pharynx. Men are affected by the disease at a higher rate than women, with an average of 7600 men and 2800 women developing it annually since 2000. Additionally, men tend to develop the disease at a younger age, with a median onset of 61 years compared to 65 years for women. Despite improved interdisciplinary diagnostic and therapeutic strategies, the 5-year relative survival rate in Germany for oral cavity and pharynx ranged from 44 to 50% for men and 55 to 72% for women according to RKI and GEKID in 2013.

2.2 Risk factors

Smoking and excessive alcohol use are major risk factors for developing oral cancer, these risk factors are generally avoidable. ¹³ , when such risk variables are discovered simultaneously, they may have an enhancing effect ¹⁴.

2.2.1 Tobacco

Cigarette smoke attenuates oral immunity by inducing gingivitis, periodontitis, and oral cancer ¹⁵. Tumor growth is induced by suppressing the tumor suppressor genes, particularly p53 and PTEN (phosphatase and tensin homolog). Smokers have a threefold increased chance of acquiring mouth cancer compared to nonsmokers ¹⁶. Further, passive or second-hand smokers have an 87% higher risk of oral cancer than never-smokers who had never been exposed ¹⁷. All tobacco products have different amounts

of cancer-causing chemicals like polycyclic hydrocarbons and tobacco-specific N-nitrosamines, which are known to play a big role in the development of cancer. Nicotine may contribute to cancer development by promoting a number of essential processes, according to evidence from experimental in vitro research on cell cultures, in vivo studies on rodents, and human investigations, including epidemiological studies ¹⁸. Nicotine binds with a greater affinity to nicotine acetylcholine receptors than acetylcholine does. Nicotine's interaction with nicotine acetylcholine receptors triggers signaling pathways, resulting in a variety of responses, including increased cell proliferation ^{19, 20}. Nicotine induces epithelial-to-mesenchymal transition, a crucial stage in the development of a malignant phenotype. This change enables the cell to develop migratory capabilities, which may promote cancer metastasis ²¹.

2.2.2 Alcohol

According to epidemiological research, drinking alcohol increases the incidence of HNSCC in a dose-dependent manner. Yet, ethanol consumption has two aspects that affect health outcomes: first, Time-related patterns of consumption, such as age at beginning of consumption and duration, may affect the association between drinking intensity and cancer risk, and second, the amount of alcohol consumed ²². Although it is unclear how alcohol contributes to oral carcinogenesis, numerous pathways have been postulated. To begin, ethanol is converted into acetaldehyde, a recognized carcinogen. Because acetaldehyde is a tumor trigger, persistent alcohol drinking promotes the development of oral cancer ²³. According to earlier studies, alcohol makes the oral mucosa more permeable, which causes epithelial atrophy. In addition, alcohol degrades the lipid composition of the oral mucosa's epithelial cell membrane, enabling carcinogens to infiltrate ²⁴. Acetaldehyde production has been linked to some *Streptococcus* species, *Neisseria* species, and other bacteria. Such bacteria have been shown to proliferate in smokers and heavy drinkers. Because these bacteria may

convert ethanol to carcinogenic acetaldehyde, they are linked to an increased risk of HNSCC ²⁵.

2.2.3 Areca Nuts

Areca-nut-containing betel quid is the fourth most commonly used addictive stimulant worldwide, behind alcohol, caffeine, and cigarettes. Researchers from King's College London were the first to identify a dependency syndrome associated with areca nut consumption ²⁶. A Taiwanese study revealed that holding and then ingesting betel-quid juice and putting unripened whole areca fruit in the quid appeared to increase the risk of oral cancer by 11-fold ²⁷. Chewing areca nuts is considered a risk factor for developing possibly malignant oral submucous fibrosis, which results in oral and esophageal squamous cell carcinoma ²⁸. Moreover, contact with its extract resulted in oral mucosa deterioration as well as genotoxic and cytotoxic consequences on oral keratinocytes and fibroblasts. The active form arecoline N-oxide is solely accountable for areca-related oral carcinogenesis ²⁸. Additionally, areca nut extract enhances the formation of prostaglandins and cyclooxygenase-2, which are essential inflammatory mediators during the process of tumorigenesis and metastasis ²⁹. In order to determine whether an educational intervention program resulted in the cessation of chewing and a decrease in incident leukoplakia, prospective cohort research was carried out in India. The key finding following a 10-year follow-up showed that the incidence of oral leukoplakia decreased significantly ³⁰.

2.2.4 Viral Infection

It is generally known that HPV contributes to pharyngeal cancer. However, further research is still being done on its impact on the oral cavity ³¹.

2.3 Histopathology

OSCC development is fostered by the accumulation of genetic transformations and epigenetic aberrations in cancer-associated signaling pathways ³². Histologically, the

lesion goes through many different stages (preneoplastic damage) before it turns into cancer ³³.

Lesions that appear in the epithelium during the process of tumorigenesis can be categorized based on their histopathological presentation as reactive epithelial changes (such as hyperkeratosis, hyperplasia, and acanthosis) or preneoplastic alteration (including mild, moderate, and severe dysplasia, before the development of an invasive carcinoma). Figure 1 depicts the progression from dysplasia to invasive cancer. OSCC begins as an epithelial dysplasia and is distinguished by the abnormal proliferation of dysplastic squamous cells on the epithelial layer's surface, which affects the subepithelial basement membrane. Basement membrane deterioration eventually causes devastation and metastasis. Local invasion of the underlying tissue also develops through epithelial cell islets and cords. Tumor cells' propensity to metastasize is strongly tied to their differentiation grade, corresponding to the architecture of neoplastic tissue and normal epithelium ³³⁻³⁷.

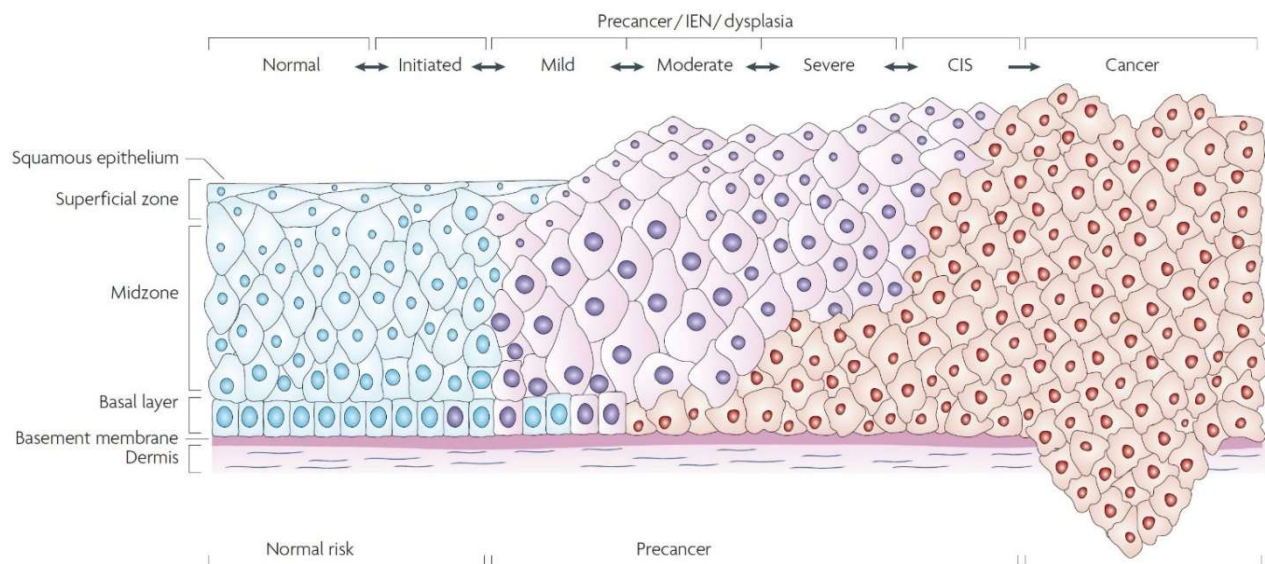


Figure 1: Schematic illustration of the development of epithelial cancer, modified according to Kelloff et al. ³⁸

2.4 Diagnosis and Staging

Despite advances in therapy over the last 20 years, there has been no significant increase in the 5-year survival rate in OSCC³⁹. Even though there are novel methods for diagnosing oral cancer, biopsy and histopathologic analysis continues to be the gold standard to diagnose OSCC⁴⁰. Premalignant or malignant oral lesions may be accurately diagnosed according to the standard of the biopsy, the availability of sufficient clinical data, and the analysis of the biopsy findings⁴¹. However, oral squamous cell carcinoma can form in intact oral mucosa, the vast majority of malignancies emerge from precursor lesions such as leukoplakia, erythroplakia, and erythroleukoplakia⁴².

Positive lymph nodes are a substantial negative predictive indicator for survival⁴³. Accordingly, a diagnostic evaluation prior to surgery is required for disease staging. To stage OSCC, computed tomography (CT) scans with contrast agent of the head, neck, and chest are routinely used to complete and optimize the staging of patients with OSCC. They are crucial in identifying tumor size, bone invasion, possible cervical node metastases, and pulmonary metastases⁴⁴.

Alternatively, ultrasound (US), magnetic resonance imaging (MRI) and in some cases, Positron emission tomography–computed tomography (PET-CT) can be employed. Especially for OSCC, an orthopantomogram and/or Cone-Beam CT (CBCT) are usually used to evaluate alveolar bone invasion and the degree of required bone resection⁴⁵. PET-CT is a type of scan in which a radioactive substance is given intravenously to the patient and is taken up by cells with a high metabolic rate, which is a characteristic of many oral cancer types. In contrast, infection and inflammation may have comparable radiologic outcomes. As a result, it is commonly utilized in advanced diseases (stage 3 or 4), salvage/recurrent cases, and metastatic disease evaluations^{46, 47}.

One of the most distinguishing clinical aspects of OSCC is its ability to locally invade surrounding tissues and metastasis. As a result, it is critical to predicting the invasive

and metastatic potential of OSCC early in the therapy process. In addition, a variety of biological markers have been developed in recent years that may improve diagnosis and provide crucial prognostic information for the management of OSCC. Molecular markers that signify a high local recurrence rate in surgically treated patients could aid in identifying patients who would benefit from postoperative radiotherapy. Furthermore, markers that indicate local recurrence in patients who have received radiotherapy could help identify highly radioresistant cancer forms. Concurrent chemotherapy, radiosensitizing drugs, adjusted radiation fractionation regimens, and other treatments targeting specific tumor molecular aberrations may represent an adjuvant therapy in patients with such cancers ⁴⁸.

2.5 Tumor Classification

Precise and reliable tumor staging is important for determining the type of treatment. According to the stage of the tumor, different aspects are decided, like curative or palliative treatment, the treatment planning (extent of resection in case of surgical treatment, or region of radiation therapy in case of primary or adjuvant radiation(chemo) therapy. The disease's staging also provides significant prognostic information to treating physicians and patients.

Staging of OSCC is performed according to the system developed by the Union for International Cancer Control (UICC) for malignant tumors" (Table 1). In addition to tumor size (T), this classification describes the existence of locoregional metastases (N) as well as distant metastasis (M) with organ involvement as seen in table 1 ⁴⁹.

Table 1: The 8th Edition of the American Joint Committee on Cancer (AJCC) TNM staging system for OSCC ⁵⁰.

TNM classification of carcinomas of the lip and oral cavity	
T	Primary tumor
TX	Primary tumor cannot be assessed
T0	No evidence of primary tumor

Tis	Carcinoma in situ
T1	Tumor 2 cm or less in greatest dimension
T2	Tumor more than 2 cm but not more than 4 cm in greatest dimension
T3	Tumor more than 4 cm in greatest dimension
T4a (oral cavity)	Tumor invades through cortical bone, into deep/extrinsic muscle of tongue (genioglossus, hyoglossus, palatoglossus, and styloglossus), maxillary sinus, or skin of face
T4b (lip and oral cavity)	Tumor invades masticator space, pterygoid plates, or skull base; or encases internal carotid artery

Note: Superficial erosion alone of bone/tooth socket by gingival primary is not sufficient to classify a tumor as T4

N - Regional Lymph Nodes

NX	Regional lymph nodes cannot be assessed
N0	No regional lymph node metastasis
N1	Metastasis in a single ipsilateral lymph node, 3 cm or less in greatest dimension
N2	Metastasis as specified in N2a, 2b, 2c below
N2a	Metastasis in a single ipsilateral lymph node, more than 3 cm but not more than 6 cm in greatest dimension
N2b	Metastasis in multiple ipsilateral lymph nodes, none more than 6 cm in greatest dimension
N2c	Metastasis in bilateral or contralateral lymph nodes, none more than 6 cm in greatest dimension
N3	Metastasis in a lymph node more than 6 cm in greatest dimension
T4b (lip and oral cavity)	Tumor invades masticator space, pterygoid plates, or skull base; or encases internal carotid artery

Note: Midline nodes are considered ipsilateral nodes.

M - Distant metastasis

M0	No distant metastasis
M1	Distant metastasis

Stage grouping

Stage 0	Tis	N0
Stage I	T1	N0

Stage II	T2	N0
Stage III	T1, T2	N1
	T3	N0, N1
Stage IVA	T1, T2, T3	N2
	T4a	N0, N1, N2
Stage IVB	Any T	N3
	T4b	Any N
Stage IVC	Any T	Any N

2.6 Management of Oral Squamous Cell Carcinoma

The majority of oral cavity cancer treatment regimens include surgery. Invasive procedures such as surgery have various advantages, including the ability to harvest a specimen for histopathologic examination and the ability to remove the cancer with a single treatment method in a single session. Although primary radiation to T1 and T2 lesions may provide comparable disease management, the negative effects of radiation to the oral environment outweigh those of surgery in the majority of cases. Additionally, it is frequently preferable to postpone radiation whenever feasible in case it becomes necessary in the future for second primary malignancies among patients with head and neck cancer. The oral cancer patient population is vulnerable to the formation of second primary malignancies. Some believe that radiation for borderline indications should be preserved for future use if the need arises.⁵¹

Patients with advanced local disease T3-T4, advanced nodal disease N2-N3, or tumors with lymphovascular and perineural invasion may benefit from radiotherapy (RT)⁵²⁻⁵⁴. Combining RCT for patients with either extracapsular extension (ECE) or positive margins enhances locoregional control and overall survival, according to two randomized studies^{55,56}. In the past, all patients at high risk of locoregional failure were included in the commonly acknowledged indications for RT. Because it is widely believed that there is no significant disadvantage to the salvage strategy in this cohort, early T1-2 and N0-1 cases are generally treated with surgical resection followed by

observation, with RT reserved as part of salvage therapy for recurrent tumor cases ^{57, 58}.

Immunotherapy is the fourth therapeutic option for HNSCC with promising outcomes. The average objective response rate for immune checkpoint inhibitors to far has only been around 15%, thus it is far from ideal. Activity levels are below expectations, with at least 80% of HNSCC patients showing no tumor size decrease. In both the CheckMate and KEYNOTE studies, there was no discernible difference between the immune checkpoint inhibitor and standard of care groups for progression-free survival. Anti-PD-1 (programmed cell death protein 1) and anti-PD-L1 therapy resulted in hyperprogression in 29% of patients with recurrent, metastatic HNSCC, shortening progression-free survival ^{51, 59}.

2.7 Recurrence of Oral Squamous Cell Carcinoma

Recurrence rates of OSCC range from 18 to 76% in individuals who received standard treatment, and it is often thought to be the primary reason for low survival rates. The previous investigations confirmed that the median time to recurrence after therapy is 7.5 months, and 86% of recurrences occur within 24 months ⁶⁰⁻⁶². The presence of cervical lymph node metastases is the major risk factor for death in individuals with OSCC. ECE is a highly predictive indicator for locoregional recurrence, distant metastasis, and disease-related death ⁶³. Local and regional recurrences are associated with post-surgery and radiation therapy failures ^{60, 64, 65}. Adjuvant Radiochemotherapy was found to reduce recurrence rates in this subset of patients compared to radiation alone ⁶⁶. Patients who are not candidates for salvage surgery or re-irradiation often get chemotherapy. However, even with the most cutting-edge medication combinations, the prognosis remains grim, and a cure is rare ⁶⁷. In terms of determining the most effective therapeutic options, patients with recurring carcinomas provide a clinical

dilemma. Salvage surgery is only appropriate for a tiny subset of patients, and roughly 30-45% of individuals have poor survival outcomes ⁶⁵.

2.8 Tumor Microenvironment (TME)

Cancer has long been thought to be a cell-autonomous mechanism in which repeated mutations in the tumor suppressor and oncogene genes cause an endless growth of malignant cells ⁶⁸. As a result, cancer therapy approaches have been focused and constrained to such tumor cell alterations ¹⁴. However, emerging data suggests that the tumor's genesis and growth are determined by tumor cells and a low TME ⁶⁹. Tumors are small organs made up of many distinct types of cells that interact to allow cancer cells to survive, develop, and spread ⁷⁰. The tumor's microenvironment, which includes immune cells (T and B lymphocytes), myeloid-derived suppressor cells (MDSCs), CAFs (or Cancer-Associated Fibroblasts), tumor-associated macrophages (TAMs), adipocytes, ECM (Extracellular Matrix) proteins, and mesenchymal stem cells, is one of the significant factors contributing to tumor development ⁷¹. Another element with a pleiotropic impact is cytokines. They generate various reactions, usually on distinct cell types, and contribute to cancer cell growth, drug resistance, initiation of EMT in cancer cells, resistance to apoptosis, and amplification of chemokine impact on recruiting immunological suppressor cells. The growing evidence of the critical involvement of various stromal components in the regulation of HNSCC development points that the tumor microenvironment plays an essential role in providing a supportive habitat, hence significantly enhancing HNSCC development and metastasis ⁷².

2.8.1 Cadherins

Cadherins have been identified as Ca^{2+} -dependent cell-cell adhesion proteins in vertebrates. They connect cells by homophilic interactions. cadherins are essential for creating and maintaining intercellular connections in healthy epithelium. Cells with fewer cadherin molecules are typically less adherent ⁷³. The classical cadherins are

single-span transmembrane cadherins with five extracellular cadherins repeat domains that collaborate with members of the catenin family to bind to the actin cytoskeleton through their cytoplasmic domains ^{74, 75}. E- and N-cadherins belong to the family of classical cadherins ⁷⁶. E-cadherin (ECAD) is found on the cell surface of all epithelial cells, whereas N-cadherin (NCAD) is present in fibroblasts, skeletal, neural tissue, cardiac muscles, and endothelial cells ⁷⁷.

2.8.2 E-cadherin

Epithelial cadherin is a 120 kDa transmembrane glycoprotein encoded by the CDH1/E-cadherin gene on chromosome 16q22.1, which is believed to be a tumor suppressor gene due to its ability to inhibit cell growth ⁷⁸. ECAD is an established member of the cadherin group and a potent tumor suppressor, since down-regulation of ECAD is typically detected in malignant epithelial tumors ⁷⁹⁻⁸¹. Due to its early identification and comprehensive characterization, ECAD is commonly considered as the model classical cadherin in mammals ⁸². It is considered a critical cell-cell adhesion protein that plays crucial roles during development and is necessary for the homeostasis of multiple organs. Multiple pathways often disturb ECAD function in cancer, making it an appealing diagnostic and prognostic candidate protein in human medicine ⁸⁰. The loss of ECAD expression in tumor tissue triggers metastatic spreading and the amplification of several EMT transcription factors ⁸³. Some investigators have referred to the loss of ECAD expression in OSCC as a high-risk marker of malignancy ^{84, 85} since it has been linked to both clinical and histological characteristics of malignancy, such as metastasis, recurrence, poor tumor differentiation, and reduced survival ⁸⁶⁻⁸⁸.

2.8.3 N-cadherin

Neural cadherin, known as cadherin 2, was discovered as the antigen identified by the mouse brain tissue-specific monoclonal antibody NCD-1, which was first developed against mouse brain tissue ⁸⁹. According to research, the structural-adhesive function

of NCAD in adult tissues is necessary for the appropriate integrity of particular tissues. It also aids in cell communication by assisting neurons form functional synapses and forming a vascular wall that is important for stabilizing blood vessels ⁹⁰⁻⁹³. The function of NCAD is diverse and varies depending on the cell environment. Any disruption in the functionality of NCAD may play a significant role in developing pathologic conditions since NCAD can influence the cytoskeleton, interact with other membrane receptors, and enhance cell attachment between cells of either the same or different types ⁹³.

2.8.4 Epithelial-Mesenchymal Transition (EMT)

EMT is a cellular process in which epithelial cells obtain mesenchymal phenotypes and behavior in response to downregulated epithelial characteristics. Historically, Elizabeth Hay developed the term "epithelial-mesenchymal transformation" in 1968 to characterize the significant cell alterations that occur during embryogenesis; later, it was termed EMT to differentiate it from malignant transformation. ⁹⁴⁻⁹⁷. The process is represented in figure 2.

EMT is induced when cells receive cues from their microenvironment. The epithelial state of cells starting EMT is denoted by stable epithelial cell–cell junctions, interactions with the basement membrane, and apical–basal polarity. These epithelial features are suppressed during EMT in favor of adopting mesenchymal traits due to modifications to gene expression and post-translational regulatory mechanisms. Cells exhibit morphology and cytoarchitecture similar to fibroblasts throughout this process and an enhanced ability to migrate. These migrating cells often have malignant characteristics ⁹⁷.

EMT is commonly believed to happen in the beginning stages of embryonic growth to promote various morphogenetic processes later in development, and it also happens during the wound-healing process ⁹⁸. Furthermore, EMT has been linked to cancer etiology and tissue fibrosis. However, during development, the opposite process,

known as a mesenchymal-epithelial transition (MET), also happens. The transition from an epithelial to a mesenchymal state is usually imperfect *in vivo*, resulting in cells having both epithelial and mesenchymal traits; based on the biological environment, these transitional phases may appear in various ways^{99, 100}. Because of the increasing complexity and diversity of the EMT literature, definitions of EMT and associated nomenclature have become unclear and frequently confusing⁹⁵. Over the last two decades, the field of EMT research has expanded rapidly. In recent years, the vast majority of published studies on EMT have focused on EMT research in tumor biology¹⁰¹⁻¹⁰⁶.

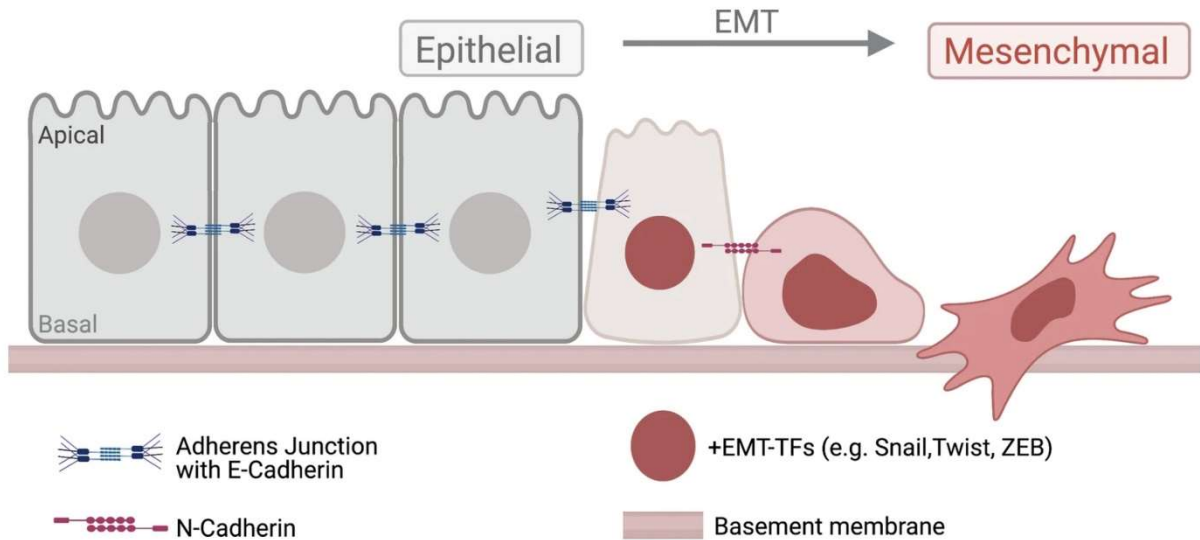


Figure 2: Schematic illustration of EMT. EMT is a complex process in which the activation of EMT-TFs causes the breakdown of cell-cell junctions, the loss of apical-basal polarity, and the upregulation of new cadherins. Modified from Amack¹⁰¹.

2.8.5 Role of EMT in Cancer

The EMT provides epithelial cells with the characteristics necessary for infiltration and metastatic potentials, such as invasiveness and the capability to degrade extracellular matrix (ECM) proteins⁹⁴. One of the challenges in understanding EMT is that the

transitions from epithelial to mesenchymal states are not binary. Instead, cancer cells frequently exhibit a mix of epithelial and mesenchymal features ^{107, 108}.

Metastatic disease, rather than primary tumors, is responsible for around 90% of cancer-related mortalities ¹⁰⁹. According to literature, the most frequent epithelial markers are cytokeratins, ECAD, and occludins, whereas the most prominent mesenchymal markers are NCAD and vimentin. ¹¹⁰. EMT has been shown to play critical roles in carcinogenesis and metastasis for decades. Most life-threatening human cancers arise from epithelial tissues, such as the liver, breast, ovary, kidney, prostate, pancreas, and colon ¹¹¹.

Circulating tumor cells (CTCs) are cancer cells discharged from a solid tumor and enter the peripheral bloodstream; they are thought to be a biomarker of the metastatic process ¹¹². Many cancer cells may not undergo full EMT but instead develop these hybrid epithelial/mesenchymal characteristics over time. Over the full EMT phenotype, hybrid E/M phenotypes provide multiple advantages. Partial EMT or hybrid E/M phenotypes have some advantages over the full EMT phenotype. It is proposed that these migratory cells are cancer cell that separated from the original tumor, infiltrated surrounding tissues, and intravasated into lymphatic and blood arteries, eventually colonizing lymph nodes and distant organs ^{94, 107}. The majority of CTCs have hybrid E/M markers, indicating incomplete EMT. Furthermore, while EMT is critical in tumor progression, its reverse phase, MET, is also important in tumor dispersion. The final stage of the invasion-metastasis cascade is known as colonization, and it is heavily reliant on MET ^{113, 114}.

2.9 Tissue microarray (TMA) as a method for histologic cancer studies

Kononen described the TMA in 1998 ¹¹⁵, as a high-throughput method for assessing histology-based laboratory procedures such as immunohistochemistry and fluorescence in-situ hybridization (FISH). Small cylindrical cores are taken from typical

formalin-fixed, paraffin-embedded tissue and organized in a matrix format within a recipient paraffin block, allowing a pathologist to quickly analyze hundreds of patient samples. TMAs have been used to investigate tumor biology, evaluate novel genetic biomarkers, and ensure laboratory quality assurance since their introduction. The TMA is also an ideal platform for validation and translation for other sorts of high-throughput molecular research. The TMA has proven indispensable for the study of tumor biology, the creation of diagnostic methods, and the analysis of oncological biomarkers ¹¹⁶.

TMAs have specific advantages over other molecular techniques, such as DNA microarrays and proteomics, in the field of molecular epidemiology. Tissues that have been formalin-fixed and paraffin-embedded serve as the basis for TMAs and are the most popular way to preserve surgical specimens. Since many hospitals must keep archival tissue blocks for at least 20 years, the source material for TMAs is widely accessible and frequently associated with long-term outcome information ¹¹⁶.

3. Aim of this study

The expression of ECAD and NCAD at the invasive front as a surrogate parameter for the epithelial-mesenchymal-transition process and its possible predictive value in terms of post-recurrence survival in recurrent OSCC have not been addressed in the existing research. In order to translate the EMT/cadherin-switch phenomenon as a histologic risk factor into a clinical setting, we needed a standardized evaluation and interpretation method of the EMT results, as well as an assignment of its degree to a specific risk profile and prognosis in an epidemiologic clinical setting.

We predicted that the cadherin switch in primary and re-OSCC specimens is an inherent characteristic of the tumor, influences its biological behavior, and further predicts post-recurrence survival outcomes in these patients.

To test this hypothesis, we analyzed the immunohistochemical expression of ECAD and NCAD in the primary and recurrent OSCC patients in a prospectively maintained, single-center cohort. We studied the post-recurrence survival of this high-risk group and correlated it with the standardized *h-score*-based immunohistochemical expression of both cadherin types with the clinical outcome (oral cancer-specific survival, overall survival, and post-recurrence disease-free survival).

To the best of our knowledge, this is the first study to investigate combined ECAD and NCAD expression as a predictor of EMT in relation to OSCC survival outcomes.

4. Material and methods

4.1 Study population

We identified all 1088 cancer patients who presented between 1992 and 2019 to the Department of Maxillofacial Surgery at the University Medical Centre of Lübeck, Germany, with primary non-metastatic OSCC. We included only patients with curative intent who underwent surgery alone or in combination with (chemo)-radiotherapy, as described in figure 3.

To ensure regular follow-up during the 5-year post-therapeutic period, all patients were enrolled in a strictly regulated recall system (every three months in the first two years and every six months after that). At the outset of the study and at each follow-up, data were accessible, including demographic information, risk factors, clinical tumor characteristics, and treatment decisions. The general condition of the patients, estimated using Charlson's comorbidity index (CCI) score ¹¹⁷, tumor stage, and other competing risk factors, were considered in the cohort analysis.

Inclusion Criteria

All OSCC patients with recurrence (N=94) were included in this study irrespective of age, sex, clinicopathological characteristics.

Exclusion Criteria

Patients with oropharyngeal carcinoma (N=87) or oral non-squamous cell carcinoma (N=131) were excluded from the study. Additionally, exclusion criteria included metastatic disease at diagnosis (N=87), patients who refused treatment or died prior to therapy (N=945), and patients who did not have a locoregional recurrence (N=815).

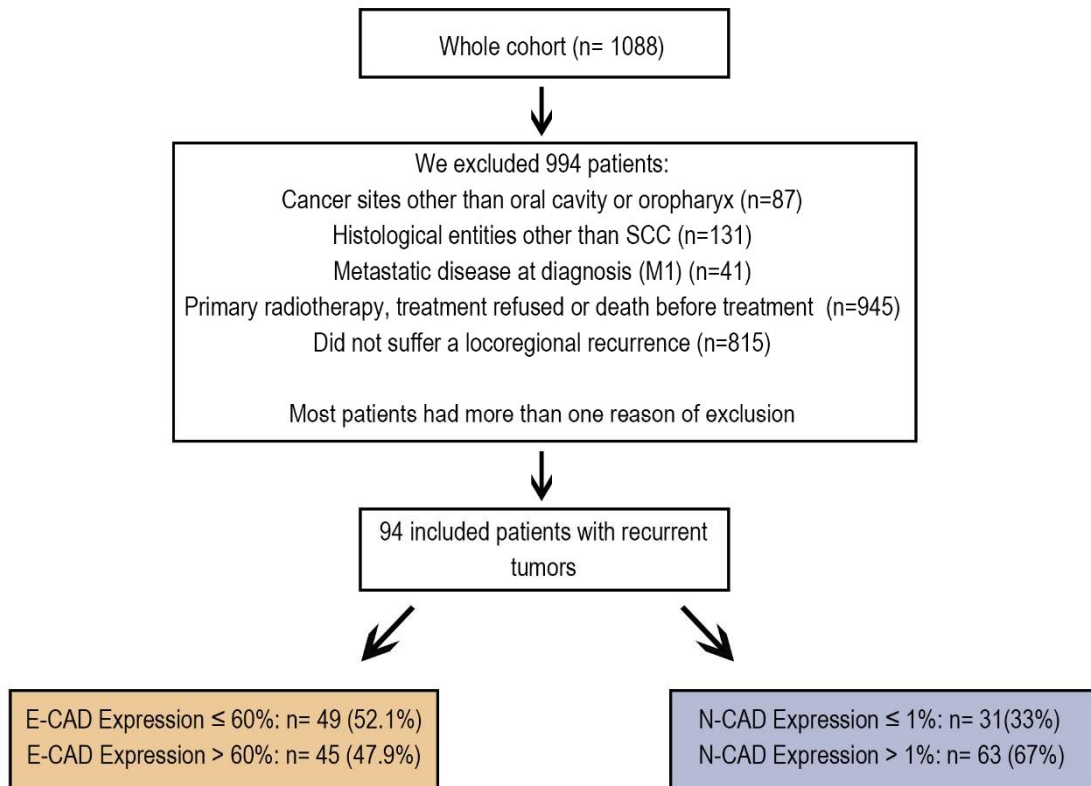


Figure 3: A flow chart depicts the inclusion/exclusion criteria for patients, as well as their assignment to the various cut-off groups for ECAD loss and NCAD de novo expression in primary and recurrent OSCC.

4.2 TMA construction and analysis

We obtained archived, formalin-fixed, paraffin-embedded tissue from surgically resected primary and recurrent oral cancer specimens from the Department of Pathology Bank at the University Hospital of Lübeck. The specimens contained tumor and adjacent normal tissues. The tissues were gathered between 1992 and 2019. For our tissue microarrays, we included 94 patients. The detailed clinical and pathologic information for these patients were available, including demographics, smoking history, clinical and pathologic TNM stage, overall survival duration, and time to recurrence.

All data concerning these patients were collected from our clinical patient database software (Agfa Orbis®). All data was re-evaluated and double-checked by two experienced maxillofacial surgeons and a pathologist.

Hematoxylin and Eosin slides and paraffin blocks of these patients were obtained from the archive. Tissue specimens were available from primary tumors, local recurrent tumors, and/or lymph node metastases (in cases with locoregional recurrence). Tissue samples were re-evaluated in order to categorize each case using the most recent TNM classification (8th edition) and UICC stages⁵⁰.

On Hematoxylin and Eosin slides, regions of interest (ROIs) were marked, and paraffin blocks were matched. Three 0.1 cm cores (triplets) were punched out of each tumor and organized in acceptor blocks as tissue microarrays (TMAs). Cores of patients that did not contain tumor tissue, had staining artifacts, or contained tissue folds were eliminated¹¹⁸.

Prostate cancer tissues were used as a positive control for ECAD, while invasive lobular carcinoma specimens were used as a negative control. Positive controls for NCAD staining were renal cell carcinoma and high-grade ovarian cancer, and negative controls were prepared by omitting the primary antibodies from the staining procedure.

TMA sections were stained with the Ventana BenchMark staining system and detected with the IViewDAB detection kit (both available from Roche, Basel, Switzerland)¹¹⁹, as mentioned in table 2.

For both ECAD and NCAD staining, heat-mediated antigen retrieval was done for 32 minutes at 92 °C with Cell Conditioning Solution 1 (CC1; #950-124, Ventana Medical Systems, Inc., Arizona, USA). The primary antibodies are mentioned in Table 1. ECAD and NCAD were used as membranous markers and cytoplasmatic markers.

Table 2: Antibodies used for the immunostaining of the Tissue microarray

Antibody	Isotype	Company	Concentration/Dilution	Clone
E-cadherin	Mouse monoclonal	Roche, Basel, Switzerland	0.314 µg/ ml	CDH1
N-cadherin	Rabbit polyclonal	Abcam, Cambridge, United Kingdom	1:100	CDH2

An excel sheet map was created before the TMA was built, as described in figure 4. This map aimed to direct assembly and subsequent scoring by designating a place within the TMA for each core sample. A tissue microarrayer was necessary for the physical fabrication of the TMA (Beecher Instruments, Sun Prairie, WI, USA). Each TMA held up to 180 tumor samples and 15 normal tissue samples as triplet cores of 1 mm² diameter.

orientation	1	1	1	2	2	2	11	11	11	12	12	12	21	21	21	22	22	22
orientation	3	3	3	4	4	4	13	13	13	14	14	14	23	23	23	24	24	24
orientation	5	5	5	6	6	6	15	15	15	16	16	16	25	25	25	26	26	26
	7	7	7	8	8	8	17	17	17	18	18	18	27	27	27	28	28	28
	9	9	9	10	10	10	19	19	19	20	20	20	29	29	29	30	30	30
	31	31	31	32	32	32	41	41	41	42	42	42	51	51	51	52	52	52
	33	33	33	34	34	34	43	43	43	44	44	44	53	53	53	54	54	54
	35	35	35	36	36	36	45	45	45	46	46	46	55	55	55	56	56	56
	37	37	37	38	38	38	47	47	47	48	48	48	57	57	57	58	58	58
	39	39	39	40	40	40	49	49	49	50	50	50	59	59	59	60	60	60

Figure 4: yellow blocks: orientation cores, green blocks: normal tissues, white blocks: tumor



Figure 5: E-Cadherin tissue microarray cores under low magnification

4.3 Immunohistochemical analysis

The slides were visualized using the Ventana iScan HT scanner (Ventana, Tuscon, AZ, USA), as seen in figure 5. For digital evaluation of the slides, the image analysis software QuPath (University of Edinburgh, UK) was used. The file extension used in QuPath was “.tif” The data were analysed using ThinkPad P1 Mobile workstation, Intel® Xeon® processor, 32GB of RAM and OLED Screen with 4K resolution ¹²⁰.

4.3.1 QuPath Software steps to analyze the antibodies:

- 1- First, a new project is created. Then, all images/data concerning this antibody are added to the selected folder. The cores are then identified using the TMA dearrayer feature. This step is essential to quickly identify, analyze, and export every core separately.
- 2- In the preprocessing phase, the stain vectors have to be estimated to improve stain separation.
- 3- Select all cores and execute the cell detection feature after adjusting all parameters based on the antibody (Threshold, Max background intensity). In our case we used the default setting for our antibodies.
- 4- The ROI are annotated and categorized into different groups: Tumor, Immune cells, Stroma, and Others
- 5- The Object Classifier feature is run to detect and analyze the number of different categories.

4.3.2 Evaluation and scoring of immunostaining results

The immunostaining evaluation based on the h-score was applied to a maximum of 300. Subsequently, this was formed by adding the percentage of strongly labeled cells (weighted 3), the percentage of moderately stained cells (weighted 2), and the percentage of weakly stained cells (weighted 1), resulting in a range of 0 to 300¹²¹. Immunostaining was performed based on the proportion of positively stained tumor cells relative to the total investigated tumor area.

The assessment of NCAD immunostaining considered the proportion of tumor cells with positive staining in relation to the total evaluated tumor area. When cytoplasmic or membrane immunostaining was observed in epithelial tumor cells, NCAD expression was considered to be positive, as shown in figures 8 and 9.

Positive ECAD expression was determined when membranous immunostaining was observed in epithelial tumor cells, as seen in figures 6 and 7. For loss of ECAD staining in epithelial cells, an inverse estimation of the h-score-based evaluation was applied as 300 minus ECAD staining in order to normalize to NCAD values and corresponding vector development in both antibodies.

The ECAD and NCAD staining scores were entered as a continuous variable, and a cutoff was established for both categories depending on the available sample size.

The appropriate cut-offs for biomarkers were calculated using the R package 'Survminer,' which employs the maximum selected rank statistics in multivariate Cox regressions to offer the value of a cut-off that most strongly correlates with the outcome. Based on the *h-score*, a cut-off value of 60% was defined for ECAD loss of expression and 1% for NCAD expression. The significance level was set at p 0.05.



Figure 6: preserved expression of E-Cadherin in a representative section

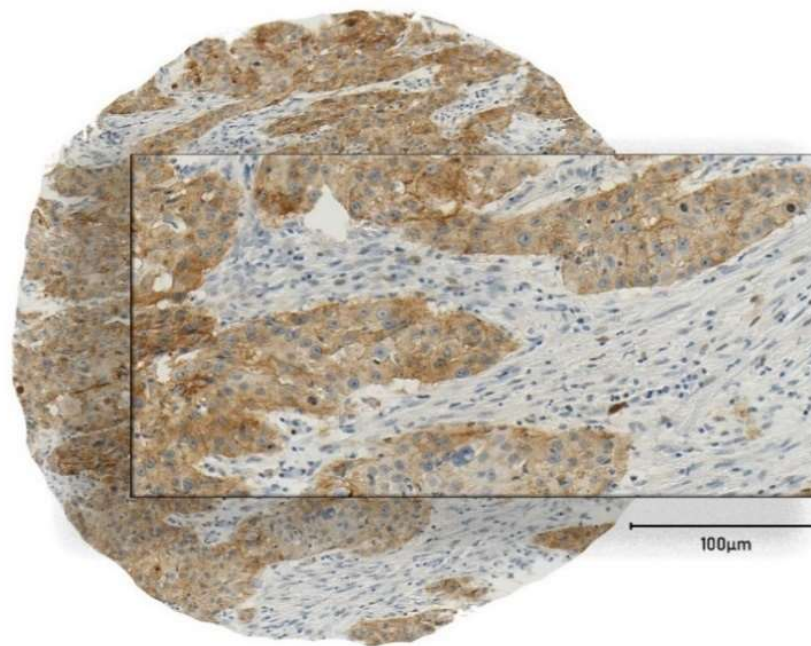


Figure 7: Loss of membrane staining in a representative section stained

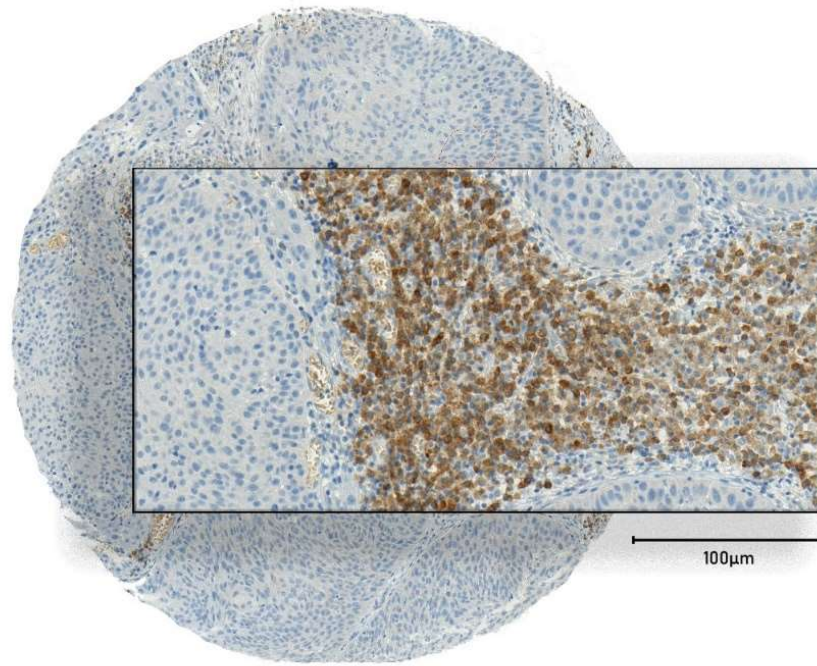


Figure 8: high expression of N-Cadherin in a representative section



Figure 9: low expression of N-Cadherin in a representative section

4.4 Statistical Analysis

From the time of recurrence, all survival outcomes were calculated. Post-recurrence disease-free survival (pr DFS) was measured by the incidence of local or regional recurrence. Overall post-recurrence survival (pr OS) was measured by death from any cause. Post-recurrence oral cancer-specific survival (pr OCSS) was measured by oral cancer mortality. Patients were censored at the most recent follow-up.

R Statistical Software was utilized for all statistical studies (version 4.0.4; R Foundation for Statistical Computing, Vienna, Austria). The proportional hazards assumption was evaluated using Schoenfeld residual plots.

4.5 Ethics

All participants signed consent forms upon admission, allowing their data to be collected and used anonymously for academic research. The study was approved by the University of Lübeck's ethics review committee (ID: 12-079A).

5. Results

5.1 Characteristics of patients

Within the investigated cohort, 94 patients with recurrent OSCC were included. The recurrence age ranged between 53 and 72 years. 68% of local recurrences were associated with cervical lymph node metastasis, while only 8.8% were associated with distant metastasis. Eleven percent of patients received radiochemotherapy as an adjuvant treatment, while fifty percent solely received radiotherapy. The remaining patients were only surgically treated (Table 3).

The majority of patients reported a positive smoking history (n = 69, 78%), as well as excessive alcohol intake (n = 52, 60%). A safe resection margin (R0) was demonstrated in 19 cases (68%). Total resection (R1) was not achieved in 9 (32%) patients. Patients with distant metastases made up 8.8% of all patients (Table 3).

Tumors were categorized as rT1 (r for recurrent) in 24 patients (26%), rT2 in 16 (18%), rT3 in 10 (11%), and rT4 in 28 (31%). There were 63 (68%) patients with rN+ nodal status and 29 (32%), with no lymph node metastases (rN0). The most common diagnosis was moderately differentiated oral squamous cell carcinoma (G2), with 40 patients (59%) falling into this category according to Table 3.

The most commonly affected areas were the floor of the mouth (n= 39, 41%), the neck solely (n= 15, 16%), the cheek/ vestibule/ retromolar (n= 13, 14%), and the anterior tongue (n=11, 12%) (Table 3).

Based on the *h-score*, appropriate cut-offs for biomarkers were established; a cut-off value of 60% was defined for ECAD loss of expression and 1% for NCAD expression.

Males were predominant in the group of patients with ECAD expression loss of less than or equal to 60%, while sex distribution in the expression of NCAD was comparable, regardless of expression levels. The CCI score was similar regardless of both NCAD and

ECAD expression. In the ECAD group with expression loss less than or equal to 60%, excessive alcohol smoking (73%) was observed. The most common site of recurrence was the floor of the mouth ranging between 40-45%, when comparing the 4 subgroups of expression of both markers. In all subgroups, between 55 and 75% of patients showed local lymph node recurrences. The resection margins were clear in 75% of patients with ECAD expression loss less than or equal to 60% and 80% of patients with NCAD expression more than 1%. In all groups, the majority of patients presented demonstrated moderate OSCC grade (Table 3).

Irrespective of NCAD and ECAD expression levels, the vast majority of patients did not have distant metastases (Table 3).

ECAD and NCAD expression among the well-differentiated OSCC tended to maintain steady levels (Table 3). Forty-three percent of re-OSCCs categorized as G3 (poorly differentiated) exhibited an E-cadherin expression loss of less than or equal to 60%, while only 21% of patients with poorly differentiated OSCC showed an ECAD expression loss of more than 60% (Table 3). On the other hand, regardless of NCAD expression, most of the patients showed moderately differentiated OSCC (NCAD \leq 1%: 62%, NCAD $>$ 1%: 57%, Table 3).

Table 3: Clinical characteristics of patients, including tumor stage and IHC expression of E-cadherin and N-cadherin in recurrent oral squamous cell carcinoma, are included in this thorough descriptive analysis.

Variable	Overall	strata by ECAD loss		strata by NCAD ²	
	N = 94 ¹	≤ 60% N=49 (52%) ¹	> 60% N=45 (48%) ¹	≤ 1% N=31 (33%) ¹	> 1% N=63 (67%) ¹
Age at recurrence diagnosis	63 (53-72)	63 (51-71)	63 (55-73)	62 (52-73)	64 (54-72)
Sex					
<i>Female</i>	30 (32%)	10 (20%)	20 (44%)	10 (32%)	20 (32%)
<i>Male</i>	64 (68%)	39 (80%)	25 (56%)	21 (68%)	43 (68%)
CCI score					
0	59 (63%)	32 (67%)	27 (60%)	21 (68%)	38 (61%)
1 ≤	34 (37%)	16 (33%)	18 (40%)	10 (32%)	24 (39%)
Missing	1	1	0	0	1
Smoking					
<i>Never</i>	20 (22%)	8 (18%)	12 (27%)	5 (17%)	15 (25%)
<i>Former or current</i>	69 (78%)	37 (82%)	32 (73%)	24 (83%)	45 (75%)
Missing	5	4	1	2	3
Alcohol consumption					
<i>None or moderate</i>	35 (40%)	12 (27%)	23 (53%)	10 (34%)	25 (43%)
<i>Excessive</i>	52 (60%)	32 (73%)	20 (47%)	19 (66%)	33 (57%)
Missing	7	5	2	2	5
Site of recurrence					
<i>Anterior tongue</i>	11 (12%)	4 (8.2%)	7 (16%)	5 (16%)	6 (9.5%)
<i>Cheek/vestibule/retromolar</i>	13 (14%)	6 (12%)	7 (16%)	3 (9.7%)	10 (16%)
<i>Floor of mouth</i>	39 (41%)	20 (41%)	19 (42%)	14 (45%)	25 (40%)
<i>Lip</i>	2 (2.1%)	0 (0%)	2 (4.4%)	0 (0%)	2 (3.2%)
<i>Neck only</i>	15 (16%)	9 (18%)	6 (13%)	7 (23%)	8 (13%)
<i>Oropharynx</i>	11 (12%)	8 (16%)	3 (6.7%)	2 (6.5%)	9 (14%)
<i>Palate</i>	3 (3.2%)	2 (4.1%)	1 (2.2%)	0 (0%)	3 (4.8%)
rT					
<i>rT1</i>	24 (26%)	12 (26%)	12 (27%)	5 (17%)	19 (31%)
<i>rT2</i>	16 (18%)	8 (17%)	8 (18%)	8 (27%)	8 (13%)
<i>rT3</i>	10 (11%)	5 (11%)	5 (11%)	3 (10%)	7 (11%)
<i>rT4</i>	28 (31%)	16 (34%)	12 (27%)	13 (43%)	15 (25%)
<i>rTx</i>	13 (14%)	6 (13%)	7 (16%)	1 (3.3%)	12 (20%)
Missing	3	2	1	1	2
rN					
<i>rN0</i>	29 (32%)	15 (31%)	14 (32%)	14 (45%)	15 (25%)
<i>rN+/x</i>	63 (68%)	33 (69%)	30 (68%)	17 (55%)	46 (75%)
Missing	2	1	1	0	2
rM					
<i>rM0/x</i>	83 (91%)	46 (98%)	37 (84%)	28 (90%)	55 (92%)
<i>rM1</i>	8 (8.8%)	1 (2.1%)	7 (16%)	3 (9.7%)	5 (8.3%)
Missing	3	2	1	0	3

Variable	Overall N = 94 ¹	strata by ECAD loss		strata by NCAD ²	
		≤ 60% N=49 (52%) ¹	> 60% N=45 (48%) ¹	≤ 1% N=31 (33%) ¹	> 1% N=63 (67%) ¹
Resection margins					
<i>R0</i>	19 (68%)	9 (75%)	10 (62%)	7 (54%)	12 (80%)
<i>R1/2/x</i>	9 (32%)	3 (25%)	6 (38%)	6 (46%)	3 (20%)
<i>Missing</i>	66	37	29	18	48
Grade					
<i>Well</i>	6 (8.8%)	2 (5.7%)	4 (12%)	2 (7.7%)	4 (9.5%)
<i>Moderate</i>	40 (59%)	18 (51%)	22 (67%)	16 (62%)	24 (57%)
<i>Poor</i>	22 (32%)	15 (43%)	7 (21%)	8 (31%)	14 (33%)
<i>Missing</i>	26	14	12	5	21

¹Median (25%-75%); n (%), ²NCAD expression, CCI: Charlson Comorbidity Index; rT: recurrent tumor; rN: recurrent regional lymph node; rM: recurrent distant metastasis.

5.2 Pattern of E-cadherin and N-cadherin immunohistological expression

Strong membranous ECAD expression was observed in intact epithelial regions close to the malignant epithelium (Figure 6), while degrees of reduced staining intensity and partial to complete absence of membranous labeling were observed in primary and re-OSCC (Figure 7). The suprabasal layer, epithelial cancer cells on the invasion front, and neoplastic epithelial nests in the stroma all showed evidence of this phenomenon.

Regardless of the tumor's degree of histological differentiation, NCAD was present in the cytoplasm and membrane of dispersed tumor cells within the invasion front and in the stroma at varying intensities (Figure 8, 9). ECAD and NCAD staining patterns were consistent across specimens and in primary and re-OSCC.

5.3 E-cadherin and N-cadherin expression in primary and recurrent malignancies

ECAD expression ranged from 43-62% (mean=53%) in primary tumors and was marginally higher in recurrent cases (mean = 57%) (Figure 10). A similar effect was seen for NCAD, which showed a non-significant increase from primary tumor tissues

(mean=2.88%, range: 0.29-4.09%) to recurrent tumor tissues (mean=5% range: 1-7%), as illustrated in Figure 10.

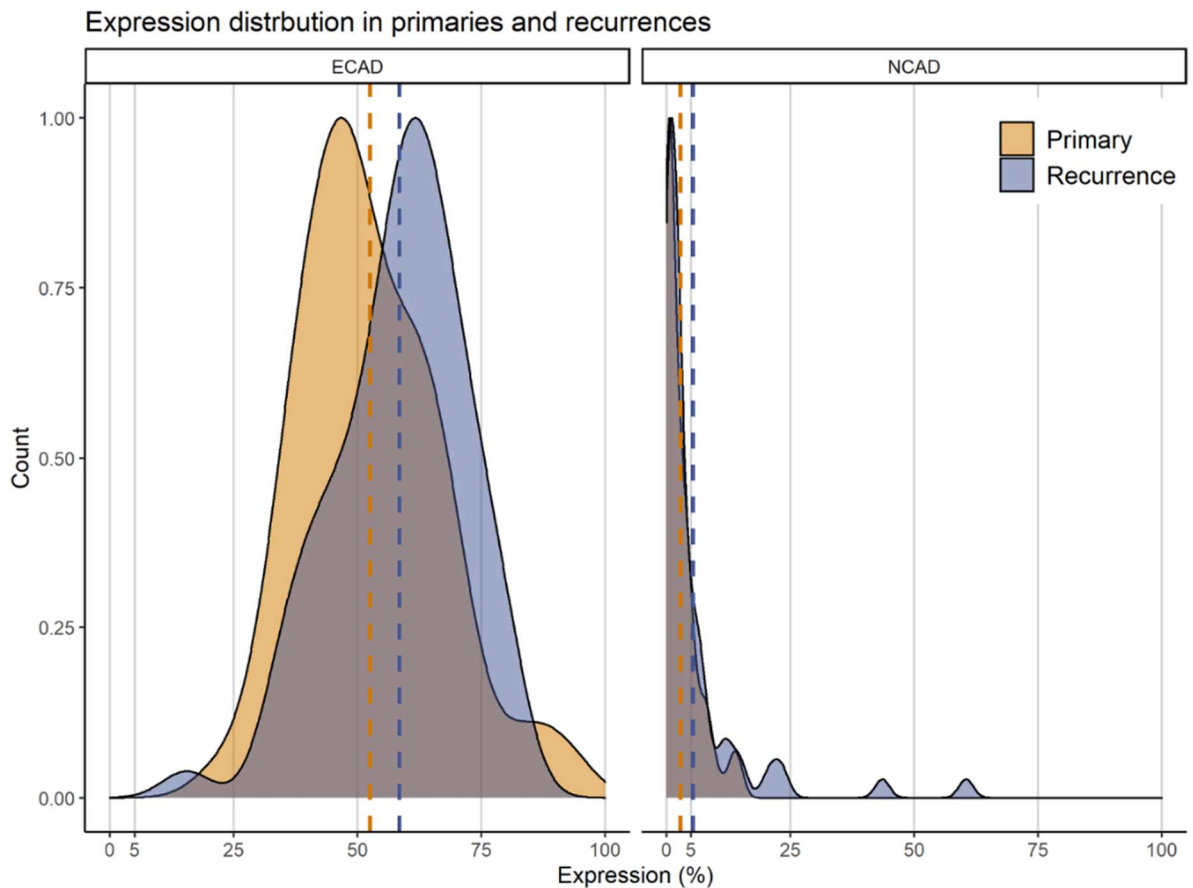


Figure 10: Histogram showing the comparison between the expression of ECAD and NCAD in primary and recurrent tumors (dotted lines represent mean value).

Overall, there was no significant difference in the expression of ECAD and NCAD between the primary and recurrent tumors, irrespective of treatment modality (Figure 11). Exceptionally, ECAD expression in the primary tumor was decreased compared to recurrent tumors in the irradiated group (Figure 11).

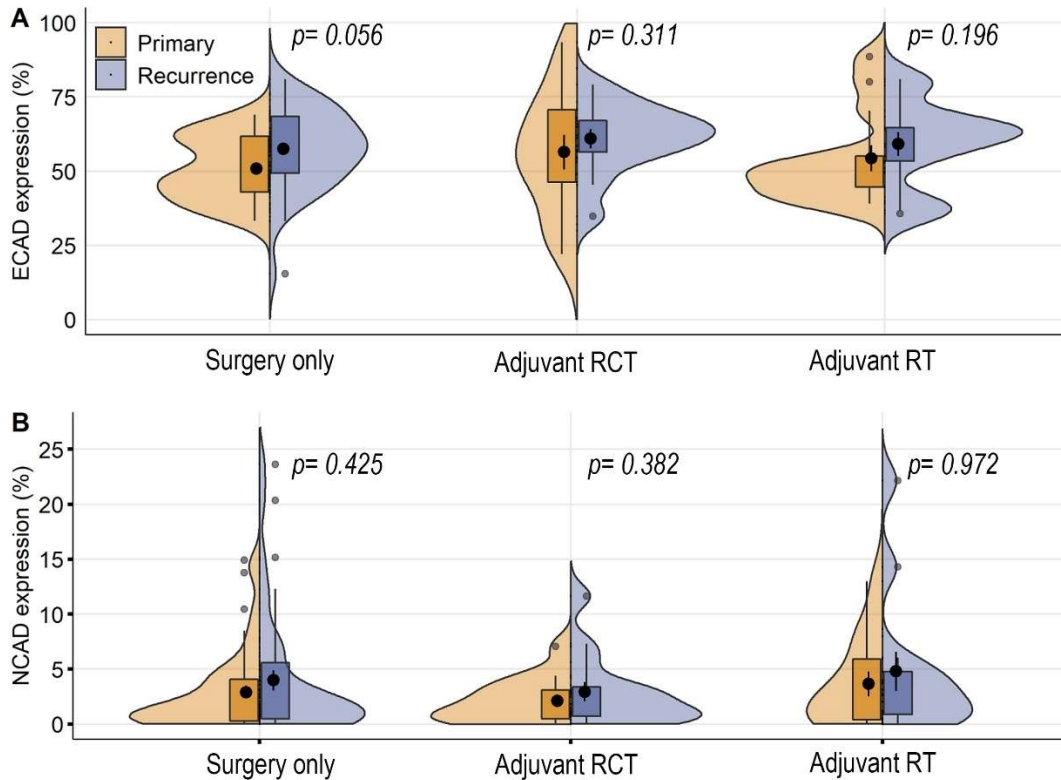


Figure 11: Distributions of the numerical data for ECAD (A) and NCAD (B) expression in primary and recurrent tumors are shown in violin plots divided into 3 subgroups (surgery only, radiotherapy, and radiochemotherapy).

These findings reveal a stable level of initial cadherin switch within the primary tumors, which does not change during the recurrence phase and appears to be irrespective of the type of adjuvant therapy (radio- or radiochemotherapy) given between the initial diagnosis and recurrence.

5.4 Survival outcomes for the E-cadherin and N-cadherin expression

Based on the *h-score* cutoff values presented in Table 5, the ECAD and NCAD survival events were evaluated. Thirty-six patients (80%) who passed away exhibited a loss of

ECAD expression greater than 60%, and of this group, 32 (74%) passed away due to complications of oral cancer. The remaining patients died from other reasons; in the group of patients with ECAD expression loss less than or equal to 60%, 31 patients (65%) died, with 25 (53%) of them died from complications related to the tumor (Table 5).

In the NCAD group, 43 (69%) of the deceased patients had an NCAD expression of more than 1%, while 24 (77%) of the deceased patients had an NCAD expression of less than or equal to 1%. Furthermore, 23 (77%) patients in the group with lower NCAD expression perished due to oral cancer, compared to 34 (57%) patients in the group with higher NCAD expression (Table 5).

Table 5: Mortality rates based on the *h-score* cutoff for E-cadherin and N-cadherin expression.

Variable	Events by ECAD loss		Events by NCAD ²	
	≤ 60% N = 49 (52%) ¹	> 60% N = 45 (48%) ¹	≤ 1% N = 31 (33%) ¹	> 1% N = 63 (67%) ¹
Death from any cause				
<i>Alive or censored</i>	17 (35%)	9 (20%)	7 (23%)	19 (31%)
<i>Dead</i>	31 (65%)	36 (80%)	24 (77%)	43 (69%)
<i>missing</i>	1	0	0	1
Cause of death				
<i>Alive or censored</i>	17 (36%)	9 (21%)	7 (23%)	19 (32%)
<i>Death from oral cancer</i>	25 (53%)	32 (74%)	23 (77%)	34 (57%)
<i>Death from other causes</i>	5 (11%)	2 (4.7%)	0 (0%)	7 (12%)
<i>missing</i>	2	2	1	3

¹n (%), ²NCAD expression

The 2-year pr OS, pr OCSS, and pr DFS rates in the ECAD group less than or equal to 60% were 41% (29-57%), 46% (34-63%) and 34% (22-52%), respectively, whereas the rates in the group greater than 60% were 24% (15-41%), 72% (60-87%), and 52% (34-78%), respectively (Table 6). The 5-year pr OS, pr OCSS, and pr DFS rates in the ECAD group

less than or equal to 60% were 30% (20-47%), 55% (42-72%) and 41% (28-61%), respectively, whereas the rates in the group greater than 60% were 17% (8.9-33%), 79% (67-93%), and 61% (42-93%), respectively (Table 6).

The 2-year pr OS, pr OCSS, and pr DFS rates in the NCAD group $\leq 1\%$ were 29% (17-50%), 70% (55-88%), and 39% (22-70%), respectively, whereas the rates in the group greater than 1% were 35% (25-49%), 53% (42-67%), and 40% (28-58%), respectively (Table 6). The 5-year pr OS, pr OCSS, and pr DFS rates in the NCAD group $\leq 1\%$ were 19% (8.9-40%), 81% (67-96%), and 39% (22-70%), respectively, whereas the rates in the group greater than 1% were 27% (18-40%), 58% (47-72%), and 51% (37-69%), respectively (Table 6).

Table 6: Survival outcomes of patients based on of E-cadherin and N-cadherin expression.

Variable	pr OS		pr OCSS		pr DFS	
	at 2 years	at 5 years	at 2 years	at 5 years	at 2 years	at 5 years
Overall	33% (25%-44%)	24% (17%-35%)	59% (49%-70%)	66% (57%-77%)	40% (30%-55%)	48% (36%-64%)
ECAD loss						
$\leq 60\%$	41% (29%-57%)	30% (20%-47%)	46% (34%-63%)	55% (42%-72%)	34% (22%-52%)	41% (28%-61%)
$> 60\%$	24% (15%-41%)	17% (8.9%-33%)	72% (60%-87%)	79% (67%-93%)	52% (34%-78%)	61% (42%-91%)
NCAD²						
$\leq 1\%$	29% (17%-50%)	19% (8.9%-40%)	70% (55%-88%)	81% (67%-96%)	39% (22%-70%)	39% (22%-70%)
$> 1\%$	35% (25%-49%)	27% (18%-40%)	53% (42%-67%)	58% (47%-72%)	40% (28%-58%)	51% (37%-69%)

²NCAD expression, pr DFS: post-recurrence disease-free survival; pr OS: post recurrence survival; pr OCSS: post-recurrence oral cancer-specific survival.

5.5 Competing Risk Analysis and Hazard Ratio for E-cadherin and N-cadherin expression

Clinicopathological variables (age, sex, CCI, smoking, tumor size, alcohol, nodal metastasis) were analyzed in Table 7. Using Cox proportional hazard regression analysis, the hazard ratios (HR) and 95% confidence intervals (CI) for overall, oral cancer-specific, and post-recurrence disease-free survival were estimated. Multivariate analysis for pr OS and pr OCSS found no statistically significant differences for potential risk factors such as sex, CCI, smoking, and alcohol, with pr OS HR values of 0.97, 1.06, 1.74, and 1.28, respectively, and pr OCSS HR values of 0.93, 1.45, 1.50, and 1.30, respectively (Table7).

In tumors larger than rT1, pr OS as well as pr OCSS were both significantly decreased. The worst prognosis for pr OS and pr DFS was for rT2 tumors (HR= 6.24, CI 2.26-17.3, $p= 0.001$; HR= 4.11, CI: 1.45-11.6, $p= 0.008$), while the worst prognosis for pr OCSS was for rT3 (HR= 8.29, CI: 2.53-27.1, $p=0.001$). Both pr OS and pr OCSS were significant in patients with positive nodal status (pr OS HR= 2.48, CI: 1.29-4.75, $p= 0.006$ and pr OCSS HR= 2.39, CI: 1.15-4.99, $p= 0.020$) (Table7).

There was a substantial association between the ECAD loss more than 60% and pr OS as well as pr OCSS (HR=2.72, CI:1.50-4.95, $p= 0.001$, HR=3.84, CI:1.93-7.63, $p= 0.001$), respectively, when comparing expression loss more than 60% to low ECAD loss in tumor cells (Table 7, Figure 11).

There was no statistically significant association between the *de novo* expression of N-cadherin and pr OS, pr OCSS, or pr DFS. The results are presented in detail in Table 7. All survival outcomes in the analysis were assessed from the first recurrence, as seen in Kaplan-Meier curves (Figure 11).

Table 7: Hazards ratios for different prognostic factors in recurrence specimens using the h-score based cut-offs of E-cadherin (60%) and N-cadherin (1%).

Characteristic	pr OS			pr OCSS			pr DFS		
	HR ¹	95% CI ¹	p-value	HR ¹	95% CI ¹	p-value	HR ¹	95% CI ¹	p-value
Age at recurrence diagnosis	1.01	0.98-1.03	0.6						
Sex									
<i>Female</i>	—	—		—	—				
<i>Male</i>	0.97	0.47-1.96	>0.9	0.93	0.43-2.00	0.9			
CCI									
0	—	—		—	—				
1 ≤	1.06	0.61-1.85	0.8	1.45	0.79-2.63	0.2			
Smoking									
<i>Never</i>	—	—		—	—				
<i>Former or current</i>	1.74	0.74-4.07	0.2	1.50	0.57-3.92	0.4			
Alcohol									
<i>None or moderate</i>	—	—		—	—				
<i>Excessive</i>	1.28	0.66-2.47	0.5	1.30	0.64-2.66	0.5			
rT									
<i>rT1</i>	—	—		—	—		—	—	
<i>rT2</i>	6.24	2.26-17.3	<0.001	6.00	1.96-18.4	0.002	4.11	1.45-11.6	0.008
<i>rT3</i>	5.85	1.98-17.3	0.001	8.29	2.53-27.1	<0.001	1.60	0.41-6.25	0.5
<i>rT4</i>	4.72	1.97-11.3	<0.001	5.15	1.85-14.3	0.002	1.43	0.51-4.00	0.5
<i>rTx</i>	4.22	1.55-11.5	0.005	7.48	2.41-23.3	<0.001	0.99	0.26-3.81	>0.9
rN									
<i>rN0</i>	—	—		—	—		—	—	
<i>rN+/x</i>	2.48	1.29-4.75	0.006	2.39	1.15-4.99	0.020	1.27	0.56-2.89	0.6
ECAD loss									
≤ 60%	—	—		—	—		—	—	
> 60%	2.72	1.50-4.95	0.001	3.84	1.93-7.63	<0.001	1.45	0.70-3.04	0.3

Characteristic	pr OS			pr OCSS			pr DFS		
	HR ¹	95% CI ¹	<i>p</i> -value	HR ¹	95% CI ¹	<i>p</i> -value	HR ¹	95% CI ¹	<i>p</i> -value
NCAD²									
≤ 1%	—	—		—	—		—	—	
> 1%	1.23	0.68- 2.21	0.5	0.90	0.47- 1.73	0.8	1.60	0.69- 3.73	0.3

¹HR = Hazard Ratio, CI = Confidence Interval, ²NCAD expression, pr DFS: post-recurrence disease-free survival; pr OS: post-recurrence survival; pr OCSS: post-recurrence oral cancer-specific survival; CCI: Charlson Comorbidity Index; rT: recurrent tumor; rN: recurrent regional lymph node; rM: recurrent distant metastasis.

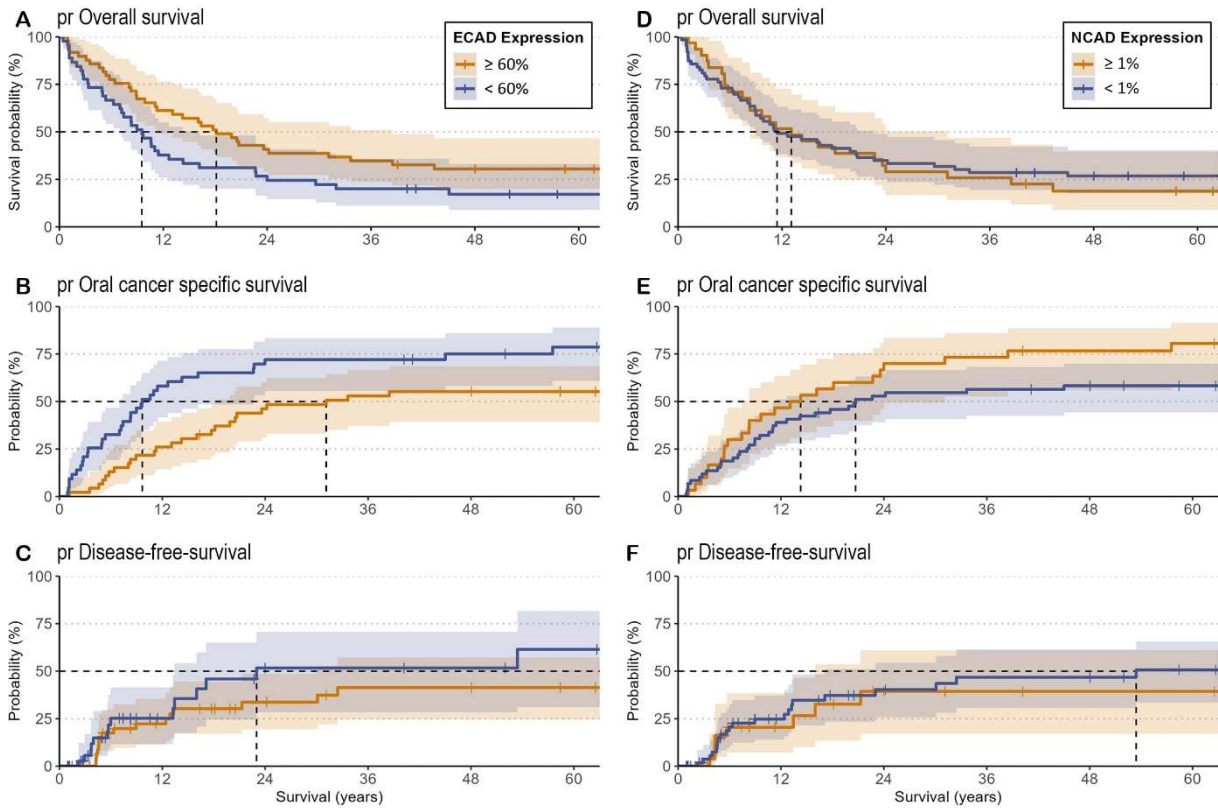


Figure 11: Separate analyses were carried out for ECAD (A-C) and NCAD (D-F). Based on the *h-score*, the cutoff was chosen at 60% for ECAD loss and 1% for N-cadherin *de novo* expression. All outcomes were assessed from the first recurrence. Kaplan-Meier curves for post-recurrence overall survival (pr OS), oral cancer-specific survival (pr OCSS), and post-recurrence disease-free survival (pr DFS) of patients with high ECAD loss and NCAD *de novo* expression (blue curves), as well as for low ECAD loss and negative NCAD expression (yellow curves).

6. Discussion

Nowadays, the way we think about malignant tumors has changed; the tumor is now seen as a complex subset of cancer cells that create TME to build a self-sufficient biological structure ⁷². OSCC is a malignant tumor that arises from the oral mucosa's stratified squamous epithelium ¹²². The lateral borders of the tongue, the floor of the mouth, as well as the lips are the most common sites for the malignant tumor ¹²³. Carcinogenesis starts with a cell being disrupted ³². The treatment of re-OSCC is a challenging inter-disciplinary endeavor. Patients who have undergone first treatment for oral cancer suffer from a variety of sequela, including altered local anatomical structures and functional impairment due to food intake, masticatory and speech deficits, and xerostomia ¹²⁴. In the event of early chemotherapy, other general problems, such as impaired kidney and bone marrow function, may exacerbate these side effects and further restrict local and systemic treatment ¹²⁵. Therefore, it is necessary to adapt the available treatment options to the general condition of these patients and to stratify therapy based on their unique needs and survival probability.

6.1 E-cadherin

ECAD is believed to be a critical tumor-suppressor protein based on the characteristic loss of ECAD-mediated cell adhesion in epithelial malignancies and its role in suppressing tumor growth ^{126, 127}. The findings from a study conducted on glioblastoma multiforme (GBM) is in line with the theory that in a small percentage of high-grade gliomas, ECAD expression plays a significant role in tumor growth and invasiveness. According to this study, patients with GBM who have an epithelial or pseudo-epithelial morphology will have a worse prognosis than those who do not express ECAD ¹²⁸.

Putzke et al. stated that ECAD overexpression is associated with aggressive disease in metastatic prostate cancer ¹²⁹. In another study, high ECAD expression was linked to

the development of ovarian epithelial cancers, as ECAD was only found in benign, borderline, and malignant ovarian epithelial tumors ¹³⁰.

The above-mentioned studies are not consistent with our findings, as we detected an inverse connection between ECAD expression and the histological grade and survival outcome of patients with recurrent OSCC tumors in the present investigation. This might be attributed to the specific type of cancer on the one hand.

On the other hand, comparable studies that focused on OSCC came in line with our observations. Specifically, Pereira et al. observed that there was a reduction in the expression of ECAD at the invasive tumor front in the cytoplasmic membrane/nucleus ¹³¹. Moreover, Peng et al. elucidated the mechanism behind ECAD loss in oral squamous cell carcinoma. They came to the conclusion that miR-134's (microRNA) oncogenic effect on oral cancer is caused by a decrease in PDCD7 (Programmed Cell Death 7) and ECAD expression ¹³². Additionally, ECAD loss was detected in the precancerous stages, which suggests that loss of expression of ECAD is related to the development of OSCC from precursor lesions, as reported by Sharma et al ¹³³.

As reported previously, we noticed a considerable degree of ECAD loss among all tumor specimens in our cohort. ECAD is largely involved in the deregulation of the extracellular matrix during EMT process, and its absence at the invasion front correlates with poor DFS and OS ¹³⁴⁻¹³⁷. ECAD functionality is not considered an easy target for tumor therapy since it appears to be affected by a multitude of mechanisms, namely genetic alterations, transcriptional repression, and suppression of the ECAD adhesion complexes via several signaling pathways ¹²⁶.

6.2 N-cadherin

The expression of NCAD is generally found in different cell types, including endothelial cells, neural cells, osteoblasts, and stromal cells ^{138, 139}. NCAD facilitates angiogenesis and mediates vascular stabilization ⁹¹.

NCAD acts as an indicator of the EMT process, and its expression has been associated with the development of numerous types of cancer¹⁴⁰⁻¹⁴³. In neuroblastoma, down-regulation of NCAD promotes metastasis¹⁴⁴. There was an increase in pluripotency-associated markers in prostate cancer cells that overexpressed NCAD¹⁴⁰.

NCAD expression, as previously indicated in the literature, plays a vital function throughout the EMT process, when it is up-regulated^{139, 145}. However, the NCAD expression in our tumor samples was less than 5%, and it did not change when comparing the expression of primary tumor to its recurrence. When we compare oral cancer to other tumor entities such as prostate cancer, we may conclude that high level of NCAD expression cannot be seen in oral cancer, which is reflective of the fact that NCAD is more frequent in non-epithelial tissues, and the increase of NCAD in normal epithelial cells implies that EMT and cancer growth are impending¹⁰⁶.

The discrepancy in ECAD loss compared to NCAD *de novo* expression raises the question of which of the Cadherins may indicate EMT and subsequently better correlates with post-recurrence survival outcome.

The key difference between the findings provided here and those in the relevant literature is the evaluation approach used for ECAD and NCAD expression. Predominantly, the threshold used to stratify patients is estimated differently. Previously, the cut-off was determined arbitrarily, mostly based on prior investigations, and was subject to multiple, not necessarily standardized immunohistological evaluation methodologies' interpretations. In addition, the available clinical data from the prospectively maintained cohort permit a valid risk-adjusted analysis, as all relevant parameters were collected at baseline and can be effectively incorporated into the applied regression model.

6.3 Effect of adjuvant treatment on the Epithelial-mesenchymal transition

Radiotherapy is used to treat more than fifty percent of cancer patients alone or in combination with other treatments. Current findings has demonstrated that RT could influence the immunobiological traits of OSCC cells. It typically works by either directly causing structural damage to DNA or indirectly causing damage via reactive oxygen species. Although RT is continuously advancing in recent years, most cancer patients continue to have poor prognosis. EMT is one of the most significant elements influencing radio-resistance in tumors, which has been recognized as a formidable hurdle to radiation efficacy ^{146, 147}.

Our findings revealed that adjuvant RT didn't impact the expression of NCAD and ECAD. Yet, several studies have focused on the relationship between EMT and radioresistance. Lin et al suggested that X-Ray irradiation induced EMT through the smad signaling pathway in colorectal carcinoma ¹⁴⁸. Various signaling pathways, such as MAPK/ERK, TGF, HIF-1, Notch and Wnt, are extensively stimulated during carcinogenesis and development, and irradiation will further trigger these pathways, resulting in malignant features, including EMT and radio-resistance. Generally, RT can enhance the production of reactive oxygen species (ROS), which can activate signaling pathways and alter the TME. ROS has been connected to irradiation induced EMT via modulating a number of EMT markers and transcriptional factors ¹⁴⁹.

One of the most popular chemotherapy drugs used now to treat HNSCC is Cisplatin. It can interact with RNA, DNA, and various proteins and induce apoptosis by triggering specific mechanisms ^{150, 151}. There are various disadvantages of utilizing Cisplatin in cancer treatment ¹⁵². One of these is tumor resistance to therapy, several studies have demonstrated that the EMT process is critical not only for cancer cell development and malignancy, but it also causes chemotherapy resistance and decreases apoptotic cell death ^{103-105, 153}. Ashrafizadeh et al. described in their review the different pathways of

EMT-mediated cancer chemoresistance through Cisplatin therapy¹⁰². In contrast to the research cited, we did not find a variation in the expression of ECAD and NCAD after Cisplatin therapy, which could be attributed to our small sample size in the presented study.

6.4 Epithelial–mesenchymal transition and survival outcome

EMT is the complex cellular program through which epithelial cells turn into mesenchymal-like cells; this process facilitates the invasion and metastasis of epithelial cancer cells. EMT is distinguished by the decrease or loss of expression of adhesion molecules and the increase of expression of mesenchymal markers.

There are many things that could control EMT. Still, these were mostly studied in cell culture models and under standard conditions, which are very different from what happens when multiple organs or tissues of a patient interact with each other. Especially, this kind of setting makes it hard to figure out how much ECAD and NCAD changes affect how long patients with re-OSCC survive^{154, 155}.

EMT process is ruled by a range of factors. To this day, these factors were predominantly studied in cell culture models and under standard conditions, which are very different from what happens when different organs or tissues of a patient interact with each other¹⁵⁶. It is especially difficult to understand how much changes in ECAD and NCAD could influence how long patients with re-OSCC survive in this environment. In this study, we investigated the predictive value of ECAD and NCAD expression as independent indicator of survival.

Our findings showed that evaluating combined protein expression of both Cadherins and correlating them to the clinical data were less accurate than evaluating single protein expression, since EMT is very complex and different regulatory components and signaling pathways are involved in the process. In our tumor samples the loss of ECAD and *de novo* expression of NCAD were detected at the tumor-stromal interface,

as mentioned above. Even the expression of the proteins in the recurrence tumor when comparing it to the primary of the same tumor didn't change. Pyo et al. investigated the relationship between the decreased ECAD expression and positive NCAD expression in OSCC, which in turn implies that cadherin switch most probably contributes significantly to the invasiveness and metastasis of oral squamous cell carcinoma ¹⁵⁷.

Few clinical studies have been conducted on the relationship between ECAD and NCAD and survival in OSCC. These studies either studied the stage of primary disease as a variable ¹⁵⁸, or compared tumor tissue with pre-cancerous lesions of oral mucosa as a control group ¹⁵⁹.

According to recent research, this is the first study to look at how between ECAD and NCAD expression affects survival after recurrence. In the risk-adjusted hazard model, we found that the post-recurrence overall survival and the post-recurrence oral cancer-specific survival both dropped significantly when the *h-score* for ECAD loss was more than 60%. Therefore, in patients with re-OSCC, this IHC score serves as an independent risk factor for poor post-recurrence survival.

6.5 Outlook and perspectives for future studies

Notwithstanding these conclusions and the findings of the current study, drug-induced EMT inhibition may present a therapeutic possibility. Since EMT plays a crucial role in tumor aggressiveness and metastasis, various medications were developed to specifically target EMT-related signaling pathways. For example, Tangeretin and Rhamnetin were reported to induce radiosensitivity by interfering with Notch pathway ^{160, 161}. A combination therapy of radiotherapy and HIF-1-targeting agents such as Sorafenib and Paclitaxel was shown to improve radio-sensitization ^{162, 163}. Furthermore, Also, By suppressing the EGFR signaling pathway, the anti-EGFR monoclonal antibody nimotuzumab was shown to increase the responsiveness of esophageal cancer KYSE-150R cells to radiation ¹⁶⁴.

Metformin was also used to test this strategy, and the results showed that EMT in oral squamous cell carcinoma might be inhibited by the mTOR/HIF-1/PKM2/STAT3 pathway¹⁶⁵. It is now necessary to conduct prospective, randomized clinical trials to determine whether this therapy is beneficial for patients with recurrent OSCC. It is becoming obvious that while previous research has mainly concentrated on alterations in gene expression and abnormal genetic and epigenetic mutations in cancer cells, examining different versions in the stromal structure of the HNSCC TME and their influence on tumorigenesis and progression may aim to grasp better the processes underlying different reactions to treatment⁷².

6.6 Limitations of the present study

One of the study's shortcomings was its small sample size. Furthermore, the distribution of patients who received different treatments (surgery alone, surgery + radiotherapy, surgery + radiochemotherapy) was inhomogeneous.

Another point to note is that the EMT process extends far beyond the cadherin switch. Cell signaling, epigenetic modification, post-translational modifications, and transcriptional control all have an impact on the process¹⁶⁶. Some studies showed that reduced ECAD expression is a signature of EMT. Still, some researchers contended that ECAD loss of expression is unnecessary during the EMT process and that restoring ECAD expression in ECAD-negative malignant cells did not affect the EMT¹⁶⁷. It was discovered that the decreased expression of ECAD was insufficient to induce EMT in a non-malignant breast cell line¹⁶⁸. The loss of ECAD expression, on the other hand, has long been associated with more aggressive, poorly differentiated malignant cells; additionally, ECAD loss has been linked to the activation of numerous EMT transcription factors. As a result, whether the loss of ECAD is a key cause or a consequence of EMT remains an open debate¹⁶⁹.

7. Summary

Previous research has indicated that the loss of ECAD and the overexpression of NCAD in tumor cells lead to metastatic dissemination and the triggering of different EMT transcription factors.

In this study the survival outcomes were determined by analyzing the immunohistochemistry expression of ECAD and NCAD in primary and recurrent OSCC patients as a marker of EMT in relation to OSCC. We correlated the post-recurrence survival of this high-risk group with the immunohistochemical expression of both cadherin types based on standardized *h-scores* and the clinical outcome (oral cancer-specific survival, overall survival, and post-recurrence disease-free survival). The patients' characteristics were scanned and digitally assessed. The evaluation of immunostaining was based on the *h-score*.

The current study reveals that Cadherin-switch appears to be an innate histological marker whose expression does not change between the primary tumor and its recurrence, regardless of the therapy used for the primary tumor. Crucial evidence was found of a proportionate correlation between ECAD loss, post-recurrence oral cancer-specific survival, and post-recurrence survival in re-OSCC using the automated evaluation of the *h-score* for IHC staining. ECAD loss of greater than 60% significantly raised the hazard ratio for post-recurrence survival outcome. Post-recurrence survival outcomes didn't correlate with the *de novo* expression of NCAD. Therefore, the loss of ECAD could be a potential biomarker for stratifying therapy and de-/escalating multimodal treatment as an independent risk factor for poor survival in patients with re-OSCC. In patients with re-OSCC, targeting EMT, may thus represent a potential adjuvant treatment option.

8. Zusammenfassung

Die epithelial-mesenchymale Transition (EMT) ist ein biologischer Mechanismus bei zahlreichen physiologischen und pathologischen Erkrankungen. Die damit verbundenen Veränderungen der Cadherin-Expression spielen eine entscheidende Rolle bei der Entstehung von Karzinomen, der Invasion und Metastasierung, der Angiogenese und der Immunantwort. EMT-Zellen weisen einen Übergang von einem epithelialen zu einem mesenchymalen Phänotyp auf (Cadherin-Switch). Dieser Prozess ist durch die *de novo*-Bildung von N-Cadherin (NCAD) gekennzeichnet, das E-Cadherin (ECAD) ersetzt und für eine erhöhte Migrationsfähigkeit und maligne Transformation der entarteten Zellen steht. Der Cadherin-Switch ist ein Charakteristikum der EMT und wurde bei verschiedenen Krebsarten beobachtet. Die vorliegende Arbeit untersucht als Hypothese den Cadherin-Switch im Gewebe von primären und rezidivierenden oralen Plattenepithelkarzinomen (re-OSCC) als inhärentes Merkmal des Tumors, das biologische Verhalten reguliert und darüber hinaus die Überlebensrate dieser Patienten nach dem Rezidiv beeinflusst. In der Methode wurden die Überlebensrate nach dem Rezidiv berechnet und die standardisierte *h-score* basierte IHC-Expression beider Cadherin-Typen mit dem klinischen Verlauf korreliert. In der zugrunde liegenden Kohorte wurden 94 Patienten mit einem Rezidiv eines OSCC im Rahmen der klinischen Routine kontrolliert. Es wurden Gewebeproben sowohl von Primär- als auch von Rezidivtumoren gesammelt und immunhistochemisch untersucht. Es wurde ein signifikanter Zusammenhang zwischen dem Verlust der ECAD-Expression und dem OSCC-spezifischen Überleben sowie dem Gesamtüberleben festgestellt (HR=2,72, CI:1,50-4,95, p=0,001) bzw. (HR=3,84, CI:1,93-7,63, p=0,001) bei einem Expressionsverlust von mehr als 60%. Es bestand kein statistisch signifikanter Zusammenhang zwischen der N-CAD *de-novo*-Expression und dem OSCC-spezifischen Überleben, dem Gesamtüberleben, oder dem krankheitsfreien Überleben nach Rezidiv.

Die aktuelle Studie zeigt, dass der Cadherin-Switch ein histologischer Marker zu sein scheint, dessen Expression sich zwischen dem Primärtumor und seinem Rezidiv nicht verändert, unabhängig von der Therapie des Primärtumors. Anhand der automatisierten Auswertung des *h-scores* für die IHC-Färbung wurde ein entscheidender Hinweis auf eine Korrelation zwischen ECAD-Verlust, OSCC-spezifischem Überleben nach Rezidiv und Gesamtüberleben nach Rezidiv gefunden. Ein ECAD -Verlust von mehr als 60 % erhöhte das Hazard Ratio für ein Sterben nach einem Rezidiv signifikant. Die Überlebensrate nach einem Rezidiv korrelierte nicht mit der *de-novo*-Expression von NCAD. Daher könnte der Verlust von ECAD ein potenzieller Biomarker für die Stratifizierung der Therapie und die De-/Eskalierung der multimodalen Behandlung sein, da er ein unabhängiger Risikofaktor für das Überleben bei Patienten mit re-OSCC ist. Bei Patienten mit re-OSCC könnte, neben der etablierten adjuvanten therapieverfahren, die gezielte Beeinflussung der EMT daher eine mögliche zusätzliche Behandlungsoption darstellen.

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