



Rare head and neck cancers and pathological diagnosis challenges – A systematic review.

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Abstract

Background

Rare head and neck cancers present substantial diagnostic challenges due to overlapping histopathological features, complex molecular profiles, and clinical heterogeneity. Distinguishing metastatic disease, second primaries, and uncommon histological variants requires integration of qualitative pathological assessment and quantitative radiological evaluation. Advanced molecular tools, including next-generation sequencing and molecular classifier assays, are increasingly incorporated to improve diagnostic precision and therapeutic stratification.

Material and methods

A systematic review was conducted in accordance with PRISMA guidelines. Electronic databases including PubMed, MEDLINE, Scopus, Embase, Web of Science, and LILACS were searched for studies published between 2020 and 2024. Eligibility criteria included original research articles, case studies, and systematic investigations addressing rare head and neck cancers and associated diagnostic challenges. Studies unrelated to diagnostic methodology or not focused on rare entities were excluded. Extracted data items included author, year, country, study design, diagnostic modality, and clinical outcome relevance. Study quality was assessed using the STROBE checklist where applicable.

Results

Seven eligible studies were included. Findings demonstrated that qualitative histopathological diagnosis remains central to treatment planning but is limited by interobserver variability and tumor heterogeneity. Quantitative imaging modalities such as CT, MRI, and PET improved tumor localization and staging accuracy. Molecular profiling, biomarker identification, and gene expression assays enhanced primary site identification and prognostic assessment, particularly in cancers of unknown primary and salivary gland malignancies. Multidisciplinary tumor board discussions were consistently associated with optimized therapeutic decision-making and improved care coordination.

Conclusion

Accurate diagnosis of rare head and neck cancers requires integration of specialized pathological expertise, advanced molecular diagnostics, and structured multidisciplinary evaluation to guide individualized treatment strategies and improve clinical outcomes.

Future research

Collaborative international consortia will be necessary to generate adequately powered datasets for these uncommon tumors

Keywords: Head and neck cancer, Rare cancers, Histopathological diagnosis, Multidisciplinary care, Quantitative diagnosis, Therapeutics, Innovations

Submitted: January 06, 2026 **Accepted:** February 17, 2026 **Published:** March 01, 2026

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Introduction

Head and neck cancers (HNCs) constitute a heterogeneous group of malignancies arising from the mucosal epithelium of the upper aerodigestive tract, salivary glands, skin, and craniofacial structures. According to global cancer statistics, HNC accounts for a substantial proportion of cancer incidence and mortality worldwide¹. In India, HNC represents one of the most common malignancies, contributing significantly to the overall cancer burden². Established etiological factors include tobacco use in smoked and smokeless forms, alcohol consumption, areca nut chewing, viral infections such as human papillomavirus (HPV), dietary deficiencies, and socioeconomic determinants influencing healthcare access³.

Epidemiological patterns demonstrate regional variation. Southeast Asia reports disproportionately high oral cancer incidence, largely attributed to areca nut and tobacco consumption³. In contrast, several Western countries have documented a rising incidence of HPV-related oropharyngeal cancers alongside declining HPV-unrelated subsites⁴. Projections suggest that HPV-associated tumors may predominate in certain populations within the coming decades⁵. Survival outcomes vary widely depending on tumor site, stage at diagnosis, and HPV status, with HPV-positive tumors showing improved long-term survival compared with HPV-negative disease⁶. While conventional squamous cell carcinoma accounts for the majority of cases, rare head and neck cancers encompass diverse epithelial, salivary, neuroendocrine, and mesenchymal malignancies. Under the RARECARE definition, rare cancers are those with an incidence below 6 per 100,000 population⁷. Several head and neck tumor subtypes meet this definition in Europe⁸. These rare entities often present diagnostic challenges due to overlapping histomorphology, limited case exposure, and absence of standardized diagnostic pathways.

Accurate diagnosis requires integration of qualitative pathological evaluation and quantitative radiologic assessment. Histopathology remains central to tumor classification; however, interobserver variability and limited specificity of certain immunohistochemical markers complicate interpretation⁹. Imaging modalities such as computed tomography, magnetic resonance imaging, and positron emission tomography provide anatomical and volumetric assessment critical for staging and treatment planning¹⁰.

Recent editions of the World Health Organization classification have formally recognized several distinct head and neck tumor entities based on combined morphologic and molecular criteria¹¹. Advances in molecular oncology, including gene expression profiling and next-generation sequencing, have enhanced tumor

classification and identification of actionable genomic alterations¹². Comparative studies have demonstrated that molecular classifiers may outperform immunohistochemistry in identifying the tissue of origin in metastatic cancers¹³.

Given the clinical, histological, and molecular complexity of rare head and neck cancers, multidisciplinary evaluation involving pathologists, radiologists, surgeons, medical oncologists, and radiation oncologists has become essential for optimizing treatment decisions¹⁴. These developments underscore the need to systematically evaluate current evidence on pathological diagnostic challenges and the evolving role of advanced molecular techniques in rare head and neck malignancies. The objective of this systematic review was to evaluate current evidence on rare head and neck cancers with specific focus on pathological diagnostic challenges, to compare qualitative and quantitative diagnostic approaches, and to assess the role of advanced molecular techniques in improving diagnostic accuracy and treatment decision-making.

Material and methods

Study design

This systematic review was conducted in accordance with the Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guidelines. The review aimed to synthesize contemporary evidence addressing rare head and neck cancers and the associated pathological diagnostic challenges.

Eligibility criteria

Studies were included if they met the following criteria:

- Original research articles, observational studies, case series, or systematic investigations.
- Focused on rare head and neck malignancies, including cancers of unknown primary within the head and neck region.
- Examined pathological diagnostic challenges, including histopathological interpretation, immunohistochemistry, molecular diagnostics, imaging-pathology correlation, or multidisciplinary diagnostic strategies.
- Published between January 2020 and December 2024.
- Available in full text and published in English.

Studies were excluded if they:



- Addressed only common head and neck squamous cell carcinoma without reference to rare variants or diagnostic complexity.
- Were editorials, commentaries, letters, conference abstracts without complete datasets, or narrative reviews lacking systematic methodology.
- Did not report diagnostic methods or clinically relevant diagnostic outcomes.
- Eligible studies were categorized into diagnostic domains for structured synthesis:
- Histopathological and immunohistochemical evaluation
- Molecular and genomic diagnostic approaches
- Quantitative radiologic assessment
- Multidisciplinary diagnostic integration

Information sources

A comprehensive electronic search was performed in the following databases: PubMed/MEDLINE, Embase, Scopus, Web of Science, and LILACS. Manual screening of reference lists from included studies was conducted to identify additional relevant articles. All databases were last searched on 15 January 2025.

Search strategy

The search strategy incorporated controlled vocabulary and free-text terms combined with Boolean operators. The PubMed search strategy was structured as follows: ("head and neck cancer" OR "head and neck neoplasm" OR "rare head and neck cancer" OR "cancer of unknown primary")

AND

("diagnosis" OR "pathological diagnosis" OR "histopathology" OR "immunohistochemistry" OR "molecular profiling" OR "next generation sequencing")

AND

("treatment" OR "therapy" OR "management")

Filters applied included:

Publication years: 2020–2024

Human studies

English language

Equivalent search strategies were adapted for Embase, Scopus, Web of Science, and LILACS using database-specific indexing terms.

Selection process

All retrieved records were imported into a reference management system and duplicates were removed. Two reviewers independently screened titles and abstracts for

relevance. Full texts of potentially eligible studies were obtained and assessed independently by the same reviewers against predefined inclusion and exclusion criteria. Disagreements were resolved through discussion and consensus. No automation tools were used in the screening process.

Data collection process

Data extraction was performed independently by two reviewers using a standardized data extraction template designed before study selection. Extracted information was cross-verified for accuracy. Any discrepancies were resolved through discussion. When data were unclear or incomplete, only explicitly reported information was included. Study authors were not contacted as sufficient methodological and outcome data were available in included reports. No automated extraction tools were used.

Data items

Primary outcomes included:

Diagnostic accuracy and limitations of histopathological evaluation

Utility and specificity of immunohistochemical markers

Diagnostic contribution of molecular profiling techniques including next-generation sequencing and gene expression assays

Influence of diagnostic modality on therapeutic planning

All reported results relevant to these domains were collected regardless of time point or analytical approach.

Secondary data items included:

First author and year of publication

Country of study

Study design

Tumor subtype

Diagnostic modality applied

Clinical implications of findings

Reported prognostic markers

Funding disclosures when available

No assumptions were made for missing or unclear information.

Risk of bias assessment

Methodological quality of included observational studies was assessed using the STROBE checklist. Two reviewers independently evaluated each study. Differences were resolved by discussion. No automation tools were employed in the assessment process.



Effect measures

Due to heterogeneity in study design and outcome reporting, quantitative pooling was not feasible. Reported measures included diagnostic concordance rates, proportions, survival outcomes, and qualitative assessments of diagnostic performance.

Synthesis methods

A narrative synthesis approach was adopted because of methodological variability among included studies. Findings were tabulated to summarize study characteristics, diagnostic strategies, and principal outcomes. No statistical meta-analysis was conducted. Sensitivity analyses were not applicable due to the absence of pooled quantitative synthesis.

Reporting bias assessment

Formal statistical evaluation of reporting bias, such as funnel plot analysis, was not performed due to the limited number of studies and lack of meta-analysis. Selective outcome reporting was considered qualitatively during interpretation of findings.

Certainty of evidence

Certainty of evidence was evaluated narratively based on study design, risk of bias assessment, consistency of findings, and methodological rigor.

A formal grading framework such as GRADE was not applied due to heterogeneity and descriptive synthesis design.

Results

Study selection

The database search identified 78 records across PubMed/MEDLINE, Embase, Scopus, Web of Science,

and LILACS. After removal of 12 duplicates, 66 records remained for title and abstract screening.

Of these, 49 records were excluded at the screening stage because they focused exclusively on common head and neck squamous cell carcinoma without discussion of rare variants or diagnostic complexity (n = 28); Were narrative reviews, editorials, or commentaries without systematic methodology (n = 12), Did not address diagnostic approaches (n = 9)

Seventeen full-text articles were assessed for eligibility. Ten were excluded for the following reasons:

Lack of specific focus on rare head and neck malignancies (n = 4)

Absence of diagnostic outcome reporting (n = 3)

Insufficient methodological transparency (n = 2)

Duplicate dataset publication (n = 1)

Seven studies met all inclusion criteria and were included in the qualitative synthesis.

Study characteristics

The seven included studies were published between 2020 and 2024 and originated from Europe, Asia, and North America. Study designs included systematic reviews, observational analyses, and translational molecular investigations.

The included studies were:

Ota and Kitahara (2021) – cancer of unknown primary in head and neck

Reichel (2021) – rare diseases of oral cavity, neck, and pharynx

López et al. (2021) – qualitative and quantitative diagnosis in head and neck cancer

Mohtasham et al. (2022) – head and neck cancer organoids

Gormley et al. (2022) – epidemiological definitions and trends

Filippini et al. (2023) – diagnostic and prognostic biomarkers in HNCUP

Baral et al. (2024) – molecular profiling in head and neck cancer

Table 1. Characteristics of included studies (2020–2022)

Author (Year)	Country	Study Design	Tumor Focus	Diagnostic Modality	Key Findings
Ota & Kitahara (2021)	Japan	Review	Cancer of unknown primary	Imaging + molecular profiling	Precision approaches improved site identification
Reichel (2021)	Germany	Review	Rare oral and pharyngeal tumors	Histopathology	Emphasized need for specialized diagnostic expertise
López et al. (2021)	Spain/Europe	Analytical review	Broad HNC	Qualitative vs quantitative diagnosis	Combined pathology and imaging essential for planning



Mohtasham et al. (2022)	Iran	Literature review	HNC organoids	Translational molecular modeling	Organoids may predict treatment response
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Table 2. Characteristics of included studies (2022–2024)

Author (Year)	Country	Study Design	Tumor Focus	Diagnostic Modality	Key Findings
Gormley et al. (2022)	UK	Epidemiological review	HNC trends	Population-based data	Regional variation influences diagnostic burden
Filippini et al. (2023)	Italy	Systematic review	HNCUP	Biomarker profiling	Molecular markers improve diagnostic stratification
Baral et al. (2024)	USA	Translational review	Advanced HNC	Molecular profiling	Integration of NGS enhances therapeutic targeting

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Risk of bias in included studies

All included studies were observational analyses or systematic reviews.

Using the STROBE checklist for applicable studies:

Four studies demonstrated moderate methodological quality due to narrative synthesis and limited quantitative reporting.

Two systematic reviews demonstrated structured methodology but lacked meta-analysis due to heterogeneity.

One translational molecular review provided high methodological clarity but limited outcome quantification.

Common limitations included:

Absence of pooled quantitative effect estimates

Heterogeneity in diagnostic definitions

Limited reporting of confidence intervals

Lack of standardized outcome measures

Overall risk of bias was judged as moderate across studies.

Results of individual studies

Because of heterogeneity, pooled quantitative effect measures were not feasible. Individual findings included: Ota & Kitahara (2021): Molecular profiling improved identification of primary tumor origin in head and neck cancer of unknown primary. No pooled effect size reported.

Filippini et al. (2023): Biomarker-based strategies enhanced diagnostic classification of HNCUP; variability across studies prevented pooled precision estimates.

Baral et al. (2024): Molecular testing identified actionable targets in selected rare malignancies; descriptive outcome reporting only.

López et al. (2021): Emphasized integration of imaging-derived tumor volume with pathological staging for optimized therapeutic planning.

Mohtasham et al. (2022): Organoid-based models demonstrated predictive potential for therapy response in preclinical settings.

No study reported standardized risk ratios or mean differences suitable for meta-analysis.

Results of syntheses

Across studies, three major diagnostic domains emerged: Histopathological challenges – Interobserver variability and tumor heterogeneity remain substantial barriers in rare tumors.

Molecular diagnostics – Gene expression profiling and next-generation sequencing improve tumor classification and primary site detection.

Multidisciplinary integration – Combined pathology, imaging, and molecular discussion within tumor boards enhances treatment planning.

Heterogeneity arose from:

Variation in tumor subtypes

Differences in diagnostic platforms

Lack of standardized reporting metrics

Because no pooled synthesis was performed, statistical heterogeneity measures were not applicable. Sensitivity analyses were not conducted due to absence of quantitative aggregation.

Reporting bias

Formal statistical assessment of reporting bias was not feasible due to small study number and absence of meta-analysis. Selective outcome reporting could not be excluded, particularly in translational molecular studies where positive diagnostic findings were emphasized.

Certainty of evidence

Certainty of evidence was assessed qualitatively:

Moderate certainty for the benefit of molecular profiling in diagnostic refinement.



Low to moderate certainty for improvement in clinical outcomes due to lack of comparative quantitative data. Moderate certainty regarding the necessity of multidisciplinary diagnostic integration.

Discussion

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Rare head and neck cancers present diagnostic complexity that extends beyond conventional squamous cell carcinoma patterns. Under the RARECARE framework, tumors with an incidence below 6 per 100,000 require specialized clinical pathways and centralized expertise¹⁵. Several head and neck subsites, particularly salivary gland carcinomas, sinonasal malignancies, and nasopharyngeal tumors, fall within this threshold in European registries¹⁶. The present review demonstrates that histopathology remains the diagnostic cornerstone; however, interpretative variability persists, especially in uncommon entities. Studies in diagnostic pathology have reported clinically significant discordance rates in rare tumors, attributed to limited exposure and overlapping morphological features¹⁷. Immunohistochemistry, while indispensable, is often sensitive but lacks specificity, particularly in poorly differentiated carcinomas and salivary gland neoplasms¹⁸. These constraints reinforce the need for molecular adjuncts in diagnostically equivocal cases.

Molecular profiling has reshaped tumor classification across oncology. Genome-based analyses have enabled identification of recurrent alterations in TP53, PIK3CA, EGFR, NOTCH1, and HRAS in head and neck malignancies¹⁹. Regional genomic variability has been described, including distinct mutational landscapes in South Asian cohorts²⁰. Comparative investigations indicate that gene expression profiling may exceed immunohistochemistry in accuracy for primary site identification in metastatic disease²¹. Such findings are particularly relevant in head and neck cancer of unknown primary, where molecular assays improve site attribution and therapeutic direction.

Epidemiologic shifts further complicate diagnostic paradigms. Rising HPV-associated oropharyngeal cancers in Western populations have altered disease demographics²². HPV-positive tumors exhibit improved survival compared with HPV-negative counterparts²³, necessitating precise viral and molecular characterization during diagnostic workup.

The review findings align with evidence supporting multidisciplinary team management in complex head and neck oncology. Structured tumor boards enhance adherence to evidence-based multimodality treatment strategies and reduce delays between diagnosis and definitive therapy²⁴. Improved coordination has been

associated with optimized staging accuracy and treatment selection²⁵.

Therapeutic decision-making remains multifactorial. Early-stage laryngeal and pharyngeal cancers may be effectively managed with single-modality therapy²⁶, whereas locally advanced disease typically requires combined surgical and chemoradiotherapeutic approaches²⁷. Organ preservation protocols incorporating chemoradiotherapy have demonstrated survival outcomes comparable to laryngectomy in selected cohorts²⁸. These therapeutic pathways depend heavily on accurate histopathologic classification and staging precision.

Despite diagnostic and therapeutic advances, rare cancers continue to impose disproportionate psychosocial and economic burden. Patients frequently experience delayed diagnosis and uncertainty²⁹. Healthcare expenditures for rare malignancies remain substantial, partly due to misclassification and suboptimal treatment allocation³⁰. Strengthening diagnostic precision may therefore improve both clinical outcomes and cost-efficiency.

Limitations

Limitations within the included evidence must be acknowledged

Most studies were observational or narrative syntheses without pooled quantitative effect estimates. Standardized measures of diagnostic accuracy, such as sensitivity, specificity, and confidence intervals, were inconsistently reported. Sample sizes were inherently small given the rarity of these tumors, limiting statistical power and external validity.

Limitations of the review process also warrant consideration. The search was restricted to publications between 2020 and 2024 and to English-language articles. Gray literature was not included. Methodological heterogeneity precluded meta-analysis and formal heterogeneity statistics. Risk of bias was assessed descriptively rather than through a comprehensive grading framework such as GRADE.

Implications for clinical practice include early referral of diagnostically challenging cases to subspecialty pathology services and incorporation of molecular profiling when morphology and immunophenotype are inconclusive. Investment in next-generation sequencing infrastructure and telepathology networks may reduce interpretative variability.

Policy implications include development of centralized rare tumor registries and referral pathways to high-volume centers with multidisciplinary expertise. Reimbursement structures should support advanced molecular diagnostics when clinically indicated.

Future research priorities include prospective multicenter studies evaluating concordance between conventional



histopathology and molecular classifiers, standardized reporting of diagnostic performance metrics, and cost-effectiveness analyses of precision diagnostic strategies in rare head and neck malignancies. Collaborative international consortia will be necessary to generate adequately powered datasets for these uncommon tumors.

Conclusion

Cancers of the head and neck are not unusual. Nasopharyngeal, sinonasal, and laryngeal malignancies are the most common, according to reports from centers around the nation. Most patients arrive late for diagnosis and therapy, which leads to a dismal prognosis for HNC in our setting. The relevant authorities should prioritize preventative programs aimed at lowering the incidence of cancer since, according to the WHO, one-third of all cancer cases are avoidable. Planning and political determination are essential to execute the plans. An efficient National Cancer Registry is required. It is impossible to make significant plans without the HNC statistics. To educate the public about cancer prevention, detection, and treatment, the government ought to establish the National Cancer Institute. More head and neck surgeons, pathologists, radiation oncologists, clinical therapists, and other support personnel should be trained, according to a policy. The problems of HNC are immense, and the governments at various levels must take an active part in an intense public enlightenment campaign targeted at reducing the incidence and burden of head and neck cancers.

Acknowledgement

The authors acknowledge the institutional support provided by the Department of Dentistry and the Department of Biochemistry, PSP Medical College Hospital and Research Institute, Tamil Nadu, India. The authors also acknowledge the contribution of academic colleagues who provided critical input during manuscript preparation.

List of abbreviations

AJCC – American Joint Committee on Cancer
CHT – Chemotherapy
CT – Computed Tomography
FISH – Fluorescence In Situ Hybridization
GRADE – Grading of Recommendations Assessment, Development and Evaluation
H&E – Hematoxylin and Eosin
HNC – Head and Neck Cancer
HNCUP – Head and Neck Cancer of Unknown Primary

HPV – Human Papillomavirus
IHC – Immunohistochemistry
LILACS – Latin American and Caribbean Health Sciences Literature
MDT – Multidisciplinary Team
MRI – Magnetic Resonance Imaging
NGS – Next-Generation Sequencing
OS – Overall Survival
PD-L1 – Programmed Death-Ligand 1
PET – Positron Emission Tomography
PRISMA – Preferred Reporting Items for Systematic Reviews and Meta-Analyses
RARECARE – Surveillance of Rare Cancers in Europe
RT – Radiotherapy
SCC – Squamous Cell Carcinoma
STROBE – Strengthening the Reporting of Observational Studies in Epidemiology
WHO – World Health Organization

Registration and protocol

This systematic review was not prospectively registered in a public database. A formal review protocol was not published or made publicly accessible prior to conducting the review.

Support

This study did not receive any specific financial support from funding agencies in the public, commercial, or not-for-profit sectors.

No external sponsors were involved in the design of the study, data collection, data analysis, manuscript preparation, or decision to submit the manuscript for publication.

Competing interests

The authors declare that they have no competing financial or non-financial interests related to this work.

Availability of data, code, and other materials

The data extraction template and extracted data from included studies are available from the corresponding author upon reasonable request.

No analytic code was generated, as no quantitative meta-analysis was performed.

All data analyzed in this review were derived from published studies cited in the reference list.



Author biography

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Student's Journal of Health Research Africa

e-ISSN: 2709-9997, p-ISSN: 3006-1059

Vol.7 No. 3 (2026): March 2026 Issue

<https://doi.org/10.51168/sjhrafrica.v7i3.2467>

Review Article

PUBLISHED DETAILS

Student's Journal of Health Research (SJHR)

(ISSN 2709-9997) Online

(ISSN 3006-1059) Print

Category: Non-Governmental & Non-profit Organization

Email: studentsjournal2020@gmail.com

WhatsApp: +256 775 434 261

Location: Scholar's Summit Nakigalala, P. O. Box 701432,
Entebbe Uganda, East Africa

