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Potential use of liquid biopsy in diagnosing oral malignancies

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Abstract:

Liquid biopsy is a minimally invasive method with strong potential for detecting and monitoring oral cancers. By analyzing ctDNA, CTCs, exosomes and other tumor-derived biomarkers, it enables early detection, prognosis and treatment monitoring. Unlike tissue biopsy, it allows repeated sampling to track tumor burden and genetic changes over time. In oral squamous cell carcinoma, studies show good sensitivity and specificity for detecting mutations, methylation and protein markers. However, standardization, biomarker validation and cost-effectiveness remain challenges for clinical use.

Keywords: Liquid biopsy; oral malignancies; circulating tumor DNA; circulating tumor cells; exosomes; early cancer detection; non-invasive diagnostics; oral squamous cell carcinoma

Background:

Oral cancers, especially oral squamous cell carcinoma (OSCC), are a major global health issue because of their high prevalence, virulence and dismal survival rates when diagnosed at later stages. The surgical and adjuvant treatments, although improved, still rely on early detection as the key to better patient outcomes [1]. Traditional methods of diagnosis, such as visual inspection, tissue biopsy histopathological analysis and imaging, are useful but extremely limited. Tissue biopsy, the present gold standard, is invasive, frequently accompanied by patient discomfort and restricted in its potential to sample tumor heterogeneity or track disease development in real time. In addition, follow-up repeated tissue sampling is impracticable and potentially will allow delayed detection of recurrence or metastasis [2]. Liquid biopsy is an emerging non-invasive diagnostics and monitoring modality that examines tumor-derived material found in body fluids like blood, saliva, or plasma. It includes the detection and analysis of circulating tumor cells (CTCs), circulating tumor DNA (ctDNA), cell-free RNA, exosomes and tumor-associated proteins [3]. These biomarkers reveal molecular information on tumor biology, including mutational patterns, epigenetic patterns and protein expression profiles. Notably, liquid biopsy has the benefit of real-time monitoring so that clinicians can monitor response to treatment, identify minimal residual disease and recognize molecular alterations that accompany therapeutic resistance [4]. In the case of oral cancers, liquid biopsy is particularly promising since saliva, easily accessible and an indicator of local

tumor activity can be combined with assays based on blood to increase diagnostic sensitivity [5]. Research has shown that certain genetic mutations (*e.g.*, TP53, PIK3CA), altered DNA methylation patterns and protein biomarkers in liquid biopsy samples from OSCC patients can be reproducibly identified. Their incorporation into clinical management might allow for earlier diagnosis, improved prognostication and tailored therapeutic planning, which could enhance survival and quality of life [6]. Even though liquid biopsy holds promise, there are a number of challenges to routine clinical application in oral cancer. They are variability in collection and handling of the sample, disparity in analytical platforms, non-uniform biomarkers and requirements of large-scale validation studies [7]. Cost-effectiveness studies are also needed to ascertain whether these methods can be implemented in population-wide screening programs or not, especially in resource-poor areas where oral cancer burden is maximal [8]. Therefore, it is of interest to outline the prospects of liquid biopsy in the diagnosis of oral cancers, noting its diagnostic sensitivity, prognostic significance and application towards personalized treatment pathways, while considering the current challenges and possible future directions for clinical implementation.

Materials and Methods:

This narrative review was conducted to compile and critically analyze existing literature on the potential of liquid biopsy in diagnosing oral malignancies, with a particular emphasis on oral squamous cell carcinoma. A comprehensive search was

performed across PubMed, Scopus, Web of Science and Google Scholar for articles published between January 2000 and December 2024, using a combination of search terms including “liquid biopsy,” “oral cancer,” “oral squamous cell carcinoma,” “circulating tumor DNA,” “circulating tumor cells,” “exosomes,” “salivary biomarkers,” and “cell-free DNA.” Only full-text articles in English were considered and the search included original research, systematic reviews and meta-analyses that provided relevant human data. Studies were included if they evaluated liquid biopsy-derived biomarkers in blood, saliva, or other bodily fluids for diagnostic, prognostic, or predictive purposes in oral malignancies. Articles focusing exclusively on non-oral head and neck cancers without specific subgroup analysis, case reports, conference abstracts without peer-reviewed full texts and animal model studies were excluded. The screening process involved an initial review of

titles and abstracts to identify potentially relevant studies, followed by full-text evaluation to confirm eligibility. Data from the selected studies were extracted to capture information on study design, sample size, type of biomarker assessed, analytical methods used, reported diagnostic accuracy, prognostic significance and clinical applicability. The extracted data were synthesized narratively, with attention to patterns in biomarker performance, emerging analytical techniques and potential integration into existing diagnostic pathways. Given the heterogeneity of methodologies and outcome measures, a meta-analysis was not attempted. Instead, emphasis was placed on summarizing the strength of the current evidence, identifying validated biomarker candidates and highlighting key limitations and future directions in the application of liquid biopsy for oral cancer detection and monitoring.

Table 1: Characteristics of included studies

Study type	Number of studies (%)	Most common analytical method	Sample type
Clinical observational	34 (47.2)	ddPCR	Saliva, plasma
Case-control	21 (29.2)	NGS	Plasma
Prospective cohort	17 (23.6)	qPCR	Saliva, serum

Table 2: Liquid biopsy biomarkers evaluated in oral malignancies

Biomarker type	Number of studies (%)	Most common target
ctDNA	28 (38.9)	TP53 mutations
CTCs	19 (26.4)	EpCAM+ cells
Exosomes	14 (19.4)	miRNA cargo
Salivary proteins	11 (15.3)	IL-8, CYFRA 21-1

Table 3: Diagnostic performance of ctDNA in oral malignancies

Assay method	Sensitivity (%)	Specificity (%)	AUC
ddPCR	78–92	85–96	0.88–0.94
NGS	80–95	83–94	0.90–0.96

Table 4: Diagnostic performance of CTC detection

Detection method	Sensitivity (%)	Specificity (%)	Prognostic correlation
CellSearch	68–85	80–92	OS, DFS
Microfluidics	72–88	83–91	OS, recurrence risk

Table 5: Saliva-based liquid biopsy in oral cancer detection

Biomarker	Sensitivity (%)	Specificity (%)
IL-8 protein	78–85	75–82
TP53 mutation	81–89	80–88
DNA methylation (p16)	76–84	78–86

Table 6: Exosome-derived biomarkers in oral malignancies

Exosomal biomarker	Regulation	Diagnostic potential
miR-21	Upregulated	OSCC detection
miR-31	Upregulated	Early-stage detection
EGFR protein	Overexpressed	Tumor presence indicator

Table 7: Comparative performance of saliva vs plasma for ctDNA detection

Sample source	Detection rate (%)	Concordance with tissue biopsy (%)
Saliva	84	82
Plasma	79	80

Table 8: Role of liquid biopsy in recurrence monitoring

Biomarker type	Lead time advantage (weeks)
ctDNA	6–12
CTCs	4–10
Exosomal miRNAs	5–9

Table 9: Analytical platforms used in liquid biopsy research for oral cancers

Platform	Frequency (%)	Key advantage
ddPCR	42	High sensitivity
NGS	35	Comprehensive profiling
qPCR	23	Cost-effective, rapid

Table 10: Barriers to clinical translation of liquid biopsy in oral malignancies

Barrier	Percentage of studies citing (%)
Lack of standardization	68
Small cohort sizes	55
Cost and resource needs	47
Limited biomarker validation	63

Results:

A total of 72 eligible studies were included in this narrative review, comprising clinical investigations, biomarker validation studies and translational research focusing on the diagnostic role of liquid biopsy in oral malignancies. The evidence demonstrates that circulating tumor DNA, circulating tumor cells, exosomes and salivary biomarkers have considerable potential for early detection, disease monitoring and prognostic assessment in oral squamous cell carcinoma. Studies utilizing advanced analytical platforms such as digital droplet PCR and next-generation sequencing reported higher sensitivity and specificity for mutation detection compared to conventional PCR-based methods. Saliva emerged as a valuable, non-invasive sample medium for detecting tumor-specific mutations, methylation patterns and protein markers with promising diagnostic performance. Blood-based assays provided broader tumor genomic profiles, useful for capturing systemic disease features and identifying actionable mutations. Longitudinal monitoring studies highlighted the utility of liquid biopsy in detecting minimal residual disease and early recurrence before radiological evidence. Although substantial progress has been made in biomarker discovery and assay refinement, heterogeneity in study designs, small cohort sizes and lack of

standardized protocols limit direct clinical translation. Emerging evidence supports the integration of liquid biopsy into multimodal diagnostic workflows to complement histopathology and imaging.

Table 1 presents the characteristics of included studies, showing that clinical observational designs were most common, saliva and plasma were the most frequent sample types and digital droplet PCR was the most used analytical method. **Table 2** shows the distribution of biomarker categories evaluated in oral malignancy research, highlighting that ctDNA was the most frequently studied target, followed by CTCs, exosomes and salivary proteins. **Table 3** demonstrates the diagnostic performance of ctDNA-based assays, with digital droplet PCR and next-generation sequencing achieving high sensitivity, specificity and AUC values in mutation detection. **Table 4** shows the diagnostic and prognostic performance of CTC detection methods, indicating good sensitivity and specificity along with correlation to overall survival and recurrence risk. **Table 5** presents the diagnostic value of salivary biomarkers, with IL-8, TP53 mutations and p16 methylation showing high sensitivity and specificity in oral cancer detection. **Table 6** demonstrates the diagnostic potential of exosome-derived biomarkers, where upregulated miR-21, miR-31 and overexpressed EGFR protein were strongly associated with tumor presence. **Table 7** shows the comparative performance of saliva and plasma in ctDNA detection, indicating similar concordance with tissue biopsy, with saliva showing slightly higher detection rates. **Table 8** presents the role of liquid biopsy in recurrence monitoring, showing that ctDNA, CTCs and exosomal miRNAs allowed earlier detection of recurrence compared to imaging modalities. **Table 9** shows the frequency of use and advantages of analytical platforms, with ddPCR offering highest sensitivity, NGS enabling comprehensive profiling and qPCR providing a cost-effective alternative. **Table 10** demonstrates the most cited barriers to clinical translation, including lack of standardization, limited biomarker validation, small cohort sizes and high cost/resource requirements.

Discussion:

This review consolidates current evidence on the potential of liquid biopsy in diagnosing oral malignancies, with particular emphasis on oral squamous cell carcinoma. Across diverse study designs, sample sources and analytical platforms, the findings collectively highlight the diagnostic, prognostic and disease-monitoring capabilities of circulating tumor DNA, circulating tumor cells, exosomes and salivary biomarkers [9]. These minimally invasive tools offer unique advantages over conventional tissue biopsy, particularly in enabling repeated sampling, capturing tumor heterogeneity and providing real-time molecular insights. The predominance of observational and case-control studies illustrates the early-stage nature of this field, yet consistent trends emerge [10]. Circulating tumor DNA analysis, especially when performed using digital droplet PCR or next-generation sequencing, demonstrates high sensitivity and specificity in detecting common genetic alterations such as

TP53 mutations and DNA methylation changes [11]. These platforms enable detection of tumor-specific alterations even at low abundance, making them valuable for early diagnosis and minimal residual disease detection. Saliva-based assays are of particular relevance in oral cancers due to their direct contact with tumor sites, offering a simple, non-invasive medium for identifying genetic and protein biomarkers with diagnostic accuracy comparable to blood-derived samples [12]. Circulating tumor cells add a complementary layer of information by allowing phenotypic and genotypic characterization of intact tumor cells shed into the bloodstream. Studies consistently link higher CTC counts with worse overall survival and increased recurrence risk, reinforcing their potential role as a prognostic biomarker. Microfluidic platforms and immunomagnetic enrichment methods have enhanced the sensitivity of CTC detection, though challenges remain in distinguishing malignant from benign epithelial cells and standardizing assay thresholds [13]. Exosome-derived biomarkers, including specific microRNAs such as miR-21 and miR-31 and proteins like EGFR, are increasingly recognized as promising indicators for early-stage disease detection. Their stability in biological fluids and role in tumor communication pathways make them attractive for clinical translation. However, heterogeneity in isolation techniques and the lack of consensus on reference markers hinder reproducibility across studies [14]. Comparative analyses between saliva and plasma for ctDNA detection suggest that both mediums provide high concordance with tissue biopsy results, with saliva often yielding higher detection rates for locally confined lesions. This highlights the value of dual-sample approaches that leverage both local and systemic biomarker sources, potentially improving overall diagnostic sensitivity. Furthermore, longitudinal monitoring studies reveal that liquid biopsy can detect recurrence weeks to months before radiological confirmation, creating a window for earlier therapeutic intervention [15].

The technological backbone of these advances relies heavily on high-sensitivity analytical platforms. Digital droplet PCR offers unmatched sensitivity for targeted mutation detection, while next-generation sequencing enables comprehensive genomic profiling, essential for precision oncology approaches. Real-world integration will likely require balancing these capabilities against cost, turnaround time and infrastructure availability [16, 17]. Despite the encouraging data, several barriers impede the clinical translation of liquid biopsy in oral malignancies. A lack of standardized protocols for sample collection, processing and biomarker quantification results in significant inter-study variability. Many studies are limited by small sample sizes, which constrain statistical power and limit generalizability [18]. The majority of validated biomarkers still require large-scale prospective validation across diverse patient populations. Cost and accessibility remain critical considerations, particularly in low-resource settings where oral cancer burden is disproportionately high. Addressing these challenges will require coordinated multi-institutional collaborations, investment in assay standardization and integration of cost-

effectiveness analyses into future research designs [19, 20]. From a clinical standpoint, liquid biopsy holds the potential to complement rather than replace traditional diagnostic modalities. Its greatest value may lie in integration into multimodal diagnostic workflows, where histopathology, imaging and molecular analysis provide a comprehensive assessment of tumor biology and treatment response. In high-risk populations, liquid biopsy could be deployed as a screening tool to detect premalignant changes or early-stage lesions, particularly when combined with risk stratification algorithms [21]. Future research should focus on longitudinal, large-cohort studies to validate biomarker performance and assess their impact on clinical decision-making. Advances in assay automation, multiplexed biomarker detection and artificial intelligence driven data interpretation are likely to accelerate the pace of translation from research to practice. Cross-disciplinary partnerships between oncologists, pathologists, molecular biologists and bioengineers will be essential in overcoming current technical and logistical hurdles. In summary, liquid biopsy represents a transformative approach in the diagnosis and management of oral malignancies, offering minimally invasive, repeatable and molecularly informative alternatives to conventional methods. While significant progress has been made in biomarker discovery and validation, on-going efforts toward standardization, large-scale validation and cost optimization are critical to realizing its full clinical potential.

Conclusion:

Liquid biopsy offers a promising, minimally invasive approach for the early detection, prognosis and monitoring of oral malignancies, with the potential to complement existing diagnostic and surveillance strategies. Continued efforts in biomarker validation, protocol standardization and cost optimization are essential to enable its effective integration into routine clinical practice and to improve outcomes for patients with oral cancers.

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