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Review

Tumor-Infiltrating Lymphocytes in Head and Neck Cancer: Ready for Prime Time?

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Simple Summary: The immune response has been shown to be a promising indicator to predict the clinical behavior of many cancers, including head and neck cancer. Tumor-infiltrating lymphocytes (TILs) were widely introduced as an important tool to reveal the status of the immune response. This review discusses the significance of TILs in head and neck cancers.

Abstract: The evaluation of tumor-infiltrating lymphocytes (TILs) has received global attention as a promising prognostic cancer biomarker that can aid in clinical decision making. Proof of their significance was first shown in breast cancer, where TILs are now recommended in the classification of breast tumors. Emerging evidence indicates that the significance of TILs extends to other cancer types, including head and neck cancer. In the era of immunotherapy as a treatment choice for head and neck cancer, assessment of TILs and immune checkpoints is of high clinical relevance. The availability of the standardized method from the International Immuno-oncology Biomarker Working Group (IIBWG) is an important cornerstone toward standardized assessment. The aim of the current article is to summarize the accumulated evidence and to establish a clear premise for future research toward the implementation of TILs in the personalized management of head and neck squamous cell carcinoma patients.

Keywords: head and neck squamous cell cancer (HNSCC); survival; tumor-infiltrating lymphocytes (TILs)



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1. Introduction

Head and neck squamous cell carcinoma (HNSCC) may manifest in all subsites of the upper aerodigestive tract, including the oral cavity, larynx, oropharynx, and hypopharynx. It is one of the most common cancers worldwide, with 878,348 new cases and 444,347 mortalities in 2020 (<https://gco.iarc.fr/today> (accessed on 21 October 2021)). Unfortunately, the incidence of HNSCC is rising [1]. The main risk factors include tobacco and alcohol consumption (for oral and pharyngo-laryngeal cancers), as well as human papillomavirus (HPV) for oropharyngeal cancer. Regarding the management of HNSCC, new therapeutic perspectives have been demonstrated with the introduction of a minimally invasive surgical

approach, using robotic surgery and deintensifying the subsequent chemoradiotherapy [2]. HNSCC is characterized by a high rate of metastatic dissemination, which is associated with worse survival compared with HNSCC cases with no metastasis [3]. While treatment planning relies on traditional criteria, including the TNM staging system (eighth edition [4]) and the status of high-risk HPV, ongoing efforts continue to improve risk stratification. Current tools for risk assessment are cancer-related parameters, while factors related to the tumor microenvironment (TME) are not yet employed in daily practice. It is important to state that the prognostic markers that are widely considered in HNSCC treatment decision making (e.g., tumor size, p16, lymphovascular invasion) are mainly based on evidence from retrospective studies [5,6].

Not surprisingly, the immune response is a major determinant influencing the survival of cancer patients. Moreover, immunotherapy is currently implemented in the standard treatment regimens of recurrent and/or metastatic head and neck cancer patients. However, the evaluation of immunological determinants of a cancer patient is not yet considered in the treatment planning of HNSCC, and there is a need to identify reliable and simple immunological biomarkers to further optimize treatment strategies [7]. Recent research in immuno-oncology and cancer biomarkers has underlined the significance of tumor-infiltrating lymphocytes (TILs) as a promising prognostic indicator in various tumor types, including HNSCC. This paper discusses recent findings regarding TILs in HNSCC and how they can be used to improve individualized treatment.

2. Accumulated Evidence on TILs as a Prognosticator: Where Are We Currently?

Evidence on the prognostic significance of TILs in patients with HNSCC has been rapidly accumulating in recent years. The selection of the molecule, e.g., CD3 (Figure 1), to be analyzed or relying on an overall assessment of TILs (Figure 2) without specific immunostaining is still a point of discussion, and there is no definitive conclusion. On the other hand, the recently published guidelines of the International Immuno-oncology Biomarker Working Group (IIBWG) [8,9] have been utilized in several studies, forming an essential step towards a standardized assessment method and implementation of TILs in routine pathology reporting. Of note, the introduced guidelines have been successfully used in head and neck cancer studies that assessed TILs in hematoxylin and eosin (HE)-stained slides [10–13].

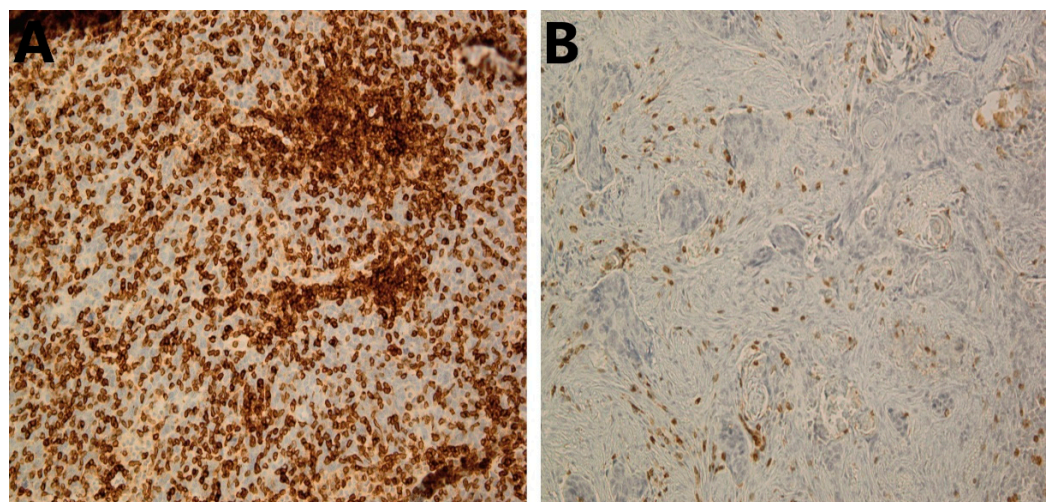


Figure 1. Immunohistochemical staining of CD3⁺ TILs in HNSCC. (A) High stromal CD3⁺ TILs in HNSCC tumor; 20×. (B) Low stromal CD3⁺ TILs in HNSCC tumor; 20×. HNSCC, head and neck squamous cell carcinoma; TILs, tumor-infiltrating lymphocytes.

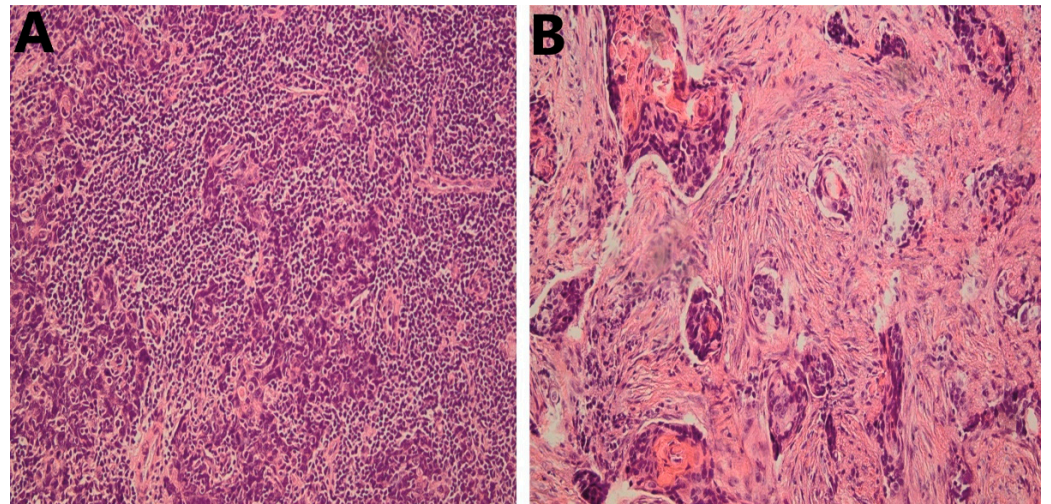


Figure 2. Representative cases of HNSCC stained with HE; 20 \times . (A) High infiltration of stromal TILs; (B) Low stromal TILs; 20 \times . HNSCC, head and neck squamous cell carcinoma; HE, hematoxylin and eosin; TILs, tumor-infiltrating lymphocytes.

In addition to many studies that have reported the significance of the overall score of TILs using HE-stained slides in head and neck cancers [10–18], the prognostic significance of cell-specific immune markers such as CD3 [19], CD4 [19,20], CD8 [20–24], CD56 [22], CD57 [25], CD163 [26], and FoxP3 [20] has been reported in these cancers. Of note, intensive research on the significance of TILs in HNSCC has been performed, which has led to the accumulation of evidence in systematic reviews and meta-analyses (Table 1). As an example, de Ruiter et al. [27], in their recent meta-analysis (2017), reported a favorable prognostic value of CD3, CD8, and FoxP3 infiltration in HNSCC based on a pooled analyses of good-quality studies with the majority of them having a low risk of bias. Furthermore, in a meta-analysis published in 2020 [28], Bisheshar et al. found a favorable prognostic value for CD56 and CD57. In another recent meta-analysis (2021) on another two TIL subsets, Borsetto et al. [29] found that high CD4 and high CD8 were associated with a reduced risk of death of HNSCC. Of note, when Borsetto et al. [29] conducted meta-analyses for subsites of HNSCC, they found that CD8 was associated with survival of oropharyngeal and hypopharyngeal cancers, but no significant association was found with oral or laryngeal SCC. Focusing on PD-L1, however, Yang et al. (2018) found no significant value for its expression in HNSCC [30]. It is important to explain that the quality of the included studies in these meta-analyses was assessed using the Quality in Prognosis Studies (QUIPS) tool in two of these meta-analyses [28,30] or by the Newcastle–Ottawa Scale in another meta-analysis [29]. Another note is that due to the overlapping time periods in the database search, some studies have been included in each of these meta-analyses [27,29].

Further, meta-analyses for specific subsites of HNSCC have also been conducted in other articles. In laryngeal SCC, for example, a recent meta-analysis (2021) found that TILs in the TME are a reliable prognostic marker [31]. This meta-analysis included studies considering the overall assessment of TILs in HE-stained sections and using subsets of TILs, including CD8 and/or CD3/CD4 [31]. However, the small number of studies (<10 studies) that were included in each meta-analysis, and the small number of cases (<100 patients) in many of the included studies, have been highlighted as a shortcoming [31].

Table 1. Summary of meta-analyses on TILs in HNSCC.

| Author (Year) | TILs Subset | No. Studies (No. Cases) | Endpoint | SCC Location | Pooled HR (95% CI), <i>p</i> |
|------------------------------|-------------|-------------------------|----------|--------------------|--|
| de Ruiter et al. [27] (2017) | CD3+ | 4 studies (904 cases) | OS | HNSCC | HR 0.64 (0.47–0.85) |
| | | 4 studies (735 cases) | DFS | HNSCC | HR 0.63 (0.49–0.82) |
| | CD8+ | 9 studies (1697 cases) | OS | HNSCC | HR 0.67 (0.58–0.79) |
| | | 7 studies (1053 cases) | DFS | HNSCC | HR 0.50 (0.37–0.68) |
| | CD4+ | 4 studies (596 cases) | LRC | HNSCC | HR 0.82 (0.70–0.96) |
| | | 4 studies (775 cases) | OS | HNSCC | HR 0.76 (0.64–0.89) |
| | FoxP3+ | 3 studies (513 cases) | LRC | HNSCC | HR 0.81 (0.68–0.96) |
| | | 6 studies (977 cases) | OS | HNSCC | HR 0.80 (0.70–0.92) |
| Yang et al. [30] (2018) | PD-L1 | 21 studies (2477 cases) | OS | HNSCC | HR 0.98 (0.71–1.37), <i>p</i> = 0.93 |
| | | 7 studies (1001 cases) | DFS | HNSCC | HR 1.07 (0.68–1.70), <i>p</i> = 0.76 |
| | | 4 studies (592 cases) | DSS | HNSCC | HR 0.90 (0.63–1.29), <i>p</i> = 0.56 |
| | | 6 studies (585 cases) | PFS | HNSCC | HR 0.71 (0.55–0.93), <i>p</i> = 0.01 |
| Bisheshar et al. [28] (2020) | CD56/CD57 | 4 studies (420 cases) | OS | HNSCC | HR 0.19 (0.11–0.35) |
| Borsetto et al. [29] (2021) | CD4+ | 3 studies (548 cases) | OS | HNSCC | HR 0.77 (0.65–0.93) |
| | CD8+ | 6 studies (1220 cases) | OS | HNSCC | HR 0.64 (0.47–0.88) |
| | CD4+ | 3 studies (498 cases) | OS | OPSCC | HR 0.52 (0.31–0.89) |
| | CD8+ | 3 studies (326 cases) | OS | OPSCC (HPV-Neg) | HR 0.39 (0.16–0.93) |
| | CD8+ | 6 studies (661 cases) | OS | OPSCC (HPV-Pos) | HR 0.40 (0.21–0.76) |
| | CD8+ | 3 studies (250 cases) | OS | Hypopharyngeal SCC | HR 0.43 (0.30–0.63) |
| Rodrigo et al. [31] (2021) | HE staining | 4 studies (719 cases) | OS | Laryngeal SCC | HR 0.57 (0.36–0.91), <i>p</i> = 0.02 |
| | HE staining | 4 studies (659 cases) | DFS | Laryngeal SCC | HR 0.56 (0.34–0.94), <i>p</i> = 0.03 |
| | CD8+ | 5 studies (536 caess) | OS | Laryngeal SCC | HR 0.62 (0.40–0.97), <i>p</i> = 0.04 |
| | CD8+ | 4 studies (574 caess) | DFS | Laryngeal SCC | HR 0.73 (0.60–0.90), <i>p</i> = 0.002 |
| | CD3+/CD4+ | 4 studies (369 caess) | OS | Laryngeal SCC | HR 0.38 (0.16–0.9), <i>p</i> = 0.03 |
| | CD3+/CD4+ | 2 studies (224 caess) | DFS | Laryngeal SCC | HR 0.23 (0.10–0.53), <i>p</i> = 0.0005 |

Abbreviations in Table 1: CI: Confidence interval; DFS: Disease-free survival; DSS: Disease-specific survival; HNSCC: Head and neck squamous cell carcinoma; HR: Hazard ratio; LRC: Locoregional control; OS: Overall survival; PFS: Progression-free survival; TILs: Tumor-infiltrating lymphocytes.

3. Clinical Scenarios of TILs in HNSCC

The immune infiltrate, due to its clinical significance, can form a useful additional prognostic parameter. In HNSCC, the assessment of TILs can aid in guiding the patient's management in two potential clinical scenarios: the first is to contribute to an improved classification of HNSCC based on the TNM–Immune staging system [32]. The currently used tumor-node-metastasis (TNM) classification has been criticized, as many cases will show variable clinical outcomes within the same stage. The addition of the p16-status, a surrogate marker for HPV-induced oropharyngeal cancer, improved risk stratification in HNSCC, but the TNM system requires further refinement. The incorporation of TILs, as an immune parameter in the TNM–Immune system could augment the prognostic performance of the classification and aid in decision making and treatment planning [33].

The second clinical scenario is to serve as an immune classifier to assess the potential need (and subsequently to predict the response) to an immunotherapy regimen [34]. This has already been introduced via the assessment of PD-L1-expressing tumor cells and immune cells in patients with recurrent/metastatic HNSCC [35]. To this extent, several clinical trials are investigating the role of immune checkpoint inhibitors (ICI) for curative approaches, such as the clinical use of neoadjuvant preoperative immunotherapy or administering ICI during concomitant radio(chemo)therapy or as a maintenance/adjuvant therapy [36,37]. To date, however, selecting HNSCC patients who might benefit from

immunotherapy remains challenging. The currently reported prognostic factors do not evaluate the full immune status of the patients, as they may be biased by pre-analytical (tissue quality) and spatio-temporal heterogeneity (specimen type and sampling time-point). Subsequently, the expected response to immune-based therapies remains unconsidered in pathology reports of HNSCC. The assessment of TILs on HE-stained slides may provide a useful parameter in addition to traditional prognostic parameters that do not assess the immune response. However, the evaluation of TILs on HE slides should be further investigated in a prospective fashion using standardized methodology, as proposed by the IIBWG, to fully comprehend their function in tailoring treatment with ICI. It will be of high clinical significance to consider the assessment of TILs following the IIBWG criteria in HE-stained sections from ongoing clinical trials of HNSCC [38]. Such a dataset is suitable to be used in the validation of TILs as a prognostic marker [39].

3.1. TILs as Indicator for Selection of HNSCC Patients for Immunotherapeutic Approaches

Numerous isolated methods for histological quantification of TILs subsets have been described, each having its own unique scoring technique, pharmacodiagnostic monoclonal antibody, and gradation or cutoff. Despite these differences in methodology, the literature concurs that TILs have an important prognostic value. Their predictive role, however, needs to be further elucidated, and few reports have made contributions regarding this topic. Essentially, tumors can be subdivided in an immune-inflamed or non-inflamed phenotype, as described in the hallmarks of cancer [40]. A preserved immunity is characterized by an adequate amount, diversity, and functioning of immune cells recruited from both the innate and adaptive immune systems, which is required to benefit from treatment with immune checkpoint inhibitors. During the process of tumoral progression, a proportion of transformed cells will not survive, allowing antigen-presenting cells to detect and pick up tumor-related antigens from dead neoplastic cells. Through human leukocyte antigen molecules, these antigens are presented to (CD4⁺ or CD8⁺) T-cells and activate the well-known cascade of tumor-cell recognition, activation, and expansion of effector cells that will induce a tumor-specific immune response. Indeed, inflamed tumor phenotypes may benefit from an improved immune-mediated elimination of tumor cells [41,42]. In anti-CTLA-4-treated melanoma patients, TILs density was significantly increased from baseline in therapy-responders, confirming their predictive significance to ICI [43]. When applied to HNSCC, Mandal et al. [44] reported that an increased density of immune-infiltrating cells, specifically CD56⁺ NK cells, was correlated with a better overall response rate (ORR) in patients treated with a variety of ICI. Furthermore, Hanna et al. [45] reported that HNSCC-patients with a high CD8⁺ lymphocyte rate and PD-1 expression were correlated with improved response rates with anti-PD-1/PD-L1 agents. At present, immunohistochemistry (IHC) for PD-L1 expression is the sole predictive biomarker to determine eligibility for treatment with ICI, yet it lacks robustness. Bearing the aforementioned theory in mind, the evaluation of TILs may provide a useful, additional parameter to assess tumor immune response, which requires further examination in prospective studies using standardized methodology.

3.2. The Role of TILs in the Elucidation of De-Escalation Therapies in HNSCC

De-escalation strategies are currently being considered in several malignancies. The goals of de-escalation are reducing therapy-related toxicity, increasing or maintaining survival outcomes, and improving patients' quality of life [46]. Indeed, the multimodal therapeutic approaches applied in HNSCC, including surgery, radiotherapy, and/or chemotherapy, are correlated with both acute and long-term toxicity. Selecting patients for de-escalation may depend on several prognostic biomarkers, such as histopathological characteristics including grade, lymphovascular, or perineural invasion; molecular markers (p16/HPV status); or clinical risk stratification (TNM, performance status) [47]. The emergence of HPV as an etiological factor has served as an important prognosticator in oropharyngeal carcinoma for many years. However, applying de-escalation strategies based on

this biomarker has not shown expected benefits with regard to survival [48,49]. The question, therefore, arises if morphological characteristics of the tumor microenvironment, i.e., the immune infiltrate, should be applied during risk stratification as an alternative for HPV status. A recent study by the group of Sylvie Rottey [13] applied the IIBWG method and reported that the increased infiltration of mononuclear cells in oropharyngeal squamous cell carcinoma (OPSCC) correlated with superior survival in comparison to OPSCC with a low TIL density. This outcome was independent of p16-status. Moreover, prognostic stage (AJCC) and stromal TILs density were considered as the two major independent prognostic factors for overall survival, indicating that OPSCC might indeed benefit from a TNM-Immune classification. The AJCC TNM, the eighth edition, deserves credit for introducing a separate classification for p16⁺ OPSCC, though it has also been subjected to criticism for failing to incorporate pivotal scientific evidence regarding the tumor's immune microenvironment. As the era of immunomodulatory agents is rapidly progressing, immune-related features such as TILs should be added to further improve the clinical and pathological staging, thus aiding physicians in clinical decision making [50]. This concept has been proposed in a similar fashion in colorectal cancer via the introduction of the Immunoscore [51]. According to this methodology, immune cell density is assessed per patient using a digital-pathology-based assay based on the quantification of CD3⁺ and CD8⁺ lymphocytes at the invasive border and tumor core. Patients with high Immunoscores in both areas were associated with better outcomes [52,53]. Although several studies have confirmed the prognostic value of this method in HNSCC, only post hoc analyses and subgroup identification will fully identify its clinical significance [12,13].

4. Automated Analysis of TILs

Digital pathology has been used in recent studies to assess markers for HNSCC. Among these, TILs identified by immunohistochemical staining of specific molecules (e.g., CD3, CD4, and CD8) were assessed using an automated method in HNSCC [21,54–57]. Similarly, the automated signature of CD8xPD-L 1 has been reported as a predictive marker in non-small-cell lung cancer patients [58]. Further research is still warranted to reach the proper application and use of digital analysis tools in our daily practices [59]. The concordance between manual and computational scoring of TILs scores was not excellent in a recent report [60]. However, the digital assessment of TILs in HE-stained slides has been successfully reported in HNSCC [61,62] and showed a superior prognostic performance in a recent large study of HNSCC [63]. These findings have been supported by studies on breast cancer [64] and colorectal cancer [65], where automated analysis showed success in the assessment of TILs.

The findings based on digital analysis can be best applied by considering a semiautomated method, where a pathologist first selects the area to be analyzed to ensure the evaluation of the representative field. The computer application/software will then identify and provide counting/estimation of TILs in the selected area, as was demonstrated in a recent study of oral SCC [55]. This will ensure correct, representative, and reproducible assessment of TILs in the stromal area adjacent to the tumor nests and thus exclude areas outside the tumor border as well as those with necrosis. Further, an open-source algorithm for the automated evaluation of TILs has been recently introduced for melanoma [66]. A similar digital evaluation approach should be considered for TILs in HNSCC to allow the comparison of different analyses.

5. Morphological Pitfalls in the Assessment of TILs in HNSCC

During the assessment of TILs, there are two major morphological pitfalls to be distinguished:

- (1). Differentiating TILs from a pre-existing immune infiltrate and other immune cells. First, this can be attributed due to the presence of pre-existing lymphoid tissue (Figure 3A), not only in the oropharyngeal/tonsillar region but in all (sub)sites of the head and neck region. Therefore, distinguishing tumor-attracted from pre-existing

mononuclear cells may be challenging. Second, regions with ulcerations (Figure 3B) or erosions are seeded with infiltrating immune cells that are predominantly polymorphonuclear cells, which can also hamper the correct assessment of TILs in the TME. Evidently, these should be excluded from evaluation.

- (2). Inadequate tumor material for evaluation. In HNSCC, available tissue specimens typically comprise diagnostic biopsies. However, these are often characterized by insufficient amounts of stroma, disabling the correct quantification of stromal lymphocytes in this compartment. Ideally, whole-tumor resected specimens are the ‘golden standard’ to perform quantification of TILs but are mostly unavailable for patients diagnosed with recurrent and unresectable or metastasized HNSCC. In addition, long time intervals may exist between the initial biopsy or resection and the decision to commence palliative treatment. The temporal variance that has occurred in the TME will ultimately have an effect on the use of TILs as a predictive biomarker in this setting.

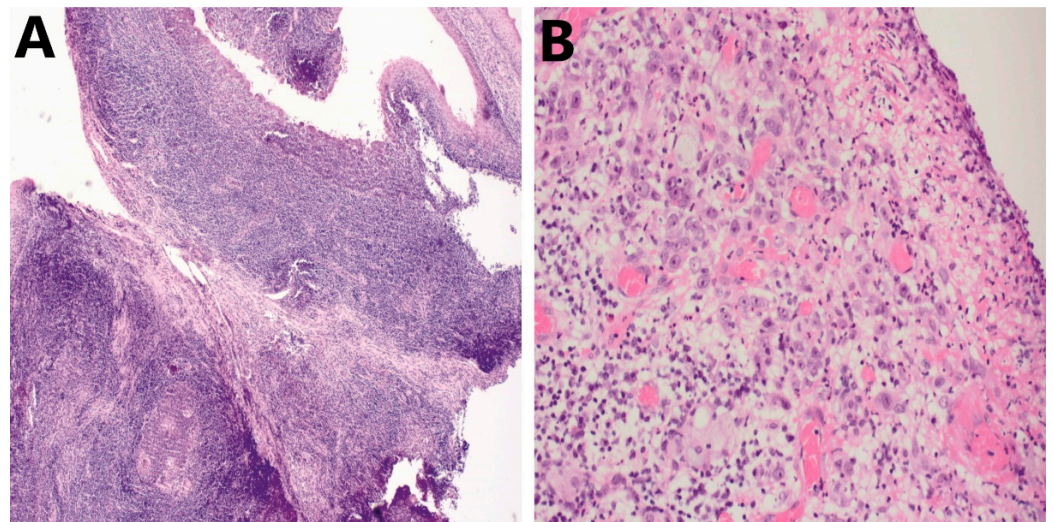


Figure 3. Examples of pitfalls during assessment of TILs in HNSCC on HE. (A) Presence of pre-existing lymphoid tissue in HNSCC; 10× (B) Presence of ulcerations seeded with polymorphonuclear cells (granulocytes); 20×. HNSCC, head and neck squamous cell carcinoma; TILs, tumor-infiltrating lymphocytes.

6. Why Is the Assessment of TILs Not Yet Used in the Daily Practice of HNSCC Pathology Reporting?

This question can be properly addressed by considering the limitations and shortcomings of the published studies. First, there are some heterogeneities, such as the variability in the chosen immunohistochemical molecules/markers to assess TILs (e.g., CD3, CD8, FOXP3) in addition to the overall assessment of TILs in HE stained slides. Of note, a recent study in breast cancer demonstrated a better inter-pathologist concordance for the overall assessment of TILs than for the quantification of cell-specific molecules [67]. This issue needs to be studied in HNSCC to find out whether the overall assessment of TILs is more advantageous compared with the identification of the best immune molecule that can be evaluated to aid in clinical decision making as an immune classifier.

Second, there are differences in the cutoff points used to define low, intermediate, and high levels of TILs. The IIBWG recommended scoring the percentage of TILs as a continuous parameter because there is no pre-determined cutoff point for each cancer type [8]. In HNSCC, this issue will need to be studied, ideally for each subsite separately. The histological location of assessment has an obvious impact: stromal TILs have been assessed in most of the published studies and were associated with survival. However, a few studies found intra-tumoral TILs to have significant clinical relevance, yet this quantification method is regarded as challenging [10,22]. The stratification of immune cells in the central area of the tumor and at

the invasive margin is therefore advised, as indicated in colon cancer [68]. Scoring TILs at the invasive front has the most relevance with a superior prognostic value compared to TILs scored in the central tumor area of oral cancer [12,22].

Finally, the prognostic significance of TILs in preoperative biopsies of HNSCC has not yet been widely studied. The potential of using TILs in preoperative prognostication to predict response to neoadjuvant therapy can be a valuable step towards personalized treatment. In some cancers, biopsies have been successfully used to assess TILs as a predictor of treatment response [69–71] and to predict lymph node metastasis [72]. Similar studies in preoperative HNSCC biopsies are warranted. Of note, a high concordance between biopsies and tumor resection samples with regard to the density of TILs assessed in HE-stained slides has been recently reported in oropharyngeal cancer [57,73]. Unfortunately, the above-mentioned morphological pitfalls that might present during the assessment of TILs (including small biopsy specimens and/or the presence of pre-existing lymphoid tissue) still form a serious obstacle that makes the evaluation of TILs sometimes challenging.

7. Future Considerations for the Inclusion of TILs in Daily Clinico-Pathological Practice

In the future, more large-scale prospective studies are needed. Firstly, registry trials using standardized protocols are recommended to optimally describe the TME at the start of systemic therapy, either (neo-)adjuvant or palliative. Based on these registry trials, validation studies in specific settings are important in order to confirm such findings, establish reference values, and ultimately implement TILs in the routine pathology of this cancer type. Subsequently, external quality control of the methodology should be arranged in all participating centers, as performed for several other biomarkers. An inter-laboratory comparison or ‘Ring’ study could therefore be invaluable in this setting: both sensitivity and specificity of the biomarker are validated in different laboratories, mostly coordinated by a main laboratory. Repeatability, inter-laboratory, and inter-observer variability of the technique are tackled in such a trial. Additionally, the dissemination of the quantification technique should be of high priority by teaching the assessors via academic courses and workshops in (inter)national congresses and symposia. Lastly, evaluating TILs as a di- or trichotomized categorical variable via universally set cutoffs would further facilitate the clinical implementation of the quantification, as these will be more easily accepted and interpreted in daily clinical practice compared to the continuous use of TILs. An overview of future steps to be tackled prior to clinical implementation is given in Table 2.

Table 2. Summary of future steps needed prior to clinical implementation.

| To do | How? | Current Status | Future Steps |
|--|--|---|---|
| Construct standardized methods | International working groups | IIBWG method Immunoscore | Further improvement upon external feedback |
| Registry trials | Large scale observational trials per subsite | Retrospective trials only | Nation-wide/international analysis using standardized protocols |
| Prospective validation (predictive/prognostic value) | Clinical trials (introduce TNM–Immune) | Retrospective trials only | Interventional clinical trials(de-escalation strategies) |
| Determining clinically valuable cutoff | Further dissemination and implementation of TIL quantification | Continuous variable only for more accurate statistical analysis | Introduce di-/trichotomized cutoffs (non vs. moderately vs. highly inflamed tumors) |
| Education | Symposia Congresses Workshops | Word-of-mouth marketing | Further implementation of the method |
| Quality control | Ring trials | Non-existing in HNSCC | Further implementation of the method |

IIBWG, International Immuno-oncology Biomarker Working Group; HNSCC, head and neck squamous cell carcinoma; TIL, tumor-infiltrating lymphocyte.

8. Conclusions

The accumulated evidence from many studies indicates that TILs are easily estimated in routine HE-stained slides of different subsites of HNSCC and therefore can pave the way towards implementation into daily practice. A recent validation study (2022) has reported that evaluation criteria from IIBWG can be easily used to score TILs in HE sections of oropharyngeal cancer and identify tumors with a high risk of poor survival [74]. Indeed, more validation studies in other subsites of the head and neck need to be considered. Meanwhile, however, the overall assessment of TILs using HE can be already reported in the daily practice of pathologists following the IIBWG method to inform clinicians about the status of the adaptive immune response and to be included in a prognostic algorithm of multiple markers (including TILs) to reach a personalized treatment strategy.

Published findings on specific molecules (e.g., CD3, CD8) are of high clinical relevance and need to be confirmed in large homogenous comparative analyses (i.e., a similar protocol of staining including concentration, cutoff points, risk categorization, etc.). Homogenous cohorts with regard to subsites and stages of HNSCC are also important to compare results. The lack of such homogenous and large cohorts for the validation of these immune molecules makes them emerging biomarkers that are not yet ready for use. Finally, the semiautomated assessment of TILs can be a step toward the precise assessment and reduction in inter-observer variation, if any.

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