Immunohistochemistry and Lung Cancer: Application in Diagnosis, Prognosis and Targeted Therapy

Aldoph B. Nanguzgambo a Rubina Razack b Mercia Louw b Chris T. Bolliger a

a Division of Pulmonology, Department of Medicine, and b Division of Anatomical Pathology, Department of Pathology, Tygerberg Academic Hospital, University of Stellenbosch, Cape Town, South Africa

Background

Lung cancer remains the commonest global cause of death from cancer [1]. Notable advances in the diagnosis of cancers have been the use of immunohistochemistry in pathological diagnosis, the characterisation of the poorly differentiated malignant tumour, or in the identification of the origin of the tumour, whether primary or metastatic, and its emerging role in lung cancer prognosis and therapy. Furthermore, lung cancer management has evolved to a multidisciplinary level involving both pathologists and clinicians, some of whom may not understand the ever increasing contribution of immunohistochemistry to diagnosis, improving targeted therapy and determining prognosis. In this review, we do not discuss the basic principles of immunostaining protocols but focus on the (a) cellular structure of the respiratory airway epithelium and lung cancer biology of the major histological subtypes; (b) definition of immunohistochemistry (IHC) and immunocytochemistry (ICC); (c) common immunomarkers used specifically for primary lung carcinomas, and (d) application of immunohistochemistry in lung cancer diagnosis, determination of primary origin, and histological subtyping for targeted therapy and prognosis.

Key Words
Lung cancer · Immunohistochemistry · Immunomarker · Targeted therapy

Abstract

Immunohistochemistry is now an established ancillary technique in lung cancer diagnosis. Not only does it help in supporting the morphological diagnosis of malignancy, but its role now extends to the determination of cell lineage, ascertaining the primary site of tumour origin and contributing to decisions on prognosis and treatment. Early detection and confirmation of lung cancer facilitate early treatment decisions. Lung cancer management now has a multidisciplinary approach which includes cytopathologists and clinicians. Some clinicians may not understand what immunohistochemistry is and what its role is in lung cancer diagnosis, prognosis and therapy. The purpose of this paper is to define immunohistochemistry, on the background of basic respiratory airway epithelial structure and cancer biology, and discuss its application in the diagnosis, treatment and determination of prognosis of lung cancer.

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Respiratory Airway Epithelial Cells and Lung Cancer Biology

The respiratory airway is composed of the conducting airways and the acinar airways. The conducting airways comprise the trachea, bronchi, bronchioles and terminal bronchioles. The acinar airway, where gas exchange takes place, is made up of respiratory bronchioles, alveolar ducts and alveolar sacs [2]. The epithelial lining of the respiratory airways is continuous but with variable composition in its cellular structure.

The bronchial epithelium is composed of the following cell types: (a) ciliated columnar cells; (b) basal cells which are cuboidal cells and in contact with the basal membrane; (c) secretory cells, namely goblet cells and serous cells, and (d) myoepithelial cells. Secretory cells produce mucus which is composed of mucins [3].

The epithelium of bronchioles is mainly made up of (a) ciliated cells as in the bronchi, and (b) Clara cells. Besides being secretory cells, Clara cells have been shown to have multiple functions, including xenobiotic metabolism via the cytochrome P450 monooxygenase system and proges-
tenicity for ciliated cells in bronchioles when there is ep-i

thelial injury [4–6].

The acinar airway epithelium has two cell types: (a) alveolar type I cells, and (b) cuboidal secretory cells, also called alveolar type II cells. Alveolar type II cells synthesise and secrete surfactant apoproteins SP-A, SP-B and SP-C, differentiated by the weight of the protein content of surfactant [7, 8].

The respiratory airway epithelium also has neuroendocrine cells. They are part of the amine precursor uptake and decarboxylation (APUD) system and also secrete the enzyme neuron-specific enolase, which is a marker for the diffuse endocrine system. Like other APUD cells, they produce various polypeptide hormones including serotonin, gastrin-releasing peptide, bombe-
sin, calcitonin and chromogranins [9–11].

The natural expectation is that external stimuli such as cigarette smoke initiate neoplastic changes at a cellular and nuclear level in the epithelial cell lines, leading to pre-malignant lesions which then evolve to lung cancer. Loss of heterozygosity, gene amplification (numeric gain in chromosomal content), chromosomal aneusomy, genetic alterations such as p53 and K-ras, epidermal growth factor receptor (EGFR), and vascular endothelial growth factor receptor mutations with overexpression of the genes have all been implicated in lung cancer evolution [12–17].

Squamous cell carcinoma is believed to be a result of proliferation of basal cells in the respiratory epithelium, which then undergo metaplastic change to squamous cells. This metaplastic cellular morphology is termed bronchial dysplasia. Based on morphological features, bronchial epithelial dysplasia has been categorised into four types, namely mild dysplasia, moderate dysplasia, severe dysplasia and carcinoma in situ [15, 18–20]. It is important to distinguish squamous cell carcinoma from squamous metaplasia and pulmonary infarct, which may present with similar cytomorphological features [21].

The precursor lesion for adenocarcinoma is thought to be atypical adenomatous hyperplasia (AAH), which is also found peripherally [22–25]. By definition, AAH is \( \leq 0.5 \) cm and comprises proliferation of atypical type II pneu-

mocytes with gaps between the cells. It is also clear that AAH represents the premalignant stage of nonmucinous bronchioloalveolar carcinoma (BAC), another periph-

erally located lesion. It must be noted that in the new pro-

posed classification of lung adenocarcinoma, the term BAC will be withdrawn and replaced by adenocarcinoma in situ (AIS) and minimally invasive adenocarcinoma (MIA), due to a 100% disease-free survival after complete resection of these preinvasive lesions [26]. In terms of size, both AIS and MIA are defined as being \( \leq 3 \) cm with lep-

idic growth pattern, but MIA has an invasive component of \( \leq 0.5 \) cm in the greatest dimension of each focus. Both lesions are usually nonmucinous. In addition to AIS and MIA, other new terms that will be used to describe BAC depending on the morphology include lepidic predominant adenocarcinoma (nonmucinous), adenocarcinoma predominantly invasive with some nonmucinous lepidic components, and invasive mucinous adenocarcinoma (formerly mucinous BAC) [26]. Using IHC, it has been demonstrated that some adenocarcinomas actually arise centrally, and that these centrally derived tumours do not derive from AAH. On the contrary, peripherally derived adenocarcinomas have similar immunohistochemical and EGFR mutational patterns to AAH [27]. These periph-

erally derived adenocarcinomas have a cellular morphol-

gy similar to that of terminal respiratory epithelial cells (type II pneumocytes, Clara cells and nonciliated bronch-

chial cells). Adenocarcinoma must be distinguished from goblet cell hyperplasia, reactive or atypical bronchial epithelial cells, granular cell tumour and hamartoma [21].

Diffuse idiopathic neuroendocrine hyperplasia (DIPNEH) is the precursor lesion of some neuroendo-

crine cancers, namely carcinoid tumourlets (small neu-

roendocrine lesions <0.5 cm extending beyond the base-

ment membrane) as well as typical and atypical carcinoid tumours. DIPNEH is a result of proliferation of the neu-

roendocrine cells already described above. Despite small
cell lung carcinoma (SCLC) and large cell neuroendocrine carcinoma (LCNEC) having neuroendocrine properties, their cell line origins are not from DIPNEH and are currently not established. There are studies that showed different immunohistochemical reactions [28, 29] and different gene expression profiles [30] between DIPNEH-derived tumours and the more aggressive neuroendocrine tumours (SCLC and LCNEC), suggesting differences in origin.

**What Is Immunohisto/Cytochemistry?**

IHC and ICC involve the study of histological (tissue) or cytological (smear) specimens, respectively, using an antibody-antigen immune reaction, the antibody being externally added to the antigen already present on the tissue or smear specimen being analysed. Immunohistochemical markers can be used to stain cell surface markers, cell membranes, cytoskeletal components, vesicles and other organelles as well as nuclear contents. This immunostaining reaction may be unique to that tissue and its site of origin. Tissue-specific monoclonal antibodies or immunomarkers are now available, and these have helped improve the identification of neoplasms.

Although morphology is still the mainstay of diagnosis in cytopathology with 85–90% of cases being diagnosed with routine stains such as Papanicolaou and Diff Quik, the remaining 10–15% require ancillary techniques, of which immunohistochemistry solves 50% of the cases [31]. Immunohistochemistry is indicated in four settings, namely (a) to help establish malignancy; (b) to determine cell lineage; (c) to ascertain the primary site of tumour origin, and (d) for prognostic and therapeutic assessment. Most of the principles which govern the application of IHC to surgical specimens can be applied to cytopathology, with a few adaptations and precautions.

Immunohistochemistry can be done on whole tissue specimens, fine needle aspiration smears, cytopsins of fluids, or cell blocks. Cell blocks are the more optimal specimen on which ICC should be performed [32]. Cell blocks meet the same conditions as tissue sections and accommodate in-built controls on the same slide. It has also been shown that ICC can successfully be applied to Papanicolaou-stained smears [33, 34]. Furthermore, cell blocks have been successfully prepared directly from scraping of already stained cytology smears, with very good preservation of the morphology and architectural structure. ICC results on these cytoscrape cell block preparations have been shown to be excellent, to be comparable to conventional cell block results and to help in achieving the final diagnosis [35].

The major advantage of performing ICC over IHC pertains to specimen acquisition. Fine needle aspiration is easy, safe and can deliver a diagnosis within minutes. However, cytological specimens have their own drawbacks (table 1). It is essential that results be interpreted in close correlation with cytomorphological, clinical and radiological findings. ICC must be seen as a complementary to reaching a definite diagnosis, never the qualifier. If any discrepancies are encountered in the above regard, a formal biopsy must be acquired and IHC must be performed. However, even IHC is ancillary and does not replace morphology. A larger repeat biopsy may need to be obtained if there is any doubt about the diagnosis.

**What Are the Useful Available Immunomarkers for Identifying the Origin of Lung Carcinomas?**

Table 2 shows the commonly used immunomarkers in lung cancer diagnosis.

**Thyroid Transcription Factor-1**

Thyroid transcription factor-1 (TTF-1) is a 38-kDa homeodomain-containing nuclear transcription protein belonging to the Nkx2 gene family. TTF-1 is involved in the transcription of thyroglobulin and thyroperoxidase genes in the follicular thyroid cells, and of the surfactant A, B, C protein genes and Clara cell secretory protein in lung epithelial cells [36, 37]. In a study by Sturm et al. [38],

<table>
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<tr>
<th>Table 1. Advantages and disadvantages of cytological specimens</th>
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<tr>
<td><strong>Advantages</strong></td>
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<tr>
<td>Easier and safer method to acquire specimens</td>
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<tr>
<td>Reduced time to diagnosis</td>
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<td>Easier access to difficult lesions</td>
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<td>Appropriate for patients not fit for biopsy</td>
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<th>Advantages</th>
<th>Disadvantages</th>
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<tr>
<td>Easier and safer method to acquire specimens</td>
<td>Numerous artefactual variables in specimens, e.g. blood or necrosis</td>
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<tr>
<td>Reduced time to diagnosis</td>
<td>Limited specimen material</td>
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<td>Easier access to difficult lesions</td>
<td>High degree of cell loss</td>
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<tr>
<td>Appropriate for patients not fit for biopsy</td>
<td>Lack of a cytology control for quality assurance</td>
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normal bronchiolar epithelia showed nuclear staining of Clara cells, whilst bronchial epithelial cells (basal cells and mucoserous glands) were negative. Normal and hyperplastic alveolar type II cells were strongly positive for TTF-1. It is expressed in all types of thyroid carcinoma except for the anaplastic type. TTF-1 is now also widely used to identify neoplasms of lung origin.

**Cytokeratins**

Cytokeratins (CKs) are water-soluble intracellular proteins found on most epithelial cells and are part of the intermediate filament family. There are at least 20 sub-classes based on their molecular weight and isoelectric pH value (Moll’s catalogue) [39]. Examples of useful CK immunomarkers used to identify the origin of lung neoplasms include CK7, CK20, 34βE12 which recognises CK1, -5, -10 and -14 of Moll’s catalogue, and CK5/6 [40]. The combination of CK7 and CK20 is commonly used to establish the pulmonary origin of a neoplasm, based on the differential diagnosis and clinical history. CK7 is a 5-kDa basic protein strongly expressed in different epithelia including the breast, endometrium, bladder, pancreas, biliary tract, stomach, and lung [40, 41]. CK20 is a 46-kDa acid protein located in the epithelium of the intestine, bladder, pancreas and biliary tract as well as in Merkel cells [42–44]. A lung tumour with CK20-positive immunoreactivity strongly suggests a metastatic carcinoma rather than a primary lung malignancy. However, mucinous adenocarcinoma of the lung may also express CK20 positivity, especially mucinous BAC [45, 46]. The 34βE12 monoclonal antibody is a highly sensitive marker of normal epithelial basal and parabasal cells and their proliferations [38].

**Neuroendocrine Immunomarkers**

Neuroendocrine immunomarkers are used for identifying tumours of neuroendocrine origin such as SCLC. They include chromogranin A, synaptophysin and neural cell adhesion molecule (NCAM). Chromogranin A is part of the chromogranin family of acidic glycoproteins. Synaptophysin is located in the neuronal presynaptic vesicles and is also positive in paraganglioma, thyroid medullary carcinoma, endocrine pancreatic tumours and carcinoid tumours. NCAM (CD56) is part of the intercellular adhesion molecule family that binds cells together. It is found on neurons, astrocytes, Schwann cells, myoblasts and natural killer lymphocytes. Among neoplasms, it is mainly expressed in neuroendocrine carcinomas [47] but also in some aggressive lymphomas such as natural killer/T-cell lymphoma with cutaneous involvement [48].

<table>
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<tr>
<th>Indication</th>
<th>Immunomarker</th>
<th>Predominant histology correlation (positive immunoreactivity)</th>
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<tr>
<td>Establishing malignancy</td>
<td>p53</td>
<td>preinvasive dysplastic lesions, e.g. autofluorescence bronchoscopic specimens</td>
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<td>Determionation of primary site of origin</td>
<td>TTF-1</td>
<td>ADC and nonmucinous BAC</td>
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<tr>
<td></td>
<td>napsin A</td>
<td>ADC</td>
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<td></td>
<td>CK7</td>
<td>ADC</td>
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<td>CK5/6</td>
<td>Sq-CC</td>
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<td></td>
<td>CK20</td>
<td>Non-pulmonary tumour</td>
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<td>34βE12</td>
<td>Sq-CC</td>
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<td></td>
<td>p63</td>
<td>Sq-CC</td>
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<tr>
<td></td>
<td>chromogranin A</td>
<td>NE, SCC and LCNEC</td>
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<td>synaptophysin</td>
<td>NE, SCC and LCNEC</td>
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<td></td>
<td>NCAM</td>
<td>NE, SCC and LCNEC</td>
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<tr>
<td></td>
<td>vimentin</td>
<td>Mesenchymal origin</td>
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<td>desmin</td>
<td>Mesenchymal origin</td>
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<tr>
<td>Determination of targeted therapy and prognosis</td>
<td>EGFR exon mutational antibodies</td>
<td>ADC (better outcome)</td>
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<tr>
<td></td>
<td>Ki-67</td>
<td>NSCLC stages I-III (poor prognosis)</td>
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ADC = Adenocarcinoma; Sq-CC = squamous cell carcinoma; SCC = small cell carcinoma; NE = neuroendocrine tumours (typical and atypical carcinoids).
Other Immunomarkers

Surfactant apoproteins A and B antibodies have been used to establish the pulmonary origin of lung cancer. Their sensitivity was only 63% in detecting primary lung cancers, with a specificity of 46% [49, 50]. p63 and p53 belong to the family of transcription factors. Early and frequent genomic amplification of p63 has been demonstrated in the development of squamous cell lung carcinoma. p63 genomic sequence amplification was seen in 88% of squamous cell carcinomas, 42% of large cell carcinomas and 11% of adenocarcinomas [51]. Napsin A is a functional aspartic proteinase expressed in normal lung parenchyma in type II pneumocytes and in the proximal and convoluted tubules of the kidneys [52]. It is strongly positive in 80% of lung adenocarcinomas. Napsin A has been reported to show a stronger, more diffuse and more sensitive expression than TTF-1 and to stain 11% more lung adenocarcinomas than TTF-1. It stains negative for squamous cell carcinoma and small cell carcinoma of the lung [49]. Ki-67 is a DNA-binding nuclear protein expressed throughout the cell cycle in proliferating cells but not in quiescent cells [53]. Ki-67 immunomarkers such as MIB-1 have been used to assess prognosis in non-small cell lung carcinoma (NSCLC), as will be discussed later.

How Are Immunomarkers Applied in Lung Cancer?

Establishing the Presence of Malignancy in a Tissue Sample

Establishing malignancy is always based on morphological features of a cytological or histological specimen. It may, however, not be feasible to morphologically diagnose lung cancer or classify lung tumours accurately due to factors such as inadequate cytology or biopsy specimens or presence of artefacts [54]. The role of immunohistochemistry in confirming malignancy is still limited. It remains an ancillary technique that supports the morphological diagnosis and does not independently establish a diagnosis of malignancy. Wang et al. [20] showed various degrees of positive immunostaining for CKs, p53 and Ki-67 in the four categories of bronchial epithelial dysplasia, with Ki-67 showing more widespread expression and stronger staining than p53. The degree and extent of dysplasia was positively correlated with the incidence of bronchogenic carcinoma (p < 0.05). Another study on p63 in 43 preinvasive lesions showed that p63 immunostaining increases progressively throughout the depth of epithelia from metaplasia to severe dysplasia [51]. In this study, however, genomic amplification of the p63 copy number gain was notable in high-grade dysplasia (n = 13), similar to the copy number gain in the adjacent invasive squamous cell carcinoma, while low-grade preinvasive lesions did not demonstrate a p63 copy number gain even if these lesions were in the vicinity of the invasive tumour. The authors concluded that due to its direct correlation with pathological dysplastic grading, p63 immunostaining is less effective than p63 genomic amplification in detecting early tumourigenesis [51]. Nowadays, autofluorescence bronchoscopy is able to detect abnormal mucosa, and biopsies can be taken from these areas; however, these biopsy specimens may not confirm overt malignancy but only show morphological dysplastic changes [55, 56]. Immunomarkers such as p53, p63 and Ki67 may be useful in such instances to establish the preinvasive process and therefore support the decision-making process regarding prognosis and therapy.

Determination of the Primary Site of Tumour Origin and the Cell Line

It must be stated that different pathology laboratories use different immunomarker panels to establish the primary site of tumour origin and the possible cell line. Rossi et al. [54] stated that the most currently validated immunomarkers for refining most diagnoses are TTF-1 and/or CK7 for adenocarcinomas and p63 and the high-molecular-weight CKs (-1, -4, -5, -6, -10, -11, -14) for squamous cell carcinomas.

Adenocarcinoma and BAC

TTF-1 is now widely used to identify adenocarcinomas of pulmonary origin, with a sensitivity of 75–94% and a specificity of 100% [57]. In one study, 23 out of 26 (88.5%) adenocarcinomas showed strong nuclear staining with TTF-1 antibody [38]. In another study, 12 out of 14 (85.7%) pleomorphic tumours with an adenocarcinomatous component showed immunoreactivity for TTF-1 [58]. TTF-1 is also positive in nonmucinous BAC. This reflects the common precursor origin from AAH, which is a terminal respiratory unit lesion. It has been shown that TTF-1 is negative in 25% of adenocarcinomas, especially in those that are centrally derived as well as in mucinous BAC [27, 59]. CK7 is also useful in identifying adenocarcinomas of pulmonary origin. Typically, an adenocarcinoma of pulmonary origin will stain positive for CK7 but negative for CK20 [41, 54]. As discussed above, CK20 is not expressed in the epithelium of the respiratory airways and is therefore used to discriminate adenocarcinomas from the gastrointestinal tract or the urothelium [43, 44, 60]. As mentioned earlier, napsin A can also
be used to identify primary pulmonary adenocarcinoma. For example, napsin A has been used to identify primary pulmonary adenocarcinomas from adenocarcinomas from other organs [61, 62].

Squamous Cell Carcinoma

As discussed above, the precursor cell of squamous cell carcinoma is believed to be the epithelial basal cell. 34βE12 is a useful antibody for identifying squamous cell tumours of basal cell origin. It is positive in all basaloid carcinomas, basaloid variants of squamous cell carcinomas and in squamous cell carcinomas. Sturm et al. [38] showed a 100% sensitivity for 34βE12 in all the above-mentioned carcinomas compared to about 31% in adenocarcinomas. The authors concluded that 34βE12 appears to be a specific marker for neoplasms arising from reserve cells of the bronchial epithelium, recognizing a wide spectrum of epithelial tumours including basaloid carcinoma, squamous cell carcinoma, basaloid variant of squamous cell carcinoma and some adenocarcinomas. Other markers used for positive identification of squamous cell carcinoma include CK5/6 and p63, although none of these have a 100% sensitivity or specificity. Squamous cell carcinoma include CK5/6 and p63, although other markers used for positive identification of squamous cell carcinoma include CK5/6 and p63, although none of these have a 100% sensitivity or specificity. Squamous cell carcinoma typically stains negative for TTF-1, surfactant apoprotein A, chromogranin A and synaptophysin.

Small Cell Carcinoma

As described above, the precursor origin of small cell carcinoma has not been established and seems different from those of other neuroendocrine tumours. However, small cell carcinoma shares common features with other neuroendocrine tumours on immunohistochemistry. It stains positive for chromogranin A, synaptophysin and NCAM CD56 [64, 65]. Morphologically, it may sometimes be difficult to distinguish between small cell carcinoma and other pulmonary neuroendocrine tumours such as carcinoid tumours due to crush artefact. Crush artefact is a known characteristic of small cell carcinomas, regardless of their origin [66]. In such cases, immunostaining with Ki-67 has helped discriminate between these two tumour groups, with small cell carcinomas showing more than 25% immunoreactivity in the crushed areas compared to less than 10% for carcinoid tumours [67]. TTF-1 has a sensitivity of about 75–80% in SCLC. In a study by Ordonez [63], 27 of 28 (96%) SCLC were TTF-1 positive compared to 4 out of 54 (7%) of non-pulmonary small cell carcinomas. CK20 which stains positive in non-pulmonary small cell carcinomas such as Merkel cell carcinomas is useful in discriminating between the two malignancies [68]. 34βE12 is always negative in small cell carcinoma and, if positive, may suggest histological heterogeneity in combination with a non-small cell tumour [29].

Large Cell Carcinoma

Large cell carcinoma is a tumour with no differentiation to allow classification into squamous cell carcinoma, adenocarcinoma or small cell carcinoma. In the WHO classification of tumours of the lung [18], several subtypes of large cell carcinoma exist. The two common subtypes are LCNEC and basaloid carcinoma. LCNEC has immunohistochemical properties similar to small cell carcinoma and other neuroendocrine tumours. In a study by Sturm et al. [38], the sensitivities in LCNEC of chromogranin A, synaptophysin and NCAM were 68.5, 84.2 and 91.2%, respectively. The sensitivity of TTF-1 expression in LCNEC ranges from 50–75% [69, 70], and therefore, TTF-1 is not helpful in differentiating LCNEC from solid adenocarcinomas in which TTF-1 also stains positive. The monoclonal antibody 34βE12 is negative in LCNEC. It is known that LCNEC can exist in pure forms or in combined forms with squamous cell carcinoma. When this happens, 34βE12 is positive. The basaloid large cell carcinoma is identified by the specific staining with 34βE12 monoclonal antibody (as discussed above for cancers of basal cell origin) and negative staining with neuroendocrine markers. One study [38] found that all 28 cases of basaloid carcinoma expressed 34βE12 but were negative for TTF-1.

Neuroendocrine Tumours

This group includes neuroendocrine hyperplasia, carcinoid tumourlets and typical and atypical carcinoid tumours. They all stain positive for the common neuroendocrine immunomarkers discussed above. Neuron-specific enolase is no longer considered useful in the identification of neuroendocrine tumours, as it has been found to be expressed in some non-neuroendocrine tumours [71]. To establish the pulmonary origin of neuroendocrine tumours, TTF-1 has been used, but the sensitivity ranges from 35 to 94% [63, 72–74]. In the study by Du et al. [57] on immunoreactivity of TTF-1 in neuroendocrine tumours from the lung, thymus, gastrointestinal tract, pancreas, and ovary, TTF-1 expression was shown in 10 out 36 (27.8%) pulmonary typical carcinoids and in 29.4% of pulmonary atypical carcinoids compared to complete none reactivity (0%) in extrapulmonary typical and atypical carcinoid tumours. On the contrary, other authors reported negative staining for TTF-1 in all carci-
noid tumourlets as well as in typical and atypical carcinoid cases [75]. Du et al. [57] argued that this observed difference in TTF-1 reactivity was multifactorial and cited reasons such as differences in methodology and antibody concentrations used, differences in scoring for intensity and extent of stain as well as possible differences in the location of the tumours analysed, that is whether central or peripherally derived.

**Sarcomatoid Carcinoma**

This is a group of carcinomas with an epithelial origin combined with varying degrees of mesenchymal differentiation [76]. The tumours have either mixed components of spindle cells or giant cells or other mesenchymal elements but can also occur in pure forms. They have been described not only in the lung but also in the skin, breast, urogenital and gastrointestinal tract [77]. The WHO histological subtypes include pleomorphic carcinoma, giant cell carcinoma, spindle cell carcinoma, carcinosarcoma and pulmonary blastoma [18]. The immunomarkers MNF116, AE1/AE3 and epithelial membrane antigen are used to confirm their epithelial origin. TTF-1 and other CKs can be used to identify the specific carcinomatous components of either adenocarcinoma or squamous cell carcinoma. Rossi et al. [58] found positive staining for TTF-1 and CK7 in 43% and 62.7%, respectively, of sarcomatoid carcinomas with spindle and/or giant cell components, whereas the pure carcinosarcoma and pulmonary blastoma had negative immunoreactivity with these immunomarkers. Immunomarkers for mesenchymal tissue such as vimentin or desmin can be used to identify the mesenchymal elements. The morphological appearance is important to identify the sarcomatous elements in the specimen, and a cytological specimen may be inadequate for such tumours.

**Therapeutic and Prognostic Applications**

It is now no longer helpful to define a lung cancer generally as NSCLC or poorly differentiated NSCLC, and efforts must be made to specify the histological subtype. Current evidence has confirmed that establishing the exact histological subtype has therapeutic and prognostic implications. Studies comparing pemetrexed-based regimens with non-pemetrexed regimens have shown improved response rates, progression-free survival and overall survival in patients with non-squamous cell histology (adenocarcinoma and large cell carcinoma) compared to patients with squamous cell carcinoma [19, 78]. Furthermore, studies have revealed that patients with lung adenocarcinomas that have EGFR mutations show better survival rates when treated with EGFR tyrosine kinase inhibitors (gefitinib and erlotinib) than with conventional cytotoxic therapy [79]. Angiogenesis inhibitors such as bevacizumab and sunitinib have been associated with an increased risk of haemorrhagic adverse effects when used in patients with squamous cell carcinoma compared to non-squamous NSCLC [19, 78]. The management of NSCLC has now evolved to optimising individually targeted therapy and addressing safety issues based on histological subtypes, and not on staging and performance status alone [78].

Although fluorescence in situ hybridization and mutational analysis are commonly used to identify which lung carcinomas, especially adenocarcinomas, are amenable to targeted therapy, immunochemistry is now being extended to this field as well, particularly in identifying EGFR mutations. The EGFR family comprises 4 transmembrane receptors belonging to the tyrosine receptor superfamily, and EGFR activation promotes tumour proliferation, angiogenesis and metastasis [80]. Twenty percent of lung adenocarcinomas exhibit EGFR mutations and 90% of EGFR mutations occur in exon 19 (15-bp/5-aa deletion) and exon 21 (L858R point mutation) [81]. Other EGFR mutations associated with sensitivity to anti-EGFR therapy are G719 mutations in exon 18 and L861 mutations in exon 21, but these are less common [81]. The gold standard method of detecting EGFR mutations is by molecular analysis with direct DNA sequencing. However, direct DNA sequencing has its own limitations which include the need for a good specimen size with a good turnover of tumour cells for DNA extraction, its high cost and a methodological complexity resulting in its use only in academic centres and commercial laboratories [82]. Specific antibody immunomarkers have now been developed targeting these EGFR exon mutants. A study of two antibodies against these exon mutants showed different sensitivities and predictive values depending on the cutoff point selected according to the immunohistochemical staining grading scale of 0 to 3. At a cutoff point of 1+, the sensitivity for the L858R antibody was 95% with a positive predictive value (PPV) of 99%, while at a cutoff value of 2+, the sensitivity decreased to 76% with a PPV of 100%. For the exon 19 deletion 15-bp/5-aa antibody, the sensitivity and PPV were 85 and 99%, respectively, at a cutoff value of 1+ and 67 and 100%, respectively, at a cutoff point of 2+ [83]. The authors, however, advised that in cases where the antibody tests were negative or inconclusive, EGFR molecular testing should still be performed. They concluded that EGFR mutant-specific antibodies should be included in the workup of...
lung adenocarcinoma to allow more rapid initiation of EGFR tyrosine kinase inhibitor therapy, and lead to a modest reduction in the volume of molecular testing [83]. A different study, however, showed a much lower overall sensitivity of 47% for these mutation-specific antibodies (L858R and del E746-A750 in exon 19), with an overall specificity of 96% [84]. The low sensitivity in this study was attributed to cross-reactivity of these antibodies with other mutations in exon 18 (G719C), exon 19 (del L747-T751 ins Q), exon 20 (A769 ins ASV) for the L858R antibody and exon 20 (D770 ins SVD) for the anti-del E746-A750 antibody. Furthermore in this study, one of two lung cancers with the EGFR wild type had positive immunoreactivity despite absence of any EGFR mutation. The authors concluded that even though these antibodies have good specificity, their low sensitivity implies that not all patients with EGFR mutations can be selected using these mutation-specific antibodies, and therefore clinical utility may be less than expected [84].

Anti-Ki-67 has been analysed in the determination of prognosis of lung cancer. In a meta-analysis of 37 studies on lung cancer, of which 29 were on NSCLC, expression of Ki-67 in resected NSCLC specimens was a poor prognostic factor for patients with stages I–III, even in subgroup analysis of histological subtypes [53]. On the contrary, Yang et al. [85] found no correlation between Ki-67 expression and prognosis in NSCLC stage I–IIIA. Such variable findings in the prognostic value of Ki-67 in NSCLC have been attributed to factors such as patient heterogeneity, retrospective nature of the study design, quantification of IHC and different cutoff values [86].

Conclusion

Understanding the basic histology of the respiratory airways as well as their cell structure and composition has led to further understanding the possible cell line origins of primary lung neoplasms. Targeting the basic cellular antigens, immunochemistry has not only improved the diagnosis of lung cancer but also shed light on the histogenesis of some of the primary lung cancers. Using immunochemistry techniques, primary lung carcinoma can now be distinguished from a metastatic carcinoma. However, further studies are required to consolidate the role of immunochemistry in guiding targeted therapy in lung cancer and determining prognosis.

Disclosure Statement

The authors have no conflicts of interest to declare.

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