

REFERENCES

1. Lee SJ, Lee JI, Nam DH, et al. Leptomeningeal carcinomatosis in non-small-cell lung cancer patients: impact on survival and correlated prognostic factors. *J Thorac Oncol* 2013;8:185–191.
2. Brem SS, Bierman PJ, Brem H, et al.; National Comprehensive Cancer Network. Central nervous system cancers. *J Natl Compr Canc Netw* 2011;9:352–400.
3. Chamberlain MC, Kormanik PA. Prognostic significance of coexistent bulky metastatic central nervous system disease in patients with leptomeningeal metastases. *Arch Neurol* 1997;54:1364–1368.
4. Chamberlain MC, Glantz M, Groves MD, Wilson WH. Diagnostic tools for neoplastic meningitis: detecting disease, identifying patient risk, and determining benefit of treatment. *Semin Oncol* 2009;36(4 Suppl 2):S35–S45.
5. Morris PG, Reiner AS, Szenberg OR, et al. Leptomeningeal metastasis from non-small cell lung cancer: survival and the impact of whole brain radiotherapy. *J Thorac Oncol* 2012;7:382–385.

Interpretation of Anti-ALK Immunohistochemistry Results

In their report,¹ Conklin et al. compared five antianaplastic lymphoma

kinase immunohistochemistry (ALK IHC) systems. I agree with their conclusion that IHC is reliable for detection of *ALK* rearrangement; however, I would like to comment on their interpretation of individual results.

Address for correspondence: Kengo Takeuchi, MD, PhD, Pathology Project for Molecular Targets, The Cancer Institute, Japanese Foundation for Cancer Research, 3-8-31 Ariake, Koto, Tokyo 135-8550, Japan. E-mail: kentakeuchi-ky@umin.net
Disclosure: The author is serving as an academic consultant for Nichirei Bioscience.

Copyright © 2013 by the International Association for the Study of Lung Cancer
ISSN: 1556-0864/13/0807-0e66

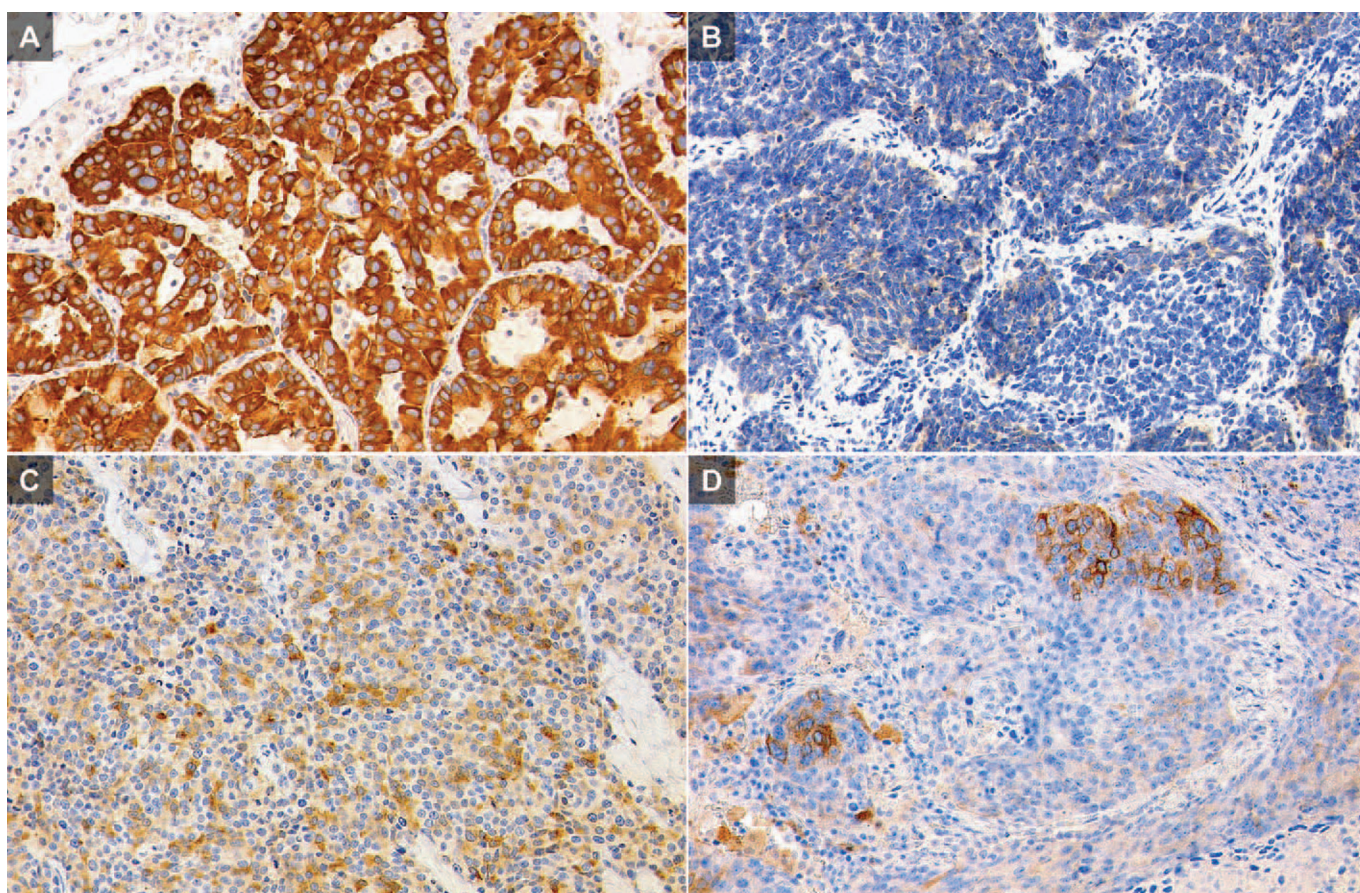


FIGURE 1. ALK iAEP IHC in lung cancers with or without *ALK* fusion genes. Expression of *ALK* fusion protein is dependent on the promoter-enhancer activity of *ALK* partner genes, including *EML4*, which are usually housekeeping genes. In *ALK* fusion-positive tumors, therefore, all tumor cells are immunostained for the *ALK* kinase domain in IHC using anti-*ALK* antibody directed to the kinase domain, when stained appropriately by using a highly sensitive method, such as iAEP. All the cancer cells in an *EML4-ALK*-positive lung cancer express *EML4-ALK* protein (A). Some lung cancer cases (<1%) may express full length *ALK*. In such a case, the staining intensity usually varies from cell to cell, showing staining heterogeneity. This heterogeneity is probably because *ALK* expression in these cases is physiological, as in normal nerve cells, and because the *ALK* promoter-enhancer activity varies among cells. *ALK* rearrangement-negative small-cell carcinoma (B), large-cell neuroendocrine carcinoma (C), and poorly differentiated carcinoma (squamous cell carcinoma in this case, D) of lung may sometimes express the full length *ALK*, and the staining pattern is usually heterogeneous. IHC, immunohistochemistry; *ALK*, anaplastic lymphoma kinase; iAEP, intercalated antibody-enhanced polymer.

They determined the sensitivity and specificity of five systems including “any ALK expression by IHC on tissue microarray (TMA) or whole section (WS),” using fluorescent in situ hybridization (FISH) on WS as the standard.¹ According to their criteria, both sensitivity and specificity of ALK1-ADVANCE should be 100%; however, these values are stated as 66% and 87.5%, respectively, in Table 3.¹

A TMA specimen corresponds to a part of the WS. Therefore, if a TMA scores 2+, the corresponding WS should score 2+ focal (heterogeneity) or higher. If a WS scores 2+ diffuse, the corresponding TMA always scores 2+. Interestingly, for the 5A4-Histofine staining in cases 10 and 11, the TMA and WS results are different by two scores (2+ or 0) (Tables 1 and 2).¹ This score difference might be because of either serious staining errors, or accidental interchanging of the results because such high difference is unlikely unless heterogeneity exists in the WS, and actually, heterogeneity was not observed in the WS. Similarly, the score difference of ALK1-ADVANCE for case 3 is also unlikely. Therefore, I would like the authors to check these results and re-stain the sections.

For ALK1 staining, the TMA and WS scores show discordance (1+ or 0) for cases 3 and 11. Such discordance may occur because the observer struggled to determine whether the faint positivity of score 1+ was real

positivity, unlike the readily detectable staining of score 2+ and 3+, as mentioned by the authors.¹ Given that, would it be appropriate to define score 1+ as positive while calculating sensitivity? From this point of view, in case 11, the best and practical sensitivity was obtained only with 5A4-Histofine staining—a readily detectable staining of score 2+—whereas with other stains, the scores were either 1+ or 0 (Table 2).¹

In my published^{2,3} and unpublished records for anti-ALK IHC of more than 4500 lung cancer cases by using the intercalated antibody-enhanced polymer (iAEP) method,³ a highly-sensitive method on which the 5A4-Histofine staining is based, almost all cancer cells were stained in more than 300 *ALK*-rearranged cases. This staining homogeneity supports the view that all tumor cells of *ALK*-rearranged tumors harbor *ALK* rearrangement.⁴ Wild-type *ALK* is weakly expressed physiologically in normal nerve cells.⁵ Therefore, lung cancers without *ALK* rearrangement sometimes show positivity in highly sensitive anti-ALK IHC, such as the iAEP method, especially in cases with neuroendocrine differentiation (small-cell, large-cell neuroendocrine, and other carcinomas with focally neuroendocrine differentiation).² However, unlike in *ALK*-rearranged cases, the staining pattern in these cases is usually heterogeneous probably because the physiological expression status varies from cell to cell (Fig. 1). In highly sensitive anti-ALK IHC for detection

of *ALK* rearrangement, therefore, a heterogeneous staining pattern should not be interpreted as *positive for ALK rearrangement*, but should be considered *probably negative for ALK rearrangement*, and then be confirmed through FISH. This anti-ALK IHC interpretation would have made the specificities of 5A4-ADVANCE and 5A4-Histofine 100% (Table 2).¹

Kengo Takeuchi, MD, PhD

Pathology Project for Molecular Targets and Division of Pathology
The Cancer Institute
Japanese Foundation for Cancer Research
Tokyo, Japan

REFERENCES

1. Conklin CM, Craddock KJ, Have C, Laskin J, Couture C, Ionescu DN. Immunohistochemistry is a reliable screening tool for identification of *ALK* rearrangement in non-small-cell lung carcinoma and is antibody dependent. *J Thorac Oncol* 2013;8:45–51.
2. Takeuchi K, Soda M, Togashi Y, et al. RET, ROS1 and *ALK* fusions in lung cancer. *Nat Med* 2012;18:378–381.
3. Takeuchi K, Choi YL, Togashi Y, et al. KIF5B-*ALK*, a novel fusion oncokine identified by an immunohistochemistry-based diagnostic system for *ALK*-positive lung cancer. *Clin Cancer Res* 2009;15:3143–3149.
4. Mano H, Takeuchi K. EML4-*ALK* fusion in lung. *Am J Pathol* 2010;176:1552–3; author reply 1553.
5. Pulford K, Lamant L, Morris SW, et al. Detection of anaplastic lymphoma kinase (*ALK*) and nucleolar protein nucleophosmin (*NPM*)-*ALK* proteins in normal and neoplastic cells with the monoclonal antibody ALK1. *Blood* 1997;89:1394–1404.