

## Immunohistochemical expression of anaplastic lymphoma kinase in neuroblastoma and its relations with some clinical and histopathological features

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**Background:** Anaplastic lymphoma kinase (ALK) mutations have been identified as a prominent cause of some familial and sporadic neuroblastoma (NB). ALK expression in NB and its relationship with clinical and histopathological features remains controversial. This study investigated ALK expression and its potential relations with these features in NB. **Methods:** Ninety cases of NB at the Department of Pathology, University of Medicine and Pharmacy at Ho Chi Minh City, Viet Nam from 01/01/2018 to 12/31/2021, were immunohistochemically stained with ALK (D5F3) antibody. The ALK expression and its relations with some clinical and histopathological features were investigated. **Results:** The rate of ALK expression in NB was 91.1%. High ALK expression (over 50% of tumor cells were positive with moderate-strong intensity) accounted for 65.6%, and low ALK expression accounted for 34.4%. All the *MYCN*-amplified NB patients had ALK immunohistochemistry positivity, most cases had high ALK protein expression. The undifferentiated subtype of NB had a lower ALK-positive rate than the poorly differentiated and differentiated subtype. The percentages of ALK positivity were significantly higher in more differentiated histological types of NB (p=.024). There was no relation between ALK expression and: age group, sex, primary tumor location, tumor stage, *MYCN* status, clinical risk, Mitotic-Karyorrhectic Index, prognostic group, necrosis, and calcification. **Conclusions:** ALK was highly expressed in NB. ALK expression was not related to several clinical and histopathological features. More studies are needed to elucidate the association between ALK expression and ALK gene status and to investigate disease progression, especially the oncogenesis of ALK-positive NB.

Key Words: Neuroblastoma; Anaplastic lymphoma kinase

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Neuroblastomas (NB) are tumors deriving from the sympathoadrenal lineage - one of the neural crest derivatives. NB represents the most prevalent extracranial solid tumor in children and accounts for 12% of all childhood cancer deaths [1]. The presence of anaplastic lymphoma kinase (*ALK*) mutations in NB has been published since 2008 [2]. Approximately 8% to 10% of NB have *ALK* mutations, and about 25% have *ALK* amplification or overexpression of the ALK protein [3]. Many studies have shown that the *ALK* gene is a significant oncogene in NB, and ALK mutations have been identified in association with familial and sporadic NB [2]. Besides, ALK is a consummate tumor antigen for targeted inhibition, due to its frequent expression in NB and restricted distribution in normal tissue [4]. These results open the door to new therapeutic strategies for treating NB with ALK inhibitors.

Several research groups have reported that ALK overexpression in NB is an adverse prognostic factor, however, the relations between ALK protein expression and clinical and histopathological features of NB are still controversial [2,5-7]. There are many methods to evaluate ALK in NB, in which immunohistochemistry (IHC) is available and capable of routine application. The clone D5F3 showed the highest sensitivity compared to other clones such as ALK 1, and 5A4. The ALK D5F3 antibody is likely the most appropriate antibody for determining ALK protein expression levels in NB, which may be used to predict patient prognosis [8,9]. In this study, ALK protein expression was studied to investigate potential relations with some clinical and pathological features of NB.

## MATERIALS AND METHODS

## Patients and specimens

Patients who were histologically diagnosed with NB (Schwannian stroma-poor) (based on International Neuroblastoma Pathology Classification [INPC]) at the Department of Pathology, University of Medicine and Pharmacy at Ho Chi Minh City, Viet Nam from January 2018 to December 2021 were enrolled in this study. All tumor specimens were obtained from the primary or metastatic tumor through surgery or biopsy before starting chemotherapy. The clinical parameters, including age at diagnosis, gender, tumor stages (based on International Neuroblastoma Risk Group Staging System [INRGSS]), primary tumor sites were collected. MYCN status (by fluorescence in situ hybridization [FISH]) was retrospectively inquired through record data from the Center for Molecular Biomedicine, University of Medicine and Pharmacy at Ho Chi Minh City. Hematoxylin a eosin slides were assessed for degree of differentiation, Mitotic-Karyorrhectic Index (MKI), and prognostic groups (according to INPC). In addition, tumor necrosis and calcification will also be noted.

Formalin-fixed, paraffin-embedded tumor tissue collected during surgery or biopsy was used. For surgical specimens, a representative area of tumor tissue will be submitted for tissue microarray, and whole biopsied specimens will be collected for IHC staining.

## IHC staining

IHC staining was performed on 4-µm-thick sections using VENTANA anti-ALK (clone D5F3, Ventana Medical Systems, Tucson, AZ, USA) rabbit monoclonal primary antibody was used in this study. The IHC staining process was fully automated using a BenchMark XT automated slide stainer (Ventana Medical Systems).

Cytoplasmic and membrane staining with any ratio was considered positive. Negative when absolutely no tumor cells were stained. ALK IHC staining was scored based on the percentage of antibody-reactive neoplastic cells and the intensity as follows: "0" (negative, no stained cells), "1+" (weak expression, <20% of cells stained), "2+" (heterogeneous weak-moderate expression, around 20%–50% of cells stained), "3+" (heterogeneous moderate-strong expression, >50% of cell stained), and "4+" (>75%, strong expression). Cases with ALK IHC score 3+, and 4+ are considered to have high ALK protein expression (i.e., more than 50% of tumor cells are stained with moderate-strong intensity). The remaining cases were considered to have low ALK protein expression (IHC ALK score 0, 1+, 2+).

## Statistical analysis

Pearson's  $\chi^2$  test (or Fisher's exact test if the sample size was small) was used to evaluate relationship between pairs of categorical variables. All statistical analyses were performed using STATA ver. 14.2 (Stata Corp., College Station, TX, USA). p-value < .05 were statistically significant.

## RESULTS

## Patient characteristics

A total of 90 patients were enrolled in this study. The clinical and pathological characteristics of the patients are summarized in Table 1. The median age of patients was 24 months old (15 days old–14 years old), and the group under 18 months old accounted for the majority. The male/female ratio is 1.3:1. In 80 cases tested for FISH *MYCN*, 11.3% had *MYCN* amplification. More than 80% of NB were poorly differentiated subtype, unfavorable histology group accounted for the majority with 57.8% (Table 1).

## ALK protein expression

ALK was positive in 91.1% of cases (82/90 cases), in which high ALK expression (3+, 4+) accounted for 65.6% (59/90 cases). The ALK IHC expression is summarized in Table 2.

ALK IHC staining was observed in the cytoplasm or membrane of NB cells and was negative for lymphocytes, endothelium, and normal adrenal gland (Fig. 1). Different intensity of ALK IHC was shown in Fig. 2.

# The relation between ALK expression and some clinical and pathological features

Our study showed all cases of differentiating NB were ALKpositive (Table 3). There was significant relation of ALK positivity with NB histological types (undifferentiated NB [9/13, 69.2%] vs. poorly differentiated NB [69/73, 94.5%] vs. differentiating NB [4/4, 100%; p = .024]). All cases with MYCN amplification were positive for ALK IHC, in which high ALK expression accounted for 7/9 cases (77.8%). No other relations between ALK expression and clinical and pathological features were observed (Table 3).

Table 1.	<ul> <li>Characteristics</li> </ul>	of patients	(n = 90)
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Characteristic	No. (%)			
Age (mo)				
<18	43 (47.8)			
18–60	31 (34.4)			
≥60	16 (17.8)			
Sex				
Male	51 (56.7)			
Female	39 (43.3)			
Primary site				
Retroperitoneum/Adrenal	74 (82.2)/47 (52.2)			
Posterior mediastinum	12 (13.3)			
Neck	3 (3.3)			
Unknown	1 (1.1)			
Stage (INRGSS)				
L1	25 (27.8)			
L2	26 (28.9)			
Ms	2 (2.2)			
Μ	37 (41.1)			
MYCN status				
Amplified	9 (11.3)			
Non-amplified	71 (88.7)			
Degree of differentiation				
Undifferentiated	13 (14.4)			
Poorly differentiated	73 (81.1)			
Differentiating	4 (4.5)			
MKI				
Low	54 (60)			
Intermediate	16 (17.8)			
High	20 (22.2)			
Prognostic group (INPC)				
Favorable	38 (42.2)			
Unfavorable	52 (57.8)			
Necrosis				
Yes	27 (30.0)			
No	63 (70.0)			
Calcification				
Yes	22 (24.4)			
No	68 (75.6)			

INRGSS, International Neuroblastoma Risk Group Staging System; MKI, Mitotic-Karyorrhectic Index; INPC, International Neuroblastoma Pathology Classification.

## Table 2. ALK IHC expression

ALK IHC score	No. (%)		
Low expression			
0	8 (8.9)		
1+	10 (11.1)		
2+	13 (14.4)		
High expression			
3+	13 (14.4)		
4+	46 (51.1)		
Total	90 (100)		

ALK, anaplastic lymphoma kinase; IHC, immunohistochemistry.

## DISCUSSION

Until now, there is no consensus method for evaluating ALK IHC staining in NB, most of the studies were based on the percentage of ALK antibody-reactive neoplastic cells and modified evaluation methods depending on the research groups [2,5,8, 10-13]. Several previous studies reported the relation between high ALK expression (> 50% of tumor cells) and poor clinical outcome [2,5,13], so a positive threshold of more than 50% of stained tumor cells was applied in many studies [6,7]. In this study, we applied the modified assessment method according to Passoni et al. [2], and investigated the relations between high/ low ALK protein expression with some clinical and histopathological factors. Our study showed that the ALK expression rate was 91.1%, similar to many other studies [2,5,9,10]. ALK protein is a single-chain transmembrane protein comprised of three regions: extracellular domain, transmembrane region, and intracellular domain [4], so ALK IHC showed cytoplasmic/membranous expression. ALK protein is expressed on the surface of most NB cells, and its expression is uncommon in normal tissue [4,9]. This makes ALK an ideal target for cancer therapy, whether the NBs exhibit ALK mutations [4,9].

In this study, ALK expression was not related to clinical features including age, sex, stage, or *MYCN* status. A previous study reported that the incidence of ALK expression increased correspondingly with more advanced tumor stages (p = .001) [7]. In another study by Passoni et al. [2], ALK protein expression was significantly up-regulated in advanced/metastatic NB, however, our study showed that ALK expression was not related to the disease stage [2]. This discrepancy may be attributed to differences in the staging system used in the two studies (INRGSS vs. International Neuroblastoma Staging System [INSS]), and the ALK antibody clone (D5F3 vs. 5A4, ALK1, SP8, each antibody binds to a different epitope, which may explain why they have different sensitivities [14]).

Our study revealed that all cases with *MYCN* amplification were positive for ALK IHC (Table 3), in which high ALK expression accounted for 7/9 cases (77.8%). Similar to our results, Lee et al. evaluated ALK expression on 70 NBs and reported that all seven cases with *MYCN* amplification were positive for ALK IHC (over 50% tumor cells expressed) [7]. In another study by Wang et al. [6], NBs with *MYCN* amplification or gain were more likely to be ALK-positive than tumors with normal *MYCN* status (p < .05) [6]. Yan et al. [8] found a strong relationship between ALK D5F3 IHC and the number of copies of the *MYCN* gene on NB. Schonherr et al. [15] reported that ALK activity was

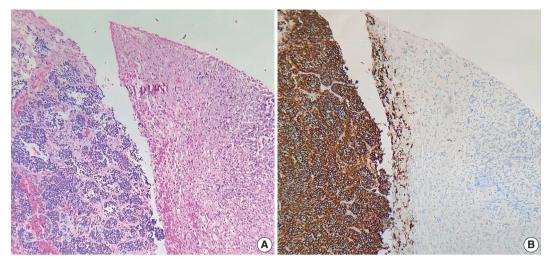


Fig. 1. (A) Neuroblastoma (left side) and normal adrenal gland (right side). (B) Strong immunohistochemistry staining of anaplastic lymphoma kinase (ALK) in tumor cells (left side), no ALK staining is found in normal adrenal tissue (right side).

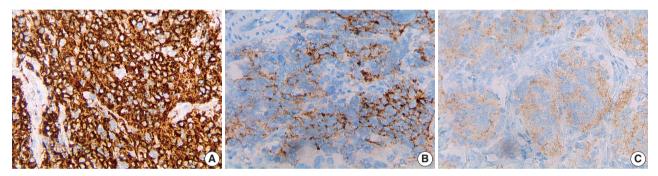


Fig. 2. (A) Immunohistochemistry (IHC) staining for anaplastic lymphoma kinase (ALK) in neuroblastoma (NB) showed strong, homogeneous cytoplasmic or membraned staining. (B) Moderate ALK IHC staining in NB. (C) Weak, heterogenous ALK IHC staining in NB.

very important in *MYCN* transcription initiation, and *MYCN* gene transcription was eliminated by using a specific ALK inhibitor. From our observations combined with several reports that high ALK expression in NB is predictive of treatment response to ALK inhibitors [2,9], we speculate that *MYCN*-amplified NB patients may benefit from ALK inhibitors treatment if high levels of ALK expression are present.

We noted higher percentages of ALK positivity in more differentiated histological types of NB (p = .024). However, there was no statistically significant correlation between ALK IHC score and NB histological types (p = .143), similar to another study [10]. ALK expression is found in neural crest cells during early development, and the involvement of ALK activation in neural crest cell migration and proliferation has been shown [16]. However, how ALK helps neural crest cells to develop in humans is poorly understood [17]. In our study, up to 4/13 cases (30.8%) of undifferentiated NB were negative for ALK IHC. As ALK protein expression was not associated with *ALK* mutations in most cases (half or more of NBs have strong ALK protein expression, but only about 10% of NBs harbor ALK aberrations) [2,5,8], so further molecular biology tests are needed to determine if these ALK IHC-negative cases have the ALK gene mutations. The D5F3 IHC test is highly concordance with ALK-FISH and approved by the U.S. Food and Drug Administration for detecting ALK rearrangements in non-small cell lung carcinoma; however, the discrepancy between these two tests has also been reported [18]. In addition, in the group of pulmonary neuroendocrine tumors, many studies have shown that there is a mismatch between ALK protein expression and ALK alterations [19-22]. Nakamura et al. [19] indicated that the immunopositivity is probably of a wild-type ALK, which may be due to epigenetic regulation or protein overstabilization. Although there are no ALK alterations detected by FISH, cases of lung cancer overexpressing ALK protein that respond positively to ALK inhibitors were announced [23,24]. Similar to NB, ALK-positive tumors may still respond to ALK inhibitor drugs even if tumors

Variable	No. of cases	IHC ALK-positive	IHC ALK-negative	p-value	Low ALK expression	High ALK expression	p-value
Age (mo)				.704ª			.928 <sup>b</sup>
<18	43	40 (93.0)	3 (7.0)		14 (32.6)	29 (67.4)	
18–60	31	28 (90.3)	3 (9.7)		11 (35.5)	20 (64.5)	
≥60	16	14 (87.5)	2 (12.5)		6 (37.5)	10 (62.5)	
Sex				.723ª			.124 <sup>b</sup>
Male	51	47 (92.2)	4 (7.8)		21 (41.2)	30 (58.8)	
Female	39	35 (89.7)	4 (10.3)		10 (25.6)	29 (74.4)	
Stage (INRGSS)				>.99ª			.631ª
L1	25	23 (92.0)	2 (8.0)		7 (28.0)	18 (72.0)	
L2	26	24 (92.3)	2 (7.7)		11 (42.3)	15 (57.7)	
Ms	2	2 (100)	0		1 (50.0)	1 (50.0)	
Μ	37	33 (89.2)	4 (10.8)		12 (32.4)	25 (67.6)	
MYCN				>.99ª			.471ª
Amplified	9	9 (100)	0		2 (22.2)	7 (77.8)	
Non-amplified	71	64 (90.2)	7 (9.8)		28 (39.4)	43 (60.6)	
Differentiation				.024ª			.143ª
Undifferentiation	13	9 (69.2)	4 (30.8)		7 (53.9)	6 (46.1)	
Poorly differentiation	73	69 (94.5)	4 (5.5)		22 (30.1)	51 (69.9)	
Differentiating	4	4 (100)	0		2 (50.0)	2 (50.0)	
MKI				.682ª			.230 <sup>b</sup>
Low	51	45 (88.2)	6 (11.8)		21 (41.2)	30 (58.8)	
Intermediate	19	18 (94.7)	1 (5.3)		6 (31.6)	13 (68.4)	
High	20	19 (95)	1 (5.0)		4 (20.0)	16 (80.0)	
Prognostic group				>.99ª			.682 <sup>b</sup>
Favorable	38	35 (92.1)	3 (7.9)		14 (36.8)	24 (63.2)	
Unfavorable	52	47 (90.4)	5 (9.6)		17 (32.7)	35 (67.3)	
Necrosis				.234ª			.735 <sup>b</sup>
Yes	27	23 (85.2)	4 (14.8)		10 (37.0)	17 (63.0)	
No	63	59 (93.7)	4 (6.3)		21 (33.3)	42 (66.7)	
Calcification				.674ª			.211 <sup>b</sup>
Yes	22	21 (95.5)	1 (4.5)		10 (45.4)	12 (54.6)	
No	68	61 (89.7)	7 (10.3)		21 (30.9)	47 (69.1)	

Table 3. The relation between ALK expression and some clinical and pathological features

Values are presented as number (%).

ALK, anaplastic lymphoma kinase; IHC, immunohistochemistry; INRGSS, International Neuroblastoma Risk Group Staging System; MKI, Mitotic-Karyorrhectic Index.

<sup>a</sup>Fisher's exact test; <sup>b</sup>χ<sup>2</sup> test.

lack *ALK* mutations [5,9], therefore, ALK-negative undifferentiated NB cases may be an indicator of unresponsiveness to ALK inhibitors. Until now, many generations of ALK inhibitors have been created and included in research. Overcoming the limitations of the pioneering ALK inhibitor crizotinib, several novel designed generations of ALK inhibitors and the combinatory therapy with either pathway inhibitors or other agents against other targets are also of interest [25,26]. The definitive efficacy of ALK inhibitors in NB remains to be elucidated upon the conclusion of ongoing clinical trials within the next few years [25].

In summary, this study highlights the characteristics of ALK expression in NB by applying both approaches to evaluating ALK expression in NB and clarifies the reason for the difference in the results of previous studies. ALK expression was not related to several clinical and histopathological features. Our study revealed a significant relation between ALK positivity with NB histological types, and all cases with *MYCN* amplification were positive for ALK, speculating that *MYCN*-amplified NB patients may benefit from ALK inhibitors. Since the role of ALK in neural crest development is still unclear, further studies are needed to elucidate the association between ALK expression and *ALK* gene status and to investigate disease progression, especially the oncogenesis of ALK-positive NB.

## **Ethics Statement**

This study was conducted in accordance with the guidelines of the Declaration of Helsinki and approved by the Institutional Review Board of Biomedical Research of University of Medicine and Pharmacy at Ho Chi Minh City (IRB No. 174/HDDD-DHYD; on February 21, 2022). Informed consent was obtained from all individual participants included in the study.

## Availability of Data and Material

The datasets generated or analyzed during the study are available from the corresponding author on reasonable request.

#### **Code Availability**

Not applicable.

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Conceptualization: TDAP, TQN, DQN. Data curation: TDAP, TQN, TLT, NTT. Formal analysis: TDAP, TQN. Funding acquisition: TDAP, TLT. Investigation: TQN, TDAP, NTT, TLT. Resources: TDAP, NTT. Methodology: TDAP, TQN. Supervision: DQN. Writing—original draft: TQN, TDAP. Writing—review & editing: TDAP, TQN, DQN. Approval of final manuscript: all authors.

#### **Conflicts of Interest**

The authors declare that they have no potential conflicts of interest.

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