

The value of cell block based on fine needle aspiration for lung cancer diagnosis

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Background: Computed tomography (CT)-guided percutaneous lung fine needle aspiration (FNA) is a convenient method to obtain samples from pulmonary lesions. FNA has a lower rate of complications than the use of a core needle biopsy, but is more difficult for the diagnosis of cytological samples. We use cell block (CB) and immunocytochemistry (ICC) to improve the accuracy of cytological diagnoses based on CT-guided percutaneous lung FNA.

Methods: We collected 526 cytological samples obtained using CT-guided percutaneous lung FNA at Shanghai Pulmonary Hospital from May 2015 to October 2015. CBs were created from these samples, and ICC was performed to help the further histological classification and confirmation of tumor as primary or metastatic. An automated Ventana *ALK* with clone D5F3 was used to identify *ALK* fusion protein.

Results: After assessment of the CBs, 32 (6.08%) diagnoses of suspected malignancy were reduced to 10 (1.90%) such diagnoses ($P<0.05$), and 161 (30.61%) cases of non-small-cell lung carcinoma (NSCLC) were reduced to 33 (6.27%) cases ($P<0.05$) after their division into specific subtypes. We also diagnosed eight (1.52%, $P<0.05$) cases of metastatic carcinoma of the lung that were difficult to diagnose by cytological smear alone. Six (3.73%) of 161 NSCLC cases exhibited *ALK* rearrangement.

Conclusions: CB and ICC are useful for accurate cytological diagnosis using CT-guided percutaneous lung FNA. These approaches are valuable for providing individualized treatment and prognostic evaluations with minor complications.

Keywords: Cytology; cell block (CB); immunocytochemistry (ICC); non-small-cell lung carcinoma (NSCLC); fine needle aspiration (FNA); *ALK* rearrangement

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Introduction

Lung cancer is currently the leading cause of cancer-related mortality worldwide (1). Obtaining an accurate diagnosis as soon as possible is very important for the lung cancer patients. Diagnosis of lung cancer can be made in two ways: histopathological diagnosis and cytopathological diagnosis.

Traditional cytopathology is used for differential diagnoses of benign and malignant lesions worldwide. However, accurately diagnosing lung cancer subtype

based only on cell morphology can be difficult due to the anaplasia of lung cancer cells. Prior to the 2004 World Health Organization (WHO) classification for lung cancer, no therapeutic implications were associated with distinguishing among histological subtypes of non-small-cell lung carcinomas (NSCLCs) (2). However, due to major therapeutic advances in the lung cancer field, pathological diagnosis has increasing implications for clinicians, particularly with respect to the differential diagnosis of squamous cell carcinoma (SCC) and adenocarcinoma (AC).

Numerous drugs focus on specific molecular targets that are primarily associated with either SCC or AC.

Core biopsy and fine needle aspiration (FNA) are the two methods of choice for diagnosing of lung cancer (3). Core biopsies yield histological specimens that can be used for immunohistochemistry (IHC), polymerase chain reaction (PCR) and fluorescence *in situ* hybridization (FISH), which are useful for precision medicine. FNA has a lower rate of complications than core needle biopsy, but making diagnoses using cytological samples obtained with FNA is more difficult (4-7). For FNA cytological samples, immunocytochemistry (ICC), which should be executed after using the cell block (CB) approach, plays an important role in the differential diagnosis of lung cancer subtypes. At present, according to the 2015 WHO classification of lung cancer, ICC is recommended for all NSCLC cases that cannot be classified as SCC or AC based on morphology alone (2). Similarly, differentiating metastatic tumors of the lung from primary tumors of the lung based only on smears is difficult, and ICC is also needed to resolve these differences.

Here, we studied the utility of CB and ICC based on FNA samples for diagnosing lung cancer.

Methods

Case collection

When CT scan reports show that the patient might have a tumor, clinicians conducted CT-guided percutaneous lung FNA. We collected 526 cytological samples from May to October of 2015 involved 362 males and 164 females, which corresponded to agender ratio of 2.21:1.00. The patient ages ranged widely from 13–92 years, with a median age of 63 years.

Each cytological sample obtained by CT-guided percutaneous lung FNA (22G) was placed into a 50-cc centrifuge tube with a ThinPrep (HOLOGIC Gen-Probe, San Diego, CA, USA) cytology test (TCT) preservation solution after the creation of conventional smears. TCT smears were obtained using the standard process. Subsequently, samples from all 526 cases were subjected to CB, and ICC was performed. The “cytology smear” group includes a TCT smear and a conventional smear from the same patient. If the TCT smear and conventional smear was positive, a positive result was obtained. The diagnosis of the “cytology smear” group was based on the TCT smear and the conventional smear.

The study was approved by the ethics committee of

Shanghai Pulmonary Hospital (No. K17-125). Written informed consent was obtained from all patients.

Diagnosis criteria

Smears with malignant cells were regarded as positive. Highly suspicious tumor cells were used to identify samples as suspicious of carcinoma. Smears with no tumor cells or only a small quantity of cells with nuclear atypia were regarded as negative. The classification of positive samples was conducted in accordance with the 2015 WHO classification criteria for lung cancer. All of the cases were diagnosed by two experienced cytopathologists, and another cytopathologist was invited to provide a diagnosis in cases of disagreement between the two primary cytopathologists.

CB and ICC

If there were difficulties making an accurate diagnosis, CBs were generated with the cytological samples obtained by CT-guided percutaneous lung FNA. On the second day, H&E-stained slides were generated, and ICC markers were evaluated. On the third day, the ICC slides were ready for assessment, and a diagnosis was made according to the ICC results. All of the slides were stained with an autostainer using an envision detection system (Leica Biosystems Melbourne Pty Ltd., Melbourne, Australia). Antibodies to the following molecules were used: P40 (P40, 1:500, OriGene), TTF-1 (8G7G3/1, 1:200, Dako), SYN (DAK-SYNAP, 1:100, Dako), CD56 {123C3[5], 1:50, Dako}, CgA (DAK-A3, 1:100, Dako), LCA (2B11 + PD7/26, 1:100, Dako), CK5/6 (D5/16B4, 1:100, Dako) and Ki-67 (MIB-1, 1:100, Dako). The negative and positive controls included in the study were napsin A (IP64, 1:100, OriGene), CDX-2 (DAK-CDX2, 1:100, Dako), villin (ID2C3, 1:1, Dako), CK7 (DV-TL, 1:100, Dako), CK20 (KS20, 1:50, Dako), ER (EP1, 1:40, Dako), PR (P9R636, 1:50, Dako), c-erBb-2 (EP3, 1:200, OriGene), and AFP (polyclonal antibodies, 1:800, Dako). *ALK* IHC was performed on a VENTANA Medical System. The primary antibody (clone D5F3, VMSI) was incubated on CB sections for 20 minutes. The Optiview DAB IHC Detection Kit (VMSI) and the Optiview Amplification Kit (VMSI) were used according to the manufacturer's protocol.

Statistical analysis

Statistical analysis, which was conducted using SPSS

Table 1 Success rate of cell block embedding

Diagnosis	Cytology smear (n)	Successful cases of CB (n)	Success rate of CB (%)
Positive	417	372	89.21
SCC	31	28	90.32
AC	158	143	90.51
SCLC	20	17	85.00
NSCLC	161	144	89.44
Low differentiated carcinoma	27	23	85.19
Others	20	17	85.00
Suspicious of malignant	32	27	84.38
Negative	77	68	88.31
Total	526	467	88.78

SCC, squamous cell carcinoma; CB, cell block; AC, adenocarcinoma; SCLC, small cell lung carcinoma; NSCLC, non-small-cell lung carcinoma.

Table 2 Value of cell block embedding for diagnosing benign and malignant tumors

Diagnosis	Cytology smear, n (%)	CB + ICC, n (%)	χ^2	P
Negative	77 (15.17)	72 (1.12)	0.195	>0.05
Positive	417 (79.28)	444 (84.41)	4.663	<0.05
Suspicious of malignant	32 (6.08)	10 (1.90)	12.003	<0.05
Total	526 (100.00)	526 (100.00)	–	–

CB, cell block; ICC, immunocytochemistry.

software (version 17.0, SPSS, Inc., Chicago, IL, USA), included χ^2 tests. P values <0.05 were regarded as statistically significant.

The sensitivity, specificity, positive predictive value (PPV), and negative predictive value (NPV) were used to quantify the ICC marker when diagnosing SCC and AC.

Results

No difference in the CB success rate was found between various specimens

Among the 526 cases, 417 cases were positive, 32 cases were suspicious of carcinoma, and 77 cases were negative. In 59 cases, no cellular component was found after CB, the success rate for CB was 88.78%. The positive specimens were diagnosed as SCC, AC, SCLC, NSCLC, not otherwise specified (NSCLC-NOS), poorly differentiated carcinoma, or other malignant tumors. The number and CB success rates for each type of specimen are summarized in *Table 1*.

No statistically significant differences with respect to the CB success rate were found among the positive specimens, the suspicious of carcinoma specimens, and the negative specimens.

CB has advantages for distinguishing between benign and malignant tumors

After the CBs were assessed, the rate of positive diagnosis increased from 79.28% to 84.41% (P<0.05); in contrast, the rate of suspected carcinoma decreased from 6.08% to 1.90% (P<0.05; *Table 2*). We also identified five false-negative cases after performing CB and ICC.

CB has advantages for determining NSCLC subtypes and identifying metastatic carcinoma of the lung

We determined final subtype diagnoses based on CB and ICC and compared these results with diagnoses based on

Table 3 Value of cell block embedding for subtyping lung cancer

Diagnosis	Cytology smear (%)	CB + ICC (%)	χ^2	P
NSCLC	161 (30.61)	33 (6.27)	23.46	<0.05
AC	158 (25.28)	253 (52.25)	53.187	<0.05
SCC	31 (4.49)	56 (21.91)	8.203	<0.05
Metastatic cancer	0 (0.00)	8 (1.52)	8.092	<0.01

CB, cell block; ICC, immunocytochemistry.

Table 4 IHC marker expression in SCC and AC

Marker	Subtype	Positive rate (%)	Sensitivity (%)	Specificity (%)	PPV (%)	NPV (%)
P40	SCC	76.79	76.79	99.60	97.73	95.09
CK5/6	SCC	85.71	85.71	93.68	75.00	96.73
Napsin A	AC	86.84	81.82	98.21	99.52	54.46
TTF-1	AC	92.59	82.21	100.00	100.00	55.45

SCC, squamous cell carcinoma; AC, adenocarcinoma; IHC, immunohistochemistry; PPV, positive predictive value; NPV, negative predictive value.

smears alone (Table 3). The use of CB and ICC decreased the NSCLC diagnoses from 30.61% to 6.27% of the cases ($P<0.05$). Additionally, in six cases, we obtained conclusions that were different from the initial diagnoses: two cases were diagnosed as AC before CB but subsequently diagnosed as SCC, and four AC cases were misdiagnosed as SCC or SCLC based on smears. We could not distinguish metastatic lung tumors from primary lung tumors based only on smears. After CB and ICC were performed, eight metastatic lung tumors were categorized as four metastatic bowel tumors, three metastatic breast tumors and one metastatic liver tumor (Table 3). We diagnosed metastatic lung tumors not only based on the morphology but also according to the ICC results. Metastatic bowel tumors were positive for CDX-2, villin, and CK20. Metastatic breast tumors were positive for ER, PR, and c-erBb-2, and the metastatic liver tumor was positive for AFP. The ICC for TTF-1 and Napsin A were negative in all of these metastatic tumors.

TTF-1, Napsin A, P40, and CK5/6 constitute reliable panel for subtyping NSCLC in CB specimens.

The positive rate, sensitivity, specificity, PPV, and NPV for each of these markers are presented in Table 4. TTF-1 is the most specific marker for diagnosing AC and has the best PPV index (specificity: 100%, PPV: 100%). P40 is the most

specific marker for diagnosing SCC (specificity: 99.60%) but is occasionally expressed in AC.

ALK fusion can be identified in CB specimens

Six (3.73%) of the 161 NSCLC patients exhibited the *ALK* fusion based on the CB specimens. Using ICC to identify *ALK* fusions is convenient and inexpensive.

Discussion

FNA and core needle biopsy are two methods used to collect pulmonary lesion specimens. However, the latter method has a higher complication rate for percutaneous transthoracic CT-guided biopsies, such as hemothorax, pneumothorax and hemoptysis (5-7). CT-guided percutaneous lung FNA is a minimally invasive procedure to diagnose lung disease that leads to few complications and relatively little damage (8). However, this procedure also has the disadvantages of providing few cellular specimens and not revealing tumor morphology, which interferes with accurate lung cancer subtyping and prognostic assessment (9). Obtaining an accurate diagnosis is extremely important for clinical treatment. In the current era of precise treatment, the lung cancer therapy chosen is determined by the pathological and molecular diagnoses.

Thus, facilitating the efficient use of cytological samples to obtain accurate pathological and molecular diagnoses is critical. Cytological samples are no longer utilized only for diagnosis but are now also used for ICC and molecular testing related to potential targeted therapy. In our hospital, we found that the complication rate from FNA was less than 10%. Clinicians recommend that the patient receive a core biopsy instead of FNA only when the lesion is near the pleura. In this situation, the complication rate of a core biopsy is approximately 25% to 30%, in our hospital, which is consistent with a previous report (5-7).

In our study, we found no statistically significant differences in the CB success rate among positive specimens, suspected carcinoma specimens and the negative specimens. The number of cells is the key factor that determines CB success. If sufficient cells are available, any FNA specimen can successfully be utilized for CB.

False-positive and false-negative diagnoses have been reported in respiratory cytopathology, even for specimens evaluated by highly experienced cytologists (10). We found five false-negative cases in this study. Highly differentiated tumors are difficult to distinguish from epithelial cells, and this difficulty can lead to misdiagnoses. Tumor cells can be squeezed into abnormal shapes, which make their diagnosis by cytopathologists challenging. Moreover, tumor cells might not be easily recognized in the presence of large quantities of blood and mucus. In this study, the utilization of CB for diagnosis increased the positive rate of lung carcinoma diagnoses and decreased the rate of suspected carcinoma diagnoses. CB can facilitate observation of the tumor cell structure in resected specimens.

Distinguishing AC from SCC is important due to clinical observations; for instance, relative to patients with AC, patients with SCC are at increased risk for hemorrhage when treated with bevacizumab. Moreover, patients with AC respond better to pemetrexed than patients with SCC. The subtyping of NSCLC based on cytological smears is possible for AC and SCC with typical cytomorphologies. For poorly differentiated tumors, ICC should be used as an adjuvant method to determine the correct diagnosis.

According to Rekhtman *et al.* (11), approximately 93% of lung cancer cases can be subtyped into AC and SCC using morphologic criteria alone; in other studies, the diagnosis of AC or SCC can be determined in 50–70% of patients based on cytological specimens (12,13). In our opinion, even experienced cytopathologists may misdiagnose based on cytomorphology alone because lung cancer cells are always poorly differentiated. The cytological features of

SCC depend on tumor grade (14). SCC may be divided into two types: keratinizing SCC and non-keratinizing SCC. However, in most cases, SCC of the lung is not keratinized. In non-keratinized cases, CB and ICC should be used for differential diagnosis. The morphology of AC cell clusters is highly variable. And reactive type II pneumocytes (RPII) undergo hyperplasia and reactive changes in response to injury resulting from various conditions, such as infections, interstitial lung diseases, organizing pneumonia, pulmonary drug toxicity, and tuberculosis (10,15-17). A lepidic pattern of AC may be observed in small biopsy specimens but is difficult to diagnose in cytological material (18,19). In the aforementioned situations, it is challenging for cytopathologists to arrive at correct diagnoses. The most important step in avoiding misdiagnosis is to correlate morphological findings with CB and ICC results.

After performing ICC, we found two cases diagnosed as SCC that were actually AC and two cases diagnosed as AC that were SCC. SCCs are known to be positive for CK5/6 and P40 (*Figure 1*), whereas ACs are positive for TTF-1 and Napsin A (*Figure 2*) (20). In the 2015 WHO classification recommends the use of ancillary techniques such as ICC to decrease the rate of NSCLC-NOS diagnoses. All these ICC markers that we used for CB specimens are regularly used for surgical specimens. In addition to these markers, we also use P63 for SCC, and SPA and CK7 for AC. In CB specimens, the quantity of cells is not sufficient for the analysis of all ICC markers; therefore, we select the most important markers to enable a diagnosis. In our study, only 33 cases were negative for CK5/6, P40, TTF-1 and Napsin A.

In our study, two other cases of AC were misdiagnosed as SCLC. In these cases, tumor cells were as large as 1.5–3 lymphocytes (12,20). Necrosis was observed in the background, and ghost cells were observed in the smears. A diagnosis can be reliably reached based on cytological morphology, but ICC may be required to confirm this diagnosis. TTF-1, SYN, CD56 and CgA are useful indices (*Figure 3*). In a small number of SCLC cases, all ICC findings could be negative (20).

Patients with *ALK* rearrangement comprise 2–5% of all NSCLC cases (21). In our study, we tested all NSCLC cases for *ALK* rearrangement using IHC. The positive rate was 3.73%, which is consistent with reports in the literature. Screening for *ALK* rearrangement using IHC (VENTANA) is convenient and economical. Cytological samples can also be screened for *ALK* rearrangement using CB.

In conclusion, CB and ICC based on CT-guided FNA can provide an accurate diagnosis. Thus, we recommended

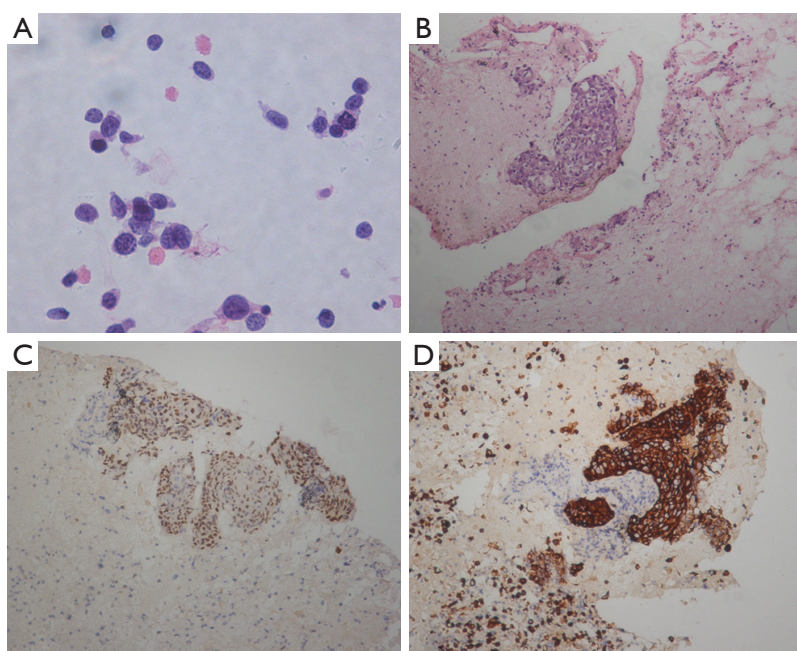


Figure 1 Image of a poorly differentiated SCC. (A) Poorly differentiated NSCLC in TCT (H & E, ×400); (B) cell block section showing a solid growth pattern without evidence of glandular or squamous differentiation (H & E, ×100); (C) tumor cells are positive for P40 (×100); (D) CK5/6 is also positive (×100). SCC, squamous cell carcinoma; NSCLC, non-small-cell lung carcinoma.

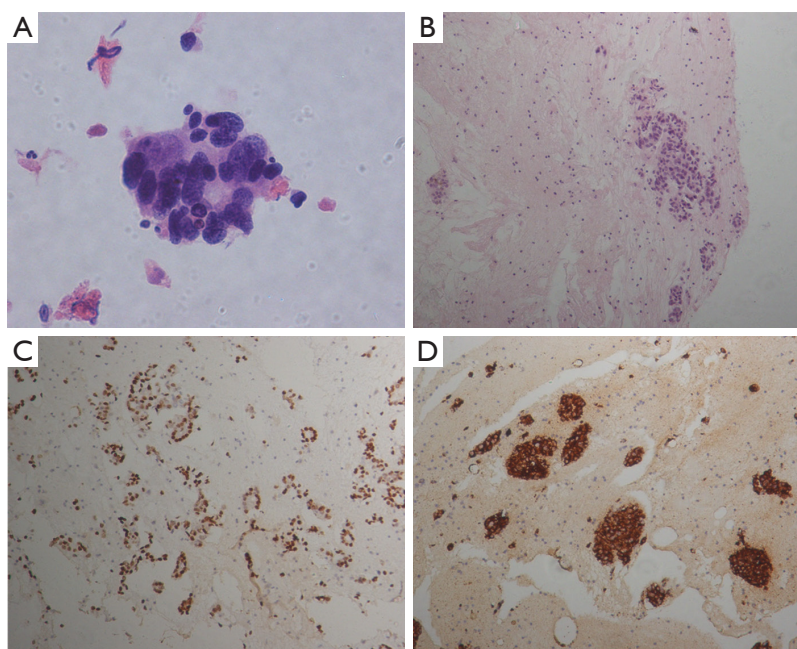


Figure 2 Photograph of a poorly differentiated AC. (A) Poorly differentiated NSCLC in TCT (H & E, ×400); (B) H & E-stained section showing a solid growth pattern without evidence of glandular or squamous differentiation (H & E, ×100); (C) tumor cells are positive for TTF1 (×100); (D) tumor cells are positive for Napsin A (×100). AC, adenocarcinoma; NSCLC, non-small-cell lung carcinoma; TCT, ThinPrep cytology test.

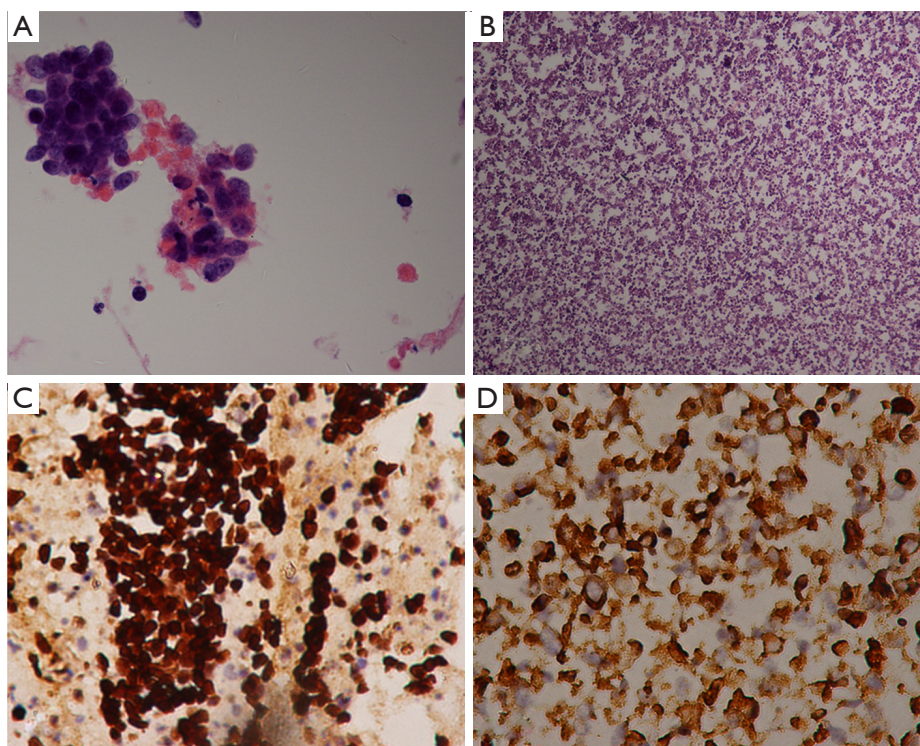


Figure 3 Immunoreactivity for TTF1 and CD56 in small cell lung carcinoma. (A) Poorly differentiated small cell lung carcinoma in TCT (H & E, ×400); (B) SCLC in cell blocks (H & E, ×100); (C) tumor cells are positive for TTF1 (×400); (D) tumor cells are positive for CD56 (×400). SCLC, small cell lung carcinoma; TCT, ThinPrep cytology test.

using CB and ICC to evaluate cytological samples derived from FNA. CB also allows for genetic testing and prognostic assessment via PCR, FISH and IHC (22-25). CB is helpful for diagnosing and subtyping advanced lung cancer, determining precise treatments, and preserving cytological specimens.

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Footnote

Conflicts of Interest: The authors have no conflicts of interest to declare.

Ethical Statement: The study was approved by the ethics committee of Shanghai Pulmonary Hospital (No. K17-125). Written informed consent was obtained from all patients.

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