

Anti-HuC and -HuD autoantibodies are differential sero-diagnostic markers for small cell carcinoma from large cell neuroendocrine carcinoma of the lung

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Abstract. Aiming to identify novel sero-diagnostic markers for neuroendocrine carcinomas of the lung, the two-dimensional gel electrophoresis-immunoblot method was used to analyze tumor-associated autoantibodies in patients with small cell lung carcinoma (SCLC) and large cell neuroendocrine carcinoma (LCNEC). Several autoantigens were revealed and anti-HuC autoantibody was detected only in sera of SCLC patients. Since Hu family proteins including HuC are well-known causes of paraneoplastic encephalomyelitis/sensory neuronopathy (PEM/SN), the expression of HuC as well as HuD mRNAs and their proteins was studied in 11 lung cancer cell lines. The expression of HuC and HuD mRNAs and proteins was only detected in SCLC- and LCNEC-derived cells. To validate the existence of anti-HuC and -HuD auto-antibodies, we studied a large number of sera including those from lung cancer patients employing dot blot analysis. Anti-HuC and -HuD autoantibodies were detected only in SCLC cases with or without PEM/SN, and not in the sera of LCNEC patients. The mechanism leading to different anti-HuC and -HuD autoantibody production between SCLC and LCNEC is unclear; however, the results from the present and previous studies suggest that anti-HuC and -HuD auto-antibodies are novel differential sero-diagnostic markers for SCLC from LCNEC.

Introduction

Lung cancer is the leading cause of cancer-related death worldwide. Based on the presence or absence of cellular

neuroendocrine differentiation, lung cancer can be grouped into non-neuroendocrine or neuroendocrine tumors. The former is roughly equal to the non-small cell lung cancer (NSCLC) largely comprising squamous cell carcinoma (SCC) and adenocarcinoma (AD). The latter ranges from low-grade typical carcinoid (TC), intermediate-grade atypical carcinoid (AC), to high-grade small cell lung carcinoma (SCLC) and large cell neuroendocrine carcinoma (LCNEC) (1,2), and the 5-year survival rates for TC, AC, LCNEC, and SCLC were 87, 56, 27, and 9%, respectively (2). Similar results were also reported by Garcia-Yuste *et al*, and the 5-year survival rates for TC, AC, LCNEC, and SCLC were 96, 72, 21, and 14%, respectively (3). These results show that both SCLC and LCNEC are highly malignant and have a similar poor prognosis. Pro-gastrin-releasing peptide (pro-GRP) is a well-known sero-diagnostic marker for SCLC; however, its positive rate is lower in stage I and II (35-45%) than in stage III (55-70%) and IV (70-80%). At present, no specific sero-diagnostic markers for distinguishing SCLC from LCNEC have been reported.

Autoantibodies are antibodies detected in the sera of patients with various autoimmune diseases. They are also frequently observed in the sera of patients with various malignancies even in the early stages, and, thus, the possibilities for them to be used as potential tumor markers have been suggested (4-10). Hanash (11) has described that harnessing the immune response to identify novel cancer biomarkers is an attractive strategy, because the immune system performs biological amplification which is equivalent to a PCR reaction by generating a detectable signal, with antigenic tumor proteins as templates, beginning at an early stage during tumor development when the tumor may be otherwise undetectable. Many tumor-related autoantibodies have been reported in pulmonary carcinomas (12-15). Thus, an exhaustive search for novel tumor-specific autoantibodies, which may serve as early sero-diagnostic markers for cancers, has commenced.

In this study, we detected tumor-associated autoantibodies by immunoblotting based on two-dimensional gel electrophoresis

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(2-DE) from the sera of patients with pulmonary neuroendocrine carcinomas. Identified antigens were further assessed to confirm their specific expressions in neuroendocrine carcinomas by RT-PCR and immunoblotting. Finally, the usefulness of the autoantigens was validated using the sera of patients with various types of pulmonary carcinoma, along with non-cancerous and healthy controls.

Materials and methods

Cell lines. Six SCLC (N230, N231, H69, H82, Lu130, and N417), two LCNEC (LCN1 and LCN2) (16), two AD (A549 and LC-2/ad), and one SCC (RERF LC-AI) cell line were used in this study. All cell lines were grown in RPMI-1640 (Sigma, Steinheim, Germany) supplemented with 10% fetal bovine serum, 100 U/ml penicillin, and 100 µg/ml streptomycin (Gibco, Auckland, NZ). After being washed with phosphate-buffered saline without bivalent ions (PBS-), harvested cells were separated into two groups, one was fixed in 10% formalin and embedded in paraffin, and the other was stored at -80°C until use for protein and total RNA extraction.

Sera. Sera from 80 pulmonary carcinoma patients (SCLC: 31, LCNEC: 7, AD: 21, and SCC: 21) and 21 non-neoplastic lung disease (interstitial pneumonia: 7, tuberculosis: 5, non-lung cancer diseases: 3, acute inflammation: 1, epithelioid granuloma: 1, cryptogenic organizing pneumonia: 1, nontuberculous mycobacteria: 1, aspergillosis: 1, inflammatory granuloma: 1) patients treated at Kitasato University Hospital. Sera from 26 healthy volunteers were also used as a normal control. All sera were kept at -80°C until use. The SCLCs were further divided into 15 limited and 12 extensive disease cases. This study was approved by the Ethics Committee of Kitasato University School of Medicine. All patients were informed of the aim of the study and gave consent to use their samples.

2D-immunoblotting (IB). Sample preparation and the two-dimensional gel electrophoresis (2-DE) used in this study were described in our previous study (17).

Proteins extracted from the mixture of two LCNECs (LCN1 and LCN2) or the mixture of three SCLCs (N231, H69, and Lu130) were separated by 2-DE. Two pieces of gel were prepared for each sample, one was transferred to a polyvinylidene difluoride (PVDF) membrane (Millipore Corp., Bedford, MA, USA) for immunoblotting and the other was visualized by coomassie brilliant blue R-350 (CBB) staining (PhastGel Blue R, GE Healthcare, Uppsala, Sweden).

Blotting membranes were blocked with 0.05% casein/TBS (0.01 mol/l Tris-HCl, pH 7.5, 150 mmol/l NaCl) for 30 min at room temperature (RT). Membranes prepared with LCNEC proteins were reacted with 100-times diluted pooled sera of five LCNEC patients and the membranes with prepared SCLC proteins were reacted with 100-times diluted pooled sera of five SCLC patients, respectively, for 1 h at RT. The dilution buffer was 0.0025% casein/TBS-T (TBS containing 0.1% Tween-20). Then, the membranes were incubated with 1,000-times diluted horseradish peroxidase (HRP)-conjugated rabbit anti-human IgG polyclonal antibody (Dako, Glostrup, Denmark) for 30 min at RT. Finally, signals were developed by stable DAB solution (Invitrogen, Carlsbad, CA, USA).

Protein identification. In brief, protein spots that reacted with patients' sera were excised from 2-DE gels and destained with 50% (v/v) acetonitrile (ACN)/50 mM NH₄HCO₃, dehydrated with 100% (v/v) ACN, and then dried under vacuum conditions. Tryptic digestion was performed for 24 h at 37°C in a minimum volume of digestion solution containing 0.5 ng/µl sequencing grade modified trypsin (Promega Corp., Madison, WI, USA) and 25 mM Tris-HCl buffer (pH 9.0). After incubation, digested protein fragments eluted in solutions were collected, and gels were washed once in 50% (v/v) ACN/5% trifluoroacetic acid (TFA) and collected in the same tube. Solutions containing digested protein fragments were measured by MALDI-TOF/TOF MS (autoflex-III; Bruker Daltonik GmbH, Bremen, Germany).

Fragment ion spectra from MS and MS/MS were submitted to MASCOT (<http://www.matrixscience.com/>) for a database search and the identification of corresponding proteins employing the following database: IPI human 20091026 (86379 sequences; 34740790 residues, <http://www.ebi.ac.uk/IPI/IPIhuman.html>).

RT-PCR. Total RNAs from the above-mentioned 11 lung cancer cell lines were extracted with Isogen (Nippon Gene, Tokyo, Japan) according to the manufacturer's instructions. Primers were designed with Oligo Primer Analysis Software, version 6.0 (Takara Bio Inc, Otsu, Japan) according to HuC and HuD mRNA sequences (18,19). HuC forward primer: 5'-TGCAAGTTGGTTCGGGACAAG-3' (582-602); reverse primer: 5'-GGCGGATGACTGGTAGAGG-3' (1031-1049). HuD forward primer: 5'-GTCTCTTCGGGAGCATTGGT-3' (415-434); reverse primer: 5'-CCTCTTATCAAAGCGGATGAA-3' (753-773). PCR was performed with pretreatment at 94°C for 2 min and 30 cycles of denaturation at 94°C for 30 sec, annealing at 60°C for 30 sec, and extension at 72°C for 30 sec. Beta-2-microglobulin was used as an internal control. PCR products were electrophoresed on 3% agarose gel and stained with ethidium bromide.

1D-immunoblotting. Proteins were extracted from lung cancer cell lines with detergent lysis buffer (20) using an ultra-sonic homogenizer (UH-50; SMT Co., Tokyo, Japan). Each extracted protein (10 µg) was boiled and separated by 10% sodium dodecyl sulfate-polyacrylamide gel electrophoresis (SDS-PAGE) with a constant current at 15 mA. The immunoblotting methods were generally the same to those used for 2D-IB with some modifications. Transferred membranes were blocked with 0.5% casein/TBS for 30 min at RT, followed by reaction with 200-times diluted HuC- and HuD-positive serum as anti-Hu protein antibody with dilution buffer for 1 h at RT, because purchased anti-HuC antibody did not show specific reactivity. Then, the membranes were incubated with 1,000-times diluted HRP-conjugated rabbit anti-human IgG polyclonal antibody (Dako) for 30 min at RT. Finally, signals were developed using Immobilon Western reagent (Millipore Corp.).

Micro-dot blot array. Anti-HuC and -HuD autoantibodies in sera were detected employing the automatic dot blot system, and the micro-dot blot array with a 256-solid pin configuration (Kakengeneq Co., Ltd., Chiba, Japan) was used. In brief, 1 µl

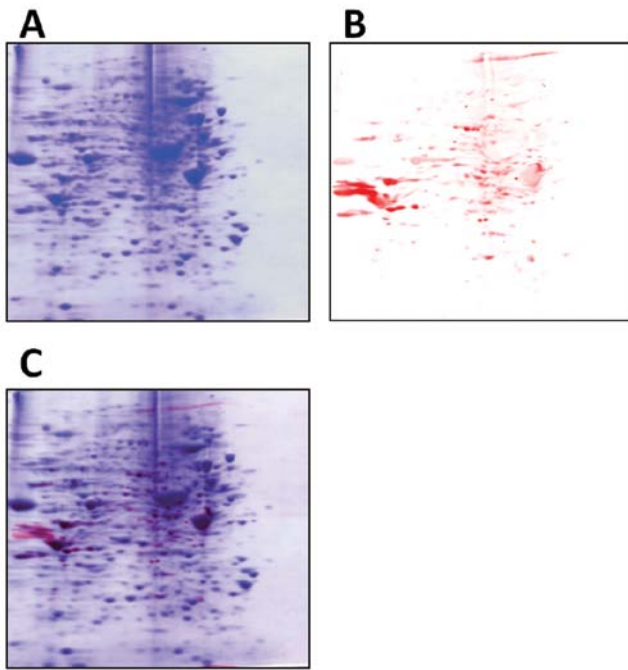


Figure 1. Detection of autoantibodies by 2D-immunoblotting. (A) Protein lysates from SCLC cell lines were separated by 2-DE and stained with CBB. (B) Immunoblot analysis was performed with mixed sera from patients with SCLC as primary antibodies, visualized with DAB solution. (C) A and B were merged.

each of proteins was spotted onto PVDF membranes, which were prepared by a wheat germ cell-free system (21). Then the membranes were blocked with 20% N101 (NOF Corp., Tokyo, Japan)/TBS for 1 h at RT. After being washed in TBS, the membranes were reacted with 100-times diluted sera with 1% N101/TBS for 30 min at RT. After TBS-T washing, the membranes were incubated with 1,000-times diluted HRP-conjugated rabbit anti-human IgG polyclonal antibody for 30 min at RT. Finally, signals were developed with Immobilon Western reagent. The data were analyzed using DotBlotChip-System software ver. 4.0 (Dynacom Co., Ltd., Chiba, Japan). Normalized signals are presented as the positive intensity minus background intensity around the spot. Statistical analysis was performed using the Mann-Whitney U test. The area under the curve (AUC) and best cut-off point were calculated employing receiver operating characteristic (ROC) analysis.

Results

Autoantigen identified by 2D-IB. The immunoreactivity of autoantibodies in sera was assessed by 2D-IB, and the representative positive protein spots on the membrane are shown in Fig. 1. Sixty-two and 63 positive spots were detected with sera from LCNEC and SCLC patients, respectively. In total, 32 proteins for LCNEC and 41 proteins for SCLC were identified as autoantigens. The identified proteins are summarized in Fig. 2. Twenty-three proteins including HuC

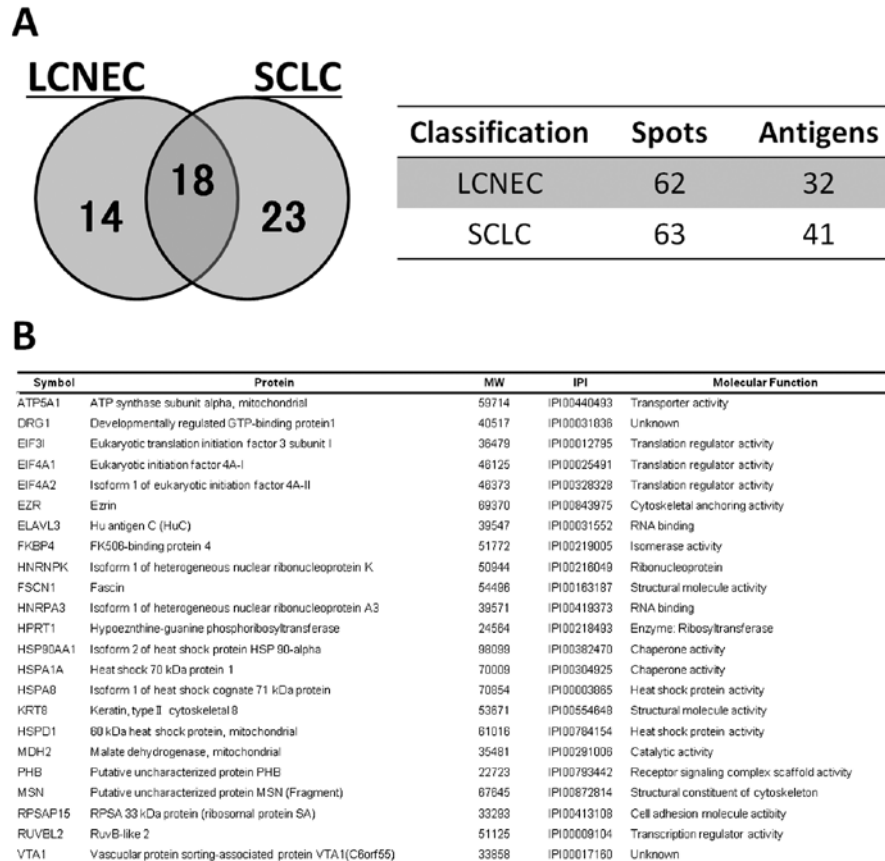


Figure 2. Identified autoantigens in LCNEC and SCLC patients. (A) Compared autoantigens identified in sera from patients with LCNEC or SCLC. Thirty-two and 41 autoantigens were identified from LCNEC and SCLC, respectively. The number of approved autoantigens identified both in LCNEC and SCLC was 18. (B) Twenty-three autoantigens including ELAVL3 (HuC) were identified only in SCLC patients.

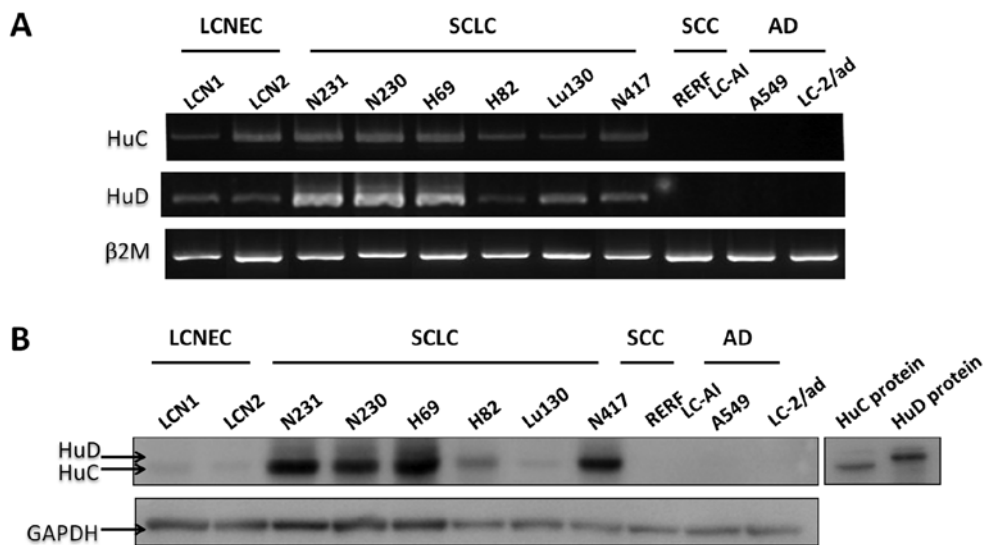


Figure 3. HuC and HuD expressions in lung cancer cell lines. (A) HuC and HuD mRNAs were detected by RT-PCR. Both mRNAs were expressed only in LCNEC and SCLC cell lines. The internal control was beta-2-microglobulin (β 2M). (B) HuC protein levels were detected by immunoblot analysis. HuC and HuD recombinant proteins were used as a positive control and GAPDH was used as an internal control. HuC protein was also expressed only in the majority of SCLC and LCNEC cell lines, and HuD protein was expressed only in a part of SCLC cell lines.

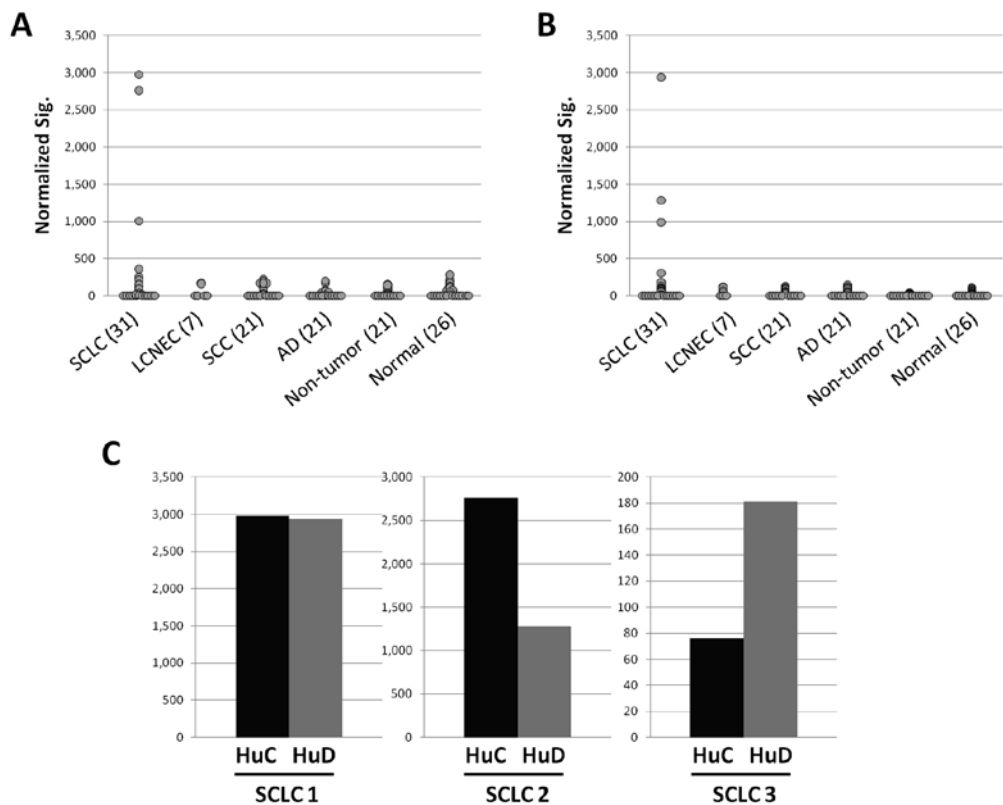


Figure 4. Anti-HuC and -HuD autoantibody levels in sera by dot blot analysis. Anti-HuC autoantibody was detected in 4/31 SCLC patients and not in the others (A). Anti-HuD autoantibody was also detected only in 6/31 SCLC patients (B). The quantitative ratio of anti-HuC and -HuD autoantibodies varied from case to case (C).

were detected only in SCLC, 14 only in LCNEC, and 18 were observed in both.

HuC and HuD expressions in lung cancer cell lines. To examine whether HuC and HuD are specifically expressed in neuroendocrine tumors of the lung, we performed RT-PCR

using 11 lung cancer cell lines (Fig. 3A). The expressions of HuC and HuD mRNAs were detected in all neuroendocrine carcinoma-derived cell lines, but not in SCC or AD cell lines.

To confirm that HuC and HuD proteins were also specifically expressed in neuroendocrine tumors, we performed immunoblot analysis using the same 11 lung cancer cell lines (Fig. 3B).

Both proteins were also detected only in neuroendocrine carcinoma-derived cell lines. HuC protein was detected in 5 of 6 SCLC and the two LCNEC cell lines. HuD protein was also detected in 4 of 6 SCLC, but not in the two LCNEC cell lines. These results were generally in accordance with those of mRNA expression analyses by RT-PCR.

Validation of anti-HuC and -HuD autoantibodies. To confirm the utility of anti-HuC and -HuD autoantibodies as potential biomarkers in neuroendocrine carcinoma of the lung, we investigated their levels in patient and control sera by dot blot analysis. Anti-HuC autoantibody was detected in 4 of 31 SCLCs (Fig. 4A), but not in other lung cancer subtypes including LCNEC, non-neoplastic lesions, or healthy controls ($p=0.003$). At a cut-off point of 360, the sensitivity for SCLC was 12.9% (95% CI: 0.036-0.298). Anti-HuD autoantibody was also detected in 6 of 31 SCLCs (Fig. 4B), but not in the others ($p<0.001$). At a cut-off of 176, the sensitivity for SCLC was 19.6% (95% CI: 0.075-0.375). When compared with the others, both anti-HuC and -HuD autoantibodies showed 100% specificity for SCLC. The AUC-ROC levels were 0.577 and 0.602, respectively.

Anti-HuC and -HuD autoantibodies are detected in SCLC patients with and without PEM/SN. In this study, these autoantibodies were detected in patients without rather than with PEM/SN. Furthermore, the quantitative ratio of anti-HuC and -HuD autoantibodies varied from case to case (Fig. 4C).

Discussion

In this study, 55 autoantigens in total were identified employing 2D-immunoblotting. Our results confirmed the utility of this approach to identify tumor-associated antigens including HuC recognized by autoantibodies in sera from patients with lung cancer (12,13).

Hu proteins are a family consisting of four RNA-binding proteins, three of which are normally expressed in the nervous system (22). All four members have three RNA-interacting domains known as RRM (RNA recognition motif) (23). PEM/SN, which occurs in less than 1% of SCLC patients, is related to high titers of autoantibodies for neuronal Hu proteins (24-26). The mechanism by which the immune system identifies Hu proteins from tumor cells as foreign proteins and generates anti-Hu autoantibodies is still unknown. A few studies have focused on the genetic causes of ectopic neuronal Hu gene (ELAV) expression in neuroendocrine tumors and their roles in the onset and progression of such tumors (27,28). Dalmaou *et al* (29) and Graus *et al* (30) reported that approximately 16% of SCLC patients without PEM/SN have detectable levels of anti-Hu autoantibodies in their sera. In this study, both SCLC patients with and without PEM/SN were detected in 12.9% for anti-HuC and 19.4% for -HuD autoantibodies in their sera. This positive rate was almost the same as those reported previously. In a mouse model study, Kazarian *et al* found that anti-Hu reactivity appeared to arise prior to chemical evidence of cancer in these mice, suggesting the possibility of using anti-Hu for the early detection of SCLC (31). The present results on anti-HuC and -HuD auto-

antibodies supported this possibility, because these autoantibodies were found from limited to extensive diseases. Although the follow-up period was short and only a few controls were used, Tsou *et al* reported that SCLC patients with high levels of anti-Hu reactivity survived for longer than those with low levels ($p=0.08$) (32). In agreement with the study by Tsou *et al* and the present study, Verschuuren *et al* also reported that anti-Hu-positive SCLC cases survived longer, and Dalmau *et al* reported that anti-Hu-positive SCLC patients have relatively limited disease (29,33). Larger scale studies are need to classify the reason for the favorable prognosis of SCLC patients with anti-Hu autoantibodies.

In this study, anti-HuC and -HuD autoantibodies were detected only in SCLC patients, and not in those with other lung cancers, non-neoplastic disease, or healthy controls. Although the quantitative ratio of anti-HuC and -HuD antibodies varied from case to case, 23.3% of SCLC patients were positive for either anti-HuC or anti-HuD or both antibodies. The positive rate may rise using a more sensitive methodology. In spite of this, we detected HuC and HuD mRNAs and proteins in LCNECs, and failed to detect anti-HuC and -HuD autoantibodies in the sera of LCNEC patients, who share many biological features with SCLC patients. Although only a small number of cases were analyzed in a previous study, somatic mutations of the HuD gene were detected in a part of SCLC, TC, and AC cases of different neuroendocrine lung tumors including LCNEC (34). Thus, genetic mutations of HuC and HuD may contribute to the production of autoantibodies.

The present results suggest that anti-Hu autoantibodies are differently expressed between SCLC and LCNEC, and they may be used as novel sero-diagnostic and differential markers for these two tumor types.

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