

# Epstein-Barr Virus-Associated Peripheral T-cell Lymphoma with Gastrointestinal Tract Involvement

WINYOU MITARNUN, MD\*,  
JINTANA PRADUTKANCHANA, MSc\*,  
SATOMI TAKAO, MSc\*\*,

VANNARAT SAECHAN, MSc\*,  
SUPAPORN SUWIWAT, MSc\*,  
TAKAFUMI ISHIDA, DSc\*\*

## Abstract

Peripheral T-cell lymphoma (PTCL) is a group of diseases which are common in Asia and areas of South and Central America. They are highly associated with the Epstein-Barr virus (EBV) infection. In the present study the authors evaluated patients with gastrointestinal involvement of PTCL with respect to clinical findings and outcome, pathologic features, and molecular analysis for EBV infection and the clonality of tumor cells. From January 1997 through December 2000, 7 patients with gastrointestinal tract involvement of PTCL were identified. The frequency of gastrointestinal tract involvement in the various types of PTCL was 5.4 per cent (7 of 129 cases). The pertinent clinical features were prolonged fever, weight loss, anemia, hepatosplenomegaly, lymphadenopathy, multiorgan involvement, and gastrointestinal bleeding. Laboratory results showed a significantly high serum level of alkaline phosphatase and lactate dehydrogenase, and abnormal coagulograms. Five patients died within 4 months after onset of illness, while two were in complete remission after chemotherapy. The tumor cell morphology was classified into three categories: small-sized cells, mixed medium- and large-sized cells, and large-sized cells. The antigenic phenotypes of the tumor cells were LCA+, CD3+, CD15-, CD16-, CD30-, CD45R0+, CD57-, CD68-, EMA-,  $\beta$ F1-, granzyme B+, TIA-1+, and p53+. The expression of CD4, CD8, CD56 and CD20 was variable. EBV-RNA expression by *in situ* hybridization (EBER-ISH) study was positive and T-cell receptor (TCR) beta and/or gamma gene rearrangements were detected in all patients. DNA sequence analysis showed high identity to the human TCR germline gene. PTCL with gastrointestinal tract involvement was associated with EBV infection. The tumor cells were mature T cells with some NK-cell antigenic expression and all demonstrated TCR gene rearrangements.

**Key word :** Peripheral T-cell Lymphoma, Epstein-Barr Virus, Gastrointestinal, T-cell Receptor Gene, Clonality

MITARNUN W, SAECHAN V, PRADUTKANCHANA J,  
SUWIWAT S, TAKAO S, ISHIDA T  
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\* Department of Pathology, Faculty of Medicine, Prince of Songkla University, Songkhla 90110, Thailand.

\*\* Department of Biological Sciences, Graduate School of Science, The University of Tokyo, Tokyo, Japan.

Peripheral T-cell proliferative diseases/lymphomas are more commonly found in Asia than in Western countries<sup>(1-4)</sup>. They also have been reported with significant frequency in Central and South America<sup>(5-7)</sup>. This suggests a racial and/or environmental connection for the disease. Peripheral T-cell proliferative disease/lymphoma exhibits considerable heterogeneity in the clinical findings, morphologic pattern, immunology, and prognosis<sup>(4,8-11)</sup>. In 1993, the International Lymphoma Study Group proposed a classification for lymphoid neoplasms, entitled, A Revised European-American Classification of Lymphoid Neoplasms, (REAL classification), by using morphologic, immunologic, and genetic techniques<sup>(12,13)</sup>. Recently, the World Health Organization (WHO) classification was proposed<sup>(14)</sup>. The mature NK/T-cell neoplasm classification in the proposed WHO scheme was largely adapted from the REAL classification. The WHO classification for predominantly extranodal malignant neoplasms was separated into four prototypes: extranodal NK/T-cell lymphoma, nasal type, enteropathy-type intestinal T-cell lymphoma, subcutaneous panniculitis-like T-cell lymphoma, and hepatosplenic  $\gamma\delta$  T-cell lymphoma<sup>(15)</sup>. The involvement of peripheral T-cell lymphomas in the gastrointestinal tract may be found in three categories: enteropathy-type intestinal T-cell lymphoma, extranodal NK/T-cell lymphoma, nasal type, and gastrointestinal  $\gamma\delta$  T-cell lymphoma<sup>(15-17)</sup>. Of these, only the enteropathy-type intestinal T-cell lymphoma is considered to be a primary gastrointestinal tract T-cell lymphoma<sup>(14)</sup>.

Gastrointestinal lymphoma accounts for 4 per cent to 20 per cent of all non-Hodgkin's lymphomas and is the most extranodal site of presentation<sup>(18)</sup>. The stomach is the major organ involved in gastrointestinal lymphoma. B-cell lymphoma of mucosal-associated lymphoid tissue (MALT) type is the commonest, and gastric MALT lymphoma may be associated with *Helicobacter pylori* and may undergo complete remission following eradication of *H. pylori* (19). In Western countries, primary gastrointestinal lymphomas account for approximately 10 per cent to 15 per cent of non-Hodgkin's lymphoma cases, and the enteropathy-type intestinal T-cell lymphoma is the most common type<sup>(19,20)</sup>.

Almost all gastrointestinal peripheral T-cell lymphoma cases which were reported from east Asia (Korea, Japan, China), Morocco, and Mexico were positive for Epstein-Barr virus (EBV) RNA by *in situ* hybridization study in most of the tumor cells<sup>(7,21-</sup>

25)

Cases of gastrointestinal T-cell lymphoma which were reported from Western Europe and the United States of America showed only 16 per cent positive for EBV RNA in the tumor cells<sup>(7,26-31)</sup>. The majority of patients in the later series were cases with enteropathy-type intestinal T-cell lymphoma.

Enteropathy-type intestinal T-cell lymphoma is a specific peripheral T-cell lymphoma that occurs in association with celiac disease<sup>(27,32,33)</sup>. Celiac disease is relatively common in Europe. It occurs worldwide, but is quite rare in Asians and black Africans<sup>(33,34)</sup>. In one study, about 7 per cent of patients with celiac disease developed enteropathy-type intestinal T-cell lymphoma in a 25-year follow-up<sup>(27)</sup>. The enteropathy-type intestinal T-cell lymphoma, to our knowledge, has not been reported from Asia.

In the present study, the authors report 7 patients with gastrointestinal tract involvement of peripheral T-cell lymphoma. All cases were studied with respect to clinical findings and outcome, histopathologic and immunohistologic features, search for an association with EBV infection, and documentation of the clonality of the neoplastic T cells.

## MATERIAL AND METHOD

The prospective study was conducted in Songklanagarind University Hospital, Songkhla, Thailand, from January 1997 through December 2000 and follow-up continued until December 2001. There were 129 patients with various types of peripheral T-cell proliferative disease/lymphoma. Seven cases were found to have gastrointestinal tract lesions, and all of them were confirmed by histologic examination.

## Immunohistochemistry

Immunohistochemical stainings for leukocyte common antigen (LCA), CD3, CD4, CD8, CD15, CD16, CD20, CD30, CD45RO, CD56, CD57, CD68, T-cell receptor  $\alpha\beta$  ( $\beta$ F1), epithelial membrane antigen (EMA), granzyme B, T-cell intracellular antigen-1 (TIA-1), and p53 protein of tumor tissues were performed on formalin-fixed, paraffin embedded tissue using the following antibodies: monoclonal mouse anti-human LCA (CD45) (Zymed, CA, USA, 1 : 100), monoclonal mouse anti-human CD3 (Novocastra, UK, 1 : 120), monoclonal mouse anti-human CD4 (Novocastra, 1 : 25), monoclonal mouse anti-human CD8 (Dako, Denmark, 1 : 20), monoclonal rabbit anti-human CD15 (Zymed, 1 : 50), monoclonal mouse anti-human CD16 (Novocastra, 1 : 30), monoclonal mouse anti-human CD20 (Dako, 1 : 70), monoclonal mouse anti-

human CD 30 (Dako, 1 : 20), monoclonal mouse anti-human CD45RO (Zymed, 1 : 70), monoclonal mouse anti-human CD56 (Novocastra, 1 : 50); neuroblastoma was used as a positive control, monoclonal mouse anti-human CD57 (Novocastra, 1 : 50), monoclonal mouse anti-human CD68 (Dako, 1 : 100), monoclonal mouse anti-human  $\beta$ 1 (Endogen, MA, USA, 1 : 10), monoclonal mouse anti-human EMA (Dako, 1 : 20), monoclonal mouse anti-human granzyme B (Novocastra, 1 : 60), monoclonal mouse anti-human TIA-1 (Coulter, France, 1 : 30), p53 protein (Dako, 1 : 250), rabbit anti-mouse immunoglobulin (Dako, 1 : 200), swine anti-rabbit immunoglobulin (Dako 1 : 25), and horseradish peroxidase-mouse antiperoxidase (Dako, 1 : 100). The staining procedure was performed as previously described(35). A tumor was considered p53 positive when more than 50 per cent of the tumor cell nuclei stained intensely with anti-p53.

### *In situ* hybridization

An *in situ* hybridization (ISH) study for the Epstein-Barr virus genomes was performed on formalin-fixed, paraffin embedded tissue using the fluorescein-conjugated EBV (EBER) oligonucleotides complementary to nuclear RNA portions of the EBER genes that were actively transcribed in latently infected cells. Briefly, tissue sections of 5 microns were deparaffinized with xylene, rehydrated in graded water-ethanol solutions, and digested with proteinase K (3 mg/L in 0.05 M Tris/HCl, pH 7.6) for 30 minutes at 37°C. After dehydration and air-drying, the fluorescein-conjugated (FITC) EBER oligonucleotide probes (Y 0017, Dako) were applied to the sections for 2 hours at 37°C. The following immunohistochemical detection system (K 046, Dako) was used: rabbit F(ab') anti-FITC/AP for 30 minutes and a solution containing 5-bromo-4-chloro-3-indolylphosphate (BCIP) and nitroblue tetrazolium (NBT) for 30-60 minutes. Then the slides were washed in running tap water and mounted in glycergel(36). Appropriate positive and negative controls were run. ISH for EBV RNA was interpreted as positive when the positively stained tumor nuclei were more than 10 per cent of the tumor cell population. The amount of positive cells between 10-25 per cent was considered +1; 26-50 per cent, +2; 51-75 per cent, +3; and more than 75 per cent was considered +4.

### DNA extraction from tissue for PCR analysis

DNA extraction was performed using QIAamp DNA Mini Kit (QIAGEN, Germany). Three to five

pieces of tissue section (3 to 5 microns) from paraffin blocks were taken, deparaffinized, resuspended in 180  $\mu$ l of buffer ATL, digested with 50  $\mu$ l of proteinase K, mixed by vortexing, and incubated at 55°C until the tissue was completely lysed. Further steps were followed as directed by the manufacturer's manual of tissue protocol(37). The presence of DNA was confirmed by agarose-gel electrophoresis or by amplification of  $\beta$ -globin gene.

### Polymerase chain reaction (PCR) analysis

For the clonality study of the tumor cells, a T-cell receptor (TCR)- $\beta$  chain and TCR- $\gamma$  chain genes were used. Amplification of the TCR- $\beta$  chain gene was carried out using the technique described by McCarthy *et al*(38). The authors used three separate reactions with the following oligonucleotide primers combinations: D2+J $\beta$ 2.3, D2+J $\beta$ 2.6, and D2+J $\beta$ 2.7. For the monoclonal rearrangement of tumor cells, one or two clearly visible band/s with the expected size range (55 to 100 bp) were identified. Amplification of the TCR- $\gamma$  chain gene was performed using the PCR technique as described by McCarthy *et al*(39,40). The authors used two separate reactions with the following oligonucleotide primers combinations: V $\gamma$ I+V $\gamma$ III/IV+J $\gamma$ 1/2 (product sizes 70-95 bp), and V $\gamma$ II+J $\gamma$ 1/2 (product sizes approximately 150-180 bp). For the monoclonal rearrangement, one or two clearly visible band/s were identified. DNA extracted from Raji and U8312 (a human T-cell lymphotropic virus type-1 positive cell line) were used as controls.

### DNA sequence analysis of TCR genes

The purified PCR products of TCR- $\beta$  chain and TCR- $\gamma$  chain genes were used as a template for automated sequencing analysis by using a ABI PRISM Dye Terminator Cycle Sequencing kit with AmpliTaq DNA polymerase, FS (ABI PRISM™ Ready Reaction, Applied Biosystems Inc., CA, USA). The purified template (approximately 500 ng) was mixed with 8  $\mu$ l of Terminator premix, 3.2 pmole of a sequencing primer and deionized water to a total volume of 20  $\mu$ l. The mixture was overlaid with one drop of mineral oil and placed in the thermal cycler preheated to 96°C. Thermal cycling was performed on a Perkin-Elmer Cetus thermal cycler model 480 under the following conditions: 96°C denaturation for 15 sec, 60°C annealing for 15 sec and 60°C extension for 4 min with 25 total cycles. After the last cycle, the extension product was precipitated and redissolved in 4  $\mu$ l of loading buffer containing 5 : 1 (vol/vol) of deionized forma-

mide with 50 mM EDTA, pH 8.0 and dextrane blue. The reaction tube was heated at 90°C for 2 min before loading onto a 6 per cent polyacrylamide sequencing gel for analysis in the Applied Biosystems Automated DNA sequencer, model 373A. The sequence obtained from each run was analysed with the assistance of SeqEd, a software program (Applied Biosystems Inc.) and compared with the sequence in Genbank using the Blast program.

## RESULTS

### Clinical features

Clinical data, pathologic findings, therapeutic interventions and outcomes of the patients are summarized in Table 1, and the pertinent laboratory findings are summarized in Table 2. Of the 129 patients with various types of peripheral T-cell proliferative disease/lymphoma, seven (5.4%) patients were found to have gastrointestinal tract lesions (4 males and 3 females, ranging in age from 23-71 years). All of these seven patients showed a prolonged fever which had lasted from one to six months. Five patients had hepatosplenomegaly; of these, four (case nos. 2, 3, 4, 5) were histologically confirmed to have peripheral T-cell lymphoma involving the liver. Three patients had peripheral T-cell lymphoma in the lymph nodes. Bone marrow involvement was found in three patients (case nos. 2, 4, 5). Case no. 7 had an extranodal NK/T-cell lymphoma, nasal type, in the nasopharynx two years before the gastrointestinal symptoms developed. Case no. 3 had peripheral T-cell lymphoma at the base of the tongue. An autopsy from case no. 5 revealed multiorgan involvement of peripheral T-cell lymphoma which included the liver, spleen, brain, and both adrenal glands.

The hemograms revealed anemia in all cases. The number of white blood cells and platelets varied. Most of these patients had high serum levels of alkaline phosphatase and lactate dehydrogenase, and abnormal prolonged coagulograms.

Of three patients who received multidrug chemotherapy, two (case nos. 2, 6) achieved a complete remission in the 46-month and 42-month follow-up; and one of them (case no. 7) died. In four patients who did not have chemotherapy, the disease followed a rapidly progressive course and all of them died within one to four months after the first onset of illness.

### Pathological features

#### *Gross pathology*

The major tumor sites in the gastrointestinal tract were the ileocaecal region and colon. Lesions in the stomach, duodenum, and jejunum were also found in some cases. In most cases, multicentric lesions involving different areas of infiltrative and ulcerative lesions were the major feature. However, two patients (case nos. 1, 6) had a single lesion. Perforation of the bowel was seen in cases 1 and 3.

#### *Histopathology*

There was a wide spectrum of cytologic morphology of tumor cells ranging from small-sized to large atypical and hyperchromatic cells. There was also an admixture of these types of cells. Some may be difficult to distinguish from an inflammatory process. Infiltrations with eosinophils, monocytes, and plasma cells were variable. The authors classified the tumor morphology into three categories: small-sized cells, mixed medium- and large-sized cells, and large-sized cells.

Small-sized cells (Fig. 1): the infiltrate was composed of small round lymphocytes, but there was no marginal zone pattern. Necrosis or angiocentric pattern were rarely observed. Infiltration with eosinophils and plasma cells varied from mild to moderate. The lesions were confined mainly to the mucosa and upper part of the muscular wall.

Mixed medium- and large-sized cells (Fig. 2): the infiltrate was composed of medium-sized cells, large atypical hyperchromatic cells, and a few small-sized cells. An angiocentric pattern and coagulative necrosis were prominent features. Infiltration with eosinophils, monocytes and plasma cells varied from mild to massive. The lesions were located mainly in the mucosa and occasionally extended to the muscular wall and adventitia.

Large-sized cells (Fig. 3): The infiltrate was mostly composed of large-size cells. The cells were pleomorphic, and had hyperchromatic nuclei and abundant mitoses. An angiocentric pattern and coagulative necrosis were very prominent features. Apoptotic cells were seen in large numbers. Infiltration of monocytes with massive hemophagocytosis was observed in the tumor areas, and in the lumen of nearby vessels. Infiltration with eosinophils and plasma varied from minimal to absent.

Table 1. Clinical and histologic findings.

Patient no.	Age (yr)/ sex	Presenting symptom and sign	Operation site	Gross and size	Histologic type	Histologic finding in other site	Post-operative therapy	Follow-up/ outcome
1	28/M	High-grade fever 2 mos, hepatosplenomegaly, lymphadenopathy, ascites, bowel perforation	Terminal ileum and right half colon resection	Bulky tumor at ileum and caecum 3 x 4 x 8 cm, caecal perforation	SC	PTCL in lymph node	NA	DOD, 4 mos
2	23/F	Low-grade fever and bloody diarrhea 1 mo, hepatosplenomegaly	Duodenum and ileum biopsy	Edema of small bowel mucosa	SC	PTCL in liver RH in lymph node	CHOP	CR, 46 mos
3	71/F	High-grade fever 1 mo, weight loss 10 kg, bloody diarrhea, hepatosplenomegaly	Terminal ileum and right half colon resection	Multiple ulcers at ileum upto 5 x 2 cm, ileal perforation	SC	PTCL in liver AILD in lymph node PTCL of the tongue	NA	DOD, 2 mos
4	28/M	High-grade fever 1 mo, bloody diarrhea 1 week, hepatosplenomegaly	Jejunum, ileum and right half colon resection	Multiple infiltrative lesions and ulcers (up to 3 cm) in jejunum, ileum and colon	MLC	PTCL in liver	NA	DOD, 2 mos
5	27/M	High-grade fever 1 mo, weight loss 5 kg, bloody diarrhea, hepatosplenomegaly	Terminal ileum and total colon resection	Multiple small infiltrative lesions and ulcers in entire colon	MLC	Autopsy showed PTCL in stomach and small bowel, liver lymph nodes, spleen, bone marrow, brain and adrenal glands	NA	DOD, 1 mo
6	34/F	Fever and abdominal pain 6 mos, weight loss 7 kg	Duodenum and portion of jejunum resection	An ulcer at duodenum, 5 x 3 cm	LC	-	CHOP	CR, 42 mos
7	51/M	High-grade fever 1 mo, maculopapular rash and skin ulcers on body and face, status post radiation therapy for NK/T in nasopharynx 2 yrs ago, bloody diarrhea 1 day	Left half colectomy, biopsy from duodenum and jejunum	Multiple small infiltrative lesions and ulcers (0.7-1.2 cm) in colon	LC	Cutaneous T-cell lymphoma RH in lymph node	Vincristine, Endoxan	DOD, 2 mos

Abbreviations: DOD = dead of disease, CR = complete remission after chemotherapy, NA = not applicable, SC = small-sized cell, MLC = mixed medium- and large-sized cells, LC = large-sized cell, PTCL = peripheral T-cell lymphoma, RH = reactive hyperplasia, AILD = angioimmunoblastic T-cell lymphoma, NK/T = extranodal NK/T-cell lymphoma, nasal type.

Table 2. Laboratory findings.

Test	Case no.						
	1	2	3	4	5	6	7
Hemoglobin, g/L	85	82	83	103	81	81	88
White blood cell count, $\times 10^9/L$	2.5	24.6	1.2	9.5	5.7	4.7	10.5
Platelets, $\times 10^9/L$	167	574	262	26	66	305	330
Total protein, g/L	45	48	50	30	46	61	57
Albumin, g/L	21	24	29	19	13	30	33
Total bilirubin, $\mu\text{mol}/L$	23.8	8.5	22.1	251.7	132.6	6.8	10.8
Conjugated bilirubin, $\mu\text{mol}/L$	19.2	3.4	13.7	161.5	107.2	5.4	3.4
Aspartate aminotransferase, U/L (0-35)	68	46	54	1970	15	15	38
Alanine aminotransferase, U/L (0-35)	27	43	27	161	14	9	35
Alkaline phosphatase, U/L (30-120)	287	218	235	266	1003	80	55
Lactate dehydrogenase, U/L (230-460)	512	280	504	ND	356	491	ND
Activated PTT, s (28-31)	69	34	ND	64.6	54	26.5	37.6
Prothrombin time, s (13-15)	25.5	13	ND	21.3	38.5	13.3	16.7
Anti HIV (by ELISA)	ND	neg	neg	neg	neg	neg	neg

Abbreviations : ND = not done, neg = negative, PTT = partial thromboplastin time.



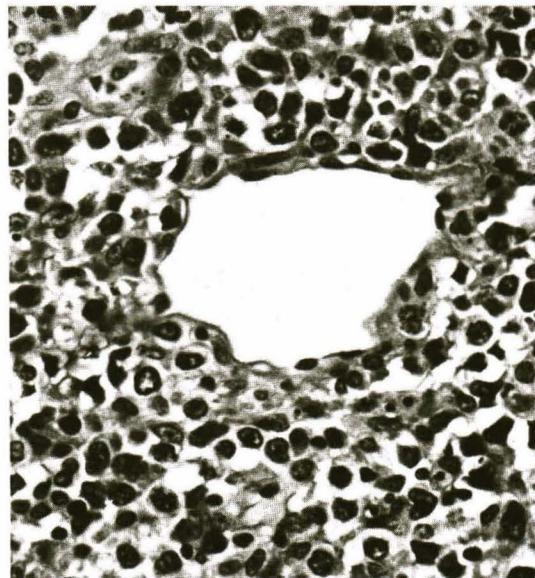
Fig. 1. Small-sized tumor cells. The majority of tumor cells are small with round to only slightly irregular nuclei and have a homogeneous pattern.



Fig. 2. Mixed medium- and large-sized tumor cells. The tumor cells are medium-sized and large-sized cells with atypical hyperchromatic nuclei. Note the angiocentric pattern.

Cases 1, 2 and 3 were classified as small-sized cells, cases 4 and 5 as mixed medium- and large-sized cells, and cases 6 and 7 as large-sized cells. In the group of small-sized cells, none showed an angio-

centric pattern and only case 1 showed coagulative necrosis. An angiocentric pattern and coagulative necrosis were found in all patients with the histologic types of mixed medium- and large-sized cells, and



**Fig. 3.** Large-sized tumor cells. Angiocentric pattern is a prominent feature. The tumor cells are pleomorphic, and have hyperchromatic nuclei and abundant mitoses.

large-sized cells. The adjacent mucosal tissue and/or the uninvolved small bowel mucosa in all cases revealed no evidence of celiac disease.

#### Immunohistochemistry

Results of the immunohistochemistry are summarized in Table 3. The phenotypes of tumor cells in these cases were LCA+, CD3+ (membrane pattern), CD15-, CD16-, CD30-, CD45RO+, CD57-, CD68-,  $\beta$ FI-, EMA-, granzyme B+, TIA-1+, and p53+. The expression of CD4, CD8, CD56 and CD20 was variable. One case expressed CD4, three cases expressed CD8, and one case expressed CD56. Case no. 4 expressed both CD20 and CD3, which are the markers of B cell and T cell, respectively.

#### *In situ* hybridization

The presence of EBV RNA (EBER) in tumor cells was detected in all cases, ranging from +2 to +4 (Table 3). Tumors of the mixed medium- and large-sized cells, and the large-sized cells morphologies, showed a strongly positive reaction.

#### Clonality study

Results of the clonality studies of these tumor cells are shown in Fig. 4 and 5. By using three sets of

**Table 3. Immunophenotypes and EBV RNA (EBER) of tumor cells in 7 patients.**

Case no.	Phenotypes										p53	EBV RNA
	LCA	CD3	CD4	CD8	CD15	CD20	CD30	CD45RO	CD56	CD57		
1	-	-	-	-	-	-	-	-	-	-	+	+2
2	+	-	-	-	-	-	-	ND	-	-	ND	+2
3	+	+	-	-	-	-	-	-	-	-	+	+3
4	+	+	+	+	+	+	+	-	-	-	+	+4
5	+	+	+	+	+	+	+	-	-	-	+	+4
6	+	+	+	+	+	+	+	-	-	-	+	+4
7	+	+	+	+	+	+	+	-	-	-	+	+4

Abbreviations : LCA = leukocyte common antigen, GrB = granzyme B, ND = not done.

oligonucleotide primers for TCR- $\beta$  gene, the monoclonal band was detected in four cases; case nos. 1, 2, 3 and 6 (Fig. 4). For the TCR- $\gamma$  gene, the monoclonal bands were detected in four cases; case nos. 4, 5, 6 and 7 (Fig. 5).

#### TCR sequence analysis

Results of the TCR gene- sequence analysis (sense strand in case nos. 1, 2, 3 and anti-sense strand in case nos. 4, 5, 6, 7) are summarized in Table 4. All of these cases confirmed the monoclonality. PCR products had a very high identity to the human TCR germline gene.

#### DISCUSSION

Peripheral T-cell proliferative disease/lymphoma has been recognized as a common group of diseases in Southern Thailand(4,41-43). In our institute, a 700-bed hospital, the authors found 129 cases in a 4-year prospective study. Of these, 7 had gastrointestinal tract involvement. The patients presented with prolonged fever, weight loss, anemia, hepatosplenomegaly, abnormal prolonged coagulograms, and significantly high serum levels of alkaline phosphatase

and lactate dehydrogenase. All four patients with liver involvement of peripheral T-cell lymphoma showed high serum levels of alkaline phosphatase. The pertinent gastrointestinal symptom was bloody diarrhea, and some patients had bowel perforation.

Gastrointestinal non-Hodgkin's lymphoma accounts for 4 per cent to 20 per cent of all non-Hodgkin's lymphoma cases, and is the most common extranodal site of presentation(18). Intestinal B-cell lymphomas appear as polypoid masses or annular lesion in the distal and terminal ileum(20). Enteropathy-type intestinal T-cell lymphomas in Western countries are mainly found in the proximal small intestine, in contrast to intestinal T-cell lymphomas in Asia and in the presented study, which tended to be located in the terminal ileum and the large intestine(20,21).

Extranodal peripheral T-cells and NK-cell neoplasms are classified into 4 major categories: nasal, intestinal, subcutaneous panniculitis-like, and hepatosplenic  $\gamma\delta$ (15). EBV-positive extranodal NK/T-cell lymphoma, nasal type, which frequently involves the intestine as a secondary site, is almost always CD56+ (70% to 100%), CD3 (membrane)-, CD3 (cytoplasm)+,



Fig. 4. PCR T-cell receptor-beta gene amplification using three pairs of primers: D2+J $\beta$ 2.3 (a), D2+J $\beta$ 2.6 (b), and D2+J $\beta$ 2.7 (c). Lanes 1 through 7 are cases 1 to 7 respectively. Lane R is Raji control, Lane N is negative control, M is ladder markers. The monoclonal band is detected in cases 1, 2, 3, and 6, (10% acrylamide gel).

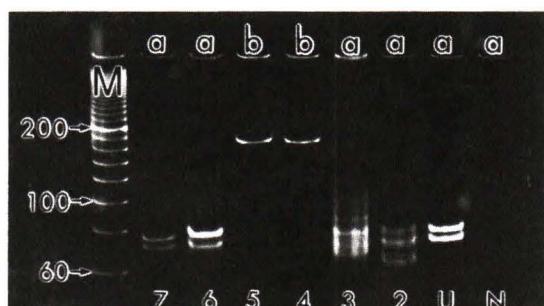


Fig. 5. PCR T-cell receptor-gamma gene amplification using two sets of primers: V $\gamma$ 1+V $\gamma$ III/IV+J $\gamma$ 1/2 (a), and V $\gamma$ II+J $\gamma$ 1/2 (b). Lanes 2 through 7 are cases 2 to 7 respectively. Lane U is U8312 positive control, Lane N is negative control, M is ladder markers. The monoclonal band is detected in cases 4, 5, 6, and 7, (case no. 1 not amplified, 10% acrylamide gel).

Table 4. Results of TCR gene-sequence analysis.

Case	Primers	PCR product (bp)	Part of sequencing product (bp)	Homology site of TCR germline gene	Identity bp	%
1	V $\gamma$ 1 + V $\gamma$ III/IV + J $\gamma$ 1/2	70-95	43	gi 5566238 gb AF 159056.1 (V $\gamma$ 1 regions)	34/34	100
2	D2 + J $\beta$ 2.3	80-90	59	gi 1197256 emb x86129.1 (71-116)	46/46	100
3	D2 + J $\beta$ 2.6	80-90	68	gi 3388552 gb M 14159.1 (2099-2148)	46/50	92
4	V $\gamma$ II + J $\gamma$ 1/2	160-180	140	gi 5566238 gb AF 159056.1 (56793-56911)	117/119	98
5	V $\gamma$ II + J $\gamma$ 1/2	160-180	141	gi 5566238 gb AF 159056.1 (56793-56924)	131/132	99
6 (upper band)	V $\gamma$ 1 + V $\gamma$ III/IV + J $\gamma$ 1/2	80-90	62	gi 3047019 gb AF 057177.1 (V $\gamma$ 1 regions)	30/30	100
6 (lower band)	V $\gamma$ 1 + V $\gamma$ III/IV + J $\gamma$ 1/2	70-80	53	gi 3047019 gb AF 057177.1 (V $\gamma$ 1 regions)	29/29	100
7	V $\gamma$ 1 + V $\gamma$ III/IV + J $\gamma$ 1/2	70-95	43	gi 5566238 gb AF 159056.1 (V $\gamma$ 1 regions)	19/19	100

high prevalence of p53 overexpression, and lacks features associated with the clonal rearrangement of TCR- $\beta$  and TCR- $\gamma$ . CD16 expression varies from 10 per cent to 60 per cent(15,44-50). Based on the morphology, the disease is characterized by a broad cytologic spectrum, ranging from cytologically benign, which cannot be distinguished from an inflammatory process, to the definite malignant morphology(16). An angiocentric pattern on histologic examination is a common feature, and necrosis is seen in most cases(15).

In gastrointestinal tract involvement of peripheral T-cell lymphomas, granzyme B-positive cells are mainly associated with enteropathy-type intestinal T-cell lymphoma. A total of 77 per cent of enteropathy-type intestinal T-cell lymphoma cases *versus* 14 per cent of the non-enteropathy-type intestinal T-cell lymphoma cases are granzyme B positive in the tumor cells(49). Although all the six cases studied were positive for granzyme B and TIA-1, the clinical history and the adjacent mucosal tissue and/or the uninvolved small intestine mucosa in all cases failed to reveal evidence of celiac disease, which is the preceding lesion of the enteropathy-type intestinal T-cell lymphoma(32,33).

The antigenic phenotype of this reported series showed T-cell markers with some NK-cell antigenic expression. The NK-cell-associated marker CD56, which has been considered as a universally positive marker in the extranodal NK/T-cell lymphoma, nasal type, was negative in 5 of 6 cases in our series. The neuroblastoma positive control and the internal positive control (neurons in tissue section) gave a strongly positive reaction. The tumor cells showed no expression of CD16, whereas the activated macrophages which infiltrated in the tumor area showed intense membrane staining. The intracellular cytotoxic molecules (granzyme B, TIA-1), and the p53 expression, were detected in all tested cases. In the present study, the cases were all positive for the EBV genome, similar to previous reports of extranodal NK/T-cell lymphoma, nasal type, but there was evidence against the diagnosis in that they all demonstrated T-cell receptor gene rearrangements, and a majority of cases failed to express CD56. The findings suggest that these cases were not extranodal NK/T-cell lymphoma, nasal type, even though case no. 1 had a preceding extranodal NK/T-cell lymphoma, nasal type, in the nasopharynx. Another important finding was the expression of CD20 in case no. 4. This concurred with our previous report on a high-grade hepatic peri-

pheral T-cell lymphoma which showed expression of both CD45R0 and CD20 in 3 of 8 cases<sup>(4)</sup>, and a recent case report of peripheral T-cell lymphoma with aberrant expression of CD20 and CD79a<sup>(51)</sup>.

All of the presented seven cases demonstrated either TCR- $\beta$  or TCR- $\gamma$  gene rearrangements. Cases 2 and 3 had only the rearrangement of the TCR- $\beta$  locus, and cases 4, 5 and 7 had only the rearrangement of the TCR- $\gamma$  locus. This finding was slightly different from a previous report of peripheral T-cell lymphomas where the TCR- $\beta$  locus was found to be rearranged in 78 per cent of cases *versus* 96 per cent of the TCR- $\gamma$  locus<sup>(52)</sup>. The TCR sequence analysis in all cases confirmed the monoclonality and had a

high identity to the human TCR germline gene. This indicated the neoplastic nature instead of an inflammatory process of peripheral (mature) T cell in the presented cases.

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## มะเร็งต่อมน้ำเหลืองชนิด เพอริเฟอร์อล ที-เซลล์ ที่มีความสัมพันธ์กับภาวะติดเชื้อไวรัสเอปสไตน์-บาร์ ในระบบทางเดินอาหาร

วิญญา มิตราనันท์, พบ\*, วรรณรัตน์ แซ่ชั้น, วทม\*, จินตนา ประดุจกาญจน์, วทม\*, สุภาพร สุวิวัฒน์, วทม\*, ชาโตมิ ทากาโอะ, วทม\*\*, ทากาพูมิ อิชิตะ, ปรด\*\*

Peripheral T-cell lymphoma (PTCL) เป็นกลุ่มโรคที่พบมากในเอเชียและกลุ่มประเทศในอเมริกาใต้ กลุ่มโรคที่มีความสัมพันธ์กับการติดเชื้อไวรัสเอปสไตน์-บาร์ (EBV) การศึกษานี้เพื่อประเมินผู้ป่วยที่มีโรค PTCL ในระบบทางเดินอาหาร โดยศึกษาอาการทางคลินิก การดำเนินโรค การเปลี่ยนแปลงทางพยาธิวิทยา ศึกษาในระดับโมเลกุลของภาวะติดเชื้อไวรัส EBV และศึกษาโคลน (clone) ของเซลล์มะเร็ง การศึกษานี้ริบดังต่อไปนี้ ตามมาตราคุณ พ.ศ. 2540 จนถึงเดือนธันวาคม พ.ศ. 2543 พบรู้ป่วยโรค PTCL ชนิดต่าง ๆ 129 ราย ในจำนวนนี้ 7 ราย (5.4%) พบรอยโรค PTCL ในระบบทางเดินอาหาร อาการสำคัญของผู้ป่วยที่มีรอยโรค PTCL ในระบบทางเดินอาหารคือไข้ร้อน น้ำหนักลด ซัด ดับและม้ามโต ต่อมน้ำเหลืองโต มีรอยโรคในหลอดอาหารและมีเลือดออกในทางเดินอาหาร การตรวจเลือดพบว่ามีการเพิ่มขึ้นของเอนไซม์ alkaline phosphatase และ lactate dehydrogenase และมีความผิดปกติในระบบการแข็งตัวของเลือด ผู้ป่วย 5 ใน 7 รายเสียชีวิตภายใน 4 เดือน หลังป่วยอาการครั้งแรก ผู้ป่วยอีก 2 ราย หายขาดจากโรคหลังการได้รับยาเคมีบำบัด ผู้วัยรุ่นได้แบ่งลักษณะของเซลล์มะเร็ง เป็น 3 ประเภทคือ ประเภทที่มีเซลล์มะเร็งขนาดเล็ก ประเภทที่มีเซลล์มะเร็งขนาดปานกลางผสมกับขนาดใหญ่ และประเภทที่มีเซลล์มะเร็งขนาดใหญ่ ผลการตรวจปรับตัวที่อยู่บนผนังเซลล์ของเซลล์มะเร็งพบว่าเป็นชนิด LCA+, CD3+, CD15-, CD16-, CD30-, CD45RO+, CD57-, CD68-, EMA-, βF1-, granzyme B+, TIA+, และ p53+, สำหรับ CD4, CD8, CD56, CD20 บางรายให้ผลบวกและบางรายให้ผลลบ ตรวจพบ EBV RNA ในนิวเคลียสของเซลล์มะเร็งของผู้ป่วยทั้ง 7 ราย การตรวจโดยวิธี PCR ของ T-cell receptor (TCR) β และ/หรือ γ gene rearrangement ปรากฏว่าให้ผลบวกในผู้ป่วยทั้ง 7 ราย และการวิเคราะห์สาย DNA ซึ่งเป็นผลผลิตโดยวิธี PCR ของ TCR ดังกล่าวข้างต้น พบว่าล้ำดับบนส่วนสาย DNA มีความคล้ายคลึงกับ human TCR germline gene มาก การวิจัยนี้แสดงว่าการเกิด PTCL ในระบบทางเดินอาหาร มีความสัมพันธ์กับภาวะติดเชื้อไวรัส EBV, เซลล์มะเร็งเป็น mature T-cell ที่แสดงโปรดีนของ NK-cell บางชนิดบนผนังเซลล์ กลุ่มเซลล์มะเร็งในผู้ป่วยแต่ละรายมาจากเซลล์เพียงหนึ่งโคลน และเป็นเซลล์ที่มีการ rearrange ของยีน TCR แล้ว

คำสำคัญ : ที-เซลล์ ลิมโฟมา, ไวรัสเอปสไตน์-บาร์, ระบบทางเดินอาหาร, ยีนของ ที-เซลล์ รีชฟเตอร์, โคลนของเซลล์มะเร็ง

วิญญา มิตราันันท์, วรรณรัตน์ แซ่ชั้น, จินตนา ประดุจกาญจน์,  
สุภาพร สุวิวัฒน์, ชาโตมิ ทากาโอะ, ทากาพูมิ อิชิตะ  
จดหมายเหตุทางแพทย์ ย 2546; 86: 816-828

\* ภาควิชาพยาธิวิทยา, คณะแพทยศาสตร์ มหาวิทยาลัยสงขลานครินทร์, สงขลา 90110

\*\* ภาควิชาชีววิทยา, มหาวิทยาลัยโตเกียว, โตเกียว, ญี่ปุ่น