

Tumors : RECIST) が発表された。その後、この RECIST は広く普及した治療効果判定法となった。2009 年に改訂が行われ、現在の最新版は RECIST 1.1 である。RECIST 1.1 では腫瘍縮小効果の評価のために、腫瘍病変を一次元的に測定する。腫瘍量の測定の精密性を向上させることそのものより、方法論の標準化と単純化が求められていたためだが、腫瘍縮小効果判定の指標とするには精度の上で難点があることは否めない。なぜなら、腫瘍は治療経過中で様々な形態をとりながら縮小することがあるため、評価時点で腫瘍サイズの測定方向が異なることがあり、測定する画像断面も全く異なることが起こり得る。また、腫瘍サイズを人が測定するため測定者間で測定誤差があり、たとえ同じ測定者であっても測定毎に誤差が生じる。現在、CT から得られるボリュームデータを 3 次的に画像解析することで、より精度高く、より客観的な指標が導出可能となった。近年特に日本においては、ヘリカル CT やマルチスライス CT の普及と高機能化が急速に進み、ボリュームデータの取得が容易となった。また、コンピュータ支援画像診断の進歩により、ボリュームデータを用いた様々な解析が可能となっている。今後、この三次元的 CT 体積測定法を WT1 がんワクチンによる肺癌の治療効果判定に応用する。

#### E. 結論

肺癌の治療効果評価において、三次元的 CT 体積測定法は有用である。

#### F. 研究発表

##### 1. 論文発表

##### 2. 学会発表

(発表誌名巻号・頁・発行年)

Yanagawa M, Tanaka Y, Morii E, Okumura M, Johkoh T, Tomiyama N, et al. Three-dimensional (3-D) quantitative analysis of preoperative CT images in pathological stage I pulmonary adenocarcinomas: Correlation of radiologic CT data with prognostic factors. RSNA 2011, Chicago, USA

#### H. 知的財産権の出願・登録状況 (予定含む)

##### 1. 特許取得

##### 2. 実用新案登録

##### 3. その他

無し

厚生労働科学研究費補助金（難病・がん等の疾患分野の医療の実用化研究事業）  
分担研究報告書

肺癌に対する治療に関する研究

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研究要旨

非小細胞肺癌切除例に対する術後補助化学療法の有用性について、カルボプラチン+パクリタクセル療法と UFT 経口投与療法の有効性を比較検討する randomized phase III study を行っている

- A. 研究目的 我が国における肺癌補助化学療法の標準治療法を探索し、WT-1 免疫療法の効果の評価にあたっての baseline 治療を確立する
- B. 研究方法 瀬戸内肺癌研究会と協力して非小細胞性肺癌症例を randomize し、補助療法におけるカルボプラチン+パクリタクセル群と UFT 群の比較を行う
- (倫理面への配慮)  
参加各施設の IRB の承認を得て臨床研究を行っている。
- C. 研究結果 既に目標症例 401 例が 2010 年末までに集積され、patients' characteristics, compliance などの情報をまとめた段階である
- D. 考察 この臨床試験によって、わが国の肺癌補助療法における WT-1 治療の best partner としての chemotherapy を選択することが可能になる。
- E. 結論 2016 年の最終生存期間の比較が待たれる
- F. 研究発表
1. 論文発表  
2011 年アムステルダムにて開催された世界肺癌学会で発表を行った
  2. 学会発表  
International Association for the Study of Lung Cancer Meeting 2011
- H. 知的財産権の出願・登録状況（予定含む）
1. 特許取得 特になし
  2. 実用新案登録 特になし
  3. その他 なし

### 別紙 3

厚生労働科学研究費補助金（難病・がん等の疾患分野の医療の実用化研究事業）  
分担研究報告書

肺癌に対する WT1 ペプチド免疫療法の開発  
分担研究「がんワクチン療法開発臨床研究における生物統計学的手法」

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研究要旨： がんワクチン療法開発において臨床試験を計画・実施するにあたって重要な点としてデータベース集積と集積したデータの有効活用である。そのための最適な統計的方法の一つがベイズ流統計手法である。そこで、本分担研究ではベイズ流統計手法の利用可能性について検討・評価を行った。

#### A. 研究目的

がんワクチン療法をはじめとする新治療法開発のためのエビデンスを得るためには前向きに臨床試験を実施するのがもっとも望ましい。しかしながら、現実の臨床現場ではエビデンスが乏しいもとでも様々な治療法が手探りの使用される側面がある。そのような場合には、日常診療で得られた情報をデータベース化し、その中からデータ解析によりエビデンスの充実化を図ることが要求される。がんワクチン療法など臨床試験のデータが相対的に十分とは言えない分野での新規治療法の評価においては、データベースから得られるヒストリカルコントロールとの比較が有用である。そのための最適な統計的方法はベイズ流統計手法であり、米国で一、二を争う M.D. アンダーソン癌センターにおいても実際の臨床試験において頻用されている[参考文献 1,2]。この方法を用いてヒストリカルコントロールとの比較解析を有効性および安全性に関して実施することで、限られたリソース（人的資源、症例

数、研究費など）を“効率的”に利用することが可能となる。がんワクチン療法開発において最適な手法の一つであると考えられる。

ベイズ流”アプローチの特徴は複雑化する様々な臨床的要求に対して柔軟に対応できる点や事前情報を積極的に活用できることから、ヒストリカルコントロールデータを取り込んだデータ解析にとっても馴染みやすく、適用事例もますます増えてきている。本研究では、データベース情報を用いた解析手法を検討するための準備として、過去に実施された臨床試験の結果を事前情報として用いるベイズ流統計手法が積極的に用いられているがん第 I 相および第 II 相臨床試験について検討する。

#### B. 研究方法

主要な臨床統計のジャーナルである、Biometrics、Statistics in Medicine、Biostatistics、Clinical Trials などに出版された論文あるいは著書を対象として、がん第 I 相および第 II 相試験に活用可能な

ベイズ流試験デザインおよびデータ解析方法を調べる。

### C. 研究結果

ベイズ流統計とは、まず、有効性や安全性に関して調べたい興味のあるパラメータ ( $\theta$ ) を考える。奏効率 (CR: complete response + PR: partial response の割合) と DLT (用量規定毒性: dose limiting toxicity) 発現率がそれぞれ代表的なものである。 $\theta$  は1つの値に決まったものとして考えず、ランダムな変数であると考え。すなわち、平均的にどれ位の値をとり、バラツキはどの程度であるかを考える。つぎに、過去の臨床的データをもとにした  $\theta$  に関する知識あるいは不確からしさを統計的な確率分布を用いて‘事前分布’として表す。新たに実施する研究/試験で観察されたデータを事前情報に加えて  $\theta$  の推定精度を高めていく。このプロセスのことを観察データで事前分布を‘更新する (update)’と呼ぶ。データで更新された後の  $\theta$  に関する情報を‘事後分布’として表す。

第 I 相試験は、第 II 相試験で用いる投与量レベルを決定するため、新試験治療の最大耐用量 (MTD: maximum tolerated dose) の推定を目的として実施される。第 I 相試験において用いられるベイズ流試験デザインが“continual reassessment method (CRM)” [3-13] である。CRM は、これまで標準的試験計画として用いられてきた“3+3”コホートデザインのように直前 3 例のデータだけでなく、その時点までに得られた全てのデータを用いて“用量-毒性カーブ”の推定を行う。また、試験開始時までに行われている情報をデータとして解析に取り込むことができるため、事前情報を効果的に活用することができる。また、用量-毒性関係に対して統計的なモデル

を仮定するため、ターゲットとする毒性発現率に相当する用量レベルの推定を行うことも可能である。

第 I 相試験で設定された用量において当該試験治療法の有効性を評価し、最終的検証ステージである第 III 相試験への移行に値するかどうかを調べるのが第 II 相試験の主目的である。試験治療群一群での評価が標準的試験デザインとなる。第 II 相試験におけるベイズ流デザインの適用事例として、米国 MD アンダーソンがんセンター (MDACC) の Estey ら [2] がある。AML 患者を対象とし、エンドポイントに CR 率を用いて、Liposomal daunorubicin (LD) + ara-C や LD + Topotecan ら 4 つの新治療を既存の標準療法と比較するランダム化第 II 相試験を実施した。MDACC でそれまでに蓄積された 591 例分の標準療法のヒストリカルデータ: CR 率 49% (291/591 例) を比較対照として、試験治療群それぞれの効果を調べた。試験治療群で 40 例の評価が完了した時点で最も成績の良い群でも CR が得られたのは 18 例であった。このデータをもとに、標準治療群より CR 率が 20%、10%以上勝る事後確率を計算したところそれぞれ 0.001 と 0.039 となった。この結果に基づいて標準療法を上回る十分な効果は期待できないと判断し、その時点で試験を中止した。さらに、有効性と毒性を一つの評価指標にまとめて同時にモニタリングを行うといったデザインも提案されている [14-18]。また、複数の治療法を同時に比較するため、“ランダム化第 II 相試験” [19-21] と呼ばれるデザインが用いられるケースが増えている。ランダム化第 II 相試験は Simon ら (1985) [19] により提案され、ランダム化によりもたらされるバイアスのない比較に基づき、次に行う検証第 III 相試験に組み込む治療群を選択することを目的とする。ランダム化第 II 相

試験においてもベイズ流統計の事前情報を活用する点は大きな魅力となる。Morita and Sakamoto (2006)はランダム化第Ⅱ相試験におけるベイズ流試験デザインの適用事例を報告しており、事前情報を積極的に用いることの有用性を議論している[21]。Neuenschwanderら[22]は、ベイズ流統計手法でもちいる事前情報分布を過去の情報に基づいてどのように構築していくかについて議論している。

#### D. 考察

事前情報を積極的に活用するベイズ流統計手法を適切に用いるためには、事前情報のもとになるデータの質の高さが鍵となる。事前情報の有効活用方法についてはいまだ研究段階であるといわざるをえず、たとえば、情報収集ルールの統一化など入手した事前情報の質を担保するための方策を厳密に議論することが重要である。

#### E. 結論

事前情報の活用の際にベイズ流アプローチの適用を考慮することは重要である。

#### F. 健康危険情報

なし

#### G. 研究発表

##### 1. 論文発表

現時点でなし

##### 2. 学会発表

なし

#### H. 知的財産権の出願・登録状況

##### 1. 特許取得

なし

##### 2. 実用新案登録

なし

##### 3. その他

なし

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### III. 研究成果の刊行に関する一覧表

研究成果の刊行に関する一覧表

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#### IV. 研究成果の刊行物・別刷

# Epstein–Barr Virus in Diffuse Large B-Cell Lymphoma in Immunocompetent Patients in Japan is as Low as in Western Countries

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According to previous reports, the frequency of Epstein–Barr virus (EBV) positivity in diffuse large B-cell lymphoma is higher in East Asia (approximately 9%) than in Western countries. The presence of the EBV genome was examined in diffuse large B-cell lymphoma patients registered with the Osaka Lymphoma Study Group (OLSG) in Osaka, Japan, situated in East Asia. The EBV-positive rate was examined with *in situ* hybridization (ISH) in 484 immunocompetent diffuse large B-cell lymphoma patients registered with OLSG. The male-to-female ratio was 1.29, with ages ranging from 16 to 95 (median, 68) years. ISH with EBV-encoded small RNAs (EBER) probes revealed positive signals in the nuclei of tumor cells: the frequency of positively stained cells among all tumor cells was almost none in 458 cases, 5–10% in 5, 10–20% in 5, 20–50% in 11, and >50% in 5. When the frequency was >20% or >50%, the EBV-positive rate in the present series (3.3% or 1.0%) was rather similar to that reported in Western cases. Careful evaluation of patient backgrounds, including age distribution, type of lymphomas, exclusion of immunocompromised patients, and establishment of definite criteria for EBV positivity (>20%, >50%, or almost all tumor cells) are essential in comparing geographical differences. *J. Med. Virol.* 83:317–321, 2011. © 2010 Wiley-Liss, Inc.

**KEY WORDS:** Epstein–Barr virus; diffuse large B-cell lymphoma; *in situ* hybridization; immunocompetence; geographical differences

## INTRODUCTION

The Epstein–Barr virus (EBV), a  $\gamma$ -herpesvirus, is transmitted by mucosal secretions among the human population. An association between EBV and human lymphomas, including endemic Burkitt's lymphoma, Hodgkin lymphoma, and non-Hodgkin lymphomas of B-, T-, or NK-cell immunophenotypes, has been reported [Shah and Young, 2009]. In the recent WHO classification (2008), the chapter for EBV-positive diffuse large B-cell lymphoma of the elderly states that the EBV-positive rate in diffuse large B-cell lymphoma in Asian countries is 8–10% [Nakamura et al., 2008]. Hematologists and hematopathologists in the Osaka Lymphoma Study Group (OLSG) in Osaka, Japan, situated in the East Asia, felt a sense of incongruity for this figure, although they had no data on this.

Abbreviations: EBV, Epstein–Barr virus; OLSG, Osaka Lymphoma Study Group; LMP, latent membrane protein; ISH, *in situ* hybridization; EBER, EBV-encoded small RNAs; MTX, methotrexate; PNA, peptide nucleic acid; FITC, fluorescein isothiocyanate; DAB, 3,3'-diaminobenzidine tetrahydrochloride; CTL, cytotoxic T-lymphocytes.

We declare that we have no conflict of interest.

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There have been two large-scale studies on the EBV positivity in diffuse large B-cell lymphoma from East Asian countries, one in Korea and one in Japan. Both studies showed a similar frequency of EBV positivity among diffuse large B-cell lymphoma, that is, 9% [Oyama et al., 2007; Park et al., 2007]. However, two points regarding these reports should be addressed. First, the age distribution was quite different between the Korean and Japanese cases: the percentage of cases older than 60 years was 36.6% in Korean and 60.6% in Japanese cases. It seems that the EBV-positive rate is higher in elder patients; thus, the difference in age distribution might affect EBV positivity. Second, the criteria for EBV positivity of diffuse large B-cell lymphoma was different: >20% of tumor cells with positive signals by *in situ* hybridization (ISH) in one report [Park et al., 2007] and >50% in another [Oyama et al., 2007]. Some Japanese investigators, adopting the criteria of almost all tumor cells with positive signals as EBV-positive, reported that 11.4% of their diffuse large B-cell lymphoma cases were EBV-positive [Kuze et al., 2000]. However, reports from Western countries apparently showed a much lower EBV-positive rate than did the East Asian ones [Gibson and Hsi, 2009; Hoeller et al., 2010]. Hoeller et al. [2010] reported that 10 (3.9%) of their 258 diffuse large B-cell lymphoma cases were EBV-positive: the percentage of positive cells was <20% in two cases, <50% in four, and >50% in four. Gibson and Hsi [2009] reported that none of their 90 diffuse large B-cell lymphoma, all older than 60 years, was EBV-positive. They stated that EBV-positive tumor cells were not identified in any of the 90 cases, and only three demonstrated very rare EBV-positive signals in small bystander lymphocytes (<1% of non-tumor cells).

In this study, we examined the presence of the EBV genome with ISH in the tumor cells in cases of diffuse large B-cell lymphoma registered with OLSG. The EBV-positive rate was then evaluated according to the criteria used in the previous and present studies, and the data were then compared with those reported from East Asia and Western countries in the light of clinical findings such as age, gender, and presence of immunodeficiencies.

## MATERIALS AND METHODS

### Patients

Between November 1999 and October 2009, in total, 4,490 cases were registered with the OLSG, Japan, in which 63 hospitals participate and register cases of malignant lymphomas and related conditions. All of the hematoxylin and eosin- and immunoperoxidase-stained sections were reviewed by one of the authors (K.A.) and classified according to the WHO classification. A diagnosis of malignant lymphoma was confirmed in 3,571 (79.5%) of 4,490 cases. Of these 3,571 cases, 3,273 (91.7%) were non-Hodgkin lymphoma and 298 (8.3%) were Hodgkin lymphoma. The number of diffuse large B-cell lymphoma cases was 1,590, which constituted 48.6% of all non-Hodgkin lymphoma. The latest 500

cases with diffuse large B-cell lymphoma registered during February 2007 to October 2009 were studied for the presence of the EBV genome. Most patients received anthracyclin-based chemotherapy and rituximab after the diagnosis. The numbers of patients who received excisional biopsy or surgical resection, punch biopsy, or needle biopsy were 384, 76, and 40, respectively. In cases receiving punch or needle biopsy, an adequate amount of material for histological analyses was obtained. The primary site was lymph node in 165 cases, extranodal organs in 204, and unknown in 131. Among them, two patients suffered from rheumatoid arthritis, one each from idiopathic thrombocytopenic purpura, interstitial pneumonia, Sjögren's syndrome, and uveitis, and were treated with prednisone, together with cyclosporine in one. These six cases were excluded for further analysis because of possible underlying immune abnormalities. During the same period, 14 cases of polymorphous lymphoproliferative disorders, 14 methotrexate (MTX)-associated lymphoproliferative disorders, five post-transplant lymphoproliferative disorders (liver, kidney, and cord blood in two, two, and one case, respectively), and one Hodgkin-like lymphoproliferative disorder were registered with the OLSG. Polymorphous lymphoproliferative disorders are defined as lymphoproliferative disorders showing a polymorphous pattern of proliferation, consisting of large lymphoid cells and various numbers of inflammatory cells [Nakamichi et al., 2010]. Polymorphous lymphoproliferative disorders usually affect individuals with a background of immunodeficiency. Clinical findings entered on the registration card for the OLSG included the following: name of hospital, age, gender, occupation, site of disease, clinical diagnosis, present illness, and peripheral blood and bone marrow findings, including differential counts, serum data, and chromosomal abnormalities of the tumor cells. History of medication (those receiving immunosuppressants) was also checked.

The Institutional Review Board for Clinical Research at Osaka University Hospital approved this study.

### Immunohistochemistry

Antibodies used for immunohistochemistry were CD20, CD79a, CD3, CD30, and anti-Epstein-Barr virus/LMP/Clones CS. 1-4 (DakoCytomation, Glostrup, Denmark, diluted at 1:400, 1:100, 1:50, 1:25, and 1:100, respectively). Immunohistochemistry was performed using an automated staining system (Dako Autostainer, DakoCytomation).

### In Situ Hybridization

ISH using the EBV-encoded small RNAs (EBER) probe was performed to examine the presence of the EBV genome in the formalin-fixed paraffin-embedded sections with the EBER DAB application kit (DakoCytomation). Briefly, sections were treated with proteinase K, diluted at 1:10 with TBS (50 mmol/L Tris-HCl buffered saline containing 150 mmol/L NaCl, pH 7.6), then hybridized with EBER PNA (peptide nucleic acid)

probe/fluorescein (DakoCytomation) at 55°C for 90 min. After blocking endogenous peroxidase activity, sections were incubated with rabbit anti-FITC (fluorescein isothiocyanate) antibody (1:50; Invitrogen, Carlsbad, CA) at room temperature for 30 min, incubated with ChemMate ENVISION/HRP polymer (DakoCytomation) at room temperature for 30 min, and colored with DAB (3,3'-diaminobenzidine tetrahydrochloride). This ISH method using an EBER probe was recently used in the author's laboratory. The EBV-positive rate in NK/T cell lymphoma by this method on paraffin-embedded sections was similar (80%) to that with the previously used method using an EBER-1 probe [Li et al., 2000].

The frequency of EBV positivity in the malignant large lymphoid cells was determined by averaging the number of positive cells from three high-power fields where the positive cells were larger in number. The tumors were then separated into five groups: tumors with almost no EBV positivity, tumors with 5–10%, 10–20%, 20–50%, and >50% of cells showing EBV positivity. According to the previous study, we adopted the criterion of >20% as EBV-positive.

#### Follow-Up

Adequate follow-up data was available in 16 EBV-positive and 204 EBV-negative diffuse large B-cell lymphoma cases. Follow-up periods for survivors in EBV-positive and -negative cases ranged from 5.2 to 38.5 (average 22.6) and 16.4–38.1 (average 25.5) months, respectively. Kaplan–Meier estimated survival rates at 2 and 3 years were calculated, and overall survival rates were compared by a log-rank test.

#### RESULTS

EBV-ISH was performed in 494 cases; however, informative data was not obtained in 10 cases because of specimen loss during staining procedures or severe non-specific staining of background/overstaining. As a result, staining results could be evaluated in only 484 cases. Positive signals were found in the nuclei of tumor cells: the frequency of positively stained cells among all tumor cells was almost none in 458 cases, 5–10% in 5, 10–20% in 5, 20–50% in 11, and >50% in 5 (Fig. 1).

Of 16 cases with >20% EBV-positive tumor cells (EBV-positive cases), large geographical necrosis was found in five cases. Angiotropism was not found. CD30 positivity was found in 6 of 15 EBV-positive cases. Tumor cells expressed latent membrane protein (LMP)-1 in 8 of 14 EBV-positive cases (Fig. 1).

In total, 40 cases with diffuse large B-cell lymphoma were excluded, due to the putative presence of immunocompromised conditions, including autoimmune diseases, receiving prednisone and cyclosporine, polymorphous lymphoproliferative disorders, MTX-associated lymphoproliferative disorders, post-transplant lymphoproliferative disorders, and Hodgkin-like lymphoproliferative disorders. The frequency of positively stained cells among all tumor cells was 20–50% in six cases and >50% in five.

The estimated survival rates at 2 and 3 years for the EBV-positive and -negative cases were 58.9% and 71.9% at 2 years and 58.9% and 69.0% at 3 years, respectively. The overall survival rates of the two groups were not statistically different (log-rank test;  $P > 0.1$ ).

#### DISCUSSION

According to previous reports, frequency of EBV positivity in diffuse large B-cell lymphoma was higher in East Asia than in Western countries (Table I) [Kuze et al., 2000; Oyama et al., 2007; Park et al., 2007; Gibson and Hsi, 2009; Hoeller et al., 2010]. However, before accepting this, a careful evaluation of the data presented in the previous studies is necessary. Regarding clinical findings, the age distribution and gender ratio varied among these reports [Kuze et al., 2000; Oyama et al., 2007; Park et al., 2007; Gibson and Hsi, 2009; Hoeller et al., 2010]. A study referenced in the WHO classification adopting >50% of the criteria included cases of immunodeficiency-associated lymphomas, although the exact number was not shown. Lymphomas known to be closely associated with EBV infection such as pyothorax-associated lymphoma and Burkitt's lymphoma were also included [Oyama et al., 2007]. These procedures inevitably increased the EBV-positive rate of the study (13.6%). The routinely registered cases with the OLSG constitute the present series; thus, collection bias in the cases could be avoided. The age distribution and gender ratio in the present series are within the spectrum of the previously reported diffuse large B-cell lymphoma cases; thus, the present findings could be adopted as standard data.

The criteria for defining EBV-positive cases vary among the previous studies; >20% of the tumor cells in one study [Park et al., 2007] and >50% in another [Oyama et al., 2007]. In particular, a study from Japan adopting the criteria of almost all EBV-positive cells reported the EBV-positive rate to be 11.4% [Kuze et al., 2000]. When adopting >20% or >50% as EBV-positive, the EBV-positive rates among diffuse large B-cell lymphoma in immunocompetent hosts in the OLSG (3.3% or 1.0%) were rather similar to those in Western series [0–2.7%, 0–1.9%; Gibson and Hsi, 2009; Hoeller et al., 2010], but much lower than that in Korean (8.9%) [Park et al., 2007] and Japanese series (11.4–13.6%) [Kuze et al., 2000; Oyama et al., 2007].

The present and previous [Park et al., 2007; Gibson and Hsi, 2009; Hoeller et al., 2010] studies have analyzed the presence of the EBV genome in diffuse large B-cell lymphoma in immunocompetent patients. When a total of 40 cases with diffuse large B-cell lymphoma, which had been excluded from the calculation of the EBV-positive rate due to the putative presence of immunocompromised conditions (such as autoimmune disease, receiving prednisone and cyclosporine, polymorphous lymphoproliferative disorders, MTX-associated lymphoproliferative disorders, post-transplant lymphoproliferative disorders, and