Concurrent Onset of Chronic Lymphocytic Leukemia and Atypical Phenotype Acute Myeloid Leukemia Revealed by Autopsy

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Case Report

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Correspondence to:
Hiroyuki Sugiura, hiroyuki.sugiura0715@gmail.com
Introduction

Myelodysplastic syndrome (MDS) or acute myeloid leukemia (AML) in patients with a previous diagnosis of chronic lymphocytic leukemia (CLL) is extremely rare, accounting for <1% of cases [1]. The development of AML in patients with previously treated CLL is diagnosed as therapy-related AML, and it generally has inferior prognosis compared to that of de novo AML [2]. Although there have been reports of therapy-related AML, those on the development of AML in patients with untreated CLL are rare [3]. In the majority of synchronous cases of CLL and AML, flow cytometry and immunohistochemistry showed characterization of 2 distinct coexisting malignant cell populations, which helped to select treatment [2, 3].

We report herein a case of CLL with concurrent onset of AML with an atypical phenotype revealed by autopsy. Concurrent onset of CLL and AML is rare, and to the best of our knowledge, no autopsy case has been reported. Notably, the diagnosis of AML was difficult during the patient’s lifetime because of the atypical phenotype of AML. However, autopsy revealed that the patient’s bone marrow, liver, and spleen were filled with large-sized blastic abnormal cells which had characteristic of AML.

Case Report

A 69-year-old man was referred to our hospital with severe anemia and thrombocytopenia. He was previously healthy and had not received any medication. Laboratory data are shown in Table 1. Splenomegaly was also found on computed tomography (Fig. 1a).

### Table 1. Laboratory data and findings of bone marrow aspiration on the admission day

| CBC and coagulation test | Biochemistry | BMA findings and tumor marker |
|-------------------------|--------------|------------------------------|
| WBC 4.240/µL            | TP 6.9 g/dL  | NCC 33,000/µL                |
| Band 1%                 | Alb 4.8 g/dL | Megakaryocyte 12/µL          |
| Neu 54%                 | T. Bil 0.6 mg/dL | Small lymphocyte 34.8%, CD5+, CD20+, CD23+, light chain κ+ |
| Mo 4%                   | AST 24 U/L   |                              |
| Lymph 41%               | ALT 18 U/L   | Blastic abnormal cell 21.4%, CD4+, CD38+, CD3–, CD20–, CD5–, CD23–, CD13–, CD34–, CD117–, MPO–, TdT– |
| RBC 246 × 10^6/µL       | LDH 282 U/L  |                              |
| MCV 87.0 fl             | γ-GTP 42 U/L |                              |
| Hb 7.4 g/dL             | UA 5.7 mg/dL |                              |
| Reti 0.24%              | Cre 0.99 mg/dL | Chromeosome analysis: 46, XY, add (1) (p36.1), del1 (1) (p?), −5, −7, −8, −10, −12, −13, −14, −16, add (19) (p13), −21, +8mar [4], 46, XY [16] |
| Plt 1.1 × 10^12/µL      | BUN 13 mg/dL |                              |
| APTT 27.6 s             | Na 141 mmol/L |                              |
| PT-INR 76%              | K 4.2 mmol/L | Tumor marker: sIL-2R 625 U/mL |
| Fib 255 mg/dL           | Cl 107 mmol/L | WT1mRNA 3,000 copy/µgRNA     |
| D-D 0.7 µg/mL           | Ca 9.3 mg/dL |                              |
| FDP <2.5 µg/mL          | IP 3.7 mg/dL |                              |

CBC, complete blood count; WBC, white blood cell; Band: band neutrophil; Neu, neutrophil; Mo, mononuclear cell; Lymph, lymphocyte; RBC, red blood cell; MCV, mean corpuscular volume; Hb, hemoglobin; Reti, reticulocyte; Plt, platelet; APTT, activated partial thromboplastin time; PT-INR, prothrombin time international normalized ratio; Fib, fibrinogen; D-D, d-dimer; FDP, fibrin and fibrinogen degradation products; TP, total protein; Alb, albumin; T. Bil, total bilirubin; AST, aspartate aminotransferase; ALT, alanine aminotransferase; LDH, lactate dehydrogenase; γ-GTP, γ-glutamyl transpeptidase; UA, uric acid; Cre, creatinine; BUN, blood urea nitrogen; Na, sodium; K, potassium; Cl, chloride; Ca, calcium; IP, inorganic phosphorus; BMA, bone marrow aspiration; NCC, nucleated cell count; CD, cluster of differentiation; MPO, myeloperoxidase; TdT, terminal deoxynucleotidyl transferase; FISH, fluorescence in situ hybridization; sIL-2R, soluble IL-2 receptor; WT1mRNA, WT1 messenger ribonucleic acid.
Bone marrow aspiration (BMA) showed 21.4% abnormal blastic cells and small lymphocyte populations (Fig. 1b, c). Flow cytometry revealed that the blastic cells did not express any lineage markers, except for CD4 and CD38. Meanwhile, small lymphocytes expressed CD5, CD20, CD23, and light chain-kappa. Immunohistochemical staining of the bone marrow clot revealed almost similar findings to those of flow cytometry, that is, infiltration of abnormal blastic cells with small lymphocytes to the bone marrow (Fig. 1d). Chromosome analysis showed complex karyotype including −5 and −7 abnormalities. The laboratory data and BMA findings are summarized in Table 1. Based on these findings, we diagnosed the small lymphocyte as CLL with Rai stage IV and Binet stage C, but we could not identify the origin of the abnormal blastic cells. Finally, we considered the abnormal blastic cells to be derived from CLL, such as Richter syndrome, which is an aggressive B-cell lymphoma transformed by CLL. The patient was started on bendamustine and rituximab (BR) treatment for CLL. However, he developed splenomegaly, and his cytopenia did not ameliorate, indicating treatment failure. We decided to start venetoclax as the second-line therapy for CLL, but it was also discontinued due to complications of tumor lysis syndrome. To evaluate the therapeutic effect of venetoclax, BMA was performed again and showed 11.8% of abnormal blastic cells with no small lymphocyte population, which seemed to be a CLL clone. However, the splenomegaly progressed, and 1–2% of abnormal blastic cells appeared in the peripheral blood. Finally, he was given palliative care because of his poor general condition and his living will, and he died 84 days after admission. We performed an autopsy with the consent of his family.

Macroscopic findings of the autopsy revealed prominent splenomegaly and hepatomegaly. In microscopy, the patient’s bone marrow, spleen, and liver were filled with large-sized abnormal...
blastic cells. These abnormal blastic cells comprised 70% nucleated cells in the bone marrow. Almost all the spleen was infiltrated by the abnormal blastic cells, and the border between white pulp and red pulp disappeared. Images of the autopsy specimen are shown in Figure 2. The abnormal blastic cells stained positively for CD4 and partially positive for MPO; negative for CD3, CD5, CD20, CD23, CD34, CD56, and c-kit; and showed p53 overexpression. Findings from flow cytometry and immunohistochemical staining of the abnormal blastic cells are summarized in Table 2. Increasing polymerase chain reaction (PCR) of WT1 mRNA suggested complications of myeloid hematologic malignancy (Table 1). Considering the partial positivity of MPO in autopsy specimens and these findings, we concluded that the abnormal blastic cell was a myeloblast and finally diagnosed the patient with AML with liver and spleen involvement. There was no CLL component. Based on these findings, the major cause of death was the progression of AML. We told the result of autopsy to the family and obtained written informed consent to publish the case from the family.

Discussion

In this case, the patient suffered from the rare combination of synchronous hematological malignancy: untreated CLL and AML. Because of the rarity and atypical phenotype, we could not diagnose AML until the autopsy proved the myeloblast in the bone marrow, liver, and spleen. Although second malignant neoplasms are more common in patients with CLL, the development of AML or MDS in untreated CLL patients is rare [4]. In Japan, Muta et al. [5] reported the first case of AML concurrently occurring with untreated CLL. Ito et al. [6] reported that the development of AML in patients with untreated CLL was associated with a poor
response to chemotherapy and extremely poor prognosis. To the best of our knowledge, this is the first autopsy case of concurrent onset of CLL and AML, and complications of AML were revealed by autopsy.

On the other hand, there are several reports about therapy-related MDS/AML developed after CLL treatment. As shown in Table 3, fludarabine-containing regimens seemed to be associated with the occurrence of MDS/AML [7–9]. In our case, the patient was previously healthy and had not received any medication including fludarabine.

Most previous studies reported that concurrent-onset CLL and AML are different clones [5, 10, 11]. However, Ito et al. [6, 12, 13] suggested the possibility of CLL and AML from the same origin because both tumors are related to TP53 alterations and aberrant Wnt signaling. In our case, we added p53 staining of the first BMA smear and PCR of IgH rearrangement in the bone marrow clot specimen for the first BMA and the autopsy specimen of the spleen. The abnormal blastic cells stained positively for p53, while the CLL component, which showed

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**Table 2.** Summary of findings from flow cytometry and immunohistochemical staining of the abnormal blastic cells

| Stain  | Pretreatment flow cytometry | Pretreatment IHC | IHC on autopsy |
|--------|----------------------------|-----------------|----------------|
| CD4    | +                          | +               | +             |
| CD38   | +                          | na              | na            |
| CD3    | -                          | -               | -             |
| CD20   | -                          | -               | -             |
| CD5    | -                          | -               | -             |
| CD23   | -                          | -               | -             |
| CD13   | -                          | na              | na            |
| CD34   | -                          | -               | -             |
| CD56   | -                          | -               | -             |
| CD117/c-kit | -                        | -              | -             |
| MPO    | -                          | -               | + (partial)   |
| TdT    | -                          | na              | na            |
| p53    | na                         | +               | Overexpression |

IHC, immunohistochemistry; CD, cluster of differentiation; na, not available; MPO, myeloperoxidase; TdT, terminal deoxynucleotidyl transferase.

**Table 3.** Summary of reports about therapy-related MDS/AML after CLL treatment

| References          | N  | CLL treatment                                                                 | Median follow-up period | Incidence of therapy-related MDS/AML (%) | Median survival after diagnosis |
|---------------------|----|-------------------------------------------------------------------------------|-------------------------|-----------------------------------------|-------------------------------|
| Morrison et al. [7]  | 521| Fludarabine alone/fludarabine and chlorambucil                                | 4.2 years               | Six patients (1.2)                       | 3.5 months                    |
| Carney et al. [8]   | 82 | Fludarabine and cyclophosphamide with or without rituximab                    | 41 months               | Five patients (6.1)                      | 11 months                     |
| Colovic et al. [9]  | 210| Fludarabine and cyclophosphamide                                              | 46 months               | Four patients (1.9)                      | 4 months                      |

MDS, myelodysplastic syndrome; AML, acute myeloid leukemia; CLL, chronic lymphocytic leukemia.
CD5 and CD23 double positivity, was negative for p53 (Fig. 3). In addition, IgH gene rearrangement was detected in the bone marrow clot specimen in the first BMA but was not detected in PCR of the autopsy specimen of the spleen. Based on these findings, we assumed that CLL and AML were different clones.

We could not identify AML before autopsy because both flow cytometry and immunohistochemical staining did not show typical antigens of AML. In addition, autopsy revealed that myeloblasts filled the bone marrow, liver, and spleen, but there were very few abnormal blastic cells in the peripheral blood during his lifetime. This is quite an atypical distribution for AML, although there have been few case reports [14]. We assumed that this unusual clinical presentation of AML was due to the progression of MDS. In the case of AML with myelodysplasia-related changes, immunophenotyping results are variable due to the heterogeneity of the underlying disease. Decreased expressions of HLA-DR, CD117, CD135, and CD38 are reported to be associated with multilineage dysplasia [15]. Kitagawa et al. [16] reported that p53-positive cells were also positive for the myeloid cell marker, and all of their 7 MDS cases that exhibited p53 expression at the time of initial diagnosis later developed overt leukemia. In our case, the autopsy revealed overexpression of p53 in the bone marrow, liver, and spleen, which helped to suspect progression of MDS to overt leukemia.

We reviewed the diagnosis of CLL because of the bone marrow infiltration of CLL cells, but the origin of the abnormal blastic cells could not be defined before biopsy. We mistook the abnormal blastic cells for cells derived from CLL, similar to Richter syndrome, and the treatment aiming for CLL seemed reasonable at that time, but the autopsy revealed the blastic cells were myeloblasts, and they seemed to be derived from different clones. Thus, BR treatment...
seemed to be effective for CLL, but not for AML. Subsequent venetoclax treatment resulted in the disappearance of CLL clones and reduced the number of myeloblasts. However, the therapeutic effect was still limited. Although venetoclax is a BCL-2 inhibitor and has been reported to be effective in not only CLL but also AML, single-agent venetoclax has been reported to have insufficient antileukemia effect in AML [17].

In conclusion, we experienced a case of concurrent onset of CLL and atypical phenotype AML with liver and spleen involvement revealed by autopsy. The case findings highlight the importance of considering the synchronous complications of AML in CLL patients, particularly in those who show an atypical clinical course.

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Statement of Ethics

Written informed consent has been obtained from the patient’s family for publication. The study protocol was approved by the Chugoku Central Hospital Ethics Committee on June 17, 2020.

Conflict of Interest Statement

The authors declare no potential conflicts of interest regarding the publication of this study.

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Author Contributions

S.K. wrote this manuscript. H.S. managed the clinical practice and authored this case study. T.K. managed and supervised the clinical practice. R.O. and T.T. managed the autopsy and supervised the pathological part of this study. T.I., S.O., N.N., T.M., and N.S. advised on the manuscript. T.K. and M.T. supervised the clinical practice.

Data Availability Statement

The data during this study are available from the corresponding author on reasonable request.
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