Cells of origin of squamous epithelium, dysplasia and cancer in the head and neck region after bone marrow transplantation

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Abstract. Secondary solid tumors that occur after hematopoietic stem cell transplantation (HSCT) are late complications of HSCT. Previously, secondary solid tumors were considered to be recipient-derived cells because transplanted cells do not contain epithelial cells. Recently, however, not only donor-derived epithelial cells but also donor-derived secondary solid tumors have also been reported in mice and humans. It means that circulating bone marrow-derived stem cells (BMDCs) including hematopoietic stem cells include the stem cells of many tissue types and the precancerous cells of many solid tumors. In most reports of donor-derived secondary solid tumors, however, tumors contained a low proportion of BMDC-derived epithelial cells in mixed solid tumor tissues. To our knowledge, there are only five known cases of completely donor-derived tumor tissues, i.e., four oral SCCs and a pharyngeal SCC. In this study, we analyzed five human clinical samples of solid tumors, i.e., two esophageal squamous cell carcinomas (SCCs), two oral SCCs and a tongue carcinoma. In the oral and tongue, completely donor-derived tissues were not observed, but in esophagus a completely donor-derived esophageal epidermis and SCC were observed for the first time. In addition, in another esophageal SCC patient, a completely donor-derived dysplasia region of esophageal epidermis was observed near recipient-derived SCC. This study suggests that BMDC-derived cells include the stem cells of esophageal epidermis and the precancerous cells of esophageal SCC and can differentiate into esophageal epithelium and esophageal SCC.

Introduction

Recent studies on regenerative medicine have indicated that mesenchymal tissues contain multipotent stem or progenitor cells that can give rise to neural and skin tissues, adipocytes, myocytes, cardiomyocytes, blood vessels, chondrocytes and hepatocytes (1). Given that bone marrow (BM) and adipose tissues are major sources of mesenchymal stem or progenitor cells (2), the observation of BM transplantation in mice and humans can provide adequate biological and biophysical information concerning cells of origin. Reportedly, bone marrow-derived cells (BMDCs) accumulate in the gastric epithelium as a result of Helicobacter pylori infection and can contribute to tumor development, indicating that infection can lead to the development of hyperplasia, metaplasia and dysplasia associated with BMDC recruitment and accumulation in the gastric epithelial mucosa, which occurs against a background of chronic inflammation (3). However, the contribution of BMDCs to head and neck cancers, including esophageal cancer, remains unknown.

GVHD is a major complication of allogeneic hematopoietic stem cell transplantation (HSCT), with significant morbidity and mortality (4). Therefore, adequate control of GVHD is critical to the continued success of transplantation. GVHD shares its molecular basis with chronic inflammation. This molecular basis includes the induction of intrinsic damage
to tissue stem or progenitor cells and deleterious effects on immune surveillance, all extensive and dynamic alterations that accumulate in the course of carcinogenesis (5). The use of immunosuppressant therapy exerts a favorable effect by ameliorating chronic GVHD, but it is associated with a higher relapse rate of hematopoietic and secondary malignancies, thus posing a major threat in the long term (6).

Cells for HSCT are obtained from BM, peripheral blood, or umbilical cord blood, which is proposed to contain mesenchymal stem or progenitor cells as well as other cell types (7,8). Secondary malignancies following HSCT are common late complications (9). With regard to the cells of origin, secondary tumors are considered to be derived from recipient-derived cells because there are very few epithelial cells in normal BM and peripheral blood (10). Nevertheless, the contribution of human BM cells, including HSCs, to epithelium, dysplasia and cancer is poorly understood (11). To study the involvement of BM cells in solid tumors developing after HSCT and distinguish the origins of epithelial cancer cells in humans, we performed highly sensitive FISH using gender chromosome-specific probes and histopathological analyses in five cases of head and neck tumors that developed subsequent to gender-mismatched BM transplantation.

Patients and methods

Patients. Patient characteristics are summarized in Table I. All patients had received gender-mismatched HSCTs and developed GVHD. Two clinical samples were obtained from patients with esophageal squamous cell carcinoma (SCC) treated at our hospital. Three clinical samples from two patients with oral SCC and one patient with tongue Diseases (Osaka, Japan). All clinical samples used in this study were acquired after obtaining written informed consent from each patient.

FISH analysis. FISH analysis was performed by Chromosome Science Labo, Inc. (Sapporo, Japan) using formalin-fixed, paraffin-embedded tissue sections as described previously (12). Briefly, 5-mm-thick sections were deparaffinized, dehydrated, microwaved (600 W) in 2X saline sodium citrate (SSC) for 10 min, cooled in PBS, digested in pepsin solution containing 0.1 N HCl at 37˚C (0.1% pepsin for 10 min for samples 4-1 and 4-2; 0.02% pepsin for 5 min for the other samples), and dehydrated. Human XY FISH probes (Chromosome Science Labo, Inc.) were applied to the pretreated sections, covered with cover slips, and simultaneously denatured at 90˚C for 13 min. Hybridization was performed at 37˚C overnight. Sections were then washed with 50% formamide/2X SSC at 37˚C for 20 min and 1X SSC for 15 min at room temperature. The slides were treated with antibodies at 37˚C for 30 min, washed three times with 0.1 % Nonidet P-40/2X SSC, counterstained with 4’,6-diamidino-2-phenylindole (DAPI), and mounted. The FISH images were captured using the CW4000 FISH application program (Leica Microsystems Imaging Solution, Ltd., Cambridge, UK) using a cooled charge-coupled device camera mounted on a Leica DMRA2 microscope (Leica Microsystems, Wetzlar, Germany). The enumeration probes for the X chromosomes were labeled with cyanine 3 (Cy3), SpectrumGold™ (Abbott Laboratories, Abbott Park, IL, USA), or Cy5, whereas the enumeration probes for the Y chromosomes were labeled with SpectrumGreen™ or SpectrumRed™ (Abbott Laboratories). The number of cells showing FISH signals was counted by the observation of at least five fields under the microscope (x100). All data were evaluated by at least three pathologists.

Histopathological analyses. Pathological diagnoses of DAPI- or hematoxylin and eosin-stained samples were performed by at least three pathologists to identify normal and malignant cells.

Results

BM significantly contributes to normal epithelium and SCC of the esophagus. Esophageal cancer developed in a male recipient with non-Hodgkin's lymphoma 115 months after

Table I. Characteristics of cases with secondary SCCs after HSCT.

<table>
<thead>
<tr>
<th>Case no.</th>
<th>Gender of donor/recipient</th>
<th>Diagnosis</th>
<th>Type of HSCT</th>
<th>GVHD</th>
<th>Location</th>
<th>Age at diagnosis</th>
<th>Time after transplantation (months)</th>
</tr>
</thead>
<tbody>
<tr>
<td>1</td>
<td>F/M</td>
<td>Non-Hodgkin's lymphoma</td>
<td>PBSCT</td>
<td>Chronic</td>
<td>Esophagus</td>
<td>42</td>
<td>115</td>
</tr>
<tr>
<td>2</td>
<td>F/M</td>
<td>Non-Hodgkin's lymphoma</td>
<td>PBSCT</td>
<td>Chronic</td>
<td>Esophagus</td>
<td>77</td>
<td>120</td>
</tr>
<tr>
<td>3</td>
<td>F/M</td>
<td>MDS</td>
<td>BMT</td>
<td>Chronic</td>
<td>Oral cavity</td>
<td>38</td>
<td>76</td>
</tr>
<tr>
<td>4-1</td>
<td>M/F</td>
<td>CML</td>
<td>BMT</td>
<td>Chronic</td>
<td>Tongue</td>
<td>45</td>
<td>150</td>
</tr>
<tr>
<td>4-2</td>
<td>F/M</td>
<td></td>
<td></td>
<td></td>
<td>Oral cavity</td>
<td>46</td>
<td>163</td>
</tr>
</tbody>
</table>

F, female; M, male; MDS, myelodysplastic syndromes; CML, chronic myelogenous leukemia; HSCT, hematopoietic stem cell transplantation; PBSCT, peripheral blood stem cell transplantation; BMT, bone marrow transplantation.
gender-mismatched HSCT using cells from a female donor (Table I, case no. 1). To assess normal epithelium in the esophagus, FISH was performed using gender chromosome-specific probes (Cy5 and SpectrumGreen for the X chromosome; SpectrumRed for the Y chromosome; Fig. 1). A comparison of FISH and histopathological analyses indicated that all infiltrating lymphocytes in an examination of 50 fields displayed green (X) signals but not red (Y) signals, suggesting that the lymphocytes were replaced with donor hematopoietic cells after HSCT (representative data in Fig. 1A-C; Tables II and III). The data indicated that both epithelial cells and basal cells were also positive for green (X) signals but not red (Y) signals, whereas mesenchymal cells in the stroma were positive for both green (X) and red (Y) signals, suggesting that the recipient cells were integrated in the mesenchymal tissues (Fig. 1A-C). The infiltration of lymphocytes with only green (X) signals in the epithelial and mesenchymal tissues was compatible with chronic GHVD. The present data propose that epithelial regions within the esophagus were replaced predominantly with donor-derived cells after HSCT (representative data in Fig. 1A-C; Tables II and III). An examination of the mesenchymal tissues indicated that stromal cells, which include fibroblasts as well as lymphocytes and coexist in flanking regions of epithelial cancer cells, displayed red (Y) signals showing the contribution of the recipient cells. On the other hand, approximately 30% of the stromal cells displayed green (X) signals but not red signals, suggesting the involvement of donor-derived cells. We found that tumor-associated fibroblasts displayed both green (X) and red (Y) signals, indicating that they were recipient cells (Fig. 1H and I).

**Significant contribution of BM to dysplasia of esophagus.** In case 2, the recipient was a male and the donor was a female. Esophageal cancer, which was characterized by dysplastic lesions, developed in the recipient 120 months after HSCT for malignant lymphoma, although the original disease
was in complete remission (Table I). As indicated in Fig. 2 (SpectrumGold for the X chromosome; SpectrumRed for the Y chromosome), FISH and histopathological analyses indicated that normal epithelial regions (lesion I in Fig. 2A) were composed predominantly of squamous cells as indicated by the yellow (X) and red (Y) signals, whereas a fraction displayed yellow (X) signals alone, suggesting that the esophageal epithelium was reconstituted at least partially with donor-derived cells in the regions examined. As indicated by the representative data in Fig. 2B-D (Tables II and III), the nuclei were similar to those of the surrounding epithelial cells in size, indicating that they were also epithelial cells. Infiltrated lymphocytes displayed only yellow (X) signals, indicating that the hematopoietic cells in the recipient were replaced by donor cells and the involvement of GVHD. Histopathological analysis of the surgical specimen from case 2 indicated dysplasia (lesion II in Fig. 2A) as a continuous lesion with the tumor (lesion III). FISH and histopathological analyses of lesion II indicated that almost all the dysplastic cells displayed only yellow (X) signals but not red (Y) signals (Fig. 2E-G; Tables II and III), suggesting that the donor-derived cells were recruited to the esophageal epithelium and transformed or that some damaged donor cells may have been incorporated into the esophagus.

Analysis of the tumor (lesion III; Fig. 2H-M; Tables II and III) showed both yellow (X) and red (Y) signals in almost all cancer cells, whereas a few cells expressed only yellow (X) signals (arrows in Fig. 2M; their nuclei were similar to those of the surrounding epithelial cancer cells in size), indicating that the tumor was composed predominantly of recipient cancer cells with a fraction of donor-derived cells. This was characterized by infiltrating lymphocytes displaying yellow (X) signals alone (arrows in Fig. 2J; the nuclei were much smaller than those of epithelial cells, indicating lymphocytes), thus being compatible with GVHD.

A mixed tumor of recipient- and donor-derived cells. In case 3, the donor was a female and the recipient was a male who developed oral SCC 76 months after HSCT for myelo-dysplastic syndrome (Table I). FISH and histopathological analyses indicated that all infiltrating lymphocytes displayed green (X) signals but not red (Y) signals, suggesting that the lymphocytes were replaced after transplantation; this was again compatible with GVHD (Cy5 for the X chromosome; SpectrumRed for the Y chromosome; Fig. 3). Analysis of the normal epithelium indicated that almost all epithelial cells displayed red (Y) signals (Fig. 3A-C; Tables II and III), whereas approximately 5% cells displayed green (X) signals alone. The nuclei of the latter cells were similar to those of the surrounding epithelial cells in size (Fig. 3C), indicating that normal epithelium in the surgical specimen of the oral cavity was composed predominantly of recipient cells. In the tumor, approximately 80% cancer cells expressed red (Y) signals (Fig. 3D-F; Tables II and III), whereas 20% cells displayed green (X) signals alone. The nuclei of the latter cells were similar size to those of the surrounding SCC cells in size (Fig. 3F), suggesting that the tumor had two different origins: a predominant contribution from the recipient cells and a partial contribution from the donor cells, which may have developed after HSCT presumably through the involvement of GVHD.

Sequentially occurring oral cancer after HSCT. We encountered a patient who developed oral SCC on different tongue regions 150 (case 4-1) and 163 months (case 4-2) after single HSCT for chronic myeloid leukemia. The recipient was a female and the donor was a male. FISH and histopathological analyses showed that all infiltrating lymphocytes displayed red (Y) signals, indicating that these lymphocytes were replaced after

Table II. Cell of origin.

<table>
<thead>
<tr>
<th>Case no.</th>
<th>Donor cells</th>
<th>Recipient cells</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>E</td>
<td>L</td>
</tr>
<tr>
<td>1</td>
<td>+</td>
<td>+</td>
</tr>
<tr>
<td>2</td>
<td>±</td>
<td>+</td>
</tr>
<tr>
<td>3</td>
<td>±</td>
<td>+</td>
</tr>
<tr>
<td>4-1</td>
<td>-</td>
<td>+</td>
</tr>
<tr>
<td>4-2</td>
<td>-</td>
<td>+</td>
</tr>
</tbody>
</table>

E, normal epithelial cells; L, peripheral mononuclear cells or lymphocytes; D, dysplasia; C, cancer cells; +, present; ±, less than 5% involved; -, absent; ND, not detected. The data summarizes the results of Table III.

Table III. Results of this study.

<table>
<thead>
<tr>
<th>Case no.</th>
<th>Normal epithelial cells</th>
<th>Cancer cells</th>
<th>Dysplasia</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Y</td>
<td>X</td>
<td>XY</td>
</tr>
<tr>
<td>1</td>
<td>0.0</td>
<td>94.3</td>
<td>0.0</td>
</tr>
<tr>
<td>2</td>
<td>36.1</td>
<td>18.1</td>
<td>41.0</td>
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<tr>
<td>3</td>
<td>43.8</td>
<td>30.8</td>
<td>24.6</td>
</tr>
<tr>
<td>4-1</td>
<td>0.0</td>
<td>57.7</td>
<td>0.0</td>
</tr>
<tr>
<td>4-2</td>
<td>0.0</td>
<td>18.9</td>
<td>0.0</td>
</tr>
</tbody>
</table>

Y, Y chromosome; X, X chromosome; XY, X and Y chromosomes; XX, two X chromosomes; ND, not detected. Scores in this table display percentages of each cell to all cells with signals. See Materials and methods. The data summarized in Table II.
transplantation and compatibility with GVHD (Cy5 for the X chromosome; SpectrumRed for the Y chromosome; Fig. 4). An analysis of normal epithelial cells and carcinoma cells revealed the expression of green (X) signals but not red (Y) signals (Figs. 4 and 5; Tables II and III). Red (Y) signals were detected only in cells with relatively small nuclei compared with those of the epithelial cells, indicating lymphocyte infiltration and compatibility with GVHD (Figs. 4C, 5C and F).

Figure 2. Recipient- and donor-derived esophageal epithelial cells and recipient-derived esophageal SCC from a male recipient (case 2) of peripheral blood stem cells from a female donor. (A) Tissue sections of the normal mucosa and solid tumor tissue are stained with hematoxylin and eosin. (B) A magnified image of normal mucosa (I). (C) Neighboring tissue sections of (B) are examined by FISH to determine the centromeres of the X (SpectrumGold, yellow) and Y (SpectrumRed, red) chromosomes. (D) A magnified image of the red-framed square in (C). A few donor-derived cells display XX signals, and their nuclei are similar to those of normal epithelial cells in size (white arrow). (E) A magnified image of the region with dysplasia (II). (F) Neighboring tissue sections of (E) are examined by FISH to determine the centromeres of the X (SpectrumGold, yellow) and Y (SpectrumRed, red) chromosomes. (G) A magnified image of the red-framed square in (F). No cell in the epithelium displays Y signals. (H) A magnified image of normal mucosa (III). (I) Neighboring tissue sections of (H) are examined by FISH to determine the centromeres of the X (SpectrumGold, yellow) and Y (SpectrumRed, red) chromosomes. (J) A magnified image of the red-framed square in (H). Infiltrating donor-derived cells display XX signals and smaller nuclei than those of normal epithelial cells (white arrow). (K) Tumor tissue sections are stained with hematoxylin and eosin. (L) Neighboring tissue sections of (K) are examined by FISH to determine the centromeres of the X (Cy5, green) and Y (SpectrumRed, red) chromosomes. (M) A magnified image of the cancer region, i.e., the red-framed square in (L). A few donor-derived cells display XX signals, and their nuclei are similar to those of SCC cells in size (white arrow).
Discussion

In this study, we utilized a highly sensitive combination of FISH using gender chromosome-specific probes and histopathological analyses to clearly demonstrate that human epithelial cancer of the head and neck can arise from donor cells after gender-mismatched hematopoietic transplantation (Tables II and III). In a case of esophageal cancer (case 1), almost all cancer cells were donor-derived in the regions examined, whereas a dysplastic lesion was composed exclusively of donor...
cells in case 2. Our observations indicated that donor BM contains various cells, including mesenchymal cells, and the collection of hematopoietic cells after HSCT caused those cells to be transferred to the recipient and contributed to the formation of epithelial structures (7,8). Alternatively, some epithelial cell types, even though normal, can free themselves from epithelial structures and migrate into the bloodstream through the well-known phenomenon of epithelial-mesenchymal transition, as occurs during the metastasis of solid tumors (10). Although the release frequency of normal epithelial cells from tissues under physiological conditions is uncertain, patients with solid tumors reportedly possess circulating epithelial cancer cells in the peripheral blood and BM (10). In the cases we examined, none of the donors developed late-onset malignancies during the observation period, which reinforced the fact that BM-derived or circulating mesenchymal stem cells, rather than apparent epithelial cells, in donors may change their phenotypes in recipient organs. This notion is compatible with recent studies in which circulating donor-derived BMSCs differentiated into epithelial cells and tumor cells after HSCT in mice and humans (11-30). Nevertheless, a mixture of recipient- and donor-derived origins was involved in cases 2 and 3; therefore, we hypothesized that the initiation and progression of esophageal carcinogenesis may occur in either recipient- or donor-derived single cells in the epithelium and that the adjacent cells may be recruited or accumulated against a background of inflammation and exposure to an immunosuppressant. However, we cannot exclude the possibility of simultaneous transformation in both recipient- and donor-derived cells.

BMSCs are known to differentiate into many types of cells. In the mouse, circulating BMSCs have been reported to differentiate into gastric mucosal cells, lung epithelial cells, renal epithelial cells, keratinocytes, hepatocytes, duct cells, astrocytes and neurons (3-7). In humans, it has been observed that BMSCs differentiate into buccal epithelial cells, keratinocytes, gastrointestinal tract cells, lung epithelial cells, hepatocytes, duct cells, astrocytes and neurons (8-11). In our cases of secondary epithelial cancers subsequent to HSCT, GVHD and exposure to immunosuppressants may have elicited the recruitment and accumulation of BMSC-derived epithelial cells in the tissues (5). The present study suggested the involvement of at least three components in the maintenance and carcinogenesis of head and neck tissues: tissue stem cells in the epithelial layer, circulating stem cells and HSCs.

In mice, BMSCs are reportedly involved in various types of solid tumor formation, including tumors of the epithelium, neural and muscle tissues, fibroblasts and blood vessel endothelium (12,14-25). In human oral SCC, BMSCs have been implicated in mucoepidermoid carcinoma of the parotid glands, invasive ductal carcinoma in the breast, papillary thyroid carcinoma, cervical carcinoma, Kaposi’s sarcoma, lung adenocarcinoma, skin SCC, glioblastoma multiforme and pharyngeal SCC (26-31). Although five previous cases suggested donor-derived tumor tissues (29,31), in the present study, we performed highly sensitive FISH using gender chromosome-specific probes and histopathological analyses in cases of head and neck tumors to clearly demonstrate the occurrence of donor-derived human esophageal cancer and dysplasia, a precancerous lesion. This is the first and definite examination to the best of our knowledge. We conclude that BMDCs can contribute to the constitution of epithelial tissues and further the occurrence of carcinogenesis stimulated by chronic inflammation and immunosuppressive conditions.

Figure 5. Recipient-derived oral epithelial cells and carcinoma cells from a female recipient (case 4-2) of bone marrow stem cells from a male donor. (A) Tissue sections of normal mucosa are stained with hematoxylin and eosin. (B) Neighboring tissue sections of (A) are examined by FISH to determine the centromeres of the X (Cy5, green) and Y (SpectrumRed, red) chromosomes. (C) A magnified image of the red-framed square in (B). A few infiltrating donor-derived cells display Y signals, and their nuclei are smaller than those of normal epithelial cells (white arrow). (D) Tumor tissue sections are stained with hematoxylin and eosin. (E) Neighboring tissue sections of (D) are examined by FISH to determine the centromeres of the X (Cy5, green) and Y (SpectrumRed, red) chromosomes. (F) A magnified image of the cancer region, i.e., the red-framed square in (E). A few infiltrating donor-derived cells display Y signals, and their nuclei are smaller than those of normal epithelial cells (white arrow).
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