CELL LINE

Establishment and characterization of NCC‑DDLPS1‑C1: a novel patient‑derived cell line of dediferentiated liposarcoma

Ryuto Tsuchiya^{1,2} · Yuki Yoshimatsu¹ · Rei Noguchi¹ · Akane Sei¹ · Fumitaka Takeshita³ · Jun Sugaya⁴ · ${\sf Suguru\ Fukushima}^4\cdot {\sf Akihiko\ Yoshida}^5\cdot {\sf Seiji\ Ohtori}^2\cdot {\sf Akira\ Kawai}^4\cdot {\sf Tadashi\ Kondo}^1$ ${\sf Suguru\ Fukushima}^4\cdot {\sf Akihiko\ Yoshida}^5\cdot {\sf Seiji\ Ohtori}^2\cdot {\sf Akira\ Kawai}^4\cdot {\sf Tadashi\ Kondo}^1$

Received: 15 July 2020 / Accepted: 13 September 2020 / Published online: 19 September 2020 © Japan Human Cell Society 2020

Abstract

Dediferentiated liposarcoma (DDLPS) is one of the four subtypes of liposarcomas; it is characterized by the amplifcation of the 12q13-15 region, which includes *MDM2* and *CDK4* genes. DDLPS has an extremely high local recurrence rate and is refractory to chemotherapy and radiation, which leads to poor prognosis. Therefore, a novel therapeutic strategy should be urgently established for improving the prognosis of DDLPS. Although patient-derived cell lines are important tools for basic research, there are no DDLPS cell lines available from public cell banks. Here, we report the establishment of a novel DDLPS cell line. Using the surgically resected tumor tissue from a patient with DDLPS, we established a cell line and named it NCC-DDLPS1-C1. The NCC-DDLPS1-C1 cells contained 12q13-15, 1p32, and 1q23 amplicons and highly expressed MDM2 and CDK4 proteins. NCC-DDLPS-C1 cells exhibited constant growth, spheroid formation, aggressive invasion, and tumorigenesis in mice. By screening a drug library, we identifed that the proteasome inhibitor, bortezomib, had inhibitory efects on the proliferation of NCC-DDLPS1-C1 cells. We concluded that the NCC-DDLPS1-C1 cell line may serve as a useful tool for basic and pre-clinical studies of DDLPS.

Keywords Sarcoma · Dediferentiated liposarcoma · Patient-derived cancer model · Patient-derived cell line

Electronic supplementary material The online version of this article [\(https://doi.org/10.1007/s13577-020-00436-5\)](https://doi.org/10.1007/s13577-020-00436-5) contains supplementary material, which is available to authorized users.

 \boxtimes Tadashi Kondo takondo@ncc.go.jp

- ¹ Division of Rare Cancer Research, National Cancer Center Research Institute, 5-1-1 Tsukiji, Chuo-ku, Tokyo 104-0045, Japan
- ² Department of Orthopaedic Surgery, Graduate School of Medicine, Chiba University, 1-8-1 Inohana, Chuo-ku, Chiba 260-8670, Japan
- ³ Department of Translational Oncology, Fundamental Innovative Oncology Core Center, National Cancer Center Research Institute, 5-1-1 Tsukiji, Chuo-ku, Tokyo 104-0045, Japan
- ⁴ Department of Musculoskeletal Oncology, National Cancer Center Hospital, 5-1-1 Tsukiji, Chuo-ku, Tokyo 104-0045, Japan
- ⁵ Department of Diagnostic Pathology, National Cancer Center Hospital, 5-1-1 Tsukiji, Chuo-ku, Tokyo 104-0045, Japan

Introduction

Dediferentiated liposarcoma (DDLPS) is one of the four subtypes of liposarcoma. Each subtype of a liposarcoma (atypical lipomatous tumor/well-diferentiated liposarcoma, dediferentiated liposarcoma, myxoid liposarcoma, and pleomorphic liposarcoma) has diferent clinical behaviors and genetic characteristics $[1, 2]$ $[1, 2]$ $[1, 2]$. DDLPS was first reported by Evans $[3]$; it is defned by the association of an atypical lipomatous tumor/ well-diferentiated liposarcoma (WDLPS) component with a non-lipogenic sarcoma of variable histological grade [\[1](#page-9-0), [2](#page-9-1)]. DDLPS is also characterized by the amplifcation of the 12q13- 15 region, which includes *MDM2* and *CDK4* genes [[4–](#page-9-3)[12](#page-9-4)]. Besides the amplification of 12q13-15, the gain of 1p32, 1q21-23, and 6q23-24 is occasionally recognized in DDLPS, which also relates with tumor transformation and progression [\[4](#page-9-3)[–8](#page-9-5), [11,](#page-9-6) [12](#page-9-4)]. Recently, several studies reported that poorly diferentiated sarcomas with MDM2 amplifcation, diagnosed earlier as malignant fbrous histiocytomas or undiferentiated pleomorphic sarcomas, are actually DDLPS, considering their clinical behaviors and molecular biology [[13](#page-9-7)[–16](#page-9-8)]. These studies suggested that sarcomas should be diagnosed by their unique genomic profling if they lack well-diferentiated components. And the latest edition of WHO classifcation of soft tissue and bone tumors mentioned that a well-diferentiated component may not be identifable [[1](#page-9-0)]. DDLPS often occurs in the retroperitoneum or deep inside the extremity muscles of patients aged > 40 years and rarely in children $[1, 2, 17]$ $[1, 2, 17]$ $[1, 2, 17]$ $[1, 2, 17]$. While WDLPS is defned as an intermediate tumor with rare metastasis, DDLPS is associated with a 15–20% metastatic rate and a nearly 50% recurrence rate [\[2](#page-9-1), [17\]](#page-9-9). Specifcally, DDLPS arising from the retroperitoneum has an extremely higher recurrence rate (80–90%), which leads to poor prognosis, than DDLPS arising from extremities [\[17](#page-9-9)[–19](#page-9-10)]. Despite over 40 years having passed since DDLPS was frst reported in 1979, there is still not enough evidence of the efectiveness of chemotherapy and radiation for the treatment of DDLPS [\[17](#page-9-9), [20](#page-9-11)]. As in the case of other sarcomas, adriamycin, docetaxel, or gemcitabinebased therapy might be used for DDLPS, but the response rates are low and such treatments rarely prolong survival [[17,](#page-9-9) [20](#page-9-11)]. Recent clinical trials have reported the efficacy of new drugs (e.g., tyrosine kinase inhibitor, MDM2 inhibitor, CDK4 inhibitor, and immune checkpoint inhibitor) [\[17](#page-9-9)]. However, considering its rarity, basic and clinical research have not been sufficient, and further studies are needed.

Patient-derived cancer cell lines are an important tool for basic and pre-clinical studies. Although long-term culturing is associated with alterations in the genetic profles of cell lines [\[21](#page-9-12)[–23\]](#page-9-13), those maintained for a short period often retain the original genetic and phenotype characteristics, which aid in examining the effects of novel drugs and evaluating the molecular mechanisms of cancer progression [[24–](#page-9-14)[28](#page-9-15)]. Therefore, there is a need to continuously establish new cell lines. Using the cell line database, Cellosaurus [\[29](#page-9-16)], we found that there were cell lines annotated as "liposarcoma." However, there were fewer cell lines annotated as "dediferentiated liposarcoma;" we identifed 23 DDLPS cell lines previously reported, but there were no DDLPS cell lines publicly available from cell banks (Supplementary Table 1). Further, because DDLPS is clinically diverse and requires more therapeutic options, there is a need to establish more patient-derived DDLPS cell lines.

Here, we report a novel DDLPS cell line, NCC-DDLPS1- C1, established from a surgically resected tumor tissue of a patient with DDLPS. We characterized NCC-DDLPS1-C1 cells and examined the antiproliferative efects of anti-cancer agents on this cell line.

Materials and methods

Patient history

The patient donor was a 57-year-old man with pleomorphic spindle cell sarcoma, most consistent with dediferentiated

liposarcoma. He visited the hospital with a major mass on his left lower abdomen. A computed tomography (CT) and a magnetic resonance imaging (MRI) detected a retroperitoneal tumor (Fig. [1](#page-2-0)a–d). A malignant tumor was suggested, and he was referred to the National Cancer Center Hospital (Tokyo, Japan). Subsequently, needle biopsy was done, which suggested dediferentiated liposarcoma. Although there was no evidence of metastatic lesions on CT and MRI scans, the tumor was too large and involved the surrounding organs. Therefore, the patient frst received neoadjuvant chemotherapy (AI; doxorubicin and ifosfamide). After three courses of chemotherapy, wide resection was performed. A part of the resected tumor was used to establish the cell line described in this study. Histological evaluation of the resected specimen demonstrated an appearance typical of dediferentiated liposarcoma (Fig. [1e](#page-2-0)–g); the spindle cells showed pleomorphic nuclei, prominent nucleoli, and frequent mitotic activity including atypical forms. MDM2/ CDK4 immunohistochemistry showed positive nuclear labeling, although cytoplasm is often co-stained for CDK4. However, there was no evidence of well-diferentiated components. These fndings led to the diagnosis of DDLPS. The resection margin was positive microscopically, and the patient received postoperative chemotherapy (AI). However, 3 months after the operation, peritoneum dissemination was detected on CT. The patient received further chemotherapy (eribulin); however, the disease progressed rapidly. Finally, the patient died 4 months after the operation. The ethical committee of National Cancer Center approved the use of clinical materials for this study, and written informed consent was obtained from the donor patient in this study.

Histological analysis

Histology examination was performed on 4 μm-thick sections from a representative parafnized tumor sample. The sections were deparaffinized and stained with hematoxylin and eosin (H&E).

Immunohistochemistry

Immunohistochemistry was also performed with the deparaffinized tumor sample. The sections were exposed to 3% hydrogen peroxide for 15 min to block endogenous peroxidase activity and then washed in deionized water for 2–3 min. Preparations were pretreated with heat-induced epitope retrieval. The primary antibodies were MDM2 (IF2, 1:100; Zymed Laboratories, San Francisco, CA, USA), and CDK4 (DCS-31, 1:200; Biosource International, Camarillo, CA, USA). The slides were incubated for 1 h at room temperature, and subsequently labeled

with peroxidase (EnVision system, Dako, Santa Clara, CA, USA).

Cell culture procedure

The surgically resected tumor tissue was used for establishing the cell line as previously described [\[30\]](#page-10-0). Fetal bovine serum was obtained from Gibco (Gibco, Grand Island, NY, USA). The cells were maintained for more than 3 months under tissue culture conditions and passaged more than 20 times.

Authentication and quality control of the established cell line

The established cell line was authenticated by examining short tandem repeats (STRs) in 10 loci by using the GenePrint 10 system (Promega, Madison, WI, USA) according to the manufacturer's instructions and the procedure described in our previous study [\[30\]](#page-10-0). Mycoplasma contamination was examined using the DNA in the tissue culture medium of the cell line as previously reported [[30\]](#page-10-0).

Single‑nucleotide polymorphism (SNP) array

SNP array genotyping was conducted with Infnium Omni-ExpressExome-8 v1.4 BeadChip (Illumina, Sn Diego, CA, USA) following the manufacturer's instructions and the procedure described in our previous study [[30](#page-10-0)]. Copy number analysis was performed using ASCAT (allele-specifc copy number analysis of tumor) (version 2.1) considering nonneoplastic cell infltration and tumor aneuploidy, and this resulted in integral allele-specifc copy number profles for the tumor cells.

Western blotting

The NCC-DDLPS1-C1 cell line, WDLPS cell lines (93T449 and 94T778; ATCC, Manassas, VA, USA), and normal human foreskin fbroblast cell line (HFF) were used for western blotting. The HFF cell line was kindly provided by Dr. Kiyono (Division of Carcinogenesis and Prevention, National Cancer Center Research Institute, Tokyo, Japan). Total proteins were extracted from each cell line using a lysis bufer (50-mM Tris–HCl at pH 7.5, 250-mM NaCl, 1% NP-40, 0.1% sodium dodecyl sulfate, 20% glycerol, 10-μM zinc chloride, 2-mM dithiothreitol, 80-mM β-glycerophosphate, 50-mM sodium fuoride, 1-mM sodium orthovanadate, and a complete protease inhibitor tablet [Roche, Basel, Switzerland]). Protein concentrations were determined using the DC Protein Assay Kit (Bio-Rad, Hercules, CA, USA). Protein lysates were electrophoresed on 12.5% SDS–polyacrylamide gels (e-PAGEL; ATTO corporation, Tokyo, Japan), and transferred to Immun-Blot PVDF

Table 1 Short tandem repeat analysis

Microsatellite (chromosome)	NCC-DDLPS1- C ₁	Tumor tissue
Amelogenin $(X Y)$	X, Y	X, Y
TH01 (3)	6	6
D21S11 (21)	32.2	32.2
D5S818(5)	12	12
D13S317 (13)	9, 12	9, 12
D7S820 (7)	9, 13	9, 13
D16S539 (16)	9, 13	9, 13
CSF1PO(5)	10	10
vWA (12)	14, 18	14, 18
TPOX(2)	8, 11	8, 11

membranes (Bio-Rad Laboratories, Inc., CA, USA). After incubation with 5% skim milk (BD Biosciences, Franklin Lakes, NJ, USA) in TBST (10-mM Tris, pH 8.0, 150-mM NaCl, 0.5% Tween 20) for 1 h, the membrane was first incubated with antibodies against MDM2 (1:100; Calbiochem, San Diego, CA, USA), CDK4 (1:1000; Santacruz, Dallas, Texas, USA) or β-actin (1:5000: abcam, Cambridge, UK) at 4 °C for 12 h. Membranes were washed 3 times for 10 min and then incubated with a 1:5000 dilution of horseradish peroxidase-conjugated anti-mouse or anti-rabbit antibodies (GE Healthcare, Chicago, IL, USA) for 1 h. After the reaction, the membranes were washed 3 times and treated with Western Lightning Plus-ECL (PerkinElmer, Waltham, MA, USA). Subsequently, the images were taken using Amershan Imager 600 (GE Healthcare). Relative expression levels of each protein were compensated by β-actin.

Cell proliferation assay

The tissue-cultured cells were seeded at 2×10^4 cells/well in 24-well culture plates at day 0, and the cell number was counted at multiple time points as previously described [\[30\]](#page-10-0). The doubling time was calculated based on the growth curve.

Spheroid formation assay

Spheroid formation was examined as previously described [[30\]](#page-10-0). Briefly, cells (1×10^5) were seeded into round-bottom 96-well ultra-low-attachment plates (96-well Clear Flat Bottom Ultra Low Attachment Microplate; Corning, Inc., Corning, NY, USA) in DMEM/F12 containing 10% FBS. The cells were maintained in a humidifed atmosphere of 5% CO₂ at 37 °C for 3 days, and the formation of spherical shaped colonies was confrmed microscopically (Keyence, Osaka, Japan). The colonies were prepared for paraffin sections using iPGell (Genostaff, Tokyo, Japan) according to the manufacturer's instruction. Cell blocks were fxed with 10% formalin neutral buffer solution and embedded in paraffin. Four-micrometer-thick paraffin sections were prepared and H&E stained. All assays were performed in triplicate.

Transwell cell invasion assay

Cell invasion assays were performed with BD BioCoat Matrigel Invasion Chambers (BD Biosciences) according to the manufacturer's instructions and the procedure described in our previous study [[30\]](#page-10-0) using NCC-DDLPS1-C1 and the osteosarcoma cell line, MG63 (JCRB Cell Bank, Ibaragi, Osaka, Japan).

Fig. 2 Study of the single nucleotide polymorphism (SNP) array. Allele-specifc copy number analysis revealed DNA copy number variations. SNP genotyping copy number profles for NCC-DDLPS1-

C1 cells showing logR and B allele frequency (BAF). The plot represents log R and BAF identifed with the ASCAT algorithm

Tumorigenesis in nude mice

All surgical procedures and care administered to the animals were in accordance with the institutional guidelines. A 100 μL volume of cells in a 1:1 mixture of Matrigel (BD Biosciences) was subcutaneously injected into female BALB/c nude mice (CLEA Japan, Inc., Tokyo, Japan) $(1 \times 10^6 \text{ cells})$. Subsequently, the tumor size was measured weekly. The tumor volume (mm³) was calculated as $L \times W^2 \times 0.52$, where *L* is the longest diameter and *W* is the shortest diameter. After 2 weeks, the tumors were surgically resected, and the specimens were stained with H&E. Three tumor samples, except one outlier, were used to describe the graph of tumor growth curve. All animal experiments were performed in accordance with the guidelines for Animal Experiments of the National Cancer Center and approved by the Institutional Committee for Ethics of Animal Experimentation.

Screening for the antiproliferative efects of anti‑cancer reagents

Screening for antiproliferative effects of anti-cancer reagents was performed according to the manufacturer's instructions and the procedure that we previously reported [[30](#page-10-0)]. We use 195 anti-cancer compounds, including FDA-approved drugs (Selleck Chemicals, Houston, TX, USA), and a list of the anti-cancer agents has been provided in Supplementary Table 2.

Dose–response experiments were also performed to validate available hits in the pilot screening according to the methods as previously reported $[30]$ $[30]$ $[30]$. IC₅₀, the sample concentration required to inhibit cell growth by 50% in comparison with the growth of the control cell, was determined from the dose–response curves.

Results

Authentication of the established cell line

The NCC-DDLPS1-C1 cell line was authenticated by analyzing the STR status of 10 microsatellites in the NCC-DDLPS1-C1 cells and the corresponding original tumor tissue. STR allele patterns were completely identical between the NCC-DDLPS1-C1 cells and the corresponding original tumor tissue (Table [1,](#page-3-0) Supplementary Figure 1). The STR patterns of NCC-DDLPS1-C1 cells did not match those of any cell lines in the public cell banks examined using a function of the cell line database, Cellosaurus [[29\]](#page-9-16). Thus, we concluded that NCC-DDLPS1-C1 was a novel cell line. We did not detect the DNA sequence unique to mycoplasma in the tissue culture medium of the NCC-DDLPS1-C1 cells (data not shown).

Fig. 3 Western blotting. **a** NCC-DDLPS1-C1 cells exhibiting overexpression of MDM2 and CDK4 proteins. **b**, **c** Graphs showing relative expression levels of each protein. Bars represent the mean standard error

Characterization of the cell line

SNP array analysis revealed multiple allelic duplications in NCC-DDLPS1-C1 cells (amplifed: 1p32, 1q23 and 12q13- 15; Fig. [2](#page-4-0), Supplementary Table 3). The amplifcation is characteristic of DDLPS, and the 12q13-15 region includes *MDM2* and *CDK4* genes. In western blotting, we used the WDLPS cell lines, which is known to highly express MDM2 and CDK4 proteins, as positive control, and the HFF cell line as negative control. We observed the overexpression of MDM2 and CDK4 proteins in NCC-DDLPS1-C1 cells by western blotting (Fig. [3a](#page-5-0)–c). These fndings demonstrate that NCC-DDLPS1-C1 cell lines maintain the unique characteristics of DDLPS and could be utilized for pre-clinical studies.

NCC-DDLPS1-C1 cells were composed of a mixed population of pleomorphic spindle and small round cells in 2D culture conditions (Fig. [4](#page-6-0)a, b). The proliferation of NCC-DDLPS1-C1 cells was monitored using a growth curve based on the number of cells. The population doubling time during the logarithmic growth phase was approximately 50 h (Fig. [4](#page-6-0)c). We confrmed that NCC-DDLPS1-C1 cells formed spheroids when they were seeded on a low-attachment microplate. The H&E stained spheroid section showed pleomorphic epithelioid cells with rhabdoid features (Fig. [4](#page-6-0)d). We also found that NCC-DDLPS1-C1 cells exhibited a more aggressive invasion ability than MG63 cells. The invasion ability of NCC-DDLPS1-C1 cells depended on the number of seeded cells and the time after seeding (Fig. [4](#page-6-0)e).

Tumorigenesis in nude mice

NCC-DDLPS1-C1 cells transplanted into BALB/c nude mice were able to undergo tumorigenesis under the

Fig. 4 Characterization of NCC-DDLPS1-C1 cells. **a**, **b** NCC-DDLPS1-C1 cells showing pleomorphic spindle appearance under 2D culture conditions. **c** Growth curve of NCC-DDLPS1-C1 cells. The *y*-axis indicates the relative cell proliferation of NCC-DDLPS1- C1 cells, and the *x*-axis represents the day after seeding. **d** The H&E stained spheroid section showing pleomorphic epithelioid cells with rhabdoid features. **e** The invasion ability of NCC-DDLPS1-C1 cells in comparison to that of MG63 osteosarcoma cells

described condition (Fig. [5a](#page-7-0)). The tumor consisted of dense proliferation of epithelioid to short spindle cells with nuclear atypia (Fig. [5b](#page-7-0)). The tumor size and volume consistently increased after the transplantation (Fig. [5](#page-7-0)c).

Sensitivity to anti‑cancer drugs

The response of NCC-DDLPS1-C1 cells to antiproliferative effects was examined using a library of 195 anti-cancer agents (Supplementary Table 2). The cells were treated with the anti-cancer agents at a fixed concentration of $10 \mu M$; the proliferation-suppressive efects are shown in Supplementary Table 4. Among the 195 anti-cancer agents examined,

4 agents that showed remarkable antiproliferative efects on NCC-DDLPS1-C1 cells were further examined to calculate their IC_{50} values. The IC_{50} values of these anti-cancer agents are listed in Table [2,](#page-7-1) and the growth curve based on which the IC_{50} values were calculated are demonstrated in Fig. [6](#page-8-0).

Discussion

DDLPS frequently relapses and has poor prognosis, and there is only limited evidence of an efective response to chemotherapy until now [[17,](#page-9-9) [20](#page-9-11)]. In particular, DDLPS arising from the retroperitoneum is extremely difficult to

Я

Fig. 5 Tumorigenesis in nude mice. **a** NCC-DDLPS1-C1 cells transplanted into BALB/c nude mice have the capability of tumorigenesis. **b** The tumor consisting of dense proliferation of epithelioid to short spindle cells with nuclear atypia. **c** The graph showing estimated tumor volume. Bars represent the mean standard error

- 50um

Table 2 Summary of half-maximal inhibitory concentration (IC50) values in the cells

be completely resected because of its anatomical position [\[17](#page-9-9)[–19](#page-9-10)]. To improve the prognosis of patients with DDLPS, establishing novel treatment strategies is urgently required. Patient-derived cell lines maintained for a short period retain their original characteristics. Therefore, these cell lines are indispensable tools to research the biology of tumors, which may lead to novel treatment methods. However, for DDLPS, only a few cell lines are available because of its rarity, and there are no means to obtain the cell lines from public cell banks. For those reasons, we established a novel DDLPS cell line, NCC-DDLPS1-C1, and demonstrated its utility.

The NCC-DDLPS1-C1 cell line was established from the retroperitoneal tumor. The site of origin of the tumor and patient's age were both typical for DDLPS [[1](#page-9-0), [2,](#page-9-1) [17](#page-9-9)]. Further, with the overexpression of MDM2 and CDK4 proteins, the tumor was considered as DDLPS. However, there was no

fnding of well-diferentiated components in the surgically resected specimen. Therefore, the tumor was diagnosed as pleomorphic spindle cell sarcoma, which is most consistent with DDLPS. In the retroperitoneum, it is difficult to distinguish between the well-diferentiated components and normal fat, which often results in inadequate resection of the tumor. Therefore, the anatomical location of the tumor may have prevented us from fnding the well-diferentiated components. Recently, because the molecular biology of DDLPS has been gradually revealed, it is often diagnosed as DDLPS by its characteristic gene amplifcation of 12q13-15 along with the overexpression of MDM2 and CDK4 proteins, if there are no well-diferentiated components [\[1,](#page-9-0) [13](#page-9-7)–[16](#page-9-8)]. In this case, the original tumor had the characteristics of DDLPS, and the NCC-DDLPS1-C1 cell line also preserved the genetic features. We, therefore, concluded that this cell line could be used as a DDLPS cell line.

Besides the amplifcation of 12q13-15, NCC-DDLPS1- C1 cells presented the amplifcation of 1p32 and 1q23, which was often observed in DDLPS patients (24%–60%) [[4–](#page-9-3)[8](#page-9-5), [11](#page-9-6)]. This feature also supported the fact that NCC-DDLPS1-C1 cells retain DDLPS characteristics. Hirata et al. reported that the amplifcation of 1p32 is associated with poor prognosis of DDLPS patients [\[4\]](#page-9-3). In this case, the patient unfortunately had rapid progression of DDLPS,

Fig. 6 Growth curves for IC_{50} value calculation of the investigated anti-cancer compounds. **a** NCC-DDLPS1-C1 cells treated with 195 anti-cancer agents at a concentration of 10 µM for 72 h. **b**–**e** Viability of the cells treated with 4 anti-cancer agents at diferent concentrations. The name of each anti-cancer agent is shown under the graph

and therefore, NCC-DDLPS1-C1 cells would be eligible for studying highly malignant DDLPS.

NCC-DDLPS1-C1 cells exhibited the typical morphology of spindle cell sarcoma, showing rapid proliferation and aggressive invasion ability, which might refect the high malignancy and poor prognosis of DDLPS in the donor patient. NCC-DDLPS1-C1 cells had the ability to form spheroids, which is useful for understanding the behavior of DDLPS in a 3D environment. The further characterization of xenografted tumor cells may be useful for the research, which requires the cell line xenografts. In addition, NCC-DDLPS1-C1 cells were adapted to nude mice under the described condition and showed similar morphology to the original tumor. Therefore, we consider that NCC-DDLPS1- C1 cells as useful both in vitro and in vivo experiments. However, the cell lines do not recapitulate all features of tumors in human body, and the complemental use of other models such as patient-derived xenografts may be required for research.

DDLPS is known to be refractory to chemotherapy, and NCC-DDLPS1-C1 cells also demonstrated resistance to most of the 195 anti-cancer compounds. In contrast, bortezomib exhibited a remarkable effect at the nanomolar level. Bortezomib is a frst-in-class proteasome inhibitor and is usually used for multiple myeloma [[31](#page-10-1)]. The efectiveness of the combination therapy of bortezomib and other anti-cancer drugs for osteosarcoma, gastrointestinal stromal tumor, and various sarcoma types was reported previously [[32,](#page-10-2) [33](#page-10-3)]. Bortezomib inhibits the 26S proteasome, and exerts its anti-tumor effects through multiple mechanisms involving NF-κβ, cell cycle proteins,

apoptosis-regulatory proteins, and endoplasmic reticulum stress [[31](#page-10-1)[–33\]](#page-10-3). In DDLPS, the amplifcation of cell cycle protein, CDK4, is related to its oncogenesis, and this fact might reflect the effectiveness of bortezomib. Furthermore, Perez et al. reported the efficacy of bortezomib in sarcomas with high expression levels of the MAP17 protein, including liposarcomas [[34](#page-10-4)]. MAP17 protein is a small non-glycosylated membrane protein and is reported to activate the Notch pathway [[35](#page-10-5)], which may also relate to the cellular toxicity of bortezomib on DDLPS. DDLPS has a lot of genetic aberrations and diversity, therefore further investigation will be required to reveal the pharmacological mechanisms on DDLPS.

The clinical features of DDLPS are highly diverse in the view of its genomic background [\[4](#page-9-3)], and the experimental fndings obtained from only a limited number of patientderived cell lines might be inadequate. Therefore, more cell lines from a larger number of patients are required for basic and pre-clinical studies. Because DDLPS is extremely rare and the means of obtaining DDLPS cell lines are limited, patient-derived cancer cell lines should be shared within the research community.

We conclude that NCC-DDLPS1-C1 may serve as a useful patient-derived sarcoma cell line and contribute to reveal further molecular backgrounds of DDLPS.

Acknowledgements We thank Drs. F Nakatani, E Kobayashi, M Nakagawa, T Komatsubara, M Saito, C Sato (Department of Musculoskeletal Oncology), and Drs. T Shibayama, and H Tanaka (Department of Diagnostic Pathology), National Cancer Center Hospital, for sampling tumor tissue specimens from surgically resected materials. We thank Dr. T Kiyono (Division of Carcinogenesis and Prevention) for kindness in providing the HFF cell line. We also appreciate the technical assistance provided by Mr. K Tanoue, T Ono, and Ms. Y Kuwata (Division of Rare Cancer Research). We appreciate the technical support provided by Ms Y Shiotani, Mr. N Uchiya, and Dr. T Imai (Central Animal Division, National Cancer Center Research Institute). We would also like to thank Editage [\(https://www.editage.jp\)](https://www.editage.jp) for their help with English language editing and their constructive comments on the manuscript. This research was technically assisted by the Fundamental Innovative Oncology Core in the National Cancer Center.

Funding This research was supported by the Japan Agency for Medical Research and Development (Grant number 20ck0106537h0001).

Compliance with ethical standards

Conflict of interest The authors declare that they have no confict of interest.

Informed consent Written informed consent for publication was provided by the patient.

Ethics approval The ethical committee of the National Cancer Center approved the use of clinical materials for this study with the approval number 2004-050. The study using animal models was approved by the ethics committee of the National Cancer Center Research Institute (T13-016).

References

- 1. The WHO Classifcation of Tumors Editorial Board. WHO classifcation of tumours of soft tissue and bone. 5th ed. Lyon: IARC; 2020.
- 2. Dei Tos AP. Liposarcomas: diagnostic pitfalls and new insights. Histopathology. 2014;64:38–52.
- 3. Evans HL. Liposarcoma: a study of 55 cases with a reassessment of its classifcation. Am J Surg Pathol. 1979;3:507–23.
- 4. Hirata M, Asano N, Katayama K, et al. Integrated exome and RNA sequencing of dediferentiated liposarcoma. Nat Commun. 2019;10:5683.
- 5. Kanojia D, Nagata Y, Garg M, et al. Genomic landscape of liposarcoma. Oncotarget. 2015;6:42429–44.
- 6. Asano N, Yoshida A, Mitani S, et al. Frequent amplifcation of receptor tyrosine kinase genes in well diferentiated/dediferentiated liposarcoma. Oncotarget. 2017;8:12941–52.
- 7. Tap WD, Eilber FC, Ginther C, et al. Evaluation of well-diferentiated/de-diferentiated liposarcomas by high-resolution oligonucleotide array-based comparative genomic hybridization. Genes Chromosom Cancer. 2011;50:95–112.
- 8. Louis-Brennetot C, Coindre JM, Ferreira C, Perot G, Terrier P, Aurias A. The CDKN2A/CDKN2B/CDK4/CCND1 pathway is pivotal in well-diferentiated and dediferentiated liposarcoma oncogenesis: an analysis of 104 tumors. Genes Chromosom Cancer. 2011;50:896–907.
- 9. Italiano A, Bianchini L, Gjernes E, et al. Clinical and biological signifcance of CDK4 amplifcation in well-diferentiated and dedifferentiated liposarcomas. Clin Cancer Res. 2009;15:5696–703.
- 10. Italiano A, Bianchini L, Keslair F, et al. HMGA2 is the partner of MDM2 in well-diferentiated and dediferentiated liposarcomas whereas CDK4 belongs to a distinct inconsistent amplicon. Int J Cancer. 2008;122:2233–41.
- 11. Snyder EL, Sandstrom DJ, Law K, et al. c-Jun amplifcation and overexpression are oncogenic in liposarcoma but not always sufficient to inhibit the adipocytic differentiation programme. J Pathol. 2009;218:292–300.
- 12. Mariani O, Brennetot C, Coindre JM, et al. JUN oncogene amplification and overexpression block adipocytic differentiation in highly aggressive sarcomas. Cancer Cell. 2007;11:361–74.
- 13. Coindre JM, Mariani O, Chibon F, et al. Most malignant fbrous histiocytomas developed in the retroperitoneum are dediferentiated liposarcomas: a review of 25 cases initially diagnosed as malignant fbrous histiocytoma. Mod Pathol. 2003;16:256–62.
- 14. Coindre JM, Hostein I, Maire G, et al. Infammatory malignant fbrous histiocytomas and dediferentiated liposarcomas: histological review, genomic profle, and MDM2 and CDK4 status favour a single entity. J Pathol. 2004;203:822–30.
- 15. Chibon F, Mariani O, Derré J, et al. A subgroup of malignant fbrous histiocytomas is associated with genetic changes similar to those of well-diferentiated liposarcomas. Cancer Genet Cytogenet. 2002;139:24–9.
- 16. Le Guellec S, Chibon F, Ouali M, et al. Are peripheral purely undiferentiated pleomorphic sarcomas with MDM2 amplification dedifferentiated liposarcomas? Am J Surg Pathol. 2014;38:293–304.
- 17. Gahvari Z, Parkes A. Dediferentiated liposarcoma: systemic therapy options. Curr Treat Options Oncol. 2020;21:15.
- 18. Gootee J, Aurit S, Curtin C, Silberstein P. Primary anatomical site, adjuvant therapy, and other prognostic variables for dediferentiated liposarcoma. J Cancer Res Clin Oncol. 2019;145:181–92.
- 19. Keung EZ, Hornick JL, Bertagnolli MM, Baldini EH, Raut CP. Predictors of outcomes in patients with primary retroperitoneal dediferentiated liposarcoma undergoing surgery. J Am Coll Surg. 2014;218:206–17.
- 20. Crago AM, Singer S. Clinical and molecular approaches to well diferentiated and dediferentiated liposarcoma. Curr Opin Oncol. 2011;23:373–8.
- 21. Ben-David U, Siranosian B, Ha G, et al. Genetic and transcriptional evolution alters cancer cell line drug response. Nature. 2018;560:325–30.
- 22. Gisselsson D, Lichtenzstejn D, Kachko P, Karlsson J, Manor E, Mai S. Clonal evolution through genetic bottlenecks and telomere attrition: Potential threats to in vitro data reproducibility. Genes Chromosom Cancer. 2019;58:452–61.
- 23. Saito S, Morita K, Kohara A, et al. Use of BAC array CGH for evaluation of chromosomal stability of clinically used human mesenchymal stem cells and of cancer cell lines. Hum Cell. 2011;24:2–8.
- 24. Ben-David U, Beroukhim R, Golub TR. Genomic evolution of cancer models: perils and opportunities. Nat Rev Cancer. 2019;19:97–109.
- 25. Hattori E, Oyama R, Kondo T. Systematic review of the current status of human sarcoma cell lines. Cells. 2019;8:157.
- 26. Barretina J, Caponigro G, Stransky N, et al. The Cancer Cell Line Encyclopedia enables predictive modelling of anticancer drug sensitivity. Nature. 2012;483:603–7.
- 27. Crystal AS, Shaw AT, Sequist LV, et al. Patient-derived models of acquired resistance can identify efective drug combinations for cancer. Science. 2014;346:1480–6.
- 28. Teicher BA, Polley E, Kunkel M, et al. Sarcoma cell line screen of oncology drugs and investigational agents identifes patterns associated with gene and microRNA expression. Mol Cancer Ther. 2015;14:2452–62.
- 29. Bairoch A. The cellosaurus, a cell-line knowledge resource. J Biomol Tech. 2018;29:25–38.
- 30. Yoshimatsu Y, Noguchi R, Tsuchiya R, et al. Establishment and characterization of NCC-CDS2-C1: a novel patient-derived cell line of CIC-DUX4 sarcoma. Hum Cell. 2020;33:427–36.
- 31. Gandolf S, Laubach JP, Hideshima T, Chauhan D, Anderson KC, Richardson PG. The proteasome and proteasome inhibitors in multiple myeloma. Cancer Metastasis Rev. 2017;36:561–84.
- 32. Deming DA, Ninan J, Bailey HH, et al. A Phase I study of intermittently dosed vorinostat in combination with bortezomib in patients with advanced solid tumors. Investig New Drugs. 2014;32:323–9.
- 33. Xian M, Cao H, Cao J, et al. Bortezomib sensitizes human osteosarcoma cells to adriamycin-induced apoptosis through ROSdependent activation of p-eIF2alpha/ATF4/CHOP axis. Int J Cancer. 2017;141:1029–41.
- 34. Perez M, Peinado-Serrano J, Garcia-Heredia JM, et al. Efficacy of bortezomib in sarcomas with high levels of MAP17 (PDZK1IP1). Oncotarget. 2016;7:67033–46.
- 35. Garcia-Heredia JM, Lucena-Cacace A, Verdugo-Sivianes EM, Perez M, Carnero A. The cargo protein MAP17 (PDZK1IP1) regulates the cancer stem cell pool activating the Notch pathway by abducting NUMB. Clin Cancer Res. 2017;23:3871–83.

Publisher's Note Springer Nature remains neutral with regard to jurisdictional claims in published maps and institutional affiliations.