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# Identification of an H2-K<sup>b</sup> or H2-D<sup>b</sup> restricted and glypican-3-derived cytotoxic T-lymphocyte epitope peptide

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**Abstract.** Glypican-3 (GPC3) is overexpressed in human hepatocellular carcinoma (HCC) but not expressed in normal tissues except for placenta and fetal liver and therefore is an ideal target for cancer immunotherapy. In this study, we identified an H2-K<sup>b</sup> or H2-D<sup>b</sup> restricted and murine GPC3 (mGPC3)-derived cytotoxic T-lymphocyte (CTL) epitope peptide in C57BL/6 (B6) mice, which can be used in the design of preclinical studies of various therapies with GPC3-target immunotherapy *in vivo*. First, 11 types of 9- to 10-mer peptides predicted to bind with H2-K<sup>b</sup> or H2-D<sup>b</sup> were selected from the mGPC3 amino acid sequence based on the binding score as calculated by the BIMAS software. We evaluated the peptide-binding affinity and confirmed that all peptides were able to bind to H2-K<sup>b</sup> or H2-D<sup>b</sup> by *in vitro* cellular binding assay. Subsequently, a mixed peptide vaccine and single peptide vaccine were given to B6 mice to evaluate immunogenic potential of the 11 selected peptides. Using the splenocytes from peptide-vaccinated mice, interferon (IFN)- $\gamma$  enzyme-linked immunospot (ELISPOT) assays showed that mGPC3-1<sub>127-136</sub> (AMFKNNYPSL) peptide was the most efficient for inducing CTLs among the 11 peptides. Next, we demonstrated that the mGPC3-1 peptide-specific CTL line could recognize mGPC3-expressing cancer cells, suggesting that mGPC3-1 peptide was an endogenously presented peptide. In conclusion, we identified mGPC3-1 as an H2-K<sup>b</sup> or H2-D<sup>b</sup> restricted, mGPC3-derived CTL epitope peptide.

## Introduction

Liver cancer ranks fifth in frequency in the world and is the third most common cause of lethal cancer (1). Liver cancer consists of hepatocellular carcinoma (HCC) and intrahepatic

cholangiocarcinoma (ICC), with HCC as the most common. Regarding HCC therapy, hepatectomy, percutaneous local therapy and transcatheter arterial embolization (TAE) are common, but the recurrence rate with conventional therapies for advanced HCC patients is still high (2). Therefore, developing a novel curative therapy or an effective adjuvant therapy for HCC is important.

Recently, immunotherapy, which consists of a peptide vaccine, protein vaccine, or DNA vaccine, has become a potentially promising option for HCC (3,4). Many tumor antigen-derived peptides recognized by cytotoxic T-lymphocyte (CTL) have been identified (5). However, to date, vaccine therapy using these peptides has not proven adequate antitumor efficacy in clinical trials for advanced HCC patients (6-8).

In HCC, glypican-3 (GPC3) is overexpressed and is not expressed in normal tissues except for the placenta and embryonic liver (9). Hence, GPC3 is a novel target molecule in HCC patients. GPC3 is a member of the heparan sulfate proteoglycan family and the glypican family regulates cell growth and division through Wnt signaling, Hedgehogs, fibroblast growth factors and bone morphogenetic proteins (10-12). We previously identified HLA-A\*24:02-restricted GPC3<sub>298-306</sub> (EYILSLEEL) and HLA-A\*02:01-restricted GPC3<sub>144-152</sub> (FVGEFFTDV) peptides and showed that both peptides can induce GPC3-specific CTLs without an auto-immune response (13,14). Clinical trials of a GPC3-derived peptide vaccine for HCC patients are currently in progress. The phase I clinical trial of a GPC3-derived peptide vaccine for advanced HCC showed safety as well as immunological evidence and potential for improving overall survival (15-17). The phase I clinical trial suggested that the GPC3-derived peptide vaccine could be an attractive approach for treatment of HCC, however, the effect of tumor reduction was limited. Therefore, further studies are needed to enhance the effect of GPC3-targeted immunotherapy and to establish a GPC3-specific CTL-inducible mouse model. We previously conducted a preclinical study of the GPC3-derived peptide vaccine using HLA-A2.1 transgenic mice (18). The treatment model experiment using HLA transgenic mice is limited.

Mice with the C57BL/6 (B6) background have been reported to spontaneously develop liver cancer (19,20). Recently, the NASH mouse model (named STAM mice C57BL/6N-NASH), which had a B6 background and spontaneously developed liver

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cancer, was exploited by Stelic Institute & Co. In this mouse model, the cancer incidence rate is high and cancer incident time is short, thus, STAM mice C57BL/6N-NASH is an attractive model for studying GPC3-targeted therapy for HCC. Therefore, identification of a mouse major histocompatibility complex (MHC) class I epitope peptide to induce GPC3-specific CTL was needed for establishment of the appropriate mouse model.

Strategies to identify epitope peptides have previously been reported (21-24). A summary of our strategy follows. First, peptides binding MHC class I epitope were predicted from antigen amino acid sequences *in silico* by prediction software and the ability of the predicted peptides to bind MHC class I was confirmed *in vitro* by a binding assay. Then, the immunogenic potential of the predicted peptides was examined by *in vivo* immunization or *in vitro* stimulation. Lastly, whether peptides that have immunogenic potential are presented by cells endogenously expressing the antigen was confirmed. In summary, we identified peptides with immunogenic potential that were presented by cells endogenously expressing the antigen. We attempted to identify H2-K<sup>b</sup> or H2-D<sup>b</sup> restricted, GPC3-derived CTL epitope peptides in C57BL/6 mice based on the above strategy.

## Materials and methods

**Mice.** C57BL/6 (B6) mice were purchased from Charles River Laboratories Japan, Inc. and STAM mice C57BL/6N-NASH were a gift from this company. Mice were maintained under the institutional guidelines set by the Animal Research Committee of the National Cancer Center Hospital East. Mice were housed in specific pathogen-free (SPF) conditions with a 12-h light cycle and food and water at *ad libitum*. Six to eight-week-old female B6 mice were used in all experiments and STAM mice C57BL/6N-NASH were provided with a very high-fat rodent diet (rodent diet with 60% kcal% fat, Research Diet Inc.). All animal procedures were performed according to the guidelines for Animal Research Committee of the National Cancer Center, Japan.

**Cell lines and transfection.** B6 thymoma RMA and RMA-S cell lines, which have H2-K<sup>b</sup> and -D<sup>b</sup> as MHC class I epitopes, were maintained in our laboratory. RMA-S is an antigen processing-defective cell line and the cells cannot present endogenous antigens with MHC class I epitopes (25). To obtain RMA transiently expressing murine GPC3 (RMA-GPC3-puro), RMA (GPC3-negative) was transfected with pCAGGS-mGPC3-internal ribosomal entry site (IRES)-puromycin-resistant (puro-R) using Lipofectamine 2000 reagent (Invitrogen Corp., Carlsbad, CA, USA) according to the manufacturer's protocols. As negative control, RMA, which was transfected with pCAGGS-IRES-puro-R in a similar way, was named RMA-puro. Expression of murine GPC3 (mGPC3) in RMA-GPC3-puro or RMA-puro was confirmed by reverse transcription polymerase chain reaction (RT-PCR). All cells were cultured in RPMI-1640 (Gibco, USA) supplemented with 10% fetal bovine serum (FBS) (Gibco).

**RT-PCR.** Total ribonucleic acid was isolated from RMA-GPC3-puro or RMA-puro homogenized with the TRIzol Reagent

(Life Technologies, Inc., Rockville, MD, USA) according to the manufacturer's protocols. The first-strand complementary deoxyribonucleic acid (cDNA) was synthesized with a PrimeScript<sup>®</sup> II 1st strand cDNA Synthesis kit (Takara Bio Inc., Japan), then mGPC3 was amplified using a Takara PCR Amplification kit (Takara Bio Inc.). The amplification protocol was as follows: 150 sec at 94°C for initial denaturation, 35 amplification cycles at 58°C for 40 sec and 72°C for 40 sec, followed by a final extension at 72°C for 5 min. The primer sequences for mGPC3 were as follows: sense, 5'-ACGGGATGGTGAAA GTGAAGA-3' and antisense, 5'-GAAAGAGAAAAGAGGGA AACA-3'. The primer sequences for  $\beta$ -actin were as follows: sense, 5'-GAGCAATGATCTTGATCTTCAT-3' and antisense, 5'-TCCATCATGAAGTGTGACGT-3'. PCR products were visualized by ethidium bromide staining after separation on a 1% agarose gel. After normalization using  $\beta$ -actin messenger ribonucleic acid (mRNA) as a control, we compared the expression of mGPC3 mRNA.

**Generation of bone marrow-derived dendritic cells (BM-DCs) from BM cells.** BM cells ( $4 \times 10^6$ ) from B6 mice were cultured in RPMI-1640 containing FBS (10%), 2-mercaptoethanol (2-ME, 50  $\mu$ M) and murine granulocyte macrophage colony-stimulating factor (mGM-CSF, 20 ng/ml) for 1 week.

**Peptides.** Eleven types of 9- to 10-mer peptides predicted to bind with H2-K<sup>b</sup> or H2-D<sup>b</sup> were selected from mGPC3 amino acid sequences (accession code AAH36126) based on the binding score as calculated by BIMAS software (Bioinformatics and Molecular Analysis Section, Center for Information Technology, NIH, Bethesda, MD, USA) and 11 synthetic peptides (custom ordered) were purchased from Scrum Inc. (Tables I and II). The 11 amino acid sequences were as follows: mGPC3-1, AMFKNNYPSL; mGPC3-2, SLFPVIYTQM; mGPC3-3, LFPVIYTQM; mGPC3-4, KSFINFYSAL; mGPC3-5, LTARLNMEQL; mGPC3-6, LGSDINVDDM; mGPC3-7, QYVQKNGGKL; mGPC3-8, YVQKNGGKL; mGPC3-9, DTLCWNGQEL; mGPC3-10, RNGMKNQFNL; mGPC3-11, MKNQFNLHEL. Each peptide was dissolved in dimethyl sulfoxide (DMSO) (Wako Pure Chemical Industries, Japan) and each peptide's density was 10 mg/ml.

**H2-K<sup>b</sup> or H2-D<sup>b</sup> binding assay.** To evaluate the binding affinity of the predicted peptides to H2-K<sup>b</sup> or H2-D<sup>b</sup> molecules, an *in vitro* cellular binding assay was performed as previously reported (23,26). Briefly, after incubation of RMA-S cells in culture medium at 26°C overnight, cells ( $1 \times 10^6$ ) were washed with PBS and suspended in 100  $\mu$ l Opti-MEM<sup>®</sup> (Invitrogen) with or without 10  $\mu$ g peptide, followed by incubation at 26°C for 3 h and then at 37°C for 3 h. After washing with PBS, H2-K<sup>b</sup> or H2-D<sup>b</sup> expression was measured with a BD FACSCanto<sup>™</sup> II flow cytometer (BD) using FITC-conjugated H2-K<sup>b</sup> (BioLegend Inc., AF6-88.5) or H2-D<sup>b</sup> (BioLegend Inc., KH95) specific monoclonal antibody and mean fluorescence intensity (MFI) was recorded. Percent MFI increase was calculated as follows: percent MFI increase = (MFI with the given peptide - MFI without peptide)/(MFI without peptide) x 100.

**Vaccination.** The mixed peptide vaccine per mouse consisted of 5  $\mu$ l mGPC3-1 to mGPC3-11 solution, 55  $\mu$ l sodium bicar-

Table I. Synthetic peptides predicted to bind with H2-K<sup>b</sup>.

	Peptide sequence (position)	Binding score <sup>a</sup>
mGPC3-1	AMFKNNYPSL (127-136)	52.8
mGPC3-2	SLFPVIYTQM (172-181)	44
mGPC3-3	LFPVIYTQM (173-181)	66
mGPC3-4	KSFINFYSAL (395-404)	40

<sup>a</sup>Binding scores were estimated by using BIMAS software ([http://www.bimas.cit.nih.gov/molbio/hla\\_bind/](http://www.bimas.cit.nih.gov/molbio/hla_bind/)).

Table II. Synthetic peptides predicted to bind with H2-D<sup>b</sup>.

	Peptide sequence (position)	Binding score <sup>a</sup>
mGPC3-5	LTARLNMEQL (82-91)	200
mGPC3-1	AMFKNNYPSL (127-136)	343.2
mGPC3-6	LGSDINVDDM (156-165)	260
mGPC3-7	QYVQKNGGKL (331-340)	720
mGPC3-8	YVQKNGGKL (332-340)	240
mGPC3-9	DTLCWNGQEL (418-127)	600
mGPC3-10	RNGMKNQFNL (437-446)	200
mGPC3-11	MKNQFNLHEL (440-449)	288

<sup>a</sup>Binding scores were estimated by using BIMAS software ([http://www.bimas.cit.nih.gov/molbio/hla\\_bind/](http://www.bimas.cit.nih.gov/molbio/hla_bind/)).

bonate solution and 110  $\mu$ l incomplete Freund's adjuvant (IFA). Single peptide vaccine per mouse consisted of 5  $\mu$ l peptide, 45  $\mu$ l sodium bicarbonate solution and 50  $\mu$ l IFA. Each vaccine solution was emulsified. The mice were immunized by intradermal injection at the base of the tail every 7 days for a total of two vaccinations. Similarly, STAM mice C57BL/6N-NASH were immunized seven times with the mGPC3-1 peptide vaccine.

**Restimulation of splenocytes obtained from immunized mice.** Seven days after the last immunization, splenocytes were collected and cluster of differentiation 8 (CD8) positive splenocytes were isolated by positive selection with anti-CD8 microbeads (Miltenyi Biotec) according to the manufacturer's protocol. CD8-positive splenocytes were cocultured with BM-DCs pulsed with each peptide as previously described (13). Seven days after coculture, the detection of antigen-specific T cells producing interferon (IFN)- $\gamma$  was performed using the BD ELISPOT kit (BD Bioscience, San Jose, CA, USA) according to the manufacturer's protocols.

**Establishment of GPC3-1-specific CTL line.** The GPC3-1-specific CTL line was established as previously described (27). Splenocytes ( $1 \times 10^4$ ) derived from B6 mice immunized with the GPC3-1 peptide vaccine were cocultured with B6-derived and irradiated (35 Gy) splenocytes ( $5 \times 10^4$ ) in RPMI-1640 contained with FBS (10%), sodium pyruvate (1 mM, Gibco),

MEM non-essential amino acid solution (1X, Gibco) and 2-ME (50  $\mu$ M). Seven days later, recombinant interleukin-2 (rIL-2, 50 U/ml, Nipro, Osaka, Japan) was added to the culture medium.

**IFN- $\gamma$  enzyme-linked immunospot (ELISPOT) analysis.** IFN- $\gamma$  ELISPOT assay was performed according to the manufacturer's protocols. Briefly, restimulated CD8-positive splenocytes ( $5 \times 10^4$ ) as target cells were added to the plate and then BM-DCs ( $5 \times 10^4$ ) pulsed with each peptide (10  $\mu$ g/ml) as effector cells or non-pulsed BM-DCs ( $5 \times 10^4$ ) as control and target cells were added to the plate, which was then incubated for 20 h at 37°C, 5% CO<sub>2</sub>. Using the GPC3-1-reactive CTL line ( $1 \times 10^5$ ) as effector cells, RMA-S ( $5 \times 10^4$ ) pulsed with each peptide (10  $\mu$ g/ml) as target cells and non-pulsed RMA-S as control and target cells ( $5 \times 10^4$ ), the plate was incubated for 20 h at 37°C, 5% CO<sub>2</sub>. Using the mGPC3-1-reactive CTL line ( $1 \times 10^5$ ) as effector cells, RMA-GPC3-puro as target cells ( $5 \times 10^5$ ) and RMA-puro ( $5 \times 10^5$ ) as control and target cells, the plate was incubated for 48 h at 37°C, 5% CO<sub>2</sub>. The number of spots was automatically counted using the Eliphoto system (Minerva Tech, Tokyo, Japan).

**Cytotoxicity assay.** Cytotoxic activity against target cells was analyzed using the Terascan VPC system (Minerva Tech) as previously described (28). Target cells were incubated with calcein AM (Dojindo, Kumamoto, Japan) solution for 30 min at 37°C and labeled. Then the labeled cells were incubated with effector cells for 4 h. Fluorescence intensity was measured before and after the culture and specific cytotoxic activity was evaluated using the following formula: % cytotoxicity =  $\{1 - [(average\ fluorescence\ of\ the\ sample\ wells - average\ fluorescence\ of\ the\ maximal\ release\ control\ wells) - (average\ fluorescence\ of\ the\ minimal\ release\ control\ wells - average\ fluorescence\ of\ the\ maximal\ release\ control\ wells)]\} \times 100\%$ .

**Statistical analysis.** Statistical analyses were performed with a Mann-Whitney U test (n=3). Significant differences were defined as \*p<0.05 or R<sup>2</sup>>0.5.

## Results

**Evaluation of selected peptide-binding affinity to H2-K<sup>b</sup> or H2-D<sup>b</sup>.** The selected 11 peptides derived from mGPC3 by the BIMAS software were evaluated by an *in vitro* binding assay to determine each peptide's binding affinity to H2-K<sup>b</sup> or H2-D<sup>b</sup>. The peptide with the highest binding affinity for H2-K<sup>b</sup> was mGPC3-2 (percent MFI, 376.6%), followed by the mGPC3-3 peptide (128.0%) and the mGPC3-1 peptide (72.7%) (Fig. 1A). That for H2-D<sup>b</sup> was mGPC3-10 peptide (539.1%) followed by the mGPC3-1 peptide (298.2%) and the mGPC3-8 peptide (191.1%) (Fig. 1B). These results show that all 11 peptides could bind H2-K<sup>b</sup> or H2-D<sup>b</sup>, although the binding score calculated by the BIMAS software did not correlate with the actual binding affinity (Fig. 1C and D).

**Induction of CTL response against mGPC3-derived peptides in B6 mice.** The vaccine schedule was performed as follows (Fig. 2A): At days 0 and 7, peptide vaccine was given. At day 14, primed mice were sacrificed and CD8-positive

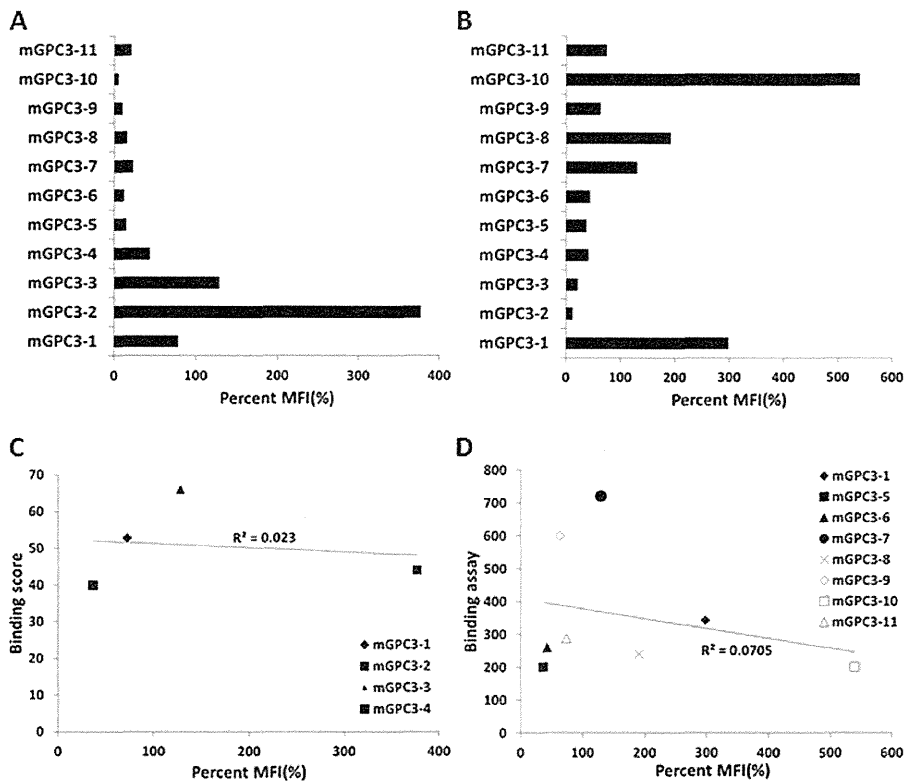


Figure 1. *In vitro* cellular peptide binding assays to H2-K<sup>b</sup> (A) or H2-D<sup>b</sup> (B) were performed using a FACS system. Comparison of BIMAS binding score with percent MFI for H2-K<sup>b</sup> (C) or H2-D<sup>b</sup> (D). Percent MFI increase = (MFI with the given peptide - MFI without peptide)/(MFI without peptide) x 100.

splenocytes were collected. CD8-positive splenocytes were restimulated with BM-DCs pulsed with each peptide. At day 21, the peptide's immunogenic potential was evaluated by IFN- $\gamma$  ELISPOT assay.

The mixed peptide vaccination was performed to evaluate immunogenic potential of the 11 peptides and IFN- $\gamma$  ELISPOT assays were performed using BM-DCs pulsed with each peptide and non-pulsed BM-DCs as target cells. The CD8-positive splenocytes from mice primed with the mixed vaccine released more IFN- $\gamma$  to BM-DCs pulsed with mGPC3-1 peptide (average number of spots,  $44.3 \pm 15.3$ ) and mGPC3-4 peptide (average number of spots,  $7.6 \pm 3.2$ ) than to non-pulsed BM-DCs (average number of spots,  $0.3 \pm 0.5$ ). These results suggest that the mGPC3-1 and mGPC3-4 peptides had immunogenic potential and were able to induce peptide-specific CTLs in B6 mice primed by the mixed vaccine system (Fig. 2B and C).

Next, to confirm whether the peptides are CTL-inducible peptides, a single peptide vaccine was given and IFN- $\gamma$  ELISPOT assays were performed using BM-DCs pulsed with either peptide and non-pulsed BM-DCs as target cells. The CD8-positive cells from mice immunized with mGPC3-1 peptide released more IFN- $\gamma$  to BM-DCs pulsed with mGPC3-1 peptide (average number of spots,  $101.0 \pm 33.2$ ) than to non-pulsed BM-DCs (average number of spots,  $2.1 \pm 3.7$ ) (Fig. 2D and E). The CD8-positive cells from mice immunized with mGPC3-4 peptide released more IFN- $\gamma$  to BM-DCs pulsed with mGPC3-4 peptide (average number of spots,  $5.3 \pm 4.0$ ) than to non-pulsed BM-DCs (average number

of spots,  $1.8 \pm 0.7$ ), but no significant differences were observed (Fig. 2F and G). These results suggest that mGPC3-1 peptide is more efficient for inducing CTLs than the mGPC3-4 peptide in a single peptide vaccine system.

Taken together, the above results suggest that mGPC3-1 peptide is the most efficient peptide for inducing CTLs among the 11 peptides.

*mGPC3-1 peptide-specific CTL line recognition of target cells endogenously expressing mGPC3.* To further investigate the ability of mGPC3-1 peptide-specific CTLs induced by peptide vaccination, we established a CTL line from immunized mice according to the above described protocol. IFN- $\gamma$  ELISPOT assays were performed using RMA-S pulsed with mGPC3-1 peptide and non-pulsed RMA-S to confirm whether the CTL line had mGPC3-1 peptide specificity. The CTL line clearly released more IFN- $\gamma$  to RMA-S pulsed with mGPC3-1 peptide than to non-pulsed RMA-S, which suggests that the CTL line is the mGPC3-1 peptide-specific CTL (Fig. 3A).

Subsequently, a cytotoxicity assay was performed to confirm whether the mGPC3-1-specific CTLs could kill RMA-S pulsed with mGPC3-1 peptide. The CTLs killed RMA-S pulsed with the mGPC3-1 peptide (16.4%) better than non-pulsed RMA-S (2.2%), suggesting that the mGPC3-1-specific CTL line could specifically recognize and kill RMA-S pulsed with the mGPC3-1 peptide (Fig. 3B).

Finally, we examined whether the CTL line could recognize RMA GPC3-puro endogenously expressing mGPC3. Expression of mGPC3 in RMA-GPC3-puro and RMA-puro

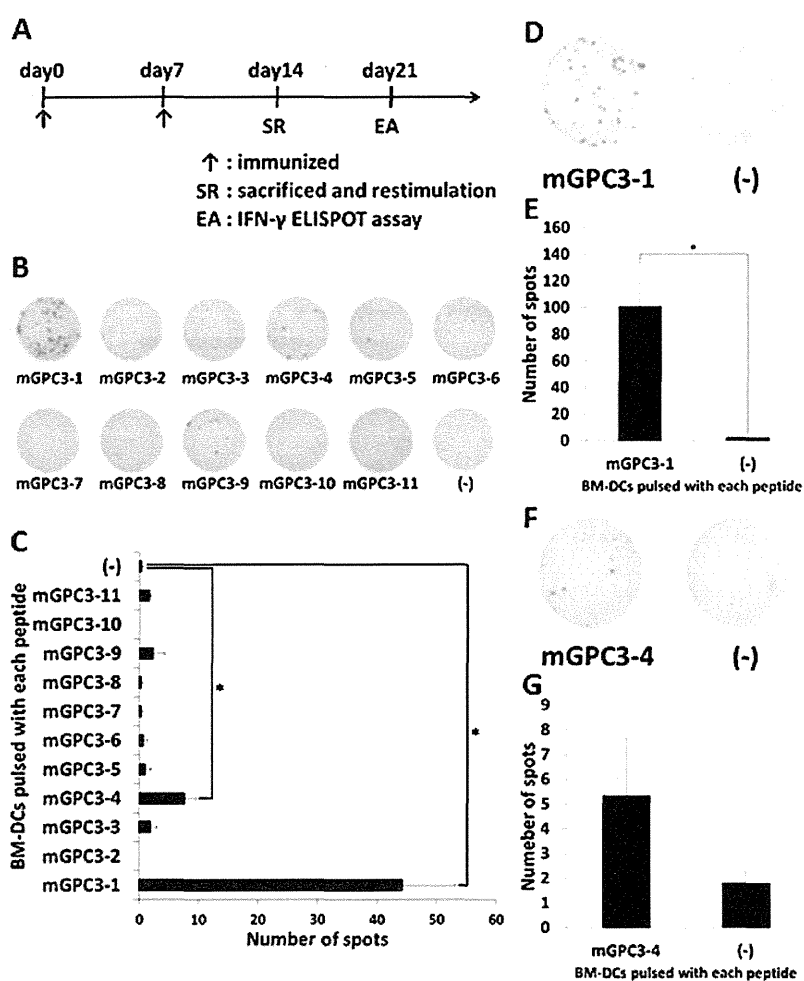


Figure 2. *In vivo* mixed peptide vaccine and single peptide vaccine. Analysis was performed for each vaccine. (A) Schedule of mixed peptide vaccine and single peptide vaccine. (B and C) The mixed peptide vaccine was given to mice and the responses of CD8-positive cells to the 11 peptides were examined. IFN- $\gamma$  ELISPOT assays were performed using BM-DCs pulsed with each peptide and non-pulsed BM-DCs as target cells (n=3, \*p<0.05). Representative data are shown (B). To confirm whether mGPC3-1 or mGPC3-4 was a CTL-inducible peptide, the single peptide vaccine was given. (D and E) mGPC3-1 peptide vaccine was given and IFN- $\gamma$  ELISPOT assays were performed using BM-DCs pulsed with mGPC3-1 and non-pulsed BM-DCs as target cells (n=3, \*p<0.05). Representative data are shown (D). (F and G) mGPC3-4 peptide vaccine was given and IFN- $\gamma$  ELISPOT assays were performed using BM-DCs pulsed with mGPC3-4 and non-pulsed BM-DCs as target cells (n=3). Representative data are shown (F).

was confirmed by RT-PCR. The results showed that RMA-GPC3-puro expressed mGPC3 and RMA-puro did not express mGPC3 (Fig. 3C). IFN- $\gamma$  ELISPOT assays were performed using RMA-GPC3-puro and RMA-puro as target cells to investigate whether the CTL line could recognize RMA-GPC3-puro expressing endogenous mGPC3. The CTL line released more IFN- $\gamma$  to RMA-GPC3-puro (average number of spots,  $32.2 \pm 5.0$ ) than to RMA-puro (average number of spots,  $18.2 \pm 6.2$ ). This result suggests that the mGPC3-1 peptide is an endogenously presented peptide (Fig. 3D).

**CTL response against the mGPC3-derived peptides induced in STAM mice.** Previously, the NASH mouse model (named STAM mice C57BL/6N-NASH) was exploited by Stelic Institute & Co. and STAM mice with a B6 background spontaneously developed liver cancer. We observed that liver cancer developed in 18-week-old STAM mice (Fig. 4A and B). Furthermore, to verify whether mGPC3-1 peptide-specific CTLs were induced in STAM mice C57BL/6N-NASH, a

mGPC3-1 peptide vaccine was given and an IFN- $\gamma$  ELISPOT assay was performed using RMA-S pulsed with mGPC3-1 peptide or non-pulsed RMA-S. The CD8-positive cells derived from immunized mice released IFN- $\gamma$  only to pulsed RMA-S (average number of spots,  $100 \pm 74.3$ ), not to non-pulsed RMA-S (average number of spots,  $0.0 \pm 0.0$ ) (Fig. 4C and E). However, the CD8-positive cells derived from unimmunized mice did not release IFN- $\gamma$  to either pulsed (average number of spots,  $0 \pm 0.0$ ) or non-pulsed (average number of spots,  $0.0 \pm 0.0$ ) RMA-S (Fig. 4D and E). These results suggest that peptide-specific mGPC3-1 could be induced in STAM mice C57BL/6N-NASH immunized with the mGPC3-1 peptide vaccine but could not be induced in un-immunized STAM mice C57BL/6N-NASH.

## Discussion

HCC is the most common liver cancer and the recurrence rate for treated HCC patients is high, thus establishment of an effective preventative method, such as a vaccination to prevent

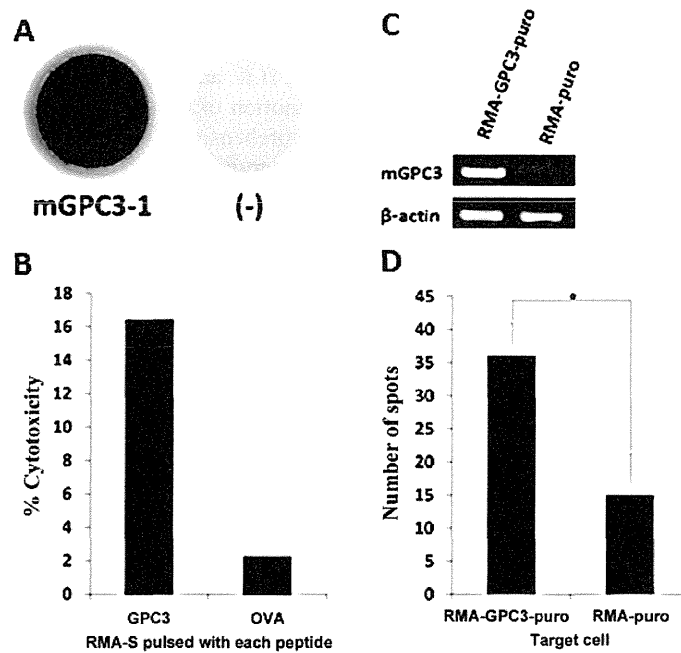


Figure 3. Analysis of established CTL line. (A) IFN- $\gamma$  ELISPOT assays were performed using GPC3-1 pulsed or non-pulsed RMA-S as target cells. (B) Cytotoxicity assays were performed using GPC3-1 pulsed or unpulsed RMA-S as target cells. Percent cytotoxicity =  $\{1 - [(average\ fluorescence\ of\ the\ sample\ wells - average\ fluorescence\ of\ the\ maximal\ release\ control\ wells) / (average\ fluorescence\ of\ the\ minimal\ release\ control\ wells - average\ fluorescence\ of\ the\ maximal\ release\ control\ wells)]\} \times 100\%$ . (C) mGPC3 expression of RMA-GPC3-puro and RMA-puro by RT-PCR. (D) IFN- $\gamma$  ELISPOT assays were performed using RMA-GPC3-puro and RMA-puro as target cells ( $n=3$ ,  $*p>0.05$ ).

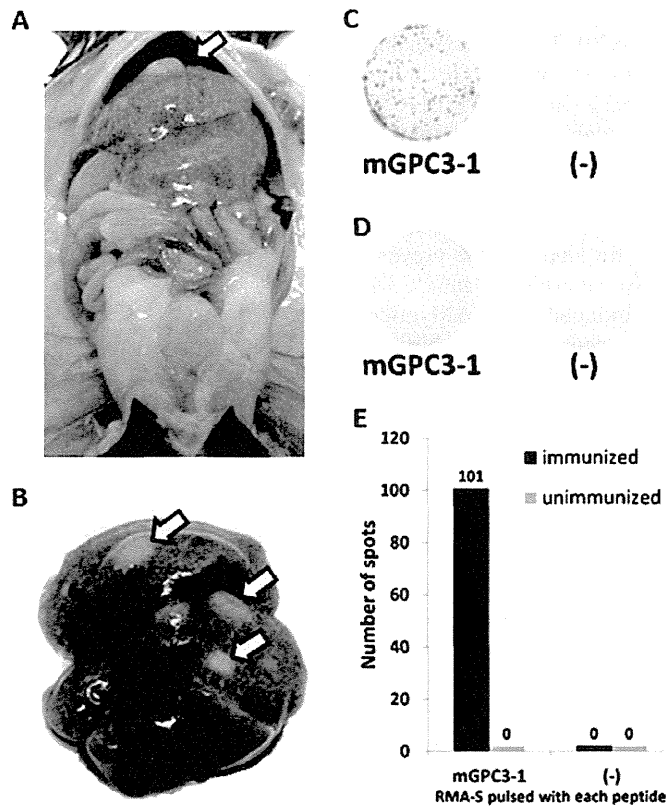


Figure 4. Analysis of STAM mice C57BL/6N-NASH. (A and B) Liver cancer was observed in 18-week-old STAM mice C57BL/6N-NASH. (C-E) To confirm whether mGPC3-1 peptide-specific CTL was induced, the mGPC3-1 peptide vaccine was given to STAM mice C57BL/6N-NASH and the IFN- $\gamma$  ELISPOT assay was performed. Arrow indicates the area of developing cancer (C and E). The CD8-positive cells derived from immunized mice released IFN- $\gamma$  to RMA-S pulsed with mGPC3-1 peptide ( $n=3$ ). Representative data are shown (C). (D and E) As a control, IFN- $\gamma$  ELISPOT assays were performed using the CD8-positive cells derived from unprimed mice. Representative data are shown (D).

the occurrence and recurrence of HCC, is needed. GPC3 is overexpressed in HCC and is not expressed in normal tissue except for the placenta and embryonic liver. Clinical trials of a GPC3-derived peptide vaccine for HCC have been performed and a phase I clinical trial has shown the safety and immunological and clinical potential of the vaccine (15,16). Moreover, to study the preventive effect as a potential of the GPC3-derived peptide vaccine, we attempted to establish a mouse model to induce GPC3-specific CTLs by the peptide vaccine.

First, mGPC3-derived peptides binding to H2-K<sup>b</sup> or H2-D<sup>b</sup> were determined *in silico* using BIMAS software. Moreover, a binding assay was performed *in vitro* and showed that all peptides predicted by the BIMAS software could bind H2-K<sup>b</sup> and H2-D<sup>b</sup>. However, the BIMAS score did not correlate with the actual binding affinity.

Peptides that can bind to MHC class I are not always able to induce peptide-specific CTLs (21,29). Therefore, to investigate actual CTL-inducible peptides among the 11 selected peptides, a mixed peptide vaccine and single peptide vaccine were given to mice. These results (Fig. 2) suggested that mGPC3-1 could induce peptide-specific CTLs. In addition, antigen-derived and CTL-inducible peptides are not necessarily presented by cancer cells endogenously expressing the antigen (23,30). Hence, we confirmed whether the mGPC3-1 peptide-specific CTL line could recognize RMA-GPC3-puro endogenously expressing mGPC3 (Fig. 3D). Furthermore, confirming whether the mGPC3-1 peptide-specific CTL line killed cancer cells presenting the mGPC3-1 peptide is important, thus a cytotoxicity assay was performed (Fig. 3B).

Mice with a B6 background that spontaneously develop liver cancer have been reported (19,20). These mice enable investigations as to whether a peptide vaccine for GPC3 has a preventive capability. Recently, the STAM mice C57BL/6N-NASH was established as a non-alcoholic-steatohepatitis (NASH) mouse model by Stelic Institute & Co. STAM mice C57BL/6N-NASH are drug-treated B6 mice and liver cancer occurs spontaneously and early in NASH mice. Therefore, this mouse is an attractive model for studying the preventive effects of a cancer vaccine. We showed that mGPC3-1 peptide-specific CTL could be induced in STAM mice C57BL/6N-NASH (Fig. 4E). Simultaneously, we established a liver cancer cell line derived from STAM mice C57BL/6N-NASH and observed the cancer cell line expressed mGPC3 (data not shown).

However, the GPC3 peptide vaccine did not prevent the occurrence of liver cancer in STAM mice C57BL/6N-NASH (data not shown). Therefore, further research to develop strong GPC3-specific immunotherapies or combinational approaches in an appropriate mouse model is needed. Identification of an H2-K<sup>b</sup> or H2-D<sup>b</sup> restricted, GPC3-derived peptide is the first step. The established cell line from STAM mice C57BL/6N-NASH, which show GPC3 expression, may help us to develop a new mouse model system for a GPC3-targeted therapy.

In conclusion, mGPC3-1<sub>127-136</sub> AMFKNNYPSL was identified as an H2-K<sup>b</sup> or H2-D<sup>b</sup> restricted, GPC3-derived CTL most-inducible epitope peptide and mGPC3-1 peptide-specific CTL can kill RMA-S pulsed with the mGPC3-1 peptide. Furthermore, we established an mGPC3-1-specific CTL-inducible model in B6 mice using an mGPC3-1 peptide vaccine.

## Acknowledgements

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# Comparison of dose intensity of vincristine, *d*-actinomycin, and cyclophosphamide chemotherapy for child and adult rhabdomyosarcoma: a retrospective analysis

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## Abstract

**Purpose** The prognosis of adult rhabdomyosarcoma (RMS) has been considered dismal. The question is raised that vincristine, *d*-actinomycin, and cyclophosphamide (VAC) chemotherapy may not be administered as per schedule for adult RMS; consequently, low dose intensity (DI) leads to poor prognosis. Herein, we examined whether the administration of VAC chemotherapy for adults and children with RMS is feasible with regard to the DIs of VAC.

**Methods** Chart review was retrospectively performed for all identified patients. The percentage of relative DI (RDI) was calculated according to the Children's Oncology Group D9803 protocol. Further, we examined the RDI in the first 6 cycles of VAC (induction phase) and the DI after the first 6 cycles of VAC (maintenance phase).

**Results** We identified a total of 27 adults and 18 children with RMS, respectively. The mean RDIs of vincristine in total phase were significantly lower in adults than that in

children ( $P = 0.04$ ). In induction phase, the mean RDIs of vincristine and cyclophosphamide were similar for both groups; however, they were dropped significantly in adults during maintenance phase ( $P < 0.05$ ). Mean RDIs of vincristine in elderly patients tended to become low. Low RDI was mainly attributable to hematologic toxicity, infection, and peripheral neuropathy. The prognosis of low versus high RDI was similar.

**Conclusions** The RDIs of vincristine and cyclophosphamide in the maintenance phase were significantly lower than that in children. VAC chemotherapy for adults was not feasible; these patients require a different regimen.

**Keywords** Rhabdomyosarcoma · Adults · Children · Chemotherapy · Soft tissue sarcoma

## Abbreviations

RMS Rhabdomyosarcoma  
VAC Vincristine, *d*-actinomycin, and cyclophosphamide  
DI Dose intensity  
RDI Relative dose intensity  
IRS The International Rhabdomyosarcoma Study  
PFS Progression-free survival  
OS Overall survival  
PCP Pneumocystis jirovecci pneumonia

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## Introduction

Recent advances in the treatment for rhabdomyosarcoma have contributed to improve survival from 55 to 73 % [1–3]. Vincristine and *d*-actinomycin with or without cyclophosphamide (VAC) regimens for rhabdomyosarcoma are considered standard options according to the previous randomized trials [4–8]. However, these trials

mainly focused on children or adolescents; therefore, the data of precious adult rhabdomyosarcoma patients were limited.

Sultan et al. [9] reported the prognosis of pediatric and adult rhabdomyosarcoma, and the data suggested that 5-year survival rate was significantly poor in adult rhabdomyosarcoma with 27 % compared to that of pediatrics with 61 %. Other facilities also reported small series of similar results [10–14]. Accordingly, the prognosis of adult rhabdomyosarcoma remained poor; however, the treatment strategy references along with childhood rhabdomyosarcoma.

There is no evidence available to uncover the reason for poor prognosis of adult rhabdomyosarcoma. The VAC chemotherapy was developed mainly for targeting pediatrics; therefore, adverse event may differ when we applied the regimen to adults. Accordingly, the VAC regimens may not be administered per schedule, which leads to poor prognosis due to low dose intensity. To our knowledge, there are no reports on dose intensity of VAC regimens in relation to prognosis. We launched this study to seek optimal treatment for adult rhabdomyosarcoma by comparing dose intensity to childhood or adolescent rhabdomyosarcoma.

## Methods

### Patients

All the patients included in this analysis meet the following criteria: histologically diagnosed with rhabdomyosarcoma, treated at the National Cancer Center Hospital in Tokyo between 1991 and 2010, and received VAC chemotherapy 6 cycles or over as a first-line treatment. The chart review was performed for all identified patients to obtain the following information: age, sex, primary tumor site, histopathology, tumor size, stage, group defined by the International Rhabdomyosarcoma Study (IRS) postsurgical grouping system [4], chemotherapy regimen, chemotherapy administration schedule, adverse event, date of radiotherapy, date of surgery, date of progression, and date of last follow-up.

### Treatment

The details of VAC chemotherapy are shown as follows. VAC consisted of vincristine at a dose of 1.5 mg/m<sup>2</sup> (up to 2 mg/body), given intravenously on day 1, 8, 15, cyclophosphamide at a dose of 2.2 g/m<sup>2</sup>, given intravenously on day 1, and *d*-actinomycin at a dose of 1.5 mg/m<sup>2</sup> (up to 2.5 mg/body), given intravenously on day 1. The treatment course was repeated every 3 weeks according to Children's

Oncology Group Study D9803 [8]. *D*-actinomycin on day 1 and vincristine on day 8 and day 15 were omitted when radiotherapy was applicable.

### Dose intensity calculation and definitions of terms

Dose intensity of vincristine and cyclophosphamide was calculated according to the following formula: Dose intensity (vincristine) = Total dose of vincristine (mg/m<sup>2</sup>) / [(last date of injection – first date of injection) + 7]/7; Dose intensity (cyclophosphamide) = Total dose of cyclophosphamide (mg/m<sup>2</sup>) / [(last date of injection – first date of injection) + 21]/7, respectively. The percentage of relative dose intensity (RDI) was calculated according to the Children's Oncology Group D9803 protocol [8]. We examined the RDI in induction phase and maintenance phase. Induction phase was defined as the first six cycles of VAC, and maintenance phase was defined as the latter cycles after induction phase. We defined cut-off points of RDI of vincristine and cyclophosphamide at 80 % and dichotomized into lower RDI (RDI < 80 %) and higher RDI (RDI ≥ 80 %).

We defined adult as the patients aged greater than or equal to 21 years, and we used the term 'children' for the others. We attempted to retrospectively define the stage of disease according to the IRS staging system, which separates patients by site of the primary tumor, tumor size, and the presence or absence of tumor-involved regional lymph nodes and of distant metastases. Adverse events were retrospectively ranked by the National Cancer Institute Common Toxicity Criteria version 3.0. As a definition of febrile neutropenia, we used the guidelines established by the Infectious Disease Society of America [15]. Treatment delay was defined as more than a week of delay to start new cycle of VAC regimen.

### Statistics

We used Student's *t* test to detect the difference of mean RDI. Progression-free survival (PFS) was defined as the time between the date of initial chemotherapy until the date of the recognition of local recurrence or distant metastases. Death was treated as an event, and the absence of disease progression was treated as a censored observation on the last day of follow-up. Overall survival (OS) was defined as the time from the date of initial chemotherapy until the date of death. Patients who were lost to follow up were treated as a censored observation on the last day of follow-up. We used the Kaplan–Meier method to draw survival curve, and significance was detected by the log-rank test. All statistical analyses were performed using SPSS ver. 19.0 (IBM Co. Inc, Okinawa, Japan).

## Results

### Patient characteristics

We identified 59 patients who received VAC regimen and of 45 patients met the eligibility criteria in the study period. Fourteen patients were excluded from this analysis by the following reasons: less than 6 cycles of VAC regimen in 12 patients, VAC chemotherapy of IRS II protocol in one, and clinical data unavailable in one. There were 27 adult patients and 18 were children.

The distribution of clinical and pathologic characteristics of these patients is listed in Table 1. The median age of adult was 28 years (range, 22–72) and that of children was 16 years (range, 2–19). Primary disease site, histopathology, and primary tumor size were similar among adult and children. Fourteen adults and 8 children had parameningeal primary site. Metastatic disease was somewhat high in adult.

Overall median follow-up period was 26 months (range, 6–114 months): 26 months in adults (range, 6–113 months) and 36 months in children (range, 6–114 months). We identified a total of 316 cycles of VAC chemotherapy in adults and a total of 209 cycles in children. The median number of chemotherapy cycles was 13 (range, 6–14) for adults and 12.5 (range, 6–14) for children.

### Relative dose intensity of VAC chemotherapy

The mean RDIs of vincristine and cyclophosphamide in total phase for adults were 76.8 and 80.8 %, respectively, and those for children were 90.2 and 86.9 %, respectively. The mean RDI of vincristine in total phase was significantly lower in adults than that in children ( $P < 0.05$ ) (Table 2).

In induction phase, the mean RDIs of vincristine and cyclophosphamide were similar for both groups; however, they were dropped significantly in adults during maintenance phase ( $P < 0.05$ ) (Table 2). We examined the RDIs of vincristine and cyclophosphamide in relation to age. Mean RDIs of vincristine in elderly patients tended to become low (Fig. 1).

### Chemotherapy efficacy by RDI

We analyzed the efficacy of RDI on PFS and OS. The PFS and OS were not statistically different in patients with high RDI and low RDI of vincristine and cyclophosphamide in total phase. For total phase of RDI for vincristine, median PFS was 28 months in patients with high RDI ( $n = 15$ ) and 24 months in patients with low RDI ( $n = 30$ ) ( $P = 0.43$ ), and median OS was 60 months in patients with high RDI ( $n = 15$ ) and 42 months in patients with low RDI ( $n = 30$ )

( $P = 0.78$ ) (Fig. 2a). For total phase of RDI for cyclophosphamide, median PFS was 28 months in patients with high RDI ( $n = 17$ ) and 24 months in patients with low RDI ( $n = 28$ ) ( $P = 0.78$ ), and median OS was 114 months in patients with high RDI ( $n = 17$ ) and 60 months in patients with low RDI ( $n = 28$ ) ( $P = 0.67$ ) (Fig. 2b). We further analyzed induction phase and maintenance phase of RDIs for vincristine and cyclophosphamide or stratified them by adults and children on PFS and OS. None of the difference regarding RDI of both agents on PFS and OS was detected (Fig. 3).

### Toxicity of VAC chemotherapy

Table 3 lists the worst degree of toxicity of VAC chemotherapy. Majority of patients experienced grade 3/4 neutropenia and anemia in both induction phase and maintenance phase. Grade 2–4 peripheral neuropathy was observed in 55.6 % of adult patients, while it was 33.3 % in children ( $P = 0.37$ ). Approximately 80 % of patients needed granulocyte colony-stimulating factor, and 73.3 % of patients received trimethoprim-sulfamethoxazole for prevention of pneumocystis jirovecii pneumonia (PCP). Two patients who did not receive trimethoprim-sulfamethoxazole developed PCP; one experienced PCP in 4th cycle of VAC, and the other did in 14 the cycle of VAC.

### Cause of dose reduction, skip, and delayed cycles

In induction phase, three of adults (11.1 %) and one of children (5.6 %) required dose reduction of vincristine due to peripheral neuropathy and myelosuppression. Eight of the patients (29.6 %) in adults and 4 of the patients (22.2 %) in children required dose reduction of cyclophosphamide, due to myelosuppression, febrile neutropenia, and liver dysfunction. In maintenance phase, four adults (16.0 %) and 2 children (11.8 %) required dose reduction of vincristine, due to peripheral neurotoxicity and myelosuppression. Nine adults (36.0 %) and 5 children (18.5 %) required dose reduction of cyclophosphamide, due to myelosuppression and febrile neutropenia.

Adult patients needed discontinuance of vincristine in total of 126 times out of 713 times (17.7 %) due to the following reasons: peripheral neurotoxicity in 75 times, myelosuppression in 35 times, and febrile neutropenia in 14 times. For these child patients, it was 55 times out of 483 times of vincristine administration (11.4 %) due to peripheral neurotoxicity in 16 times, myelosuppression in 16 times, and febrile neutropenia in 9 times.

In induction phase, 14 adult patients (51.9 %) and 9 child patients (50.0 %) experienced cycle delay. In maintenance phase, 17 adult patients (68.0 %) and 9 child patients (52.9 %) required cycle delay. The major causes of

**Table 1** Patient characteristics

	Adults (N = 27)	Children (N = 18)
Age, median (range)	28 (22–72)	16 (2–19)
Gender		
Female	8 (29.6 %)	9 (50.0 %)
Male	19 (70.4 %)	9 (50.0 %)
Primary tumor site		
Head-neck	3 (11.1 %)	3 (16.7 %)
Parameningeal primary site	14 (51.9 %)	8 (44.4 %)
Unfavorable other	5 (18.5 %)	1 (5.6 %)
Extremity	3 (11.1 %)	5 (27.8 %)
Genitourinary primary site	2 (7.4 %)	1 (5.6 %)
Histopathology		
Alveolar	15 (55.6 %)	10 (55.6 %)
Embryonal	10 (37.0 %)	7 (38.9 %)
Others	2 (7.4 %)	1 (5.6 %)
Tumor size		
≥5 cm	10 (37.0 %)	8 (44.4 %)
<5 cm	15 (55.6 %)	9 (50.0 %)
Not available	2 (7.4 %)	1 (5.6 %)
IRS group		
I	3 (11.1 %)	2 (11.1 %)
II	3 (11.1 %)	1 (5.6 %)
III	12 (44.4 %)	12 (66.7 %)
IV	9 (33.3 %)	3 (16.7 %)
IRS pre-treatment stage		
Stage 1	4 (14.8 %)	4 (22.2 %)
Stage 2	4 (14.8 %)	1 (5.6 %)
Stage 3	10 (37.0 %)	10 (55.6 %)
Stage 4	9 (33.3 %)	3 (16.7 %)
Complete surgical resection		
Pre-chemotherapy	7 (25.9 %)	4 (22.2 %)
Post-chemotherapy	4 (14.8 %)	6 (33.3 %)
No surgery	16 (59.3 %)	8 (44.4 %)
Radiation therapy		
Pre-chemotherapy	2 (7.4 %)	2 (11.1 %)
Post-chemotherapy	20 (74.1 %)	14 (77.8 %)
No radiation therapy	5 (18.5 %)	2 (11.1 %)

IRS Intergroup Rhabdomyosarcoma Study

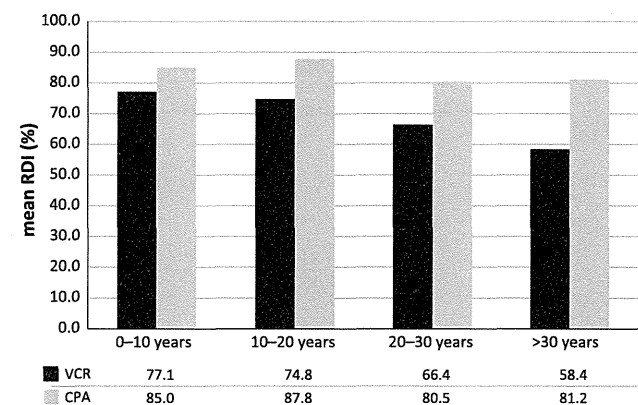
cycle delay were myelosuppression, febrile neutropenia, and patient preferences.

## Discussion

In our analyses, 27 adults and 18 children of rhabdomyosarcoma who received VAC chemotherapy at a single institution were evaluated for RDIs of VAC chemotherapy. Adult patients had significantly lower RDIs of vincristine and cyclophosphamide in the maintenance phase compared

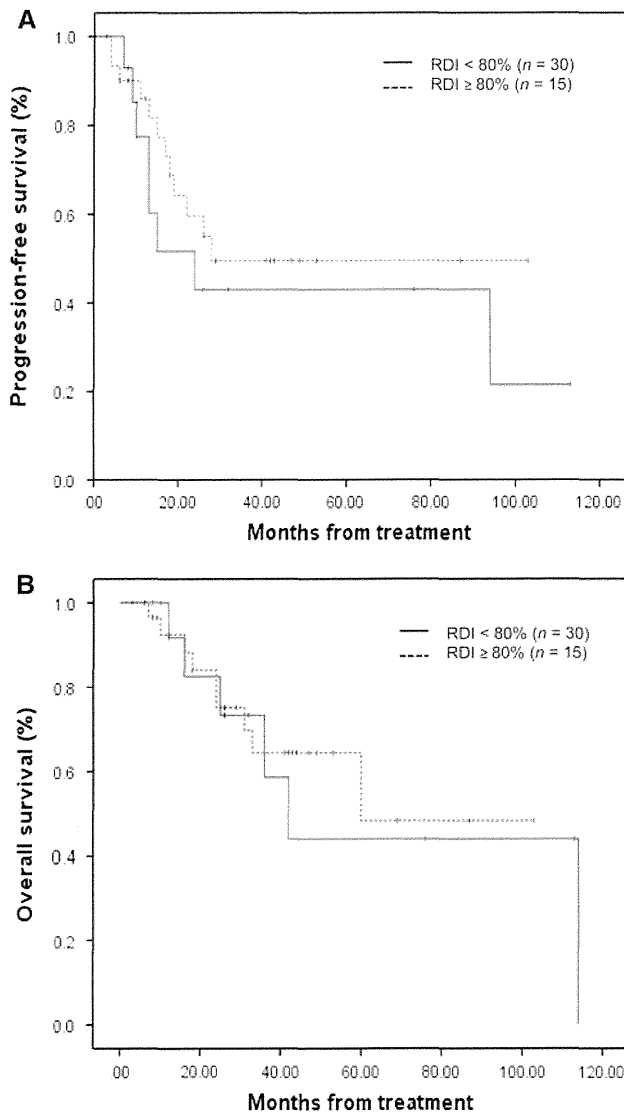
**Table 2** Mean relative dose intensity in patients with adult and child rhabdomyosarcoma

	Adult (%)	Child (%)	P value
Vincristine			
Induction phase	77.7	86.4	0.109
Maintenance phase	71.9	100.1	0.042
Total phase	76.8	90.2	0.040
Cyclophosphamide			
Induction phase	87.1	88.2	0.820
Maintenance phase	69.7	86.4	0.011
Total phase	80.8	86.9	0.156

**Fig. 1** Relative dose intensity of vincristine and cyclophosphamide in relation to age. VCR vincristine, CPA cyclophosphamide

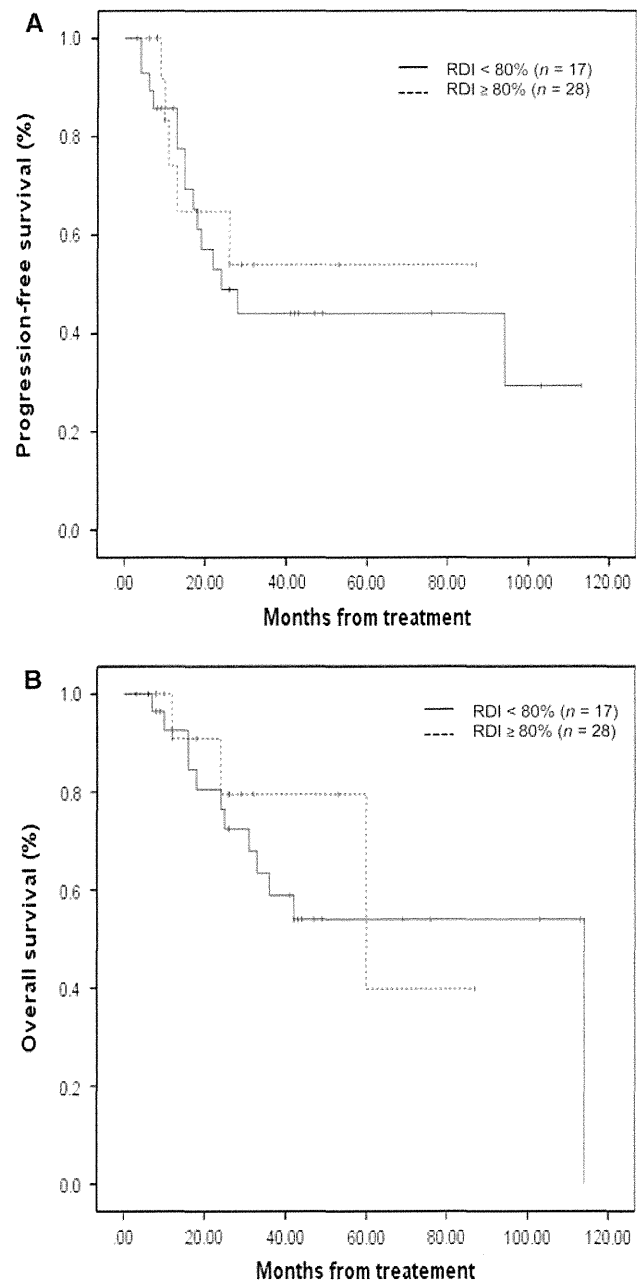
to those of children. Further, the mean RDIs of vincristine tended to become low depending on patients' ages. The reasons for dose reduction, discontinuance, and treatment delay of vincristine and cyclophosphamide were generally due to myelosuppression, infection, and peripheral neurotoxicity. We revealed that the adult rhabdomyosarcoma patients significantly received lower RDIs of vincristine and cyclophosphamide than those of children; however, the PFS and the OS were not statistically different in patients with high RDI and low RDI in total phase and maintenance phase.

The prognosis of adult rhabdomyosarcoma remains poor, with approximately 30 % at 5-year survival [9–14, 16]. The reason for the poor outcome of adults remains unclear. Sultan et al. [9] reported the comparison of pediatric and adult rhabdomyosarcoma; they reported that the typical pediatric rhabdomyosarcoma variants (embryonal and alveolar subtypes) occurred less frequently in adult rhabdomyosarcoma and that the most common primary sites in adult were extremities and unfavorable sites and that radiation therapy was performed less frequently in adult rhabdomyosarcoma than in pediatric patients. In our study, adult rhabdomyosarcoma patients had lower RDIs in



**Fig. 2** **a** Kaplan–Meier curve of progression-free survival in patients with high relative dose intensity of vincristine (*dash line*) and low relative dose intensity of vincristine (*black line*). **b** Kaplan–Meier curve of overall survival in patients with high relative dose intensity of vincristine (*dash line*) and low relative dose intensity of vincristine (*black line*)

VAC chemotherapy, especially in maintenance phase. The reason for the low RDI was not apparent, but we demonstrated that VAC chemotherapy for adult rhabdomyosarcoma patient may not be feasible as VAC chemotherapy was developed mainly targeting pediatric patients. We also suggested that none of the differences regarding RDIs of both vincristine and cyclophosphamide on PFS and OS were detected. However, in our previous data focusing on childhood RMS and adult RMS survival suggested significant differences in survival in localized tumor [17]. Although we could not observe significant differences in survival in metastatic tumors for both childhood RMS and adult RMS, controversy remains as to the standard



**Fig. 3** **a** Kaplan–Meier curve of progression-free survival in patients with high relative dose intensity of cyclophosphamide (*dash line*) and low relative dose intensity of cyclophosphamide (*black line*). **b** Kaplan–Meier curve of overall survival in patients with high relative dose intensity of cyclophosphamide (*dash line*) and low relative dose intensity of cyclophosphamide (*black line*)

treatment for patients with adult rhabdomyosarcoma. Recently, several types of trials for these patients have demonstrated to show additional benefit of adding active new agents to standard VAC chemotherapy, which failed to show survival benefit [18–20].

In IRS-V study, the dosage of cyclophosphamide was designed 1.2 g/m<sup>2</sup>/cycle. High-dose therapy with stem cell transplantation for high-risk metastatic RMS did not

**Table 3** Adverse event in patients who received VAC regimen

	Adults			Children		
	Induction phase (n = 27)	Maintenance phase (n = 25)	Total phase	Induction phase (n = 18)	Maintenance phase (n = 17)	Total phase
Neutropenia Gr. 4	27 (100 %)	23 (92.0 %)	27 (100 %)	18 (100 %)	17 (100 %)	18 (100 %)
Anemia Gr. 3/4	24 (88.9 %)	21 (84.0 %)	27 (100 %)	16 (88.9 %)	14 (82.6 %)	16 (88.9 %)
Thrombocytopenia Gr. 4	12 (44.4 %)	14 (56.0 %)	19 (70.3 %)	13 (72.2 %)	10 (58.8 %)	13 (72.2 %)
FN Gr. 3/4	23 (85.2 %)	19 (76.0 %)	26 (96.3 %)	17 (94.4 %)	14 (82.6 %)	18 (100 %)
Peripheral neuropathy $\geq$ Gr. 2	11 (40.7 %)	12 (48.0 %)	15 (55.6 %)	6 (33.3 %)	5 (29.4 %)	6 (33.3 %)

VAC vincristine, *d*-actinomycin, cyclophosphamide, FN febrile neutropenia, Gr grade

improve outcome significantly [21]. Accordingly, reduced dose of VAC chemotherapy for adults might not be inferior to standard VAC dose regarding survival.

This study has several limitations as of the nature of retrospective design and small sample size. More patients may be needed to detect the small differences regarding survival in low RDI patients and high RDI patients. Decision of administration of vincristine on day 8 and 15 may vary by physicians as evaluation of peripheral neuropathy by vincristine is not integrated. However, considering poor prognosis of adult RMS with highly toxic regimen of VAC, new strategy to treat this rare cohort must be developed urgently.

In summary, adult rhabdomyosarcoma is a rare malignancy with poor prognosis. The treatment strategy has been performed along with child rhabdomyosarcoma; however, the RDIs of vincristine and cyclophosphamide in the maintenance phase were significantly lower than that in children. Causes of low RDIs were generally myelosuppression, infection, and peripheral neurotoxicity in both groups. VAC chemotherapy for adults may not be feasible; thus, the prospective trial for adult rhabdomyosarcoma should be designed.

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**Conflict of interest** None declared.

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## Clinical outcomes of adult and childhood rhabdomyosarcoma treated with vincristine, *d*-actinomycin, and cyclophosphamide chemotherapy

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### Abstract

**Background** Outcomes in adult patients with rhabdomyosarcoma are poor, with a 5-year survival rate of approximately 30 %. The current study aimed to compare the clinical outcomes of adult and childhood rhabdomyosarcoma patients with local and metastatic disease and to examine the impact and timing of local therapy on metastasis.

**Methods** Clinicopathological features and patient outcomes were reviewed retrospectively for rhabdomyosarcoma patients receiving chemotherapy between 1981 and 2010 at our institution. Adults were defined as those aged 21 years or older.

**Results** Of the 98 patients identified, 36 were adults (median age, 29; range, 21–72) and 62 were children

(median age, 11; range, 0.6–20). Median progression-free survival of localized and metastatic disease for children and adults was as follows: localized disease, 166.9 versus 22.4 months ( $p = 0.005$ ), and metastatic disease, 13.3 versus 13.3 months ( $p = 0.949$ ), respectively. Multivariate regression analysis revealed that older age ( $\geq 21$  vs.  $< 21$ ) was a significant poor prognostic factor in localized disease. Conversely, age was not related to survival in metastatic disease. Receiving radiotherapy to the primary site was an independent factor indicating a better prognosis. An analysis of the optimal timing of local therapy was performed for 53 patients; however, its significance on survival could not be determined.

**Conclusions** Age was a negative prognostic factor in rhabdomyosarcoma patients with localized disease, but it did not affect the survival in metastatic disease. For metastatic disease, although local therapies may be effective for survival, the timing of such therapies should be determined individually.

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**Keywords** Adult · Child · Rhabdomyosarcoma ·  
Clinical outcome

### Introduction

Rhabdomyosarcoma (RMS) is the most common soft tissue sarcoma in children, but it is far less common in adults (Ferrari et al. 2003). Improvements in multimodal treatment of RMS have improved survival in children from 25 to approximately 70 % over the past 40 years (Pappo et al. 1997; Raney et al. 2001; Breitfeld and Meyer 2005; Leaphart and Rodeberg 2007). In contrast, the prognosis for adult RMS remains poor, with a 5-year survival rate of approximately 30 % (Sultan et al. 2009; Ferrari et al. 2003;

La Quaglia et al. 1994; Little et al. 2002; Esnaola et al. 2001; Hawkins et al. 2001).

Metastatic RMS affects approximately 15 % of all children with RMS (Breneman et al. 2003). According to the risk categories identified by the Intergroup Rhabdomyosarcoma Study (IRS), low- and intermediate-risk patients have improved outcomes, with 96–97 % of low-risk patients achieving 5-year survival (Meza et al. 2006) and 79 % of intermediate-risk patients attaining 4-year survival (Arndt et al. 2009). However, survival for high-risk patients remains poor, with 34 % achieving 3-year survival (Oberlin et al. 2008) and 24 % achieving 5-year survival (Carli et al. 2004). Consequently, standard treatment for high-risk patients remains controversial (Klingebiel et al. 2008; Pappo et al. 2007; Lager et al. 2006).

The treatment strategy for RMS requires multidisciplinary therapy, including chemotherapy, surgery, and radiation therapy. Vincristine and *d*-actinomycin, with or without cyclophosphamide (VAC) regimens, are considered the standard option for RMS (Maurer et al. 1988, 1993; Crist et al. 1995, 2001; Meza et al. 2006; Arndt et al. 2009), with wide surgical resection of tumor and postoperative radiation therapy required for local control of localized RMS (La et al. 2011; Rodeberg et al. 2011; Schuck et al. 2004). The timing of radiation therapy is critical: for localized RMS, radiation is appropriate to start during weeks 9–12, and for parameningeal RMS with intracranial extension, local radiation treatment should begin during the first 1–2 weeks of chemotherapy (Michalski et al. 2004; Raney et al. 2002). However, the impact and optimal timing of local therapy for metastatic disease is unknown. Therefore, the purpose of the current study was to compare the clinical outcomes of local or metastatic adult and childhood RMS and to examine the impact and timing of local therapies on metastatic disease.

## Patients and methods

### Patients

All patients included in this analysis met the following criteria: histologically diagnosed with RMS, treated at the National Cancer Center Hospital in Tokyo between 1981 and 2010, and received VAC or VAC-like chemotherapy. Medical records were then retrospectively reviewed to obtain the following information: date of birth, gender, date of diagnosis, primary tumor site, histopathology, initial tumor size, presence of central nervous system (CNS) invasion, clinical stage, group category as defined by the IRS (Breitfeld and Meyer 2005), date of treatment initiation, chemotherapy regimen, best response, chemotherapy administration schedule, date of radiotherapy, date of

surgery, date of progression, date of last follow-up, and survival status.

### Treatment

The VAC regimen after the year of 2000 consisted of vincristine at a dose of 1.5 mg/m<sup>2</sup> (up to 2 mg/body), given intravenously on days 1, 8, and 15; cyclophosphamide at a dose of 2.2 g/m<sup>2</sup>, given intravenously on day 1; and *d*-actinomycin at a dose of 1.5 mg/m<sup>2</sup> (up to 2.5 mg/body), given intravenously on day 1. The treatment course was repeated according to IRS-IV or Children's Oncology Group (COG) Study (D9803) protocols (Arndt et al. 2009). Before the year of 2000, the VAC-like regimen includes the following regimens: vincristine, *d*-actinomycin, and either ifosfamide, etoposide, or doxorubicin. The treatment course was administered according to IRS-II or III protocols.

Local therapy includes surgery, radiation therapy, or both. Surgical resection defined in this manuscript includes only total gross resection. Microscopic complete resection (Group I) was confirmed microscopically later on. Radiation therapy was delivered to each primary tumor site and regional lymph nodes where applicable. The total dose ranged from 30 to 56.3 Gy, with a median dose of 45 Gy.

### Definitions of terms

We defined adults as patients aged 21 years or more, and considered the remaining patients to be children. The induction phase was defined as the first six cycles of the VAC/VAC-like regimen, and the maintenance phase was defined as the later cycles following the induction phase. Response to chemotherapy was compared with baseline status. A complete response (CR) was defined as the disappearance of tumors with no evidence of disease. A partial response (PR) was a 50 % or greater decrease in the sum of tumor diameters. Stable disease (SD) was a less than 50 % decrease in the sum of tumor diameters. Progressive disease (PD) was defined as a 25 % or greater increase in the sum of tumor diameters and/or the appearance of new lesions.

Surgery or radiation therapy in this study means resection or radiation therapy to the primary site during primary treatment, respectively. Resection or radiation therapy to a relapsed site or metastatic site is not included in this category; likewise, biopsy only is not included in this definition.

### Statistical analyses

Progression-free survival (PFS) was defined as the time from the date of initial chemotherapy to the date when

local recurrences or distant metastases were recognized. Overall survival (OS) was defined as the time from the date of initial chemotherapy to the date of death due to any cause. Patients who survived were treated as censored observations on the last day of follow-up. PFS and OS were estimated using the Kaplan–Meier method, and survival curves were compared using the log-rank test. Multivariate Cox regression analysis was used to estimate the hazard ratios and 95 % confidence intervals (CI). A two-sided  $p < 0.05$  was considered statistically significant. All statistical analyses were performed using SAS (version 9.1.3; SAS Institute Inc., Cary, NC, USA).

**Results**

**Patient characteristics**

We identified 98 patients who met the eligibility criteria. The distributions of clinical and pathologic characteristics of patients are listed in Tables 1 and 2. There were 36

**Table 1** Characteristics of patients with localized rhabdomyosarcoma ( $n = 73$ )

	Adults ( $n = 22$ )		Children ( $n = 51$ )	
	No.	%	No.	%
<b>Gender</b>				
Female	5	22.7	24	47.1
Male	17	77.3	27	52.9
<b>Tumor size</b>				
<5 cm	9	40.9	23	45.1
≥5 cm	13	59.1	28	51.9
<b>Site</b>				
Favorable	5	22.7	22	43.1
Unfavorable	17	77.3	29	56.9
<b>Histology</b>				
Alveolar	9	40.9	18	35.3
Embryonal	12	54.5	29	56.9
Other	1	4.5	4	7.8
<b>Group</b>				
1	3	13.6	9	17.6
2	3	13.6	5	9.8
3	16	72.7	37	72.5
<b>Stage</b>				
1	4	18.2	21	41.2
2	5	22.7	8	15.7
3	13	59.1	22	43.1
<b>CNS invasion</b>				
No	20	90.9	46	90.2
Yes	2	9.1	5	9.8

CNS central nervous system

adults (age: median, 29; range 21–72) and 62 children (age: median, 11; range, 0.6–20). Seventy-three patients had localized disease (22 adults and 51 children), while 25 patients had metastatic disease (14 adults and 11 children). The most common histology was embryonal in localized disease (56.2 %) and alveolar in metastatic disease (68.0 %). Botryoid was found in only one patient. Common primary sites in localized disease were parameningeal (38.4 %), head and neck (21.9 %), and extremity (15.1 %), while common primary sites in metastatic disease sites were parameningeal (32.0 %), extremity (28.0 %), and other (24.0 %) (Table 3). The most common metastatic

**Table 2** Characteristics of patients with metastatic rhabdomyosarcoma ( $n = 25$ )

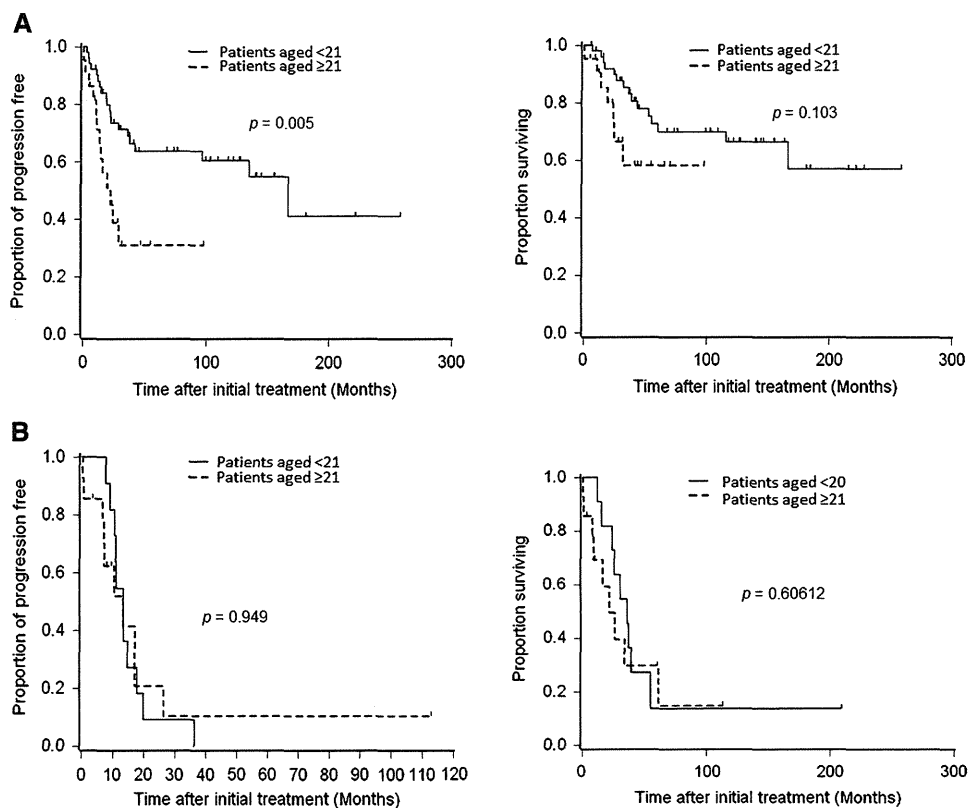
	Adults ( $n = 14$ )		Children ( $n = 11$ )	
	No.	%	No.	%
<b>Gender</b>				
Female	6	42.9	6	54.5
Male	8	57.1	5	45.5
<b>Tumor size</b>				
<5 cm	2	14.3	4	36.4
≥5 cm	12	85.7	7	63.6
<b>Site</b>				
Favorable	1	7.1	1	9.1
Unfavorable	13	92.9	10	90.9
<b>Histology</b>				
Alveolar	11	78.6	6	54.5
Embryonal	2	14.3	5	45.5
Other	1	7.1	0	0
<b>CNS invasion</b>				
No	14	100	10	90.9
Yes	0	0	1	9.1

CNS central nervous system

**Table 3** Primary site of rhabdomyosarcoma in localized and metastatic disease

	Localized disease No. (%)	Metastatic disease No. (%)
Extremity	11 (15.1)	7 (28.0)
Parameningeal site	28 (38.4)	8 (32.0)
Head and neck	16 (21.9)	1 (4.0)
Genitourinary primary site (non-bladder/prostate)	6 (8.2)	0 (0.0)
Orbit	5 (6.9)	1 (4.0)
Genitourinary primary site (bladder/prostate)	2 (2.7)	2 (8.0)
Unfavorable other	5 (6.9)	6 (24.0)

**Fig. 1** **a** Kaplan–Meier curve of progression-free survival and overall survival in patients with localized disease of adults (*dashed line*) and children (*solid line*). **b** Kaplan–Meier curve of progression-free survival and overall survival in patients with metastatic disease of adults (*dashed line*) and children (*solid line*)



sites were bone ( $n = 17$ ), bone marrow ( $n = 7$ ), and lung ( $n = 6$ ). Eleven patients had metastases to multiple sites.

Thirty-one patients (31.6 %) underwent surgical resection at RMS diagnosis. Ten patients were classified as IRSG Group I, 7 as Group II, and 14 as Groups III/IV. Surgery (primary tumor resection at diagnosis or second-look surgery) was performed in 43 patients (58.9 %) with localized disease and in 5 patients (20.0 %) with metastatic disease. We identified 59 patients (60.2 %) who had received VAC regimens and 39 patients (39.8 %) who had received VAC-like regimens. Radiation therapy was performed in 52 patients (71.2 %) with localized disease and in 18 patients (72.0 %) with metastatic disease during the course of treatment.

#### Patient outcomes in adults and children

The best responses to chemotherapy were as follows: among those with localized disease, 65 patients (89.0 %) achieved CR/PR, 5 patients (6.9 %) achieved SD/PD, and the data are not available for 3 patients (4.1 %); among patients with metastatic disease, 22 patients (88.0 %) achieved CR/PR and 3 patients (12.0 %) achieved SD/PD ( $p = 1.000$ ). The overall median follow-up period was 37 months (range, 0–263 months); 37 months in metastatic disease (range, 0–213 months); and 43 months in localized disease (range, 0–263 months). At the time of analysis, 50

patients (51.0 %) experienced recurrence, and 41 of these patients (41.8 %) later died. Sites of first recurrence/progression were locoregional in 26 patients and distant metastases in 24 patients. Seven patients in whom the sites of first recurrence/progression were locoregional achieved CR and were still alive following second-line treatment.

Adult patients with localized disease had a significantly greater probability of poor outcome compared with children. The median PFS times for localized and metastatic disease for children and adults were as follows: localized disease, 166.9 versus 22.4 months ( $p = 0.005$ ) (Fig. 1a), and metastatic disease, 13.3 versus 13.3 months ( $p = 0.949$ ) (Fig. 1b), respectively. Median OS times were not statistically different in patients with metastatic disease for both adults and children.

#### Analyses of prognostic factors in localized and metastatic disease

To determine the independent predictors of survival, we used a multivariate Cox regression model. The results of multivariate analysis for PFS and OS in localized and metastatic disease are shown in Tables 4 and 5. According to Table 4, age (<21 vs. ≥21) was the only statistically significant negative predictor of PFS for patients with localized disease ( $p = 0.018$ ). In contrast, for metastatic disease, age was not significantly different with respect to