

## PolyI:C and mouse survivin artificially embedding human 2B peptide induce a CD4<sup>+</sup> T cell response to autologous survivin in HLA-A\*2402 transgenic mice

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### ARTICLE INFO

#### Article history:

Received 28 March 2014

Received in revised form 4 August 2014

Accepted 6 August 2014

Available online 23 August 2014

#### Keywords:

Survivin

PolyI:C

CD4 epitope

Peptide vaccine

Th1 response

Interferon- $\gamma$

Tumor immunity

### ABSTRACT

CD4<sup>+</sup> T cell effectors are crucial for establishing antitumor immunity. Dendritic cell maturation by immune adjuvants appears to facilitate subset-specific CD4<sup>+</sup> T cell proliferation, but the adjuvant effect for CD4 T on induction of cytotoxic T lymphocytes (CTLs) is largely unknown. Self-antigenic determinants with low avidity are usually CD4 epitopes in mutated proteins with tumor-associated class I-antigens (TAAs). In this study, we made a chimeric version of survivin, a target of human CTLs. The chimeric survivin, where human survivin-2B containing a TAA was embedded in the mouse survivin frame (MmSVN2B), was used to immunize HLA-A-2402/K<sup>b</sup>-transgenic (HLA24<sup>b</sup>-Tg) mice. Subcutaneous administration of MmSVN2B or xenogeneic human survivin (control HsSNV2B) to HLA24<sup>b</sup>-Tg mice failed to induce an immune response without co-administration of an RNA adjuvant polyI:C, which was required for effector induction *in vivo*. Although HLA-A-2402/K<sup>b</sup> presented the survivin-2B peptide in C57BL/6 mice, 2B-specific tetramer assays showed that no CD8<sup>+</sup> T CTLs specific to survivin-2B proliferated above the detection limit in immunized mice, even with polyI:C treatment. However, the CD4<sup>+</sup> T cell response, as monitored by IFN- $\gamma$ , was significantly increased in mice given polyI:C + MmSVN2B. The Th1 response and antibody production were enhanced in the mice with polyI:C. The CD4 epitope responsible for effector function was not Hs/MmSNV<sub>13-27</sub>, a nonconserved region between human and mouse survivin, but region 53-67, which was identical between human and mouse survivin. These results suggest that activated, self-reactive CD4<sup>+</sup> helper T cells proliferate in MmSVN2B + polyI:C immunization and contribute to Th1 polarization followed by antibody production, but hardly participate in CTL induction.

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### Introduction

Dendritic cells (DCs) present exogenous antigens (Ags) to cells in the major histocompatibility complex (MHC) class I-restricted Ag-presentation pathway and cause the proliferation of CD8<sup>+</sup> T

cells specific to the extrinsic Ag. When tumor cells have soluble and insoluble exogenous Ags, MHC class I Ag presentation is mainly transporter associated with antigen processing (TAP)- and proteasome-dependent, suggesting the pathway is partly shared with the pathway for endogenous Ag presentation. The delivery of exogenous Ag by DCs to the pathway for MHC class I-restricted Ag presentation is called cross-presentation (Bevan 1976).

PolyI:C is a double-stranded RNA analog that activates RNA-sensing pattern-recognition receptor pathways (Matsumoto and Seya 2008; Seya and Matsumoto 2009). PolyI:C is an efficient trigger of cross-presentation, and facilitates cross-priming of CD8<sup>+</sup> T cells in the presence of Ag. Tumor-associated antigens (TAAs) usually expressed in low levels are thought to need support from pattern-recognition receptor activation to induce TAA-specific cytotoxic T lymphocytes (CTLs) (Seya et al. 2013).

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Many TAAs have been identified and tested for tolerability to patients and for ability to suppress tumor progression. Peptide vaccine immunotherapy against cancer has been studied clinically (Rosenberg et al. 2004). Survivin (SVN) is a TAA that generates CTLs in cancer patients (Schmitz et al. 2000; Andersen et al. 2001). Human survivin (HsSVN) is a 16.5 kDa cytoplasmic protein that inhibits caspase 3 and 7 in cells stimulated to undergo apoptosis (Altieri 2001). SVN is a member of the inhibitor of apoptosis protein family associated with fetal development. Therefore, except for testis, thymus and placenta, normal tissues express little SVN (Ambrosini et al. 1997; Altieri 2001). SVN is required in early thymocyte development from CD4/CD8-double-negative cells to CD4/CD8-double-positive lymphocytes (Okada et al. 2004). SVN is expressed in a wide variety of malignant cells (Altieri 2001; Fukuda and Pelus 2006). There are several splicing variants including a variant HsSVN2B with a cryptic epitope for MHC class I in humans. An HsSVN2B peptide (AYACNTSTL: 80–88) is an HLA-A\*2402-restricted peptide recognized by CD8+ CTLs (Hirohashi et al. 2002). Some cancer cells have higher mRNA levels of the HsSVN splice variant 2B, but whether this splice variant functions in tumorigenesis is unknown (Li 2005).

Several trials have studied the SVN2B peptide in cancer patients (Tsuruma et al. 2008; Honma et al. 2009; Kameshima et al. 2013). Although CTLs specific for SVN were detected in peripheral blood mononuclear cells of most cancer patients, as determined by HLA-A\*2402/SVN2B tetramer assays, no substantial therapeutic effect on cancer is seen in most clinical studies. A phase I clinical study found that vaccination with SVN2B peptide combined with IFN- $\alpha$  had significant therapeutic benefits in advanced pancreatic cancer patients, in spite of IFN-mediated side effects. Thus, an IFN-inducing adjuvant, that simultaneously up-regulates Ag-presentation and IFN-inducible genes, might more efficiently contribute to the clinical benefits of SVN for cancer patients.

PolyI:C is an analog of virus double-stranded RNA with IFN-inducing adjuvant properties. To test the effect of polyI:C on survivin-derived CTLs, we used a mouse model expressing human HLA-A24 that presents the SVN2B peptide (Gotoh et al. 2002). Mice have no splice counterpart for HsSVN2B and therefore mouse survivin (MmSVN) lacks the 2B portion of HsSVN, although the mouse ortholog is 84% homologous to HsSVN (Kobayashi et al. 1999). When BALB/c mice are injected intraperitoneally with HsSVN2B + RNA adjuvant, high levels of CD4+ T cells are induced in splenic T cells, as determined by IFN- $\gamma$ , TNF- $\alpha$ , and IL-2 production, as well as development of lytic MHC class II-restricted T cells and memory (Charalambous et al. 2006).

The N-terminal sequence of HsSVN, which includes amino acids 13–27 (FLKDHRISTFKNWPF), differs from that of MmSVN (YLKNIYRIATFKNWPF) (Charalambous et al. 2006). Therefore, high frequencies of self-reactive CD4+ T cells specific for a tumorigenic protein might be elicited in mice with xenogeneic HsSVN. However, self-reactive CD4+ T cells can be induced toward syngeneic or nonmutated CD4 epitopes in cancer patients (Topalian et al. 1996; Osen et al. 2010). To test the possibility that sub-derived self-CD4 epitopes participate in CD8+ CTL proliferation, we made a chimeric survivin protein (MmSVN2B), where the human 2B exon sequence was embedded into MmSVN. We immunized HLA-A-2402/K<sup>b</sup>-transgenic (HLA24<sup>b</sup>-Tg) B6 mice with MmSVN2B. The results indicated that the CD8+ CTL response to a self-tumor Ag (2B peptide) was barely enhanced by treatment of HLA24<sup>b</sup>-Tg mice with MmSVN2B in the presence of polyI:C. However, CD4+ T cell immune responses to the CD4 epitope of MmSVN2B and HsSVN2B were significantly enhanced in HLA24<sup>b</sup>-Tg mice with SVN2B proteins + polyI:C. The CD4 epitopes were not the N-terminal HsSVN<sub>13–27</sub> and MmSVN<sub>13–27</sub> sequences, but the Hs/MmSVN<sub>53–67</sub> (DLAQFFCFKELEGW) sequence, which is identical in HsSVN2B and MmSVN2B and thus a nonmutated CD4 epitope.

PolyI:C was required for proliferation of self-reactive CD4+ Th1 cells that recognized the syngeneic epitope. We discuss how RNA adjuvant might induce CD4+ Th1 cells and act in the antitumor immune response.

## Materials and methods

### Bioinformatics analysis

Ensembl databases (<http://asia.ensembl.org/index.html>) were used to investigate human and mouse SVN genomic structure. Primate and rodent short interspersed nuclear elements (SINES) were predicted using the Repeat Masker program (<http://www.repeatmasker.org/>). Results from databases were confirmed by comparison to previous reports (Mahotka et al. 1999).

### Expression analysis

Total RNA was extracted from tissues from C57BL/6 mice and murine cell lines using RNeasy Mini Kits (Qiagen) following the manufacturer's instructions. RT-PCR used High Capacity cDNA Reverse Transcription Kits (Applied Biosystems) according to the manufacturer's instructions. Primer pairs were designed to span separate exons to avoid amplifying other genomic DNA. Primers were 5'-ACTACCGCATCGCCACCT-3' (forward) and 5'-GCTTGTGTTGGTCTCCTTTG-3' (reverse) for detection of the murine SVN gene (MmSVN) and 5'-TGTAACCAACTGGGACGATAT-3' (forward) and 5'-CTTTTCACGGTTGGCCTTAG-3' (reverse) for murine *Gapdh*. PCR conditions for mSVN were 94 °C 3 min; 35 cycles of 94 °C 30 s, 65 °C 30 s, 72 °C for 30 s; and 7 min 72 °C. *Gapdh* PCR conditions were 94 °C 3 min; 30 cycles of 94 °C 30 s, 65 °C 30 s, and 72 °C 30 s; and 7 min at 72 °C.

### Antigens

The HsSVN2B-coding sequence was amplified using primers 5'-CGGGATCCATGGGTGCCCGACG-3' (underline: *Bam*HI site) and 5'-GGAATTCATCCATGGCAGC-3' (underline: *Eco*RI site). To construct the mSVN 2B gene (MmSVN2B), we used two-step PCR to make a chimeric gene of the mSVN gene and the human 2B exon (Fig. 2). In the first PCR, two fragments containing exon 1–2 and exon 3–4 were amplified using primers 5'-CCGCTCGAGATGGGAGCTCCGGCGCT-3' (underline: *Xho*I site) and 5'-ACCGTGCCCGGCCAATCGGGTTGTCA-3' (italics: 5'-end of exon 2B of the HsSVN2B gene) for exon 1 and exon 2 and 5'-GGGCGGATCACGAGAGAGGAGCATAGAAAGCA-3' (italics: 3'-end of exon 2B) and 5'-CGGGATCCTTAGGCAGCCAGCTGCTCAAT-3' (underline: *Bam*HI site) for exon 3 and exon 4. The exon 2B fragment was amplified using primers 5'-CGATGACAACCCGATTGGGCCGGGCACGG-3' (italics: 3'-end of exon 1 and exon 2 of MmSVN) and 5'-TTTCTATGCTCTCTCTCGTGATCCGCC-3' (italics: 5'-end of exon 3 and exon 4 of MmSVN). In the second PCR, the three templates from the first PCR were mixed in equal amounts and amplified using primers 5'-CCGCTCGAGATGGGAGCTCCGGCGCT-3' (underline: *Xho*I site) and 5'-CGGGATCCTTAGGCAGCCAGCTGCTCAAT-3' (underline: *Bam*HI site). The pCold vector II (TaKaRa) and SVN fragments were restriction digested and ligated overnight with T4 ligase (Promega) at 4 °C. Ligation mixtures were transformed into competent *Escherichia coli* strain BL21 (DE3) cells. After preculturing for 2 h at 37 °C, cells were cooled on ice. Recombinant protein expression was induced with isopropyl-1-thio- $\beta$ -D-galactopyranoside at a final concentration of 1 mM and cultured for 24 h at 16 °C. N-His-tagged survivin proteins were purified using a Profinia protein purification system (Biorad). Buffer of

purified SVN proteins was sequentially exchanged with PBS containing 2 M urea. To rule out lipopolysaccharide contamination, we treated survivin proteins with 200 µg/ml of polymixin B (Sigma) for 30 min at 37 °C before use. OVA (ovalbumin) (Sigma) was similarly treated with polymixin B as an Ag.

### Mice

C57BL/6 (H-2b) mice were from Clea Japan (Tokyo). HLA24<sup>b</sup>-Tg was from SLC Japan (Gotoh et al. 2002). Mice were maintained in the Hokkaido University Animal Facility (Sapporo, Japan) in specific pathogen-free conditions. All experiments used mice that were 8–12 weeks old at the time of first procedure. All mice were used according to the guidelines of the institutional animal care and use committee of Hokkaido University, which approved this study (ID number: 08-0243, “Analysis of immune modulation by toll-like receptors”).

### Reagents, antibodies and cells

PolyI:C and OVA<sub>323–339</sub> peptide (ISQAVHAAHAEINEAGR) were from Sigma. OVA<sub>257–264</sub> peptide (SIINFEKL: SL8), OVA (H2K<sup>b</sup>-SL8), HLA-A\*2402 survivin-2B and HIV tetramer were from MBL. SVN2B peptide (AYACNTSTL) and HLA-A\*2402/2B peptide-restricted human T cell clones (Ikenoue et al. 2005) were kindly provided by Dr. Noriyuki Sato (Department of Pathology, School of Medicine, Sapporo Medical University). Human and murine-specific helper peptides (Charalambous et al. 2006) MmSVN<sub>13–27</sub> (YLKNYRIATFKNWPF) and Hs SVN<sub>13–27</sub> (FLKDHRISTFKNWPF) and the common helper peptide Hs/Mm SVN<sub>53–67</sub> (DLAQCFKCFKELEGW), were synthesized by Biologica Co. Ltd (Nagoya). Peptide purity was >95%. To eliminate lipopolysaccharide contamination, all peptides were treated with 200 µg/ml polymixin B (Sigma) for 30 min at 37 °C before use (Nishiguchi et al. 2001). Anti-CD3 $\epsilon$  (145-2C11), anti-CD8 $\alpha$  (53-6.7) and anti-IFN $\gamma$  (XMG1.2) antibodies (Abs) were from BioLegend. Anti-CD4 Ab (L3T4) was from eBiosciences and ViaProbe was from BD Biosciences. Dendritic cells were prepared from spleens of mice as described previously (Azuma et al., 2012).

### Antigen-specific T cell expansion in vivo

HLA24<sup>b</sup> Tg mice (Gotoh et al. 2002) were subcutaneously immunized with 100 µg of each antigen and 100 µg poly I:C once a week for 4 weeks. After 7 days from the last immunization, spleens were extracted, homogenized and stained with FITC-CD8 $\alpha$  and PE-OVA (Azuma et al. 2012) or PE-SVN2B tetramer for detecting antigen-specific CD8<sup>+</sup> T cells (Tsuruma et al. 2008). For intracellular cytokine detection, splenocytes were cultured with 100 nM SL8 or survivin 2B peptide for 6 h with 10 µg/ml brefeldin A (Sigma–Aldrich) added in the last 4 h. For intracellular cytokine detection of antigen-specific CD4<sup>+</sup> T cells, splenocytes were cultured with 100 nM OVA<sub>323–339</sub> peptide or SVN helper peptide for 6 h with 10 µg/ml brefeldin A (Sigma–Aldrich) added in the last 5 h. Cells were stained with PE-anti-CD8 $\alpha$ /FITC-anti-CD3 $\epsilon$  for CD8<sup>+</sup> T cells or PE-anti-CD4/FITC-anti-CD3 $\epsilon$  for CD4<sup>+</sup> T cells. After cell-surface staining, cells were fixed and permeabilized with Cytofix/Cytoperm (BD Biosciences) according to the manufacturer's instruction. Fixed and permeabilized cells were stained with APC-anti-IFN $\gamma$ . Stained cells were analyzed with FACSCalibur (BD Biosciences) and FlowJo software (Tree Star) (Azuma et al. 2012).

### ELISA

Sera were collected from immunized mice once a week for 4 weeks and 96-well plates were coated with 10 µg/ml OVA,

MmSVN2B and HsSVN2B in ELISA/ELISPOT coating buffer (eBioscience) and incubated overnight at 4 °C. ELISA diluent solution (eBioscience) was used for blocking and antibody dilution. PBS with 0.05% Tween 20 was used for washes. Anti-OVA or anti-SVN in sera was assessed by ELISA using antiserum for IgG2a/b and IgG1 diluted 1000-fold and 10,000-fold and incubated for 2 h at room temperature. After washing, isotype IgGs were detected using goat anti-mouse total IgG, IgG1, or IgG2a conjugated to HRP (Southern Biotechnology Associates). After washing, plates were stained with 1XTMB ELISA substrate solution (eBioscience) and reactions stopped with 2 N H<sub>2</sub>SO<sub>4</sub> before measuring absorbance.

### Statistical analyses

For comparison of two groups, *P*-values were calculated with a Student's *t*-test. For comparison of multiple groups, *P*-values were calculated with one-way analysis of variance (ANOVA) with Bonferroni's test. Error bars are SD or SEM between samples.

## Results

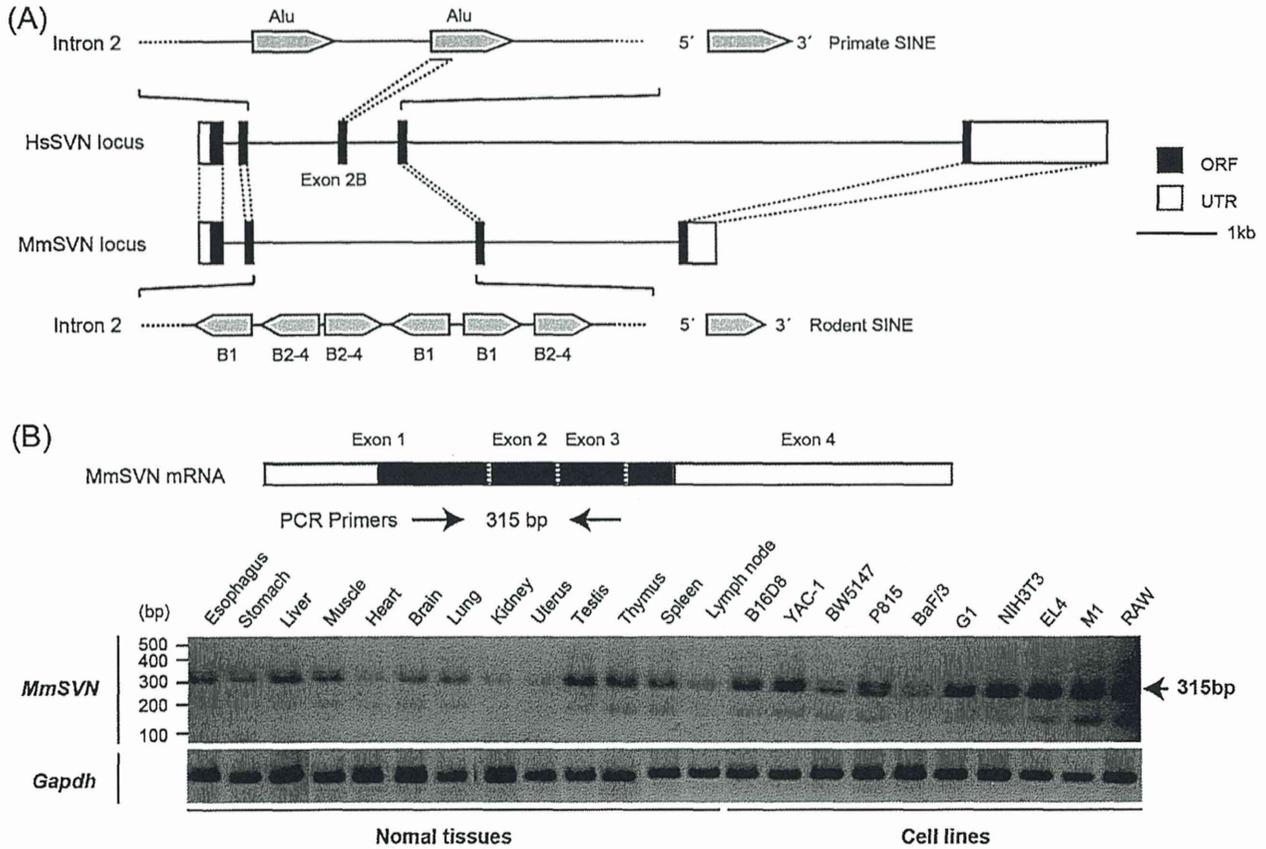
### Origin of human SVN exon 2B

The HsSVN gene has four conserved and two cryptic exons (Mahotka et al. 1999). The authentic HsSVN gene encode 142 amino acids in exons 1–4. On the other hand, the HsSVN2B product is 165 amino acids encoded by exons 1, 2, 2B, 3 and 4. Exon 2B is hidden within intron 2, which is spliced into mature HsSVN2B mRNA in-frame between exons 2 and 3 (Mahotka et al. 1999). Exon 2B is followed by the GT-AG rule and expressed in many tumor cells and tumor cell lines, suggesting that splicing predominantly occurs in malignantly transformed cells (Mahotka et al. 2002). According to the Ensembl database, HsSVN intron 2 had two Alu sequences (Fig. 1A), and exon 2B resulted from the second Alu. In contrast, the MmSVN gene had four exons separated by three introns with no Alu sequence in intron 2; instead, MmSVN had several SINE sequences characteristic of rodents in intron 2 (Fig. 1A). Although the exon sequences were conserved in human and mouse SVNs, two intron sequences diverged between human and mouse (Fig. 1A). These results suggested that integration of exon 2B was evolutionarily new and formed after an Alu insertion. Although the SVN gene is conserved in yeast and humans, exon 2B was established after the divergence of human and mouse.

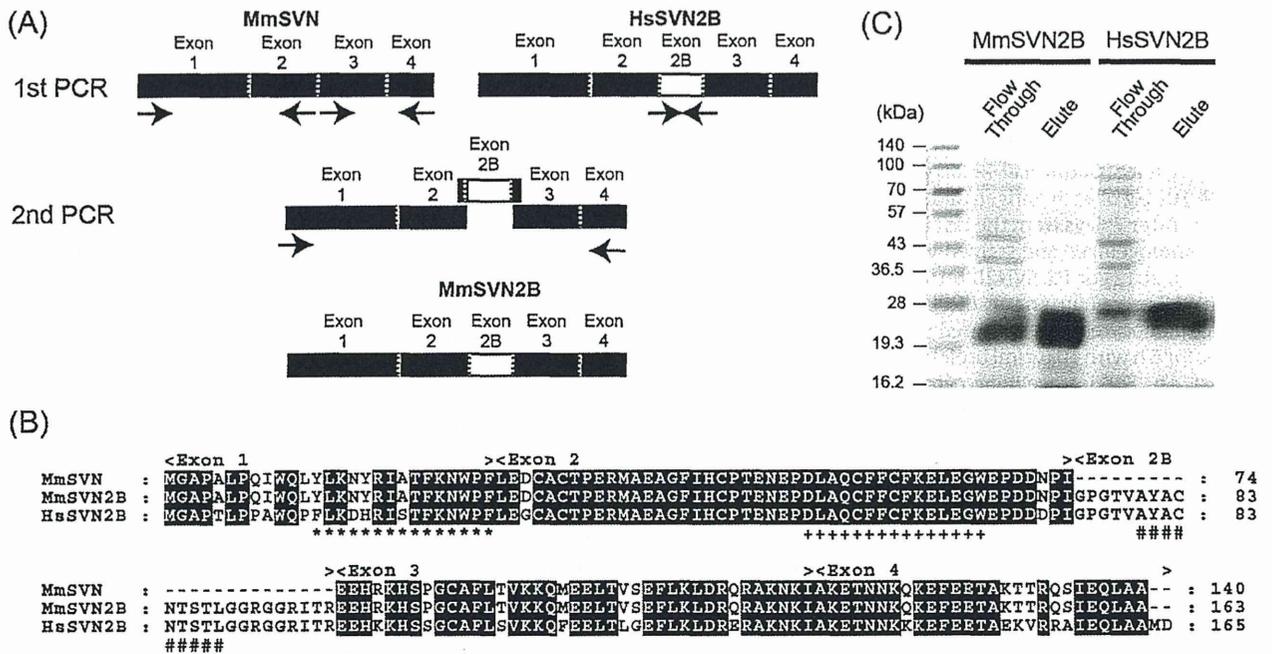
We used RT-PCR to investigate transcripts resulting from splicing other exons around exon 2 into the MmSVN mRNA. Results of mRNAs from mouse organs and cell lines are in Fig. 2B. The results suggested that no alternative exons around exon 2 in the MmSVN gene. We detected a ~200 bp product in most organs and cell lines tested (Fig. 2B), but this was not an MmSVN transcript.

### Generation of a mmSVN2B construct

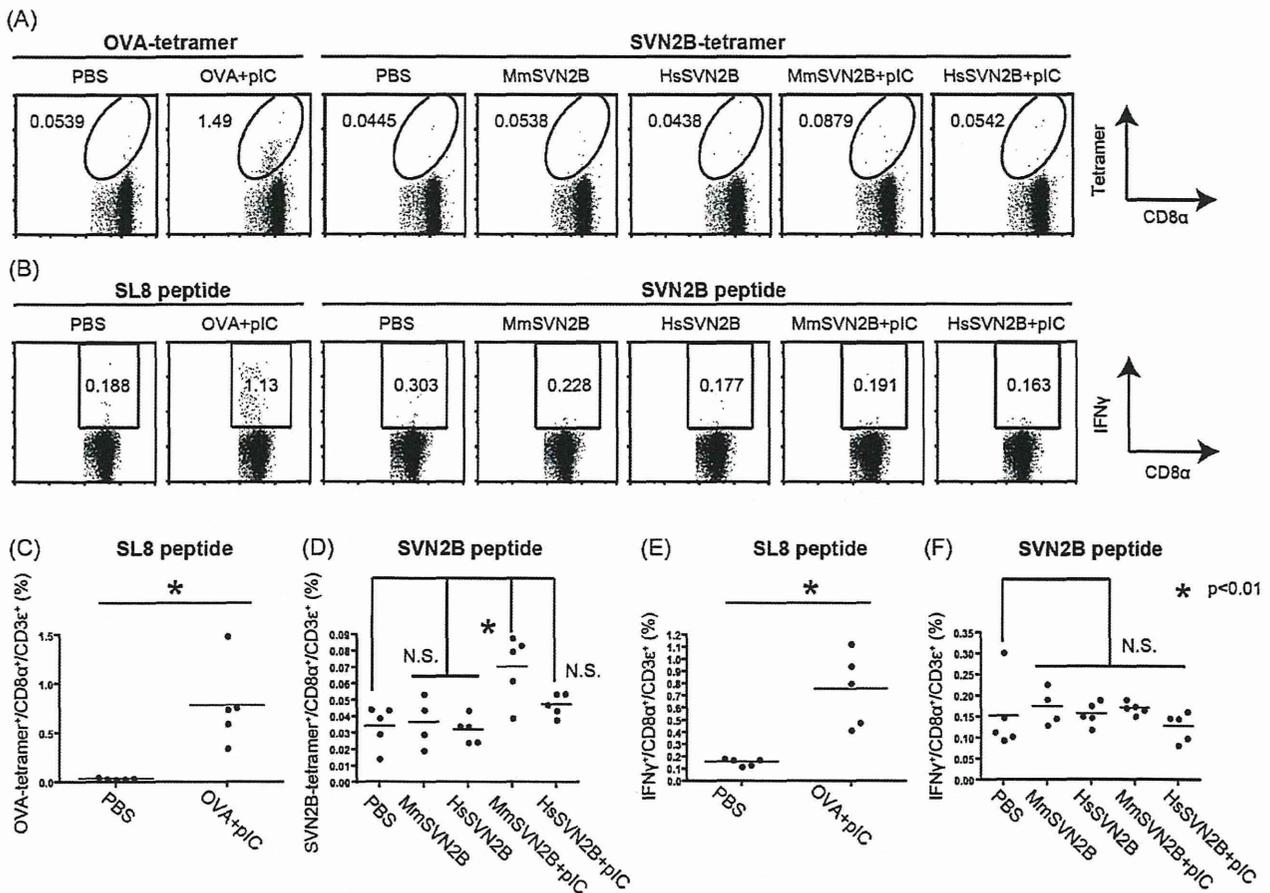
A SVN2B peptide derived from the HsSVN2B gene that contained the exon 2B sequence was recognized by CTLs in cancer patients (Hirohashi et al. 2002; Tsuruma et al. 2008; Honma et al. 2009) and a CTL clone was established from patients (Ikenoue et al. 2005). We artificially constructed an MmSVN2B with a xenogeneic human exon 2B inserted into the boundary between exon 2 and 3 of SVN (Fig. 2A and B). Prominent amino acid substitutions between MmSVN2B and HsSVN2B were concentrated in the N-terminal region encoded by exon 1 (Fig. 2B), and a CD4 epitope is in this region (Li 2005; Mahotka et al. 2002). In an earlier paper, this HsSVN<sub>13–27</sub> region, but not MmSVN<sub>13–27</sub>, was an effective CD4 epitope that promoted HsSVN<sub>13–27</sub>-specific CD4<sup>+</sup> T cell proliferation



**Fig. 1.** Genome structure and expression of human and murine SVN gene. (A) Comparison of human and murine survivin gene structure. Survivin gene structures were defined by the Ensembl genome browser. Primate and rodent SINES were predicted using Repeat Masker program. Filled boxes, coding regions; open boxes, 5'- and 3'-untranslated regions. (B) Structure of murine survivin transcript and RT-PCR analysis of organs and cell lines. Arrows, survivin-detecting PCR primers.



**Fig. 2.** Structure and purification of chimeric MmSVN2B protein. (A) Strategy for constructing chimeric MmSVN2B protein. Human exon 2B was inserted into MmSVN by PCR. (B) Alignment of murine and human SVN sequences. Black shaded area, residues conserved between human and murine SVN; Hs, human; Mm, mouse. \*, MmSVN<sub>13-27</sub>/HsSVN<sub>13-27</sub> peptide; +, Hs/Mm SVN<sub>53-67</sub> peptide; #, SVN2B peptide. (C) Purification of N-His-tagged MmSVN2B and HsSVN2B proteins. N-His-tagged SVN proteins were purified using a Profinia protein purification system from BL21 (DE3) competent cells. Purified SVN protein buffer was sequentially exchanged to PBS containing 2 M urea.



**Fig. 3.** Expansion of OVA and SVN-specific CD8<sup>+</sup> T cells. (A) HLA24<sup>b</sup>-Tg mice were immunized with 100 μg antigen and 100 μg polyI:C once a week for 4 weeks. After 7 days from the last immunization, spleens were homogenized and stained with FITC-CD8α and PE-OVA or PE-survivin tetramer to detect antigen-specific CD8<sup>+</sup> T cells. (B) Splenocytes were cultured *in vitro* in the presence of SL8 or SVN2B peptides for 6 h and IFN-γ production was measured by FACS. (C, D) Average percentages of OVA-positive and SVN2B-tetramer positive CD8<sup>+</sup> T cells shown in (A). (E, F) Average percentages of IFN-γ producing CD8<sup>+</sup> T cells specifically in response to SL8 or SVN2B peptide in (B). \**p* < 0.01.

(Charalambous et al. 2006). His-tagged MmSVN2B and HsSVN2B proteins were purified and used as Ags (Fig. 2C).

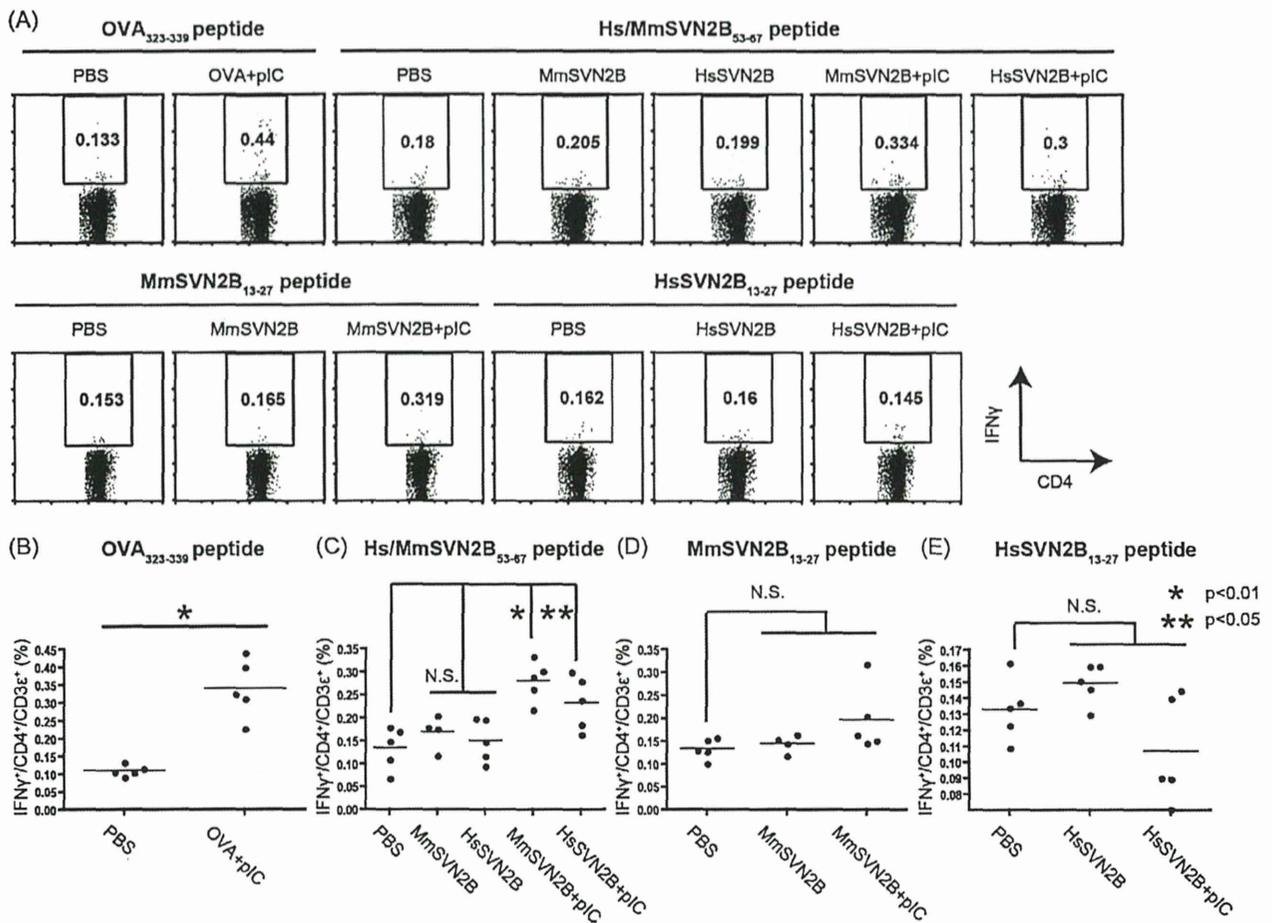
#### CD4<sup>+</sup> and CD8<sup>+</sup> T cells that react to MmSVN2B plus polyI:C

We examined the ability of MmSVN2B to induce IFN-γ and CD8<sup>+</sup> T cell proliferation by immunizing HLA24<sup>b</sup>-Tg mice with MmSVN2B or HsSVN2B with or without polyI:C (Fig. 3). SVN2B-specific CTLs were probed by SVN2B-tetramer (Fig. 3A) and IFN-γ staining (Fig. 3B). SVN2B-specific human CD8<sup>+</sup> T cells were detected with SVN2B-tetramer (Fig. S1), which enabled us to search for SVN2B-specific CTLs in HLA24<sup>b</sup>-Tg mice (Ikenoue et al. 2005). Expression of CD40 was up-regulated in CD8α<sup>+</sup> conventional DCs to a similar extent with MmSVN2B or HsSVN2B (Fig. S2), consistent with a report on CD40 that promotes cross-priming by Ahonen et al. (J Exp Med, 2004). OVA and polyI:C were used as positive controls (Fig. 3A, B left panels), and SL8 (SIINFEKL)-specific CTLs were monitored with OVA tetramer (Azuma et al. 2012). Both OVA-tetramer-positive and IFN-γ-producing CD8<sup>+</sup> T cells were detected in mice immunized with OVA and polyI:C (Fig. 3C, E). Without polyI:C stimulation, only small number of OVA-tetramer-positive cells were upregulated compared to controls (Azuma et al. 2012; Azuma & Seya unpublished data).

When HLA24<sup>b</sup>-Tg mice were immunized with MmSVN2B or HsSVN2B without polyI:C, no significant induction of SVN2B-tetramer-positive (Fig. 3D) or IFN-γ-inducing cells was observed

(Fig. 3F). When polyI:C was included, only a small increase in SVN2B-tetramer-positive cells was detected in mice given MmSVN + polyI:C with no significant increase in IFN-γ (Fig. 3F). Mice receiving HsSVN + polyI:C (Fig. 3D) or polyI:C alone (not shown) showed no significant increase in SVN2B-specific CD8<sup>+</sup> T cells. Consistent with the lack of tetramer-positive CTL induction, MmSVN2B treatment failed to regress MmSVN2B-transfected tumor cells implanted into HLA24<sup>b</sup>-Tg mice. In EG7 tumor-bearing mice, administration of polyI:C alone (without Ag) induces tumor-growth retardation due to the contribution of endogenous Ag (Azuma et al. 2012), but in this case with tumor-unloaded mice polyI:C exhibited no tumor-regressing activity (data not shown), possibly due to the lack of Ag.

Next, we determined the amounts of CD4<sup>+</sup> T cells that reacted with MmSVN2B. The positive control group received OVA Ag and polyI:C (Fig. 4A, B). The negative control group received PBS without Ag and polyI:C, but basal frequencies of IFN-γ-producing CD4<sup>+</sup> T cells were detected in this group even in the absence of polyI:C or Ag (Fig. 4). When MmSVN2B or HsSVN2B only was used to immunize mice, no significant response was seen in CD4<sup>+</sup> T cells compared to PBS controls (Fig. 4A, C–E). When polyI:C was included, IFN-γ-producing CD4<sup>+</sup> T cells restimulated with Hs/MmSVN<sub>53–67</sub> peptide increased significantly in mice that received MmSVN and HsSVN (Fig. 4C, D). The sequence of MmSVN<sub>53–67</sub> was identical to the sequence of HsSVN<sub>53–67</sub> (Fig. 2B). However, we did not detect a significant increase in IFN-γ-producing CD4<sup>+</sup> T cells in mice



**Fig. 4.** Expansion of OVA and SVN-specific CD4<sup>+</sup> T cells. (A) HLA24<sup>b</sup>-Tg mice were immunized with 100 μg each antigen and 100 μg poly I:C once a week for 4 weeks. After 7 days from the last immunization, splenocytes were cultured with 100 nM OVA<sub>323-339</sub> peptide or SVN helper peptide for 6 h, and 10 μg/ml brefeldin A (Sigma–Aldrich) was added in the last 5 h. After cell surface and intracellular staining, IFN-γ production of CD4<sup>+</sup> T cells was measured by FACS. Average percentages of IFN-γ-producing CD4<sup>+</sup> T cells in response to (B) OVA<sub>323-339</sub> peptide; (C) Hs/Mm SVN<sub>53-67</sub> peptide; (D) MmSVN<sub>13-27</sub> peptide; (E) HsSVN<sub>13-27</sub> peptide. \**p* < 0.01, \*\**p* < 0.05.

restimulated with MmSVN<sub>13-27</sub> or HsSVN<sub>13-27</sub> peptide (Fig. 4D, E). Differences in these two CD4 epitope sequences are in Fig. 2B.

*Ab production by immunization with MmSVN2B with polyI:C*

Activation of Th1 cells is essential for B cell antibody class switching. Therefore, we examined production of SVN-specific Ab in Tg mice that did or did not receive polyI:C. Serum was collected from HLA24<sup>b</sup>-Tg mice immunized with different Ags and polyI:C. OVA and polyI:C were the positive control and resulted in a significant increase in OVA-specific IgG1, IgG2a and IgG2b by ELISA (Fig. 5 left panels). When HLA24<sup>b</sup>-Tg mice were immunized with MmSVN2B or HsSVN2B without polyI:C, no significant production of any isotypes was observed (Fig. 5 center and right panels). When polyI:C was included, MmSVN2B or HsSVN2B-specific isotypes increased significantly.

**Discussion**

We demonstrated that HLA24<sup>b</sup>-Tg mice induced Hs/MmSVN<sub>53-67</sub>-specific CD4<sup>+</sup> T cells and SVN-specific Ab followed by Th1 cell activation in response to injection of polyI:C and MmSVN2B protein. This result was partly inconsistent with a previous report (Charalambous et al. 2006) using Balb/c mice and HsSVN conjugated to Dec205 mAb. That is, our study with C57BL/6 mice and MmSVN2B did not detect significant increases

in MmSVN<sub>13-27</sub>-specific CD4<sup>+</sup> T cells after subcutaneous injection of MmSVN2B with polyI:C. Thus, the xenogeneic differences in sequence between HsSVN and MmSVN did not always contribute to generating effective CD4<sup>+</sup> T cells specific for a tumorigenic protein in C57BL/6 mice. The haplotype of the MHC class II proteins between Balb/c (having H-2d) and C57BL/6 mice (having H-2b) and Dec205 mAb conjugation (Charalambous et al. 2006) might be the reason for these different results. However, no CD8<sup>+</sup> CTLs against the 2B peptide were detected even when using a specific tetramer for detection of CD8<sup>+</sup> CTLs (Fig. S1). Hence, polyI:C was required for proliferation of self-reactive CD4<sup>+</sup> Th1 cells that recognized the syngeneic epitope without proliferation of SVN2B peptide-specific CTLs.

OVA were used as positive controls (Fig. 3A, B left panels), and SL8 (SIINFEKL)-specific CTLs were monitored with OVA tetramer (Azuma et al. 2012). Here, T cell activation by polyI:C + MmSVN2B is a focus in this study. However, there is a lot-to-lot difference of T cell-activating activity in polyI:C + OVA as in our present and previous studies (Azuma et al. 2012). This difference of T cell activation may be attributable to the fact that polyI:C consists of a variety of length of polyI chains and polyC chains with a lot-to-lot heterogeneity. In addition, the amounts of Ags in Azuma's experiment are higher than those in the present experiment (Azuma et al. 2012). CD40 stimulation by specific Ab results in high enhancement of cross-priming of CD8 T cells (Charalambous et al. 2006) and CD40 was up-regulated in CD8α<sup>+</sup> DCs by polyI:C treatment, but the CD40

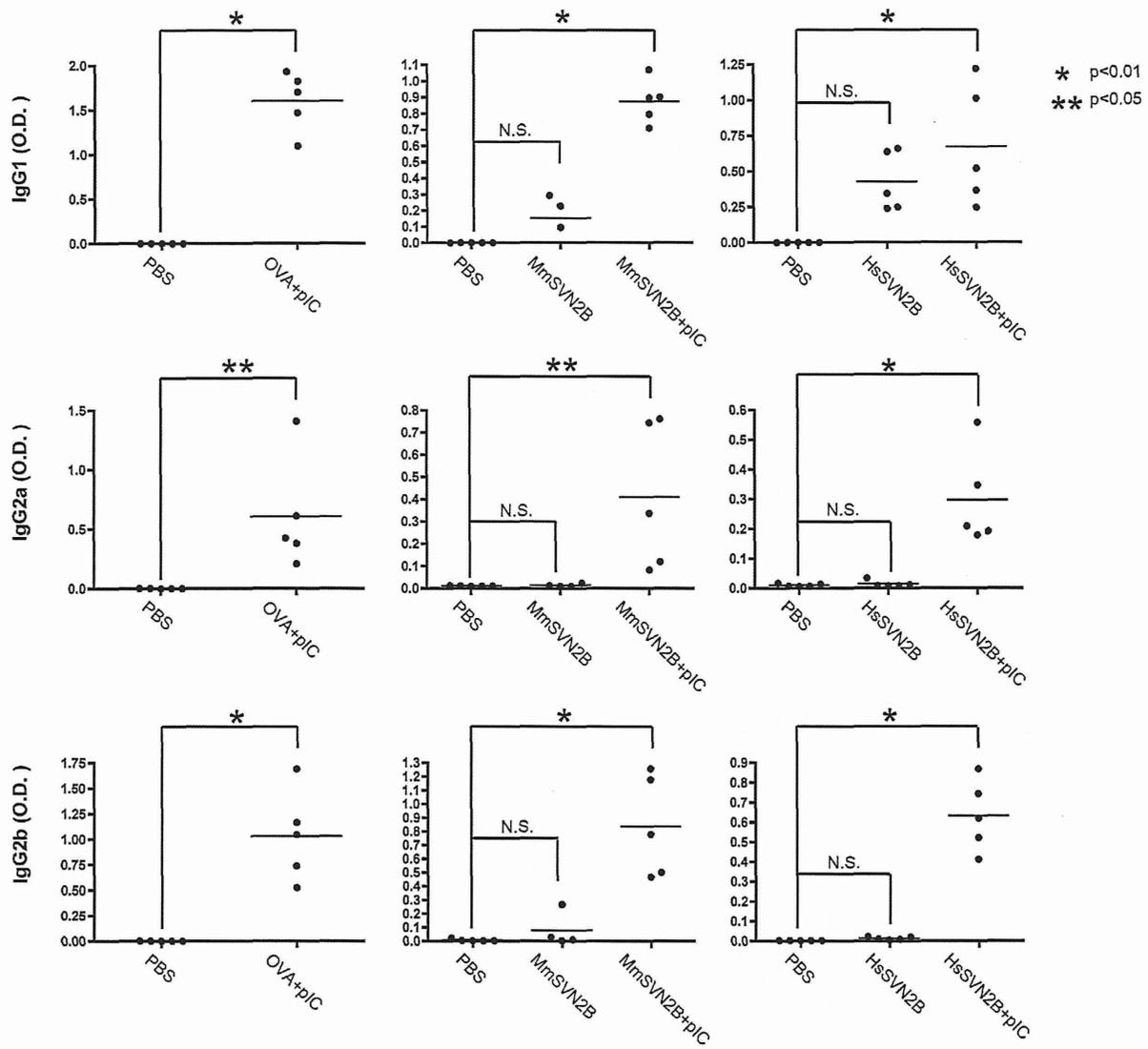


Fig. 5. Production of OVA and SVN-specific antibodies. Sera were collected from immunized mice at once a week for 4 weeks. Anti-OVA or anti-SVN in sera was assessed by ELISA using antiserum for IgG2a/b and IgG1. \* $p < 0.01$ , \*\* $p < 0.05$ .

levels were also variable depending upon the polyI:C lots. Development of a synthesizing method for defined length of RNA duplex will settle the issue.

Two points are noted. First, polyI:C, an RNA adjuvant, induces CD4<sup>+</sup> T cells in addition to the reported cases of CD8<sup>+</sup> T cells. The factors that participate in polyI:C-mediated CD4<sup>+</sup> T cell proliferation and the kind of CD4<sup>+</sup> T subsets that are predominantly induced by polyI:C remain unknown. PolyI:C is primarily a potential activator of the IFN-inducing pathways RIG-I/MDA5 and TLR3 (Matsumoto & Seya 2008). These pathways allow host immune cells to produce type I/III IFNs and cytokines and are soluble effectors against cancer. TLR3 preferentially induces cross-presentation in CD8 $\alpha^+$  DC in response to dsRNA including polyI:C (Schulz et al., 2005; Azuma et al. 2012) and causes proliferation of CD8<sup>+</sup> T cells including cells that respond to TAAs *via* cross-priming (Azuma et al. 2012). CD4<sup>+</sup> T cells that are likely evoked by polyI:C stimulation function in antitumor immunity since their helper function is usually suppressed in tumor-bearing mice and can be relieved by innate immune response (Lee et al. 2013). Stimulation with polyI:C + SVN Ag might change a tumor-derived suppressive environment to an environment suitable for primary activation and maintenance of

Ag-specific cytotoxic CD8<sup>+</sup> T cell responses (Ridge et al. 1998; Janssen et al. 2003).

According to a recent report, however, adoptively transferred CD4<sup>+</sup> T cells induce tumor rejection independently of CD8<sup>+</sup> T cells (Corthay et al. 2005; Perez-Diez et al. 2007). This rejection is apparently based on cytokines released from CD4<sup>+</sup> T cells (Corthay et al. 2005) and on interaction with CD4<sup>+</sup> T cells and other immune cells such as macrophages (Mfs) and natural killer (NK) cells (Perez-Diez et al. 2007). DCs stimulated with polyI:C also result in NK cell activation after DC-NK cell-to-cell contact (Akazawa et al. 2007). Mfs in tumors might be a direct target of dsRNA, which converts tumor-supporting Mfs into tumoricidal Mfs (Shime et al. 2012). IL-12p40 is preferentially produced *via* the TICAM-1/Batf3 pathway in response to dsRNA (Azuma et al. 2013). Thus, a variety of cellular effectors can be triggered as antitumor agents by administration of dsRNA with TAA peptides or proteins. We found that CD4<sup>+</sup> T cells with Th1 properties were effectors induced by polyI:C possibly acting as an antitumor agent in SVN-responding tumor cells. Although epitope sequence and hydrophobicity might affect Th1 polarization in mice, CD4<sup>+</sup> T effectors are successfully induced in tumor-bearing or tumor-implanted mice by stimulation with MmSVN2B + polyI:C.

Hence, *in vivo* administration of an RNA adjuvant with Ag proteins induce CD4<sup>+</sup> helper T cells secondary to class II presentation in DCs, together with induction of type I IFNs and cytokines. CD4<sup>+</sup> T cells also facilitate Ab production caused by stimulation of B cell development (Mak et al. 2003).

Notably, this is a specific feature of RNA adjuvants, since TLR2 agonist Pam2 lipopeptides such as Pam2CSK4 and MALP2s induce antitumor CTLs with sufficient potential (Chua et al. 2014) but fail to induce DC-mediated antitumor NK cell activation (Yamazaki et al. 2011; Sawahata et al. 2011). CD4<sup>+</sup> T cells with regulatory modes such as Tregs and Tr-1 cells and IL-10 were induced by Pam2 peptide in the presence of Ag (Yamazaki et al. 2011). Nevertheless, robust proliferation of antitumor CTL is induced by Pam2 lipopeptides (Chua et al. 2014). Thus, the mode of CD8<sup>+</sup> T cell proliferation is differentially modulated between TLR2 and TLR3/MDA5 agonists.

The other point is how self-Ag-reactive CD4<sup>+</sup> T cells that act as Th1 effectors in SVN-based immunotherapy are generated. Proliferation of self-reactive T cells is prevented in normal mice, so the levels of self-reactive T cells are usually lower than the detection limit of assays (Gebe et al. 2003). Self-reactive CD4<sup>+</sup> T cells might be positively regulated by polyI:C in the presence of protein antigen, since mice, when exposed to DNA/RNA, harbor autoimmune diseases against the protein (Mills 2011). However, even with Ag proteins, polyI:C induced minimal cross-priming of CD8<sup>+</sup> T cells in our setting, as with previous reports (Charalambous et al. 2006). In this and other studies, both syngeneic and xenogeneic CD4 epitopes prime CD4<sup>+</sup> T cells, stimulating Ab production and Th1 polarization with antitumor activity, but with little association with CTL induction (Charalambous et al. 2006). Our SVN results suggested that self-responsive CD4 epitopes that are identical in sequence in human and mouse SVN have a conserved function as a Th1 skewer, albeit modest, in mice by stimulating DCs and Mfs to prime T and B cells. In this context, however, a question remains to be settled about why the insertion of the 2B sequence in MmSVN caused induction of auto-reactive CD4<sup>+</sup> T cells secondary to the class II presentation of the common SVN sequence (53–67) rather than the reported uncommon 13–27 region.

Generally, the presence of Tregs and regulatory cytokines such as IL-10 usually suppresses the function of self-reactive CD4<sup>+</sup> T effectors, so an autoimmune response cannot be detected (Danke et al. 2004; Quezada et al. 2010). In tumor-bearing mice, polyI:C releases the restriction of T cell autoreactivity by Tregs to enhance CD4<sup>+</sup> T function in a tumor microenvironment. Although the level of Treg cells increases in MALP2s-stimulated tumor-bearing mice (Yamazaki et al. 2011), the amount of Treg cells is not affected by polyI:C injection (Chua et al. 2014). Signs of autoimmune diseases have not yet been observed in mice that received intermittent administration of polyI:C under our conditions. Further studies on the function of regulatory factors in tumor-bearing mice after treatment with various adjuvants are needed to determine the balance between CD4<sup>+</sup> T effector functions and regulatory factors including Tregs (Quezada et al. 2010; Corthay et al. 2005).

It has been reported that treatment of murine glioma with DCs loading MmSVN long overlapping peptide covering CD4 and CD8 epitopes (DC therapy) conferred good prognosis on tumor-bearing mice (Ciesielski et al. 2008). In previous trials on peptide vaccine therapy, SVN2B peptide + IFN- $\alpha$  resulted in clinical improvements and enhanced immunological responses of patients (Kameshima et al. 2013). Treatment with SVN2B peptide alone did not result in good prognosis or effective tumor regression in late stage patients with cancer, however (Tsuruma et al. 2008; Honma et al. 2009). These results suggest that both killer and helper T cells are required for *in vivo* induction of tumor regression, as previously suggested (Perez-Diez et al. 2007). NK cells, Mfs, and soluble and angiogenic factors might be involved in tumor rejection (Shime et al. 2012; Müller-Hermelink et al., 2008; Coussens and Werb 2002) in

addition to Ag + polyI:C. According to the study with Ag and polyI:C, a protein or long peptide Ag containing CD4 epitopes, adjuvant RNA and additional factors that disable immunoregulatory factors, are required to effectively induce TAA-specific killer and helper T cell proliferation and subsequent tumoricidal activity in future studies (Casares et al. 2001). Ag peptides should be designed to present both class I and class II peptides on DCs to facilitate proliferation of CD4<sup>+</sup> T cells and Ab production. Methods for inducing potential CD8<sup>+</sup> CTLs against tumors still need to be considered.

### Competing interests

The authors have declared that no competing interest exists.

### Acknowledgements

We are grateful to members in our laboratories. JK is a Research Fellow of the Japan Society for the Promotion of Science. This work was supported in part by Grants-in-Aid from the Ministry of Education, Science, and Culture and the Ministry of Health, Labor, and Welfare of Japan, and by a MEXT Grant-in-Project 'the Carcinogenic Spiral', 'the National Cancer Center Research and Development Fund (23-A-44)'. Financial support by the Takeda Science Foundation, the Yasuda Cancer Research Foundation and the Ono Foundation are gratefully acknowledged.

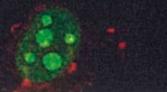
### Appendix A. Supplementary data

Supplementary data associated with this article can be found, in the online version, at <http://dx.doi.org/10.1016/j.imbio.2014.08.017>.

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**INAM Plays a Critical Role in IFN- $\gamma$  Production by NK Cells Interacting with Polyinosinic-Polycytidylic Acid-Stimulated Accessory Cells**

This information is current as of February 5, 2015.

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*J Immunol* 2014; 193:5199-5207; Prepublished online 15 October 2014;

doi: 10.4049/jimmunol.1400924

<http://www.jimmunol.org/content/193/10/5199>

**Supplementary Material** <http://www.jimmunol.org/content/suppl/2014/10/15/jimmunol.1400924.DCSupplemental.html>

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