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2 **MHC class I molecules are incorporated into human herpesvirus-6 viral particles**  
3 **and released into the extracellular environment**

4 Running title: MHC class I expresses in HHV-6 virions

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17

18    **Abstract**

19    Human herpesvirus-6 (HHV-6) belongs to the betaherpesvirus subfamily and mainly  
20    replicates in T lymphocytes. Here, we show that major histocompatibility complex  
21    (MHC) class I molecules are incorporated into HHV-6 viral particles and released into  
22    the extracellular environment. In addition, HHV-6A/B-infected T cells showed reduced  
23    surface and intracellular expression of MHC class I molecules. The cellular machinery  
24    responsible for molecular transport appears to be modified upon HHV-6 infection,  
25    causing MHC class I molecules to be transported to virion assembly sites.

26

27

28    **Key words**

29    HHV-6A/B, MHC class I, viral particles

30

## 31 **Introduction**

32 Human herpesvirus 6 (HHV-6), which belongs to the betaherpesvirus subfamily (1)  
33 was first isolated from peripheral blood lymphocytes obtained from patients with  
34 lymphoproliferative disorders (2). HHV-6 isolates are classified as HHV-6A and  
35 HHV-6B (see the Virus Taxonomy List 2011), based on genetic and antigenic  
36 differences and their cell tropism (3, 4, 2, 5). Primary infection with HHV-6B causes  
37 exanthem subitum (6). The diseases caused by HHV-6A have been unknown. HHV-6B  
38 mostly infects infants and remains latent in more than 90% of the population (7).

39 In general, herpesviruses use several strategies to evade host immune responses. For  
40 example, viruses may inhibit major histocompatibility complex (MHC) class  
41 I-associated antigen presentation to escape detection by cytotoxic T lymphocytes  
42 (CTLs). Several proteins expressed by herpesviruses block the transport of antigenic  
43 peptides from the cytosol to the endoplasmic reticulum (ER) (8-11), whereas others  
44 retain (12-14) or destroy class I molecules, or deliver them to lysosomes for degradation  
45 (15-18). The result is reduced surface expression of MHC class I molecules, enabling  
46 the virus to evade host immune surveillance.

47 HHV-6A, but not HHV-6B, downregulates the expression of MHC class I in dendritic  
48 cells (19). HHV-6 U21 binds to and diverts MHC class I molecules to an endolysosomal  
49 compartment, effectively removing them from the cell surface and providing a possible  
50 means of immune escape (20).

51 Here, we show that the expression of MHC class I molecules by infected cells was  
52 downregulated with incorporation into HHV-6 viral particles, suggesting a possible  
53 mechanism by which the virus escapes host immune surveillance.

54

55

## 56 **Materials and Methods**

### 57 **Cells and viruses**

58 Umbilical cord blood mononuclear cells (CBMCs) were prepared as described  
59 previously (21). CBMCs were provided by K. Adachi (Minoh Hospital, Minoh, Japan)  
60 and H. Yamada (Kobe University Graduate School of Medicine, Kobe, Japan) and  
61 purchased from the Cell Bank of the RIKEN BioResource Center, Japan. Virus stocks  
62 were also prepared as described previously (22, 21). HSB-2 and MT-4 cell lines were  
63 used in this study (23). HHV-6A (GS strain) and HHV-6B (HST strain) were prepared as  
64 previously described (21).

65

### 66 **Antibodies**

67 Monoclonal antibody (Mab) OHV-1 (24) and a polyclonal antibody against gB (23, 25)  
68 have been described previously. Other Mabs were purchased: MHC class I (clone:  
69 W6/32; Bio Legend), CD63 (clone: CLB-gran/12, 435; Sanquin), and  $\alpha$ -tubulin (clone:  
70 B-5-1-2; Sigma). The following secondary antibodies were used: Alexa Fluor 488- or  
71 594-conjugated F(ab')<sub>2</sub> fragment of goat anti-mouse or rabbit immunoglobulin G (IgG)  
72 (Invitrogen) and anti-mouse IgG, horseradish peroxidase-linked whole antibody (from  
73 sheep) (GE Healthcare).

74

### 75 **Virion and exosome isolation**

76 Virions and exosomes were purified as previously described (26, 23). The collected  
77 fractions were used for Western blotting, electron microscopy, or LC-MS/MS.

78

79 **Liquid chromatography-tandem mass spectrometry**

80 The fractions described above were analyzed by LC-MS/MS. Proteins were 10-times  
81 diluted with 9.8M urea. The solutions were adjusted to pH 8.5, reduced with 13 mM  
82 DTT at 37 °C for 1.5 h and alkylated with 27 mM iodoacetamide in the dark for 1 h.  
83 The protein mixtures were further diluted with 100mM Triethylammonium bicarbonate  
84 (TEAB) pH8.5 to reduce urea to 1 M, and digested with 4 µl of 1mg/ml trypsin-TPCK  
85 solution. Samples were digested overnight at 37°C. Following digestion, lysates were  
86 acidified by adding 10% TFA. The samples were desalted using peptide cleanup C18  
87 spin tubes (Agilent Technologies) and vacuum-dried. NanoLC-MS/MS analyses were  
88 performed on an LTQ-Orbitrap XL mass spectrometer (Thermo Fisher Scientific,  
89 Waltham, MA, USA) as described previously (27), while spray voltage was changed to  
90 1800 V. Peptides and proteins were identified by automated database searches using  
91 Proteome Discoverer v.1.1 (Thermo Fisher Scientific) against all entries of the Swiss  
92 Prot protein database (version 3.26) with a precursor mass tolerance of 10 p.p.m., a  
93 fragment ion mass tolerance of 0.8 Da, and strict trypsin specificity, allowing for up to  
94 two missed cleavages. Cysteine carbamidomethylation was set as a fixed modification,  
95 and methionine oxidation was allowed as a variable modification.

96

97 **Western blotting**

98 Western blotting was performed as described previously (28, 29).

99

100 **Electron microscopy**

101 Electron microscopy was performed as described previously (30).

Briefly, the virion-containing pellet was resuspended in 2% (wt/vol) paraformaldehyde solution buffered with 0.1 M phosphate (pH 7.2). Next, 5  $\mu$ l of the resuspended pellet was loaded onto Formvar–carbon-coated grids to adsorb the virions. Immunostaining was then performed. The virions were incubated with mouse anti-gB, anti-MHC class I, or anti-CD63 antibody for 1 h at room temperature, followed by goat anti-mouse IgG conjugated to 10 nm colloidal gold particles (GE Healthcare) for a further 1 h at room temperature. After immunolabeling, the samples were washed in distilled water, stained for 5 min with uranyl oxalate, pH 7.0, washed again, embedded in a mixture of 1.8% methylcellulose and 0.4% uranyl acetate, pH 4.0, at 4°C, air-dried, and observed under a Hitachi H-7100 electron microscope. For the control experiments, samples were incubated with the secondary antibody alone.

#### **Flow cytometry**

MT-4 cells were infected with HHV-6B. At 72 h post-infection, the cells were fixed with 4% (wt/ vol) paraformaldehyde at room temperature for 15 min and incubated with the anti-MHC class I Mab at 37°C for 1 h. Cells were then stained with an appropriate secondary antibody at 37°C for 30 min. For the control experiments, samples were incubated with the secondary antibody alone. Stained cells were analyzed using a flow cytometer (ec800; SONY;).

#### **Immunofluorescence assay**

The immunofluorescence assay was performed as described previously (28). Briefly, MT-4 cells were infected with HHV-6B. At 72 h post-infection, the cells were fixed with cold acetone-methanol (7:3) and incubated at 37°C for 1 h with an anti-HHV-6 gB

126 rabbit antibody or an anti-MHC class I Mab. After washing for 10 min with PBS  
127 containing 0.02% Tween-20, the cells were incubated with an appropriate secondary  
128 antibody at 37°C for 30 min, followed by Hoechst33342 at 37°C for 40 min. After  
129 washing as described above, signals were detected by a confocal laser-scanning  
130 microscope (Olympus FluoView FV1000).

## 134 **Results**

### 135 **Virion and exosome isolation**

136 The extracellular viral particles containing exosomes were purified from the culture  
137 supernatant of HHV-6A (strain GS)-infected HSB-2 or HHV-6B (strain HST)-infected  
138 MT-4 cells. The particle-containing fractions were confirmed by Western blotting with  
139 an anti-gB antibody (23, 25). Next, the particle-containing fractions were analyzed by  
140 liquid chromatography-tandem mass spectrometry (LC-MS/MS) (27), which detected  
141 many cellular proteins (unpublished data). Of the host proteins detected, we focused our  
142 analyses on MHC class I molecules.

### 144 **Virion- or exosome-associated fractions contain MHC class I molecules**

145 To verify the expression of MHC class I within viral particles, the proteins in fractions  
146 3–10 were separated by SDS-PAGE and analyzed by Western blotting with anti-gB  
147 rabbit, anti-MHC class I, or anti-CD63 antibodies. As shown in Figure 1, gB protein  
148 was detected in fractions 5–6 and MHC class I was detected primarily in fractions 6–8.  
149 We previously reported that the multivesicular body (MVB) marker, CD63, was

incorporated into virions and exosomes (23); therefore, we also examined the expression of CD63. As expected, CD63 was detected in fractions 5–10 (Fig. 1c). To confirm the expression of MHC class I within both virions and exosomes, we performed negative staining of fractions 6 and 7 followed by electron microscopy (30). Fraction 6 contained mainly viral particles of approximately 200 nm in diameter. Both MHC class I (Fig. 1e) and gB protein (Fig. 1d) were present in these particles. Fraction 7 contained mainly exosomes of approximately 50–100 nm in diameter (Fig. 1f). These exosomes contained MHC class I, which conformed the results of the Western blotting experiments. Taken together, these results indicate that MHC class I molecules are present in exosomes and virions released from HHV-6B-infected cells.

#### **Downregulated expression of MHC class I molecules on the surface of HHV-6B-infected cells**

Downregulation of MHC class I occurs in many different virus-infected cells (31–37). Because MHC class I molecules were incorporated into virions, HHV-6-infected MT-4 cells might show an apparent downregulation in cell surface expression. To confirm this, HHV-6B- or mock-infected cells, which were harvested at 72 h post-infection, were fixed and then stained with an anti-MHC class I antibody. The surface expression of MHC class I was then analyzed by flow cytometry. As expected, HHV-6B-infected cells showed downregulated cell surface expression of MHC class I when compared with mock-infected cells (Fig. 2a). This reduced expression was confirmed by Western blot analysis (Fig. 2b), indicating that the expression of MHC class I molecules within HHV-6-infected cells (not just expression on the cell surface) was also downregulated. We next observed the localization of MHC class I molecules in these cells after they

were fixed and co-stained with anti-MHC class I and gB antibodies. MHC class I in infected cells was localized mainly within intracellular compartments, and colocalized with the envelope glycoprotein gB during the later stages of infection; however, MHC class I was mainly localized to the plasma membrane in mock-infected cells (Fig. 2c).

## **Discussion**

Here, we used mass spectrometry-based proteomics analysis to show that MHC class I molecules are incorporated into HHV-6 viral particles. The downregulation of MHC class I molecules in virus-infected cells is an important mechanism by which viruses evade immune surveillance (31-37). We showed that downregulation of MHC class I molecules occurs in T cells infected by HHV-6. MHC class I molecules are incorporated into viral particles and exosomes and then released into the extracellular environment, suggesting a possible strategy for escaping host immune responses. In addition, MHC class I molecules incorporated into virions and exosomes may assist viral entry. Further studies are needed to address this question.

We previously reported that immature HHV-6 particles bud into TGN or TGN-derived vesicles (which are produced in HHV-6B-infected cells), that vesicles containing mature virions become MVBs, and that virions and exosomes are released into the extracellular environment via an exosomal secretory pathway (23). It is possible that MHC class I molecules are transported into the TGN-derived membranes from which the virions bud and then incorporated into virions within infected cells without being recycled (Fig. 3).

The MHC class I molecules within infected cells colocalized with the gB protein in the cytoplasm, indicating that, like viral glycoproteins, they are sorted into vesicles. The

reduction in the total (both cell surface and intracellular) expression of MHC class I in HHV-6-infected cells suggests that some of them may be transported to lysosomes and degraded, as this route is the same as that used to transport particles to MVBs.

Several host proteins are expressed in the same intracellular compartments incorporating viral particles, although they are usually expressed on the surface of uninfected cells. Newly formed compartments within HHV-6-infected cells may show the combined characteristics of early and late endosomes. Recycling to early endosomes in HHV-6-infected cells may be modified or defective; therefore, several cellular proteins that use the same recycling system may be incorporated into virions and exosomes.

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217     **Competing interests**

218     The authors have declared that no competing interests exist.

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222     **References**

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## Figure Legends

**Figure 1. MHC class I molecules are incorporated into virions and exosomes and released from HHV-6B-infected cells.** Virions and exosomes were collected from the culture medium of HHV-6B-infected cells by sucrose density gradient centrifugation and examined by Western blotting (a, b, and c) and electron microscopy (d, e, and f). Western blots with anti-gB rabbit (a), anti-MHC class I (W6/32) (b), or anti-CD63 (CLB-gran/12, 435) (c) antibodies are shown. The same amount of each protein fraction was added to each well of the gel. Immunogold labeling of gB (d) in Fraction 6 and of MHC class I in Fractions 6 and 7 (e and f). The Fraction number is shown at the top of each panel. The fractions were collected from the bottom of tube. Labeled virions (empty arrowheads) and exosomes (arrowheads). Scale bars: 200 nm (d-f).