

Innate Immune Response Induced by Baculovirus Attenuates Transgene Expression in Mammalian Cells

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The baculovirus *Autographa californica* nucleopolyhedrovirus (AcNPV) has been widely used to achieve a high level of foreign gene expression in insect cells, as well as for efficient gene transduction into mammalian cells without any replication. In addition to permitting efficient gene delivery, baculovirus has been shown to induce host innate immune responses in various mammalian cells and in mice. In this study, we examined the effects of the innate immune responses on gene expression by recombinant baculoviruses in cultured cells. The reporter gene expression in IRF3-deficient mouse embryonic fibroblasts (MEFs) infected with the recombinant baculovirus was shown to be enhanced in accordance with the suppression of beta interferon (IFN- β) production. Furthermore, efficient gene transduction by the recombinant baculovirus was achieved in MEFs deficient for stimulator of interferon genes (STING), TANK binding kinase 1 (TBK1), IFN regulatory factor 3 (IRF3), or IFN- β promoter stimulator 1 (IPS-1), but not in those deficient for IRF7, MyD88, or Z-DNA binding protein 1 (ZBP1)/DAI. Enhancement of gene expression by the recombinant baculovirus was also observed in human hepatoma cell lines replicating hepatitis C virus (HCV), in which innate immunity was impaired by the cleavage of IPS-1 by the viral protease. In addition, infection with the recombinant baculovirus expressing the BH3-only protein, BIM_S, a potent inducer of apoptosis, resulted in a selective cell death in the HCV replicon cells. These results indicate that innate immune responses induced by infection with baculovirus attenuate transgene expression, and this characteristic might be useful for a selective gene transduction into cells with impaired innate immune nity arising from infection with various viruses.

he baculovirus Autographa californica nucleopolyhedrovirus (AcNPV) is an enveloped, double-stranded-DNA (dsDNA) virus that is primarily pathogenic to insects. Although AcNPV has long been used as an efficient gene expression vector in insect cells (1, 2), recombinant baculoviruses (rBVs) have also been shown to be capable of entering into various mammalian cells without any replication and of expressing foreign genes under the control of mammalian promoters (3-7). Therefore, baculovirus is now recognized as a useful viral vector not only for abundant gene expression in insect cells but also for gene delivery into mammalian cells. Recent studies suggest that dynamin- and clathrin-dependent endocytosis, macropinocytosis, and cholesterol in the plasma membrane play crucial roles in the internalization of baculovirus; however, the mechanisms of entry of baculovirus into insect and mammalian cells have not been well characterized yet. In contrast to the efficient transgene expression in vitro by the recombinant baculoviruses, the in vivo gene delivery is still unsatisfactory. Several obstacles, such as serum complement and the acute inflammatory response induced by inoculation of baculovirus, might be implicated in the inactivation of viral particles (8-12). In vivo foreign gene expression has been achieved by introducing the baculovirus vectors into rabbit endothelial cells lining the artery through collar-mediated delivery (13), mouse skeletal muscle cells in the quadriceps by intramuscular injection (14), neural or choroid plexus cells in the rodent brain by intracranial injection (15, 16), mouse retinal pigment epithelial cells following subretinal injection (17), and the cerebral cortex and testis of mice by direct inoculation (4). The level of foreign gene expression has been no more than satisfactory, mainly due to the brief duration of transgene expression in vivo following local administration. In addition

to achieving gene delivery of recombinant baculovirus vectors, the wild-type (WT) baculovirus has also been shown to stimulate host antiviral immune responses in mammalian cell lines (8–10, 18–23). These observations suggest possible disadvantages to use of the baculovirus for sustained transduction of gene expression *in vivo*. Moreover, AcNPV was shown to possess a strong adjuvant activity to promote humoral and cellular immune responses against coadministered antigens, maturation of dendritic cells (DCs), and production of proinflammatory cytokines, chemokines, and type I interferons (IFNs) (24). We have previously shown that baculovirus induced the expression of proinflammatory cytokines and type I IFNs through the Toll-like receptor (TLR)-dependent and -independent pathway in mouse embryonic fibroblasts (MEFs), respectively (19). However, the precise mechanisms of the innate immune response and their influence

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Induction of type I IFN by pathogens is crucial for innate immunity, and it is mediated by the activation of pattern recognition receptors (PRRs), such as TLRs and retinoic-acid-inducible protein I (RIG-I)-like receptors (RLRs), including RIG-I and melanoma differentiation-associated gene 5 (MDA5) (25). The type I IFN induction is primarily controlled at the gene transcriptional level, and a family of transcription factors, IFN regulatory factors (IRFs), plays a pivotal role in this regulation (26). IRF3 and IRF7 are now known to be essential for the RIG-I, MDA5, and TLRmediated type I IFN production pathway. IRF3 is induced primarily by a response to initiate beta interferon (IFN- β) production, whereas IRF7 is induced by IFN- β and participates in the late phase of IFN-α induction (26). All TLRs, except for TLR3, activate the MyD88-dependent pathway, whereas TLR3 and TLR4 activate the TRIF-dependent pathway. A novel TLR-independent cytosolic surveillance system for transfected dsDNA that elicits type I IFN induction through the TANK binding kinase 1 (TBK1)/IKB kinase-related kinase (IKKi)-IRF3 pathway has been discovered (27, 28). Recently, a cytoplasmic recognition receptor, ZBP (Z-DNA binding protein 1), was shown to be activated by dsDNA from a variety of sources and to produce type I IFN through an IRF3 pathway and probably through an IRF7 pathway (29). More recently, it has been shown that an endoplasmic reticulum (ER)associated multiple transmembrane protein, termed "stimulator of interferon genes" (STING; also referred to as TMEM173, MPYS, MITA, and ERIS) is an essential innate immune signaling adaptor involved in the cytosolic sensing of dsDNA ligands in the TBK1/IRF3 axis (30–33).

In this study, we examined the mechanisms of induction of the innate immune response and their influence on the transgene expression in MEFs upon inoculation with the recombinant baculovirus. The efficiency of transgene expression by the recombinant baculovirus was enhanced in IRF3^{-/-} MEFs compared to the wild-type, IRF7^{-/-}, or MyD88^{-/-} MEFs. These results are consistent with the abrogation of IFN-B production via the STING/ TBK1/IRF3 pathway. Importantly, IFN-β promoter stimulator 1 (IPS-1), which is an adaptor molecule of the RLR signaling pathway, was also suggested to be involved in the efficient gene transduction by baculovirus. This was consistent with the enhancement of transgene expression by baculovirus in human liver cell lines replicating hepatitis C virus (HCV), in which innate immunity was impaired by the cleavage of IPS-1 by the viral protease. Finally, infection with a recombinant baculovirus expressing BIM_s, a potent inducer of apoptosis, selectively induced cell death in cells replicating HCV. Collectively, these results suggest that an innate immune signaling pathway attenuates the transgene expression in mammalian cells by recombinant baculovirus.

MATERIALS AND METHODS

Mice and cells. All animal experiments were conducted in accordance with the Animal Care and Use Committee of the Research Institute for Microbial Diseases, Osaka University, Japan. C57BL/6 mice were purchased from CLEA Japan (Tokyo, Japan). MyD88-, IRF3-, IRF7-, or IPS-1-deficient mice were described previously (19). TBK1^{+/-} IKK $\beta^{+/-}$ (WT) and TBK1^{-/-} IKK $\beta^{+/-}$ (TBK1^{-/-}) MEFs or ZBP1-deficient MEFs were derived from mice that were established as described previously (34, 35). STING-deficient MEFs were provided by G. Barber. MEFs from the deficient mice were obtained from day-12.5 to -13.5 embryos and maintained in Dulbecco's modified Eagle's medium (DMEM) (Sigma, St.

Louis, MO) supplemented with 10% heat-inactivated fetal calf serum (FCS), 1.5 mM L-glutamine, 100 U/ml penicillin, and 100 μ g/ml streptomycin. In some cases, MEFs were immortalized by a lentivirus carrying the simian virus 40 (SV40) large T antigen. Huh7OK1 cells, which exhibit high susceptibility to propagation of cell culture-adapted HCV of the genotype 2a JFH1 strain (HCVcc) (36), Huh7 cells, and HCV subgenomic RNA replicon (SGR)-cured cells were maintained in DMEM supplemented with 10% FCS. Huh7 cells harboring SGR of genotype 1b (Con1 strain) (37), 1a (RMT strain) (provided by M. Kohara), and 2a (JFH1 strain) were cultured in DMEM supplemented with 10% FCS and 1 mg/ml G418 (Nacalai Tesque, Kyoto, Japan). The cells were cultured at 37°C in a humidified atmosphere with 5% CO₂.

Plasmids, reagents, and viruses. The cDNA fragments encoding murine IRF3 and IPS-1 were amplified by reverse transcription (RT)-PCR from total RNA of MEFs and cloned into a retroviral vector, pMIH (provided by D. Huang), and the resulting plasmids were designated pMIH FLAG-IRF3 and pMIH FLAG-IPS-1, respectively. Retroviral vectors for expressing short hairpin RNA (shRNA) targeting human IRF3, IPS-1, or STING were purchased from TaKaRa Bio (Shiga, Japan). The pFG EF BIM_s2A GFP and pFG EF BIM_s4E 2A GFP constructs were generated by PCR from pEF BIM_s2A GFP and pEF BIM_s4E 2A GFP, respectively, and cloned into pFastBac VSVG (Invitrogen, Tokyo, Japan) for the construction of recombinant baculoviruses by using an In-Fusion cloning kit (Ta-KaRa Bio). All PCR products were confirmed by sequencing by using an ABI PRISM 3100 genetic analyzer (Applied Biosystems, Tokyo, Japan). The generation of recombinant vesicular stomatitis virus (VSV) possessing a luciferase gene, VSV-luc, was described previously (38). The recombinant baculoviruses were generated by a Bac-to-Bac system (Invitrogen) following the manufacturer's protocol. For efficient gene transduction to MEFs, we utilized the baculovirus possessing the cDNA of VSV envelope G protein (VSVG) under the control of the polyhedrin promoter and bearing VSVG on the virion surface (3, 4). Recombinant baculoviruses carrying cDNAs of luciferase and green fluorescence protein (GFP) under the CAG promoter consist of chicken β -actin promoter (39), and those carrying cDNAs of BIM_S and BIM_S4E under the human elongation factor (EF) promoter (40) were designated rBV-luc, rBV-GFP, rBV-BIM_s, and rBV-BIM_s4E, respectively. These recombinant baculoviruses and wildtype baculovirus (AcNPV C6 strain) were propagated in Spodoptera frugiperda (Sf-9) cells in Sf-900II insect medium (Invitrogen) supplemented with 10% FCS and purified as previously described (18). The infectious titers of recombinant baculovirus are expressed in PFU (41). The retroviruses expressing FLAG-IRF3, FLAG-IPS-1, or shRNAs against human IRF3, IPS-1, or STING were generated by PlatE cells as described previously (42). The culture supernatants containing retroviruses were passed through a 0.45-µm-pore filter, inoculated into MEFs, and incubated with hygromycin (50 µg/ml), and then stable cell lines were established. The VSV variants GLPLF and NCP12.1, derived from Indiana strains, were provided by M. Whitt. Human recombinant IFN-α was purchased from PBL Biomedical Laboratories (New Brunswick, NJ).

Antibodies. Antibodies to GFP (polyclonal), FLAG tag (M2), mouse IPS-1 (no. 3013), TBK1/NAK1 (no. 4983), HCV NS5A (HCM-131-5), calnexin (sc-70481), and β -actin (A8481) were purchased from Clontech (Mountain View, CA), Sigma, Cell Signaling (Tokyo, Japan), Austral Biologicals (San Ramon, CA), and Santa Cruz Biotechnology (Santa Cruz, CA), respectively. The BIM antibody (rat monoclonal antibody 3C5) was a gift from A. Strasser (43).

Immunoblotting. Cells were washed three times with ice-cold phosphate-buffered saline (PBS) and suspended in 0.4 ml lysis buffer containing 20 mM Tris-HCl (pH 7.4), 135 mM NaCl, 1% Triton X-100, 1% glycerol, and protease inhibitor cocktail tablets (Roche Molecular Biochemicals, Mannheim, Germany). Cell lysates were incubated for 30 min at 4°C and centrifuged at 14,000 × g for 15 min at 4°C. The proteins were boiled in 20 μ l of sample buffer, subjected to sodium dodecyl sulfate-12.5% polyacrylamide gel electrophoresis (SDS-PAGE), and transferred to polyvinylidene difluoride membranes (Millipore, Tokyo, Japan). The

membranes were then blocked with Tris-buffered saline containing 20 mM Tris-HCl (pH 7.4), 135 mM NaCl, 0.05% Tween 20, and 5% skim milk and incubated with primary antibody at room temperature for 1 h and then with horseradish peroxidase (HRP)-conjugated secondary antibody at room temperature for 1 h. The immune complexes and cell lysates were visualized with Super Signal West Femto substrate (Pierce, Rockford, IL) and detected by an LAS-3000 image analyzer system (Fujifilm, Tokyo, Japan).

Immunofluorescence assay. MEFs were infected with recombinant baculovirus expressing luciferase (rBV-luc) at a multiplicity of infection (MOI) of 100 and fixed with methanol-acetone (1:1) for 10 min at 6 h postinfection after being washed twice with PBS. Cells were incubated for 1 h at 4°C with 1 μ g/ml of primary antibody in PBS containing 10% FCS (PBSF) and then incubated at room temperature for 1 h with 1 μ g/ml of Alexa Fluor-conjugated secondary antibody (Invitrogen) after three washes with PBSF. Cell nuclei were stained with 4',6-diamidino-2-phenylindole (DAPI). After an extensive wash with PBSF, the samples were examined with a FluoView FV1000 laser scanning confocal microscope (Olympus, Tokyo, Japan).

Reporter assays. Cells were inoculated with rBV-luc or VSV-luc, and the relative light units (RLU) of luciferase were determined by using the Promega luciferase assay system (Promega, Madison, WI) at 24 h postinfection according to a protocol provided by the manufacturer. The expression of GFP in various cells inoculated with rBV-GFP was examined by fluorescence microscopy (Olympus).

Production of cytokines. MEFs prepared from each of the deficient mouse strains were seeded into 96-well plates at a concentration of 2×10^4 cells/well and infected with rBV-GFP at an MOI of 100, and then the production of IFN-β in the culture supernatants was determined at 24 h postinfection by enzyme-linked immunosorbent assay (ELISA) kits purchased from PBL Biomedical Laboratories (Piscataway, NJ).

Real-time PCR. Baculoviral genomes attached on the cell surface were quantified by SYBR green-based quantitative PCR (qPCR) using a virusspecific primer as described previously (44). In brief, MEFs derived from WT and IRF3^{-/-} mice were inoculated with rBV-GFP at an MOI of 100, incubated at 4°C for 30 min, and washed three times with PBS. The total DNA was prepared from cells by using QIAamp DNA minikit (Qiagen, Valencia, CA), and the amounts of baculovirus genome in the total DNA (20 ng) were determined by using Platinum SYBR green qPCR SuperMix UDG (Invitrogen) using GP64-specific primers and normalized with GAPDH (glyceraldehyde-3-phosphate dehydrogenase) according to the manufacturer's protocol. Fluorescent signals were analyzed by an ABI PRISM 7000 (Applied Biosystems). To determine the gene expression, total RNA was prepared from cells by using an RNeasy minikit (Qiagen, Valencia, CA), and the first-strand cDNA was synthesized by using a ReverTra Ace (Toyobo, Osaka, Japan) and oligo(dT)₂₀ primer. The mouse IFN-β, IFN-γ-induced protein 10 (IP-10), GFP, and GAPDH genes were amplified using the primer pairs 5'-ACACCAGCCTGGCTTCCATC-3' and 5'-TTGGAGCTGGAGCTGCTTATAGTTG-3', 5'-TGAATCCGGA ATCTAAGACCATCAA-3' and 5'-AGGACTAGCCATCCACTGGGTAA AG-3', 5'-ACACCAGCCTGGCTTCCATC-3' and 5'-TTGGAGCTGGA GCTGCTTATAGTTG-3', and 5'-ACCACAGTCCATGCCATCAC-3' and 5'-TCCACCACCCTGTTGCTGTA-3', respectively. The human IP-10, ISG15, and interleukin-8 (IL-8) genes were amplified using the primer pairs 5'-GGCCATCAAGAATTTACTGAAAGCA-3' and 5'-TCTGTGTG GTCCATCCTTGGAA-3', 5'-AGCGAACTCATCTTTGCCAGTACA-3' and 5'-CAGCTCTGACACCGACATGGA-3', and 5'-ACACTGCGCCAA CACAGAAATTA-3' and 5'-TTTGCTTGAAGTTTCACTGGCATC-3', respectively. The amount of each cDNA was determined by using Platinum SYBR green qPCR SuperMix UDG (Invitrogen). Fluorescent signals were analyzed by an ABI PRISM 7000 (Applied Biosystems). The human IRF-3, IPS-1, and STING genes were amplified by using the TaqMan probes (IRF3, Hs01547283_m1; IPS-1, Hs00920075_m1; STING, Hs00736958). The amount of each cDNA was determined by using Taq-Man Fast advanced master mix (Applied Biosystems). Fluorescent signals

were analyzed by a ViiA7 PCR system (Applied Biosystems). The expression of mRNA of each of the cytokines was normalized with that of GAPDH mRNA.

Cell viability. Cells in the culture supernatants and adherent cells detached from the plates by trypsin were collected by centrifugation at 1,500 rpm for 5 min. Cell pellets were resuspended in 0.25 ml of PBS containing 2% FCS and 100 μ g/ml of propidium iodide (PI), and cell viability was determined by flow cytometry using FlowJo software (Tree Star, Ashland, OR).

Statistical analysis. Results are expressed as means \pm standard deviations (SD). The significance of differences in the means was determined by Student's *t* test.

RESULTS

IRF3-dependent innate immune response suppresses transgene expression by recombinant baculovirus in MEFs. We have previously shown that baculovirus induced proinflammatory cytokines and type I IFNs in MEFs through the TLR-dependent and TLR-independent pathways, respectively (19). Although the recombinant baculovirus was shown to be capable of internalizing into mammalian cells without any replication, the precise mechanisms of induction of an innate immune response that induces suppression of the transgene expression in MEFs remain unclear. To elucidate the involvement of innate immunity in the influence of gene transduction by recombinant baculovirus, MEFs derived from wild-type and IRF3-, IRF7-, or MyD88-deficient mice were inoculated with rBV-GFP possessing the GFP genes under the control of the CAG promoter, and the transgene expression was evaluated by microscopic observation. Although GFP expression in the MEFs derived from wild-type, MyD88-, and IRF7-deficient mice was low, IRF3-deficient cells exhibited efficient GFP expression upon infection with rBV-GFP (Fig. 1A). The finding that the transgene expression was not enhanced by rBV-GFP in MyD88deficient MEFs suggests that TLR-mediated innate immunityexcept for that involving the TLR3-dependent pathway-was not involved in the suppression of transgene expression by recombinant baculovirus in MEFs. In addition, there was no difference in the amounts of viral particles attached to the cell surface between wild-type and IRF3-deficient MEFs (Fig. 1B). Immunoblotting and quantitative RT-PCR analyses also confirmed the enhancement of gene transduction in IRF3-deficient MEFs upon infection with rBV-GFP (Fig. 1C, left panel, and D). Furthermore, the production of IFN-B and IP-10 in the IRF3-deficient MEFs decreased significantly compared with that in the wild-type cells (Fig. 1C, middle and right panels), suggesting that the enhancement of GFP expression in IRF3-deficient MEFs was due to the suppression of innate immune responses. To further confirm that the transgene expression was enhanced by the recombinant baculovirus through the suppression of IRF-3, FLAG-IRF3 was exogenously expressed in IRF3-deficient MEFs by a retroviral vector (Fig. 2A). trans-complementation of FLAG-IRF3 in the IRF3-deficient MEFs induced suppression of the luciferase expression by rBV-luc (Fig. 2B). Furthermore, confocal microscopy observation revealed that exogenous expression of FLAG-IRF3 in the IRF3-deficient MEFs was translocated into the nucleus upon infection with rBV-luc (Fig. 2C), suggesting that the FLAG-IRF3 exogenously expressed in the IRF3-deficient MEFs was functional. Collectively, these results suggest that the innate immune response through the IRF3-dependent pathway attenuates the transgene expression by recombinant baculovirus in MEFs.

Type I IFN production through the STING/TBK1/IRF3 axis is

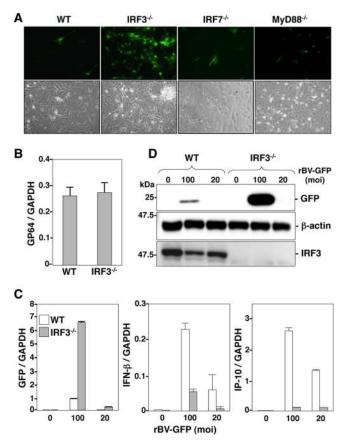


FIG 1 Efficient gene transduction by recombinant baculovirus in the IRF3^{-/-} MEFs. (A) MEFs derived from wild-type (WT) and IRF3^{-/-}, IRF7^{-/} , or MyD88^{-/-} mice were inoculated with rBV-GFP (MOI of 100). At 24 h after inoculation, GFP signal was determined by microscopic observation after fixation in 4% paraformaldehyde. (B) MEFs derived from WT and IRF3^{-/} mice were inoculated with rBV-GFP at an MOI of 100, incubated at 4°C for 30 min, and washed three times with PBS. The total cellular DNA was extracted, and the amounts of baculovirus genome were quantified by real-time PCR. Data represent means \pm SD from 2 independent experiments. (C) MEFs derived from WT and IRF3^{-/-} mice were inoculated with 2 doses of rBV-GFP (MOI of 100 or 20). At 24 h after inoculation, total RNA was extracted, and the expression of GFP, IFN-β, and IP-10 mRNAs was determined by real-time PCR. (D) MEFs derived from WT and IRF3-deficient mice were inoculated with 2 doses of rBV-GFP (MOI of 100 and 20). At 24 h after inoculation, cell extracts were subjected to SDS-PAGE and immunoblotted with antibodies against GFP, IRF3, or β -actin, respectively.

involved in the suppression of transgene expression by recombinant baculovirus in MEFs. A TLR-independent cytosolic surveillance system against exogenous dsDNA that induces type I IFN production through a TANK binding kinase 1 (TBK1)/I κ B kinase-related kinase (Ikki)/IRF3 pathway has been identified (27, 28). To determine the involvement of the TLR-independent cytosolic sensor to detect exogenous dsDNA on the transgene expression by recombinant baculovirus, we examined the involvement of TBK1 in the induction of the innate immune response in MEFs upon infection with rBV-GFP. Immunofluorescence, immunoblotting, and quantitative RT-PCR analyses revealed that TBK1deficient MEFs exhibited enhancement of the transgene expression by rBV-GFP, as seen in the IRF3-deficient MEFs (Fig. 3A, B, and C, left panel). In addition, significant reductions in IFN- β and IP-10 production were observed in the TBK1-deficient MEFs

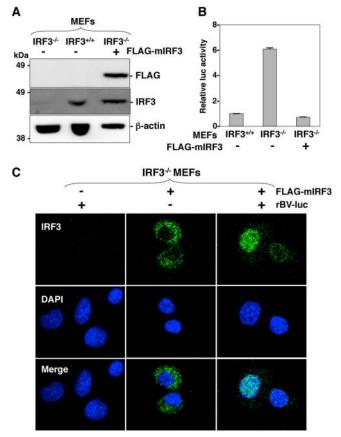


FIG 2 IRF3-dependent enhancement of gene transduction by recombinant baculovirus. (A) Cell extracts of MEFs derived from WT or IRF3^{-/-} mice and FLAG-mIRF3-transduced IRF3^{-/-} MEFs were subjected to SDS-PAGE and immunoblotted with antibodies against FLAG, IRF3, or β -actin, respectively. (B) MEFs derived from WT or IRF3^{-/-} mice and FLAG-mIRF3-transduced IRF3-deficient MEFs were inoculated with rBV-luc at an MOI of 100. At 24 h after inoculation, the luciferase activity was determined. Data represent the means \pm SD from 3 independent experiments. (C) MEFs were inoculated with rBV-luc at an MOI of 100. At 6 h after inoculation, cells were fixed in 50% methanol–50% acetone for 10 min. IRF3 (green) was stained with the appropriate antibodies, followed by staining with Alexa Fluor 488-conjugated secondary antibodies. Nuclei were stained by DAPI.

compared with the levels in the wild-type cells (Fig. 3C, middle and right panels), suggesting that the enhancement of the transgene expression in the TBK1-deficient MEFs was due to suppression of the innate immune response. Recently, the ER-associated membrane protein STING was identified as a candidate DNA sensor for production of type I IFN in response to virus infection (30-33, 45). On the other hand, the Z-DNA binding protein 1 (ZBP1; also referred to as DAI/DLM-1) has also been reported as a candidate DNA sensor (29), while the involvement of these DNA sensors in the type I IFN production is still controversial (35). We have previously shown that the induction of an innate immune response upon infection with baculovirus was dependent on viral DNA (19). To examine the roles of STING and ZBP1 in the induction of innate immune responses that lead to the suppression of transgene expression by the recombinant baculovirus, STINGor ZBP1-deficient MEFs were inoculated with either rBV-GFP or rBV-luc. As shown in Fig. 4A and B, STING-deficient MEFs exhibited the enhancement of GFP and luciferase expression upon

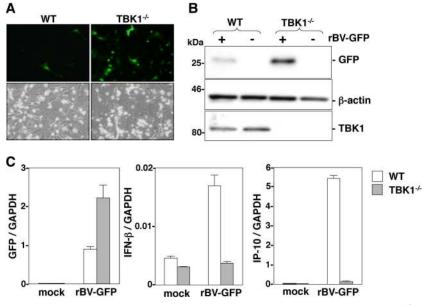


FIG 3 Involvement of TBK1 in efficient transgene expression by recombinant baculovirus. (A) MEFs derived from TBK1^{+/-} IKK $\beta^{+/-}$ (WT) and TBK1^{-/-} IKK $\beta^{+/-}$ (WT) ince were inoculated with rBV-GFP (MOI of 100). At 24 h after inoculation, the GFP signal was determined by microscopic observation after fixation in 4% paraformaldehyde. (B) MEFs derived from WT and TBK1^{-/-} mice were inoculated with rBV-GFP at an MOI of 100. At 24 h after inoculation, cell extracts were subjected to SDS-PAGE and immunoblotted with antibodies against GFP, TBK1, or β -actin, respectively. (C) MEFs derived from WT and TBK1-deficient mice were inoculated with rBV-GFP (MOI of 100). At 24 h after inoculation, total RNA was extracted, and the expression of GFP, IFN- β , and IP-10 mRNA was determined by real-time PCR. Data from the real-time PCR were normalized to the amount of GAPDH mRNA.

inoculation with rBV-GFP and rBV-luc, respectively, while there was no such enhancement in ZBP1-deficient MEFs. STING-deficient MEFs, but not ZBP1-deficient MEFs, exhibited a significant reduction of IFN-B production upon infection with rBV-GFP (Fig. 4C). It has previously been shown that STING is dynamically translocated from the ER to the perinuclear region in cells stimulated with exogenous DNA ligands or infected with DNA virus (30, 46). We confirmed that endogenous STING was accumulated in the perinuclear region upon infection with AcNPV in MEFs (Fig. 4D), suggesting that baculovirus DNA participates in the induction of innate immune response through the translocation of STING to the perinuclear region. These results suggest that recombinant baculovirus induces type I IFN production through the STING/TBK1/IRF3 axis in MEFs, and those signaling pathways may be involved in the suppression of transgene expression by recombinant baculoviruses.

RLR signaling pathways participate in the suppression of transgene expression in MEFs by recombinant baculovirus. RIG-I and MDA5 have been identified as TLR-independent cytoplasmic RNA detectors and have been shown to induce type I IFN production through an adaptor molecule, IPS-1 (also referred as MAVS/VISA/Cardif), that localizes in mitochondria (25). We have previously shown that AcNPV produced substantial amounts of the inflammatory cytokines, chemokines, and type I IFNs in IPS-1-deficient MEFs in spite of a slight reduction in IFN- β and IL-6 (19). Recently, it was shown that the DNA virus is also capable of stimulating the RLR signaling pathways to induce type I IFNs through the participation of an adaptor molecule, IPS-1 (47, 48). To determine the involvement of RLR signaling pathways in the transgene expression by recombinant baculovirus, IRF3-, TBK1-, IPS-1-, or ZBP1-deficient MEFs were inoculated with rBV-luc. Interestingly, not only MEFs deficient in IRF3

or TBK1 but also those deficient in IPS-1 showed significant enhancement of luciferase expression (Fig. 5A). In addition, IPS-1deficient MEFs exhibited enhancement of GFP expression and a significant reduction of IFN-B and IP-10 production compared with those in wild-type cells upon infection with rBV-GFP, as seen in IRF3-deficient MEFs (Fig. 5B and C). These results suggest that the RLR signaling pathway also participates in the suppression of the gene transduction by recombinant baculovirus in MEFs. To confirm the enhancement of the transgene expression by the recombinant baculovirus through the suppression of IPS-1, FLAG-IPS-1 was exogenously expressed in IPS-1-deficient MEFs by a retroviral vector (Fig. 5D). trans-complementation of FLAG-IPS-1 in the IPS-1-deficient MEFs induced suppression of luciferase expression by rBV-luc (Fig. 5E). These results suggest that an IPS-1-dependent RLR-signaling pathway also participates in the suppression of the transgene expression by recombinant baculovirus in MEFs.

RLR signaling pathways also participate in the suppression of transgene expression in human hepatoma cells by recombinant baculovirus. The NS3/4A protease of HCV participates not only in the processing of the precursor polyprotein of HCV but also in the circumvention of the induction of the host innate immune response through the cleavage of adaptor molecules, such as TRIF and IPS-1, in the human hepatoma cell lines (49). Therefore, we hypothesized that transgene expression by recombinant baculovirus might be enhanced in the HCV replicon cells, in which the innate immune response is impaired by the cleavage of IPS-1 by HCV NS3/4A protease. Reporter gene expression by infection with rBV-luc and VSV-luc was higher in the replicon cells harboring the HCV genome derived from genotypes 1a, 1b, or 2a than in those in which the viral RNA (genotype 1b) was eliminated by the treatment with IFN- α (cured) (Fig. 6A). The reporter gene expres-

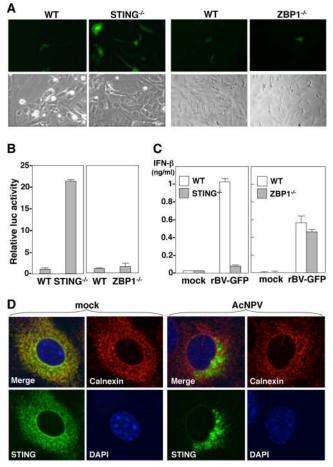


FIG 4 Efficient gene transduction by recombinant baculovirus vectors through the STING/TBK1/IRF3 axis in MEFs. (A) MEFs derived from wild-type (WT) and STING- or ZBP1-deficient mice were inoculated with rBV-GFP (MOI of 100). At 24 h after inoculation, the GFP signal was determined by microscopic observation after fixation in 4% paraformaldehyde. (B and C) MEFs derived from WT and STING- or ZBP1-deficient mice were inoculated with rBV-luc (MOI of 100) or rBV-GFP (MOI of 100). At 24 h after inoculation, the luciferase activity of cell lysates was determined by sandwich ELISA (C). (D) The MEFs were inoculated with wild-type baculovirus (AcNPV) (MOI of 100). At 6 h after inoculation, cells were fixed in 50% methanol–50% acetone for 10 min. STING (green) and ER-calnexin (red) were stained with the appropriate antibodies, followed by staining with Alexa Fluor 488- or Alexa Fluor 555-conjugated second antibodies, respectively. Nuclei were stained by DAPI.

sion was increased in a dose-dependent manner upon infection with rBV-luc of the HCV replicon cells of genotype 1b (Fig. 6B). Although levels of production of IL-8 in cells infected with either rBV-luc or VSV-GFP were comparable, no induction of IP-10 and ISG15 was observed in cells infected with rBV-luc, in contrast to those infected with VSV-GFP (Fig. 6C). These data suggest that IP-10 and ISG15 are not involved in efficient gene transduction by rBV-Luc. Furthermore, the enhancement of reporter gene expression in cells infected with HCVcc was canceled by the elimination of viral replication by the treatment with IFN- α (Fig. 6D), suggesting that an IPS-1-dependent RLR-signaling pathway is involved in suppression of the transgene expression by recombinant baculovirus in human hepatoma cells, as seen in MEFs. To further confirm that IPS-1 is implicated in the suppression of the transgene

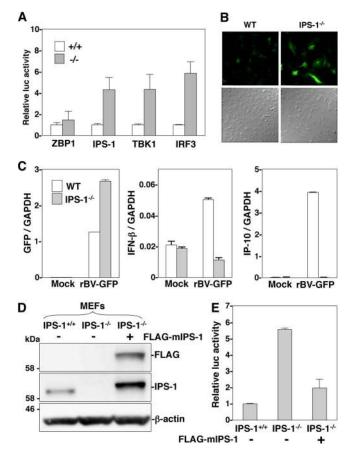


FIG 5 RLR signaling pathways participate in the suppression of gene transduction in MEFs upon infection with recombinant baculovirus. (A) MEFs derived from wild-type (WT) and IPS-1-, TBK1-, ZBP1-, or IRF3-deficient mice were inoculated with rBV-luc at an MOI of 100. At 24 h after inoculation, the luciferase activity was determined. Data represent means \pm SD from 3 independent experiments. (B) MEFs derived from WT and IPS-1^{-/-} mice were inoculated with rBV-GFP at an MOI of 100. At 24 h after inoculation, GFP expression was detected by microscopic observation after fixation in 4% paraformaldehyde. (C) MEFs derived from WT or IPS-1-deficient mice were inoculated with rBV-GFP at an MOI of 100. At 24 h after inoculation, total RNA was extracted, and the expression of GFP, IFN-β, and IP-10 mRNAs was determined by real-time PCR. Data from the real-time PCR were normalized to the amount of GAPDH mRNA. (D) Cell extracts of MEFs derived from WT or IPS-1^{-/-} mice and FLAG-mIPS-1-transduced IPS-1^{-/-} MEFs were subjected to SDS-PAGE and immunoblotted with antibodies against FLAG, IPS-1, or $\beta\text{-actin},$ respectively. (E) MEFs derived from WT or $\text{IPS-1}^{-\prime-}$ mice and FLAG-mIPS-1-transduced IPS-1-deficient MEFs were inoculated with rBVluc at an MOI of 100. At 24 h after inoculation, the luciferase activity was determined. Data represent means \pm SD from 3 independent experiments.

expression by recombinant baculovirus, we established IPS-1, IRF3, or STING knockdown Huh7 cell lines by using shRNA (Fig. 6E, left). The highest enhancement of luciferase expression by rBV-luc was achieved in Huh7 cells with knockdown of STING, followed by those with knockdown of IPS-1 and IRF3 (Fig. 6E, right), suggesting that IPS-1 also participates in the suppression of transgene expression by recombinant baculovirus in human hepatoma cells in a manner similar to its participation in MEFs.

Induction of apoptosis in cells replicating HCV RNA by a recombinant baculovirus expressing a proapoptotic protein. The results in this study suggest that recombinant baculovirus is capable of achieving an enhanced transgene expression in cells

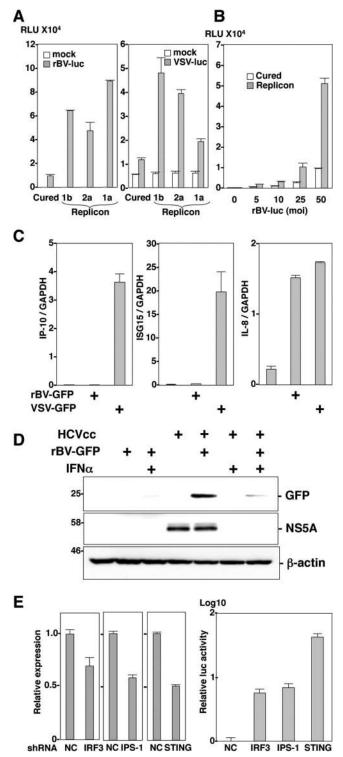


FIG 6 Efficient gene transduction by recombinant baculovirus in HCV replicon-harboring cells. (A) Huh7OK1 cells (cured) and the HCV replicon-harboring cells derived from genotype 1a (RMT strain), 1b (con1 strain), and 2a (JFH1 strain) were inoculated with rBV-luc (MOI of 100) or VSV-luc at an MOI of 5 and (B) with rBV-luc at MOI of 5, 10, 25, and 50. At 24 h after inoculation, the luciferase activity of cell lysates was determined. (C) Huh7 cells were inoculated with rBV-GFP (MOI of 100) or VSV-GFP (NCP mutant) at an MOI of 0.05. At 24 h after inoculation, total RNA was extracted, and the expression of IP-10, ISG15, and IL-8 mRNAs was determined by real-time PCR. Data from the real-time PCR were normalized to the amount of GAPDH

with an impaired host innate immune response. Therefore, we thought that it might be feasible to eliminate cells replicating the HCV genome by infection with a recombinant baculovirus encoding one of the proapoptotic Bcl-2 family proteins, BIM_s, which induces apoptosis through activation of Bax and Bak. We generated two recombinant baculoviruses, rBV-BIM_S and rBV-BIM_s4E, carrying cDNAs of either BIM_s or BIM_s4E (an inactive mutant of BIM₂), 2A peptides of *Thosea asigna* virus, to express multiple proteins as a polyprotein that dissociates into each protein upon translation (50), and enhanced GFP under the control of the EF promoter (Fig. 7A), and exogenous expression of both BIM_s and BIM_s4E was detected in the HCV replicon cells infected with the recombinants in the presence of an apoptosis inhibitor (Fig. 7B). HCV replicon cells infected with $rBV-BIM_s$, but not with rBV-BIM_s4E, exhibited an efficient cell death (Fig. 7C), and expression of rBV-BIMs induces cell death at 24 h postinfection in the replicon cells but not in the cured cells (Fig. 7D). Furthermore, cell death induced in HCV replicon cells by the infection with rBV-BIM_s was inhibited by the treatment with a caspase inhibitor, qVD-Oph (Fig. 7E), suggesting that infection with rBV-BIM_s selectively induced apoptosis in cells replicating HCV RNA through the activation of Bcl-2 family proteins. These results suggest that infection with rBV-BIM_s might eliminate HCV through the induction of cell death by the enhanced expression of the Bcl-2 family protein in HCV-infected cells in which innate immunity is impaired by the NS3/4A protease.

DISCUSSION

Viral vectors have great advantages for efficient and sustained gene delivery to target cells, but the induction of immune responses against viral components and products, including viral proteins and nucleic acids, is a great concern (51). Production of proinflammatory cytokines and type I IFNs upon delivery of the viral vectors may abrogate transgene expression and induce cytotoxicity to the target cells and tissues (52–54). In fact, it has been shown that the removal of viral elements responsible for replication and pathogenicity, together with the provision of a large-capacity space to accommodate foreign genes and broader cell tropism by bearing envelope proteins that have high affinity to various tissues, makes the viral vectors amenable for use in gene therapy (3–7, 55–57).

Recently, the application of baculovirus vectors to gene therapy has also been shown to have several advantages: (i) low cytotoxicity in mammalian cells even at a high multiplicity of infection, (ii) inherent inability to replicate in mammalian cells to induce pathogenicity, and (iii) the absence of preexisting antibodies. In addition to the efficient gene delivery into mammalian cells,

mRNA. (D) Huh7 cells infected with HCVcc at an MOI of 1 and incubated for 72 h were inoculated with rBV-GFP (MOI of 100) in the presence or absence of human recombinant IFN- α (rIFN- α) (100 U/ml). At 24 h after inoculation, the cell extracts were subjected to SDS-PAGE and immunoblotted with antibodies against GFP, NS5A, or β -actin, respectively. (E) Relative luciferase activity in Huh7 cells with IRF3, IPS-1, or STING knocked down. Cells were inoculated with rBV-luc at an MOI of 100. At 24 h after inoculation, the luciferase activity of cell lysates was determined (right panel). Luciferase activity is normalized to control shNC cells, and data represent the means \pm SD from 2 independent experiments. Total RNA was extracted, and the expression of IRF3, IPS-1, or STING mRNA was determined by real-time PCR (left panel). Data from the real-time PCR were normalized to the amount of GAPDH mRNA.

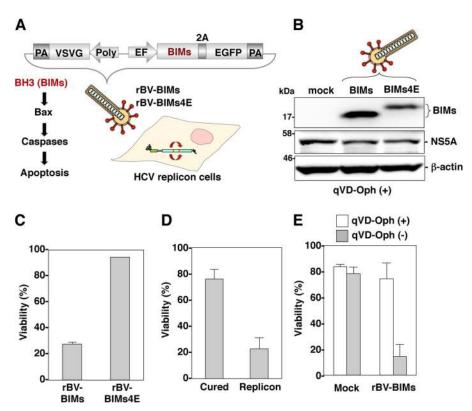


FIG 7 Induction of apoptosis in cells replicating HCV RNA by a recombinant baculovirus expressing a proapoptotic protein. (A) Structure of recombinant baculoviruses, rBV-BIM_S, and rBV-BIM_S4E carrying cDNAs of either BIM_s or BIM_s4E 2A peptides, and enhanced GFP under the control of the EF promoter. The apoptosis pathway was induced by BIM_s. (B) HCV replicon cells were infected with either rBV-BIM_S or rBV-BIM_S4E at an MOI of 500 in the presence of the caspase inhibitor qVD-Oph (20 μ M). Cells were subjected to immunoblotting by appropriate antibodies. (C) HCV replicon cells and cell viability was determined by PI exclusion at 24 h postinfection. (D) HCV replicon cells and cured cells were infected with rBV-BIM_S at an MOI of 500, and cell viability was determined by PI exclusion at 24 h postinfection. (E) HCV replicon cells and cured cells were infected with rBV-BIM_S at an MOI of 500 in the presence of 20 μ M qVD-Oph, and cell viability was determined by PI exclusion at 24 h postinfection. (E) HCV replicon cells are cells were infected with rBV-BIM_S at an MOI of 500 in the presence of 20 μ M qVD-Oph, and cell viability was determined by PI exclusion at 24 h postinfection. (E) HCV replicon at 24 h postinfection. Diffection at 24 h postinfection. (E) HCV replicon cells and cured cells were infected with rBV-BIM_S at an MOI of 500 in the presence of 20 μ M qVD-Oph, and cell viability was determined by PI exclusion at 24 h postinfection. (E) HCV replicon at 24 h postinfection.

the baculovirus has been shown to stimulate a host innate immune response in mammalian cells and to confer protection from lethal virus infection in mice (8, 10, 18, 19). Furthermore, it has been reported that the baculovirus possesses a strong adjuvant activity to promote humoral and cellular immune responses (24). Although these characteristics might be disadvantageous for the transgene expression, the baculovirus may have potency for use as a vaccine vector possessing adjuvant activity. Recombinant baculoviruses bearing the chimeric GP64 with Plasmodium berghei circumsporozoites (58, 59), Plasmodium falciparum circumsporozoites (60), Toxoplasma gondii (61), rinderpest virus (62), footand-mouth disease virus (63), pseudorabies virus (64), and avian influenza virus (65) have been shown to elicit an efficient antibody response. These results suggest that recombinant and pseudotyped baculoviruses encoding or transiently bearing foreign epitopes may induce immune responses in the antigen-presenting cells. Furthermore, gene delivery by using recombinant baculoviruses bearing specific ligands of interest may elicit more potent immunogenicity (66-68).

In this study, we have shown that reporter gene induction by recombinant baculovirus was enhanced in STING-, TBK1-, or IRF3-deficient MEFs. Moreover, an endogenous STING was translocated to the perinuclear region upon infection with recombinant baculovirus, suggesting that STING plays a crucial role in the induction of the innate immune response via baculovirus infection. Although a direct interaction of viral and synthetic DNAs with STING is not known, baculoviral DNA may be involved in the STING-associated signaling complex, including TBK1 and IRF3. Baculovirus has a circular dsDNA that contains bioactive CpG motif sequences, and the immune responses induced by the baculovirus DNA produce inflammatory cytokines and type I IFNs through TLR-dependent and -independent pathways, respectively (18, 19). Previously, Zhang et al. reported that the helicase DDX41, which is a member of the DExD/H-box helicase superfamily, acts as a cytosolic DNA sensor in immunocompetent cells (69). However, the involvement of DDX41 in type I IFN production in other cell types has not been demonstrated. Recently, Chen and colleagues have identified a cyclic GMP-AMP (cGAMP) synthase (cGAS) as a candidate for a cytosolic DNA sensor (70, 71). cGAMP produced by cGAS works as an endogenous second messenger and activates STING in cells stimulated by a broad species of DNAs. Although further studies are needed, it might be feasible to speculate that cGAS participates in the baculovirus-mediated innate immune response in a cell-type-specific manner.

IPS-1^{-/-} MEFs exhibit enhanced gene transduction by recombinant baculovirus, which is consistent with the report that murine gammaherpesvirus induces cytokine production in an IPS-1-

dependent manner (72), suggesting the involvement of the RLR pathways in the innate immune responses against DNA viruses. Although the evidence is not yet conclusive, Choi et al. suggested that IPS-1 may exclusively participate in a branch of the B-DNA/ RLR/IRF3 signaling pathway (47). On the other hand, the small RNAs encoded by adenovirus (73) and the cytosolic DNA-dependent RNA polymerase III (74) are suggested to be involved in the RLR signaling pathway (75). Furthermore, transcripts of several early viral genes were detected in the baculovirus-transduced mammalian cells (76, 77). Collectively, these data suggested that viral genomes and RNA transcripts of baculovirus and cytosolic DNA-dependent RNA polymerase III may participate in the IPS-1-dependent IFN-B production in cells infected with recombinant baculovirus. Recently, it has been shown that HCV infection suppresses not only IPS-1-dependent IFN signaling by NS3/4A but also STING-mediated IFN response by NS4B (78, 79). Therefore, suppression of STING by HCV replication may participate in the efficient gene transduction in the HCV replicon cells by baculovirus.

Chronic infection with HCV is a major risk factor for liver diseases, including chronic hepatitis, cirrhosis, and hepatocellular carcinoma (80). Newly developed directly acting antivirals targeting NS3/4A protease, NS5A, and NS5B polymerase have achieved a sustained virological response in over 80% of individuals infected with HCV, but drug-resistant breakthrough viruses have been reported to emerge during the treatment (81). In this study, we have raised the possibility of selective elimination of HCVinfected cells by the enhanced gene transduction of recombinant baculoviruses expressing proapoptotic Bcl-2 family proteins. Thus, our notion indicates that combinations of baculovirus-mediated gene therapy with current chemotherapy would be useful methods for eliminating HCV.

In conclusion, we have demonstrated that baculovirus induces innate immune responses that attenuate transgene expression. Recently, it was reported that baculovirus vectors exhibited an effective transgene expression in pluripotent stem cells, without elicitation of immune response, apoptosis, or pluripotency differentiation (82), supporting our notion that the induction of an acute immune response attenuates transgene expression by baculovirus. Although further studies are needed to clarify the induction of immune responses in more detail, these characteristics of baculovirus might be advantageous for vaccine vehicles possessing adjuvant activity and vectors capable of selective gene transduction into cells with impaired innate immunity due to infection with various viruses.

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