

FIG. 2. (A) Genetic and transcriptional map of type 5 AAV. The p7 and p19 promoters drive Rep78 and Rep52, respectively. The p41 promoter transcribes two mRNAs. One expresses VP1, and the other is for VP2 and VP3. The initiation codon for VP2 is ACG, which is poorly utilized for translation, leading to production of a smaller amount of VP2 polypeptides than VP3 polypeptides. The ITRs are indicated by black boxes. pA, polyadenylation signal sequence. (B) Recombinant baculoviruses (rBac) constructed. An rBac for Rep utilizes a truncated promoter for the immediate-early 1 gene of *Orgyia pseudotsugata* nuclear polyhedrosis virus (Δ IE) for type 5 Rep78, and another RepBac expresses Rep78 under the control of a truncated p10 promoter (Δ p10). See Fig. 4A for details. Either RepBac uses the polyhedrin promoter (polh) for Rep52. For expression of type 5 capsid proteins, a recombinant baculovirus that harbored a VP gene on which the initiation codon for VP1 is mutated to ACG was constructed (VP5Bac). Another series of VPBacs that had the type 5 VP1 gene partially replaced by the corresponding portion of type 2 VP1 at the N terminus was generated. (C) Western analysis of Sf9 cells infected with VP5Bac. The initiation codon for VP1 was mutated to an ACG codon, which enabled synthesis of VP1, -2, and -3 from a single VP mRNA. The amount of VP1 synthesized was extremely small compared to that in 293 cells. rAAV5-GFP generated with VP5Bac was used for the infection of COS cells at 10^5 vg per cell. The number of GFP-positive cells was 10% of the number of positive cells obtained with rAAV5-GFP produced in 293 cells.

peptides produced with VP251Bac through VP256Bac. Each VPBac produced chimeric VP1 at levels comparable to those of VP2. Formation of empty capsids was confirmed by CsCl density gradient analysis of Sf9 cell lysate infected with VP254Bac, as shown in Fig. 3C. The peak of VP polypeptides came to the fraction of 1.31 g/cm³, a buoyant density of empty capsids. The GFP gene between the type 5 ITRs could be

packaged into each type of chimeric capsid, and all of the chimeric rAAV5-GFPs except VP251 could transduce COS cells with efficiency similar to that of 293 cell-produced rAAV5-GFP (data not shown). The yields of rAAV5-GFP produced with VP253Bac or VP254Bac were approximately 1.2 times higher than others, although the difference was not statistically significant. We thus used VP254Bac to produce rAAV5 for the next experiments.

The initial Rep baculovirus for type 2 rAAV production drove type 2 Rep72 expression with a truncated promoter for the immediate-early 1 gene of *Orgyia pseudotsugata* nuclear polyhedrosis virus (Δ IE) and type 2 Rep52 under the control of the polyhedrin promoter (31) (Fig. 2B). The AAV5 genome encodes nonstructural proteins Rep78 and Rep52 (Fig. 2A). Similarly, we constructed a Rep baculovirus that expressed type 5 Rep78 and Rep52 under the control of the Δ IE promoter and the polyhedrin promoter, respectively. The titers of the type 2 or type 5 Rep baculoviruses, however, were lower than those of other recombinant baculovirus vectors (e.g., VPBac, GFPBac). The immediate-early 1 gene promoter becomes active at the early stage of baculovirus infection, and we thought that early expression of Rep78 in insect cells might negatively affect the yields of recombinant baculoviruses. The very late p10 promoter, which is widely used for recombinant protein production, is active at the latest stage of baculovirus infection. Thus, to delay and suppress the expression of Rep78, we tested a series of truncated p10 promoters. First, we screened the truncated p10 promoters for production of type 2 rAAV and selected one that generated high-titer rAAV2. Figure 4A shows the map of the p10 promoter and the truncated p10 promoter we constructed. The upstream TAAG sequence does not affect the activity of the p10 promoter (32). The sequence between the TAAG sequence and the p10 protein initiation codon at +72 (where the transcription start site is defined as +1) is called the burst sequence and is required for the "burst" of expression of the p10 protein at the very late stage of baculovirus infection. The *vlf-1* transactivator interacts with the burst sequence and strongly stimulates the transcription from the p10 promoter (35). To construct a weak p10 promoter (Δ p10), we removed the burst sequence between positions +39 and +72 from the original p10 promoter. The Δ p10 promoter was best for the production of rAAV2 among a series of truncated p10 promoters we examined. The titers of recombinant baculoviruses with the Δ p10 promoter were comparable to those of other recombinant baculoviruses. The Δ p10 promoter was transferred to express type 5 Rep78 (Fig. 2B). Figure 4B compares the time courses of type 5 Rep expression by Δ IE and Δ p10 promoters over 72 h after infection, indicating that the Δ p10 promoter-driven Rep78 expression was detected at 24 h after infection while the Δ IE promoter expressed Rep78 as early as 12 h after infection. To examine whether this modest difference in the levels of Rep78 affected replication of the AAV vector DNA, we isolated the low-molecular-weight DNA from the Sf9 cells infected with hGFP baculovirus and a Rep baculovirus (Fig. 4C). A ladder of replicative forms (RF) of rAAV5 DNA began to appear at 36 h postinfection in either case. The expected size of rAAV5-hGFP or monomer RF is 2.4 kb and the sizes of dimer and trimer RF are 4.8 and 7.2 kb, which is consistent with the result of the agarose gel electrophoresis.

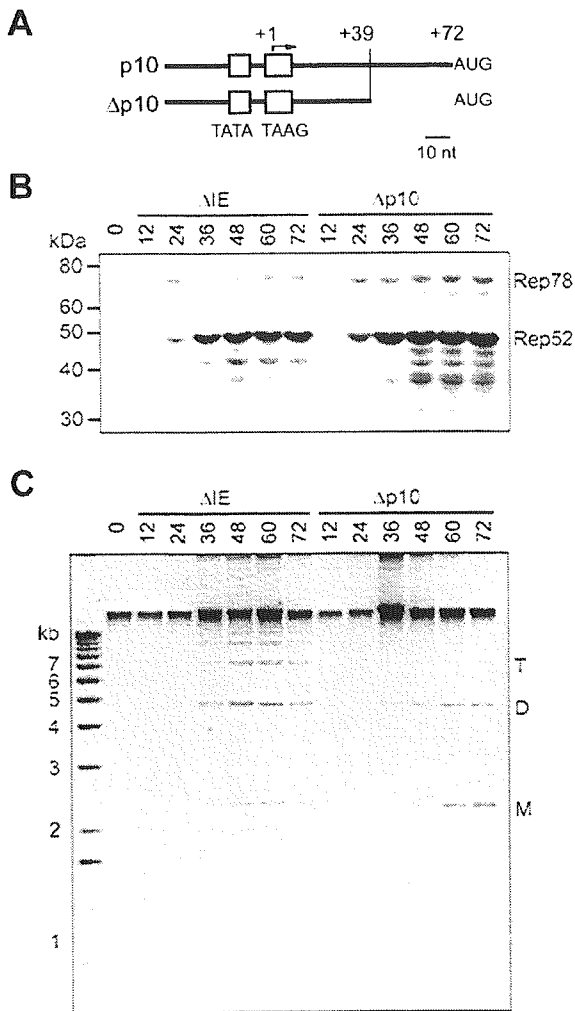


FIG. 4. (A) Map of the $\Delta p10$ promoter used for Rep78 expression. The sequence between positions +39 and +72 is deleted in the $\Delta p10$ promoter, where the T of the TAAG sequence or the transcription start site (marked with a bent arrow) is defined as +1 and the A of the p10 protein AUG codon is defined as +72. The original AUG codon for the p10 protein was mutated to ACT with pFastBac Dual (Invitrogen). The positions of the TATA box and the TAAG sequence are indicated. (B) Time course of Rep78 expression by ΔE or $\Delta p10$ promoter. Sf9 cells were infected with a Rep baculovirus, and the cells were harvested at the times indicated (in hours) for Western analysis with a monoclonal anti-Rep antibody. (C) Replication of hGFP vector DNA in insect cells. Sf9 cells were coinfecting with a Rep baculovirus and an hGFP baculovirus at 1 PFU per cell and incubated for the times indicated (in hours). Low-molecular-weight DNA was isolated, and DNA equivalent to 10^6 cells was resolved onto a 1% agarose gel. T, trimer; D, dimer; M, monomer.

der electron microscopy, showing typical rAAV particles of a diameter of 20 nm in addition to empty capsids (Fig. 5C). According to the staining pattern, approximately 30% of capsids contained vector genomes. In another experiment, rAAV5-hGFP was purified with two rounds of CsCl ultracentrifugation and the titers of rAAV5-hGFP were determined by real-time PCR using a pair of CMV-specific primers. Figure 5D summarizes the yields of rAAV5-hGFP with the use of different serotypes of small Rep. The titer of rAAV5-GFP

produced with type 1, 2, 3, or 4 small Rep was $56,000 \pm 3,200$ ($n = 4$), $41,000 \pm 18,900$ ($n = 4$), $42,000 \pm 7,300$ ($n = 3$), or $39,000 \pm 3,500$ ($n = 3$) particles per Sf9 cell, respectively, while that of rAAV5-GFP produced using AAV5 Rep52 was $13,500 \pm 3,200$ ($n = 5$). The rAAV5-hGFP particles produced with the indicated serotype Rep52 were further purified by anion-exchange column chromatography, and a total of 3×10^9 vg of either rAAV5-hGFP were then fractionated by sodium dodecyl sulfate-polyacrylamide gel electrophoresis and examined by silver staining along with 293 cell-produced rAAV5-hGFP (Fig. 5E). Densitometric analysis indicated that the intensities of the VP3 bands were almost equal to one another.

Type 5 vector DNA was packaged into type 5 capsids consisting of chimeric VP1 between types 2 and 5 in the baculovirus system. To examine the possible effect of the chimeric VP1 on packaging of type 5 vector DNA with heteroserotypic Rep52, we tested the production of rAAV5-hGFP by using either Rep5/1Bac or Rep5/5Bac and VP5Bac or VP254Bac. Interestingly, the yields of rAAV5 produced with type 5 Rep52 and type 2/5 chimeric capsids were constantly lower than yields produced with other combinations (Fig. 5F). Type 1 Rep52 was capable of packaging type 5 vector DNA into type 5 capsids and type 2/5 chimeric capsids with similar levels of efficiency. Although the result was not conclusive, the presence of a type 2 VP1-unique portion might interfere with type 5 Rep52 packaging rAAV5 DNA into type 5 capsids in insect cells.

Insect cell-produced rAAV5 infects cells via an $\alpha 2$ -3 sialic acid receptor. AAV2 capsids utilize HSPG as a primary coreceptor to infect target cells (30), whereas AAV5 capsids require $\alpha 2$ -3 sialic acid for efficient uptake (14). rAAV5 capsids generated in Sf9 cells are composed of VP1 partially replaced with type 2 VP1. The domains involved in receptor binding are within the VP3 portion (16), and the type 2 VP1-unique portion does not appear to be involved in attachment to target cells (19). To determine whether rAAV5 chimeric capsid particles infect cells via sialic acid and not via HSPG, we performed competition experiments with receptor analogs. The results of the heparin competition study show that rAAV2-GFP failed to transduce COS cells in the presence of heparin, an analog of heparan sulfate, as expected (Fig. 6A, top panels). By contrast, rAAV5-GFP produced in 293 cells (Fig. 6A, middle panels) or insect cells (Fig. 6A, bottom panels) was able to express GFP in COS cells irrespective of the presence of heparin, suggesting that Sf9 cell-produced rAAV5-GFP did not utilize HSPG as a primary coreceptor. The number of GFP-expressing cells was counted by flow cytometry, and the percent change in transduction compared to transduction in the absence of heparin was calculated, which clearly corroborated the observation with fluorescent microscopy. We next examined whether insect cell-produced rAAV5-GFP infects cells via $\alpha 2$ -3 sialic acid. As shown in Fig. 6B, COS cells were infected with rAAV5 generated in 293 cells (middle panels) or Sf9 cells (bottom panels) in the presence or absence of an analog of $\alpha 2$ -3 sialic acid, 3'-SLN. The analog inhibited GFP expression in COS cells by both 293 cell- and Sf9 cell-produced rAAV5-GFP, suggesting that rAAV5-GFP produced in insect cells infected cells via $\alpha 2$ -3 sialic acid as did 293 cell-produced rAAV5. To confirm that rAAV5-GFP derived from insect cells utilized sialic acid as a cell attachment receptor, we infected cells denuded of sialic acid by neuraminidase treatment. The

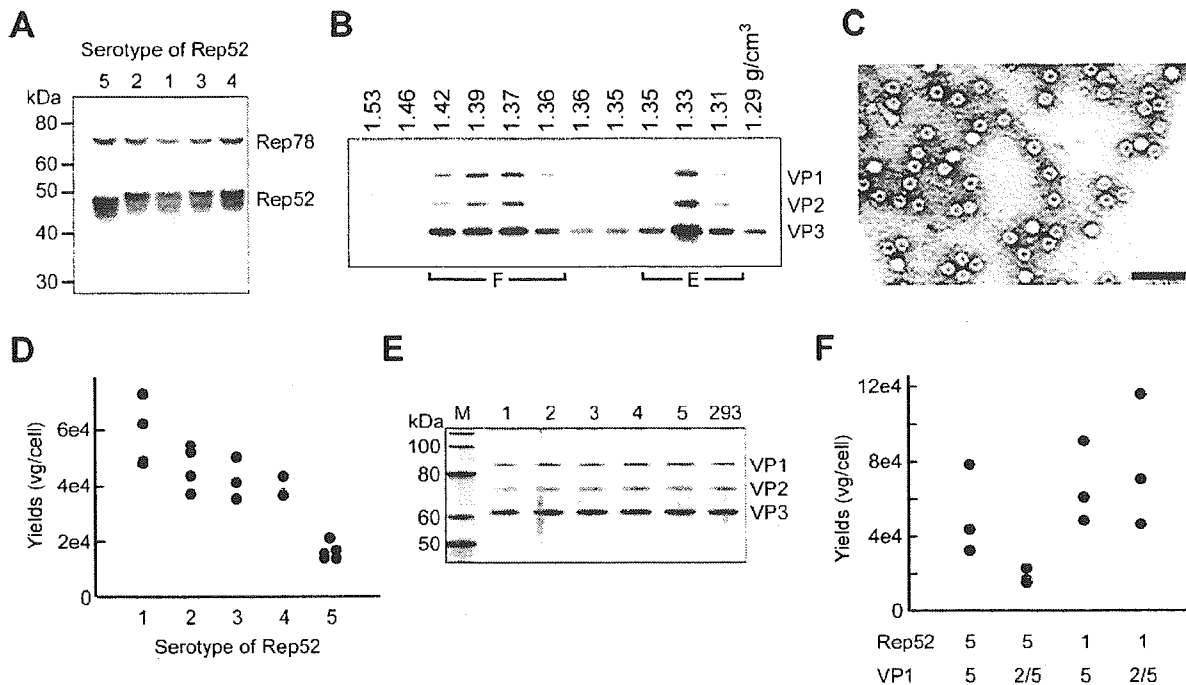


FIG. 5. (A) Western analysis of RepBacs expressing type 5 Rep78 and type 1, 2, 3, 4, or 5 Rep52 with an anti-Rep antibody. (B) Analysis of Sf9 cells coinfecting with Rep, VP254, and hGFP baculoviruses by CsCl density gradient ultracentrifugation. Three days after infection, the cells were lysed and subjected to ultracentrifugation. F, filled, or containing rAAV particles; E, empty capsids. (C) Negative staining of rAAV5-hGFP particles purified with ion-exchange column chromatography alone. Particles were stained with 2% uranyl acetate. Magnification, $\times 100,000$. Bar, 100 nm. (D) Generation of rAAV5-hGFP produced with different serotypes of Rep52. The yield of rAAV5-hGFP produced with type 1, 2, 3, 4, or 5 small Rep was $56,000 \pm 3,200$ ($n = 4$), $41,000 \pm 18,900$ ($n = 4$), $42,000 \pm 7,300$ ($n = 3$), $39,000 \pm 3,500$ ($n = 3$), or $13,500 \pm 3,200$ ($n = 5$) particles per Sf9 cell, respectively. (E) Analysis of rAAV5-hGFP produced in insect cells or 293 cells by silver staining. rAAV5-hGFP (3×10^8 particles) produced with serotype 1, 2, 3, 4, or 5 and that produced in 293 cells were resolved onto a 4 to 12% NuPAGE Bis-Tris gel (Invitrogen). Lane M, molecular size markers. (F) Comparison of the yields of rAAV5-hGFP produced with type 1 or type 5 Rep52 and VP5Bac or VP254Bac. Sf9 cells were coinfecting with hGFPBac, Rep5/1Bac or Rep5/5Bac, and VP5Bac or VP254Bac at an MOI of 1 in each of three independent experiments. The rAAV5-hGFP produced was purified by two rounds of CsCl density gradient ultracentrifugation, and the genomic titer was determined by real-time PCR.

result shows that prior incubation with neuraminidase significantly inhibited the transduction of COS cells mediated by rAAV5-GFP produced in 293 cells and Sf9 cells (Fig. 6C).

Comparison of transduction efficiencies with rAAV5 in cultured cells. We next compared the efficacy of rAAV5-GFP/Neo produced in Sf9 cells to that for a mammalian-cell-produced counterpart. COS cells were infected with either Sf9-produced or 293-produced rAAV5-GFP/Neo at 1×10^5 through 1×10^2 vg per cell for 1 day, and the number of GFP-positive cells was counted by flow cytometry. As shown in Fig. 7A, both Sf9-produced and 293-produced rAAV5-GFP/Neo showed similar dose-response curves. In addition, the vector genome-to-transducing unit ratio was calculated based on the number of GFP-positive cells at 3×10^3 vg per cell. Three independently produced samples were examined, and the vector genome-to-transducing unit ratio for Sf9-produced rAAV5-GFP was $3.9 \times 10^4 \pm 1.6 \times 10^4$ (mean \pm standard deviation), while the ratio for 293-produced rAAV5-GFP was $3.6 \times 10^4 \pm 1.2 \times 10^4$. These results indicated that insect cell-generated rAAV5-GFP/Neo had a similar ability to transduce COS cells. Although the capsids produced in Sf9 cells contain type 2/5 chimeric VP1 and those produced in HEK293 cells were composed of original type 5 VP1, rAAV5-GFP/Neo de-

rived from Sf9 cells and that derived from HEK293 cells did not show any significant difference in GFP expression in COS cells, suggesting that the difference in the VP1-unique portion did not impact the expression of the transgene or affect the intracellular processing of type 5 capsids in COS cells. We also compared transduction efficiencies of rAAV5-hGFP generated in Sf9 cells and rAAV5-hGFP generated in HEK293 cells. Surprisingly, the dose-response curve obtained by Sf9-produced rAAV5-hGFP shifted to the right and the number of GFP-positive cells at the dose of 3×10^3 vg per cell was five times larger than that for 293-produced rAAV5-hGFP (Fig. 7B). Since the substitution of the type 5 VP1-unique portion with the equivalent portion of type 2 did not impact the GFP expression in COS cells (Fig. 7A), we explored the rAAV genomes packaged into vector capsids. Virion DNA was isolated and analyzed on an alkaline gel. After electrophoresis, the DNA was transferred to a nylon membrane and hybridized with a ^{32}P -labeled CMV-specific probe. The GFP/Neo DNA packaged into AAV5 capsids is essentially the same in size and amount as expected (Fig. 7C). We next analyzed virion DNA isolated from rAAV5-hGFP produced with the indicated serotype Rep52 in insect cells, as well as 293-produced rAAV5-hGFP (Fig. 7D). The encapsidated hGFP DNA is present as

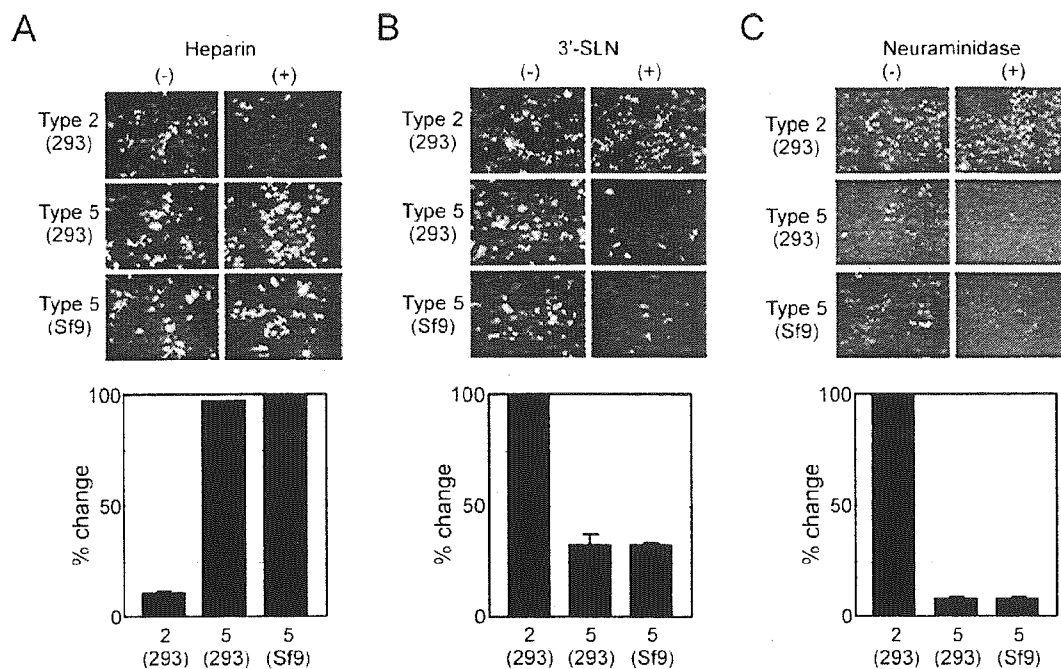


FIG. 6. (A) Heparin, an analog of HSPG, does not inhibit transduction of COS cells infected with rAAV5-GFP/Neo produced in insect cells. Cells were infected with adenovirus at 3 PFU per cell and incubated at 37°C for 2 h. After being washed with medium, the cells were infected with rAAV2-GFP/Neo produced in 293 cells at 10^4 vg per cell or rAAV5-GFP/Neo generated in 293 cells or Sf9 cells at 10^5 vg per cell in the presence or absence of 20 μ g per ml of heparin in triplicate. One day after infection, the cells were observed under a fluorescent microscope. The number of GFP-positive cells was also counted by flow cytometry. Data are presented as percent change in transduction compared to transduction in the absence of heparin. (B) An analog of α -2-3 sialic acid inhibits both 293 cell- and Sf9 cell-produced rAAV5-GFP/Neo. COS cells were infected with an adenovirus at 3 PFU per cell and incubated at 37°C for 2 h. After being washed with medium, the cells were infected with rAAV2-GFP/Neo at 10^4 vg per cell or rAAV5-GFP at 10^5 vg per cell for 1.5 h in the presence of 0 or 0.5 mM 3'-SLN (Sigma-Aldrich). The cells were washed twice with medium and further incubated for 1 day. The number of GFP-expressing cells was measured by flow cytometry ($n = 3$). Data are presented as percent change in transduction compared to transduction in the absence of the analog. (C) Neuraminidase treatment of COS cells inhibits transduction with rAAV5-GFP generated in 293 cells or Sf9 cells. COS cells were infected in triplicate with adenovirus at 3 PFU per cell for 1 h at 37°C, treated with 0.08 U per ml of neuraminidase (*Vibrio cholerae*, type III; Sigma-Aldrich) for 1 h, and infected with 10^5 vg per cell of rAAV5-GFP produced in 293 cells or Sf9 cells for 2 h. COS cells were similarly treated and also infected with 10^4 vg per cell of rAAV2-GFP/Neo. The infected cells were then washed twice with medium and incubated for an additional day. After incubation, the cells were observed under fluorescent microscopy and the number of GFP-positive cells was counted by flow cytometry.

two DNA species. The higher-mobility virion DNA corresponds with 2.4-kb hGFP vector DNA or a single-stranded monomer, which is confirmed by comigration with a 2.4-kb vector DNA obtained by treatment with a restriction enzyme of the hGFP vector plasmid, pSR485hGFP. The lower-mobility DNA is the same in size as the monomer RF or duplex form of hGFP DNA (Fig. 7D) isolated from Sf9 cells coinfecting with RepBac and hGFPBac (Fig. 4C). The intensity of the larger virion DNA, which was quantified with an imaging analyzer, was roughly double that of shorter DNA for each rAAV5 produced in Sf9 cells. If the larger virion DNA is a monomer duplex form and thus has two CMV promoter sequences hybridizing to a CMV probe, then we estimated that the quantity of the double-stranded monomeric form was equal to that of the single-stranded monomer form in the rAAV5-hGFP virion produced in 293 cells is 1 to 3.5. AAV particles have been shown to package two copies of vector genomes that are less than 50% of the 4.8-kb AAV genome, and the packaged vector DNA appeared to be monomeric single-stranded and double-stranded RF (6). For gene expression, the single-stranded vector genome has to be

converted to a double-stranded form by either second-strand synthesis (8, 9) or annealing of complementary strands (23). The monomeric duplex vector DNA, on the other hand, can function directly as a template for mRNA synthesis. Thus, the more potent gene expression mediated by rAAV5-hGFP generated in Sf9 cells is probably due to the presence of the encapsidated monomer duplex form.

Comparison of efficacies of rAAV5 in vivo. To compare the efficacies of rAAV5 produced in 293 cells and rAAV5 produced in Sf9 cells, we constructed a type 5 vector that expressed human SEAP. rAAV5 particles produced in Sf9 cells consisted of chimeric VP1 between type 2 and type 5. To eliminate the possible difference in intracellular processing of rAAV5 particles due to replacement of the type 5 VP1-unique portion with the equivalent one of type 2, we compared the in vivo activities of rAAV5 particles containing type 2/5 VP1 polypeptides produced in insect and mammalian cells. Five mice each intramuscularly received a total of 10^{11} vg of rAAV5-SEAP generated in either 293 cells or Sf9 cells, and serum SEAP levels were monitored. As shown in Fig. 8A, the expression profile of the Sf9-produced type 5 SEAP vector differed from that of the 293-produced one. The rAAV5-SEAP

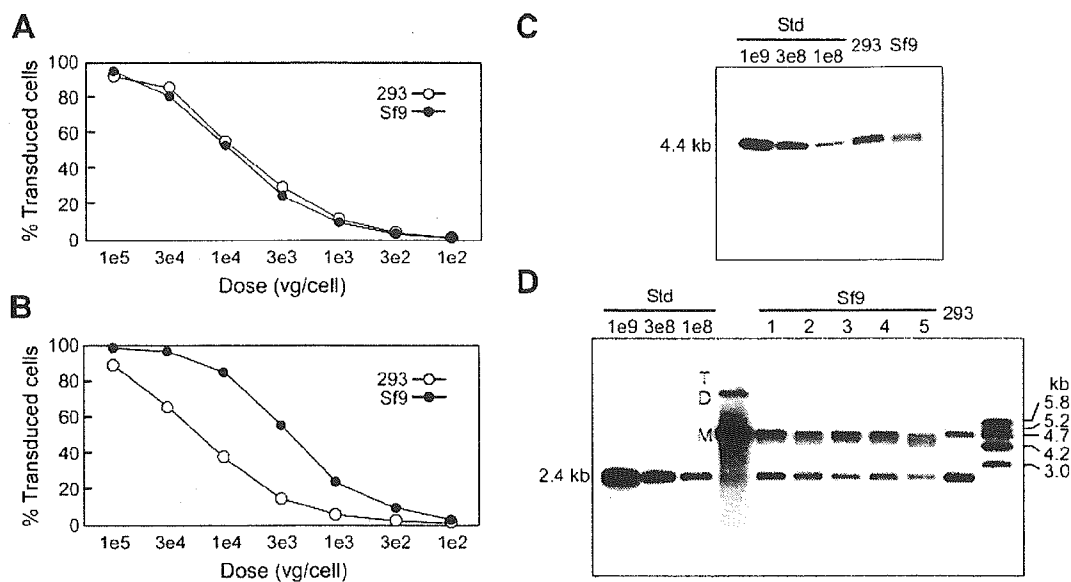


FIG. 7. Comparison of infectivities of rAAV5 produced in 293 cells and rAAV5 produced in Sf9 cells. (A) COS cells were infected with rAAV5-GFP/Neo produced in 293 cells or Sf9 cells at the doses indicated, ranging from 1×10^5 through 1×10^2 vg per cell. One day after infection, the cells were examined for GFP fluorescence by flow cytometry in triplicate. (B) COS cells were infected with rAAV5-hGFP produced in HEK293 cells or Sf9 cells at a dose of 1×10^5 through 1×10^2 vg per cell for 1 day. The number of GFP-positive cells was counted cytometrically. (C) Analysis of GFP/Neo vector virion DNA on an alkaline gel. Virion DNA was isolated from rAAV5-GFP/Neo produced in HEK293 cells or Sf9 cells by treatment with proteinase K, and samples equivalent to 3×10^8 vg were resolved onto a 0.8% alkaline gel. The DNA was then transferred to a nylon membrane and hybridized to a 32 P-labeled CMV-specific probe. A 4.4-kb-size copy number standard (Std) (1×10^9 , 3×10^8 , and 1×10^8 copies) was loaded, which was derived from the GFP/Neo vector plasmid, pSR485a, with a restriction enzyme that cut out the vector portion. (D) Alkaline agarose gel electrophoresis of virion DNA isolated from rAAV5-hGFP. Vector DNA isolated from rAAV5-hGFP particles produced with type 1, 2, 3, 4, or 5 Rep52 was analyzed along with 293-produced rAAV5-hGFP. Low-molecular-weight DNA isolated from insect cells infected with RepBac and hGFPBac (Fig. 4C) serves as a reference for monomer (M), dimer (D), and trimer (T) replicative forms.

generated in HEK293 cells showed a gradual increase in serum SEAP activity over 1 month after injection, which is a typical expression pattern by rAAV-mediated transduction. The Sf9-produced rAAV5-SEAP induced levels of SEAP activity at 1 or 2 weeks after injection that were more than 30 or 10 times higher, respectively, than those of the 293-produced rAAV5-SEAP, and the serum SEAP activity by Sf9 produced rAAV5-SEAP decreased at 4 weeks after injection. There was no significant difference between the two groups after 4 weeks following administration. We also analyzed the SEAP vector DNA on an alkaline gel (Fig. 8B). The expected size of rAAV5-SEAP vector genomes is 3.4 kb. The majority of 293-produced rAAV5-SEAP DNA is single-stranded monomer in both type 5 capsids and type 2/5 chimeric capsids. In addition to the 3.4-kb single-stranded vector genome, DNA extracted from Sf9 cell-produced rAAV5 particles contained an additional DNA of approximately 4.7 kb. One model for AAV packaging proposes that when the size of vector DNA is larger than the size of the wild-type AAV, 4.7 kb, the vector DNA is cleaved to 100% of the AAV genome during packaging into virion (6). The 4.7-kb virion DNA may be a cleaved product of duplex multimers synthesized in Sf9 cells.

DISCUSSION

Recent advances in understanding of biology of AAV and in production of rAAV have facilitated the use of rAAV as a gene transfer vector. A human clinical trial with rAAV2 expressing

a coagulation factor IX has shown that intramuscular delivery of more than 10^{15} rAAV2 particles would be required for amelioration of hemophilia B (15). Currently, the widely employed method for production of rAAV is transfection of packaging cells, such as HEK293 cells, with plasmids carrying AAV and adenovirus genes. Plasmid transfection is more easily adaptable to packaging different serotype AAV vectors than establishing a packaging cell line. However, the transfection process is the limiting step in rAAV production, which requires adherent HEK293 cells on a two-dimensional surface for efficient production of rAAV.

The production of other AAV serotype-derived vectors has been described previously (26) and follows the strategy developed for rAAV2 (20). Some modifications have been reported, such as lipofection of 293 cells in suspension culture in serum-free media, which makes the handling of the cells and the purification step easier (28). However, the use of a lipid reagent for transfection may be neither cost-effective nor scalable. A recombinant herpes simplex virus harboring type 5 *rep* and *cap* genes was created to eliminate the transfection process (33), although the yields of rAAV5 were low. The baculovirus/insect cell-based rAAV5 production system presented here does not require plasmid transfection and is scalable. By extrapolation from culture volume, we expect to obtain more than 10^{14} particles of rAAV5-GFP from a 1-liter culture. This is consistent with yields of rAAV1 or rAAV2 produced in Sf9 cell cultures (20a, 31).

To produce infectious rAAV5 particles in insect cells, we

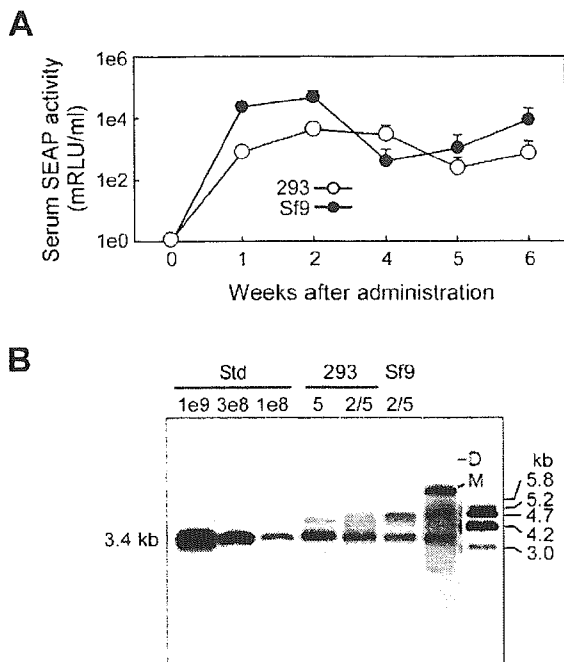


FIG. 8. (A) Serum SEAP activity following intramuscular injection of rAAV5-SEAP. Five mice each received a total of 10^{11} vg of pseudotyped rAAV5-SEAP produced in 293 cells or Sf9 cells into tibialis anterior muscles. Blood was taken at the indicated weeks after injection. The serum SEAP activity was measured by a SEAP report gene assay (Roche Diagnostics, GmbH, Penzberg, Germany). RLU, relative light units. (B) Molecular analysis of SEAP vector DNA. Vector genomes were isolated from type 5 or type 2/5 SEAP vector particles produced in HEK293 cells or from Sf9-produced rAAV5-SEAP. Extrachromosomal low-molecular-weight DNA isolated from Sf9 cells infected with RepBac and SEAPBac was also analyzed. Copies (10^9 through 10^5) of copy number standard (Std) vector DNA derived from SEAP vector plasmid, pAAVSEAP, were also loaded. M, monomer replicative form of SEAP vector genomes; D, dimer replicative form.

inserted an N-terminal portion of type 2 VP1 into the corresponding site of type 5 VP1. The N termini of VP1 polypeptides contain the phospholipase A₂ motif and are essential to viral infectivity (36). Electron microscopy indicated that the VP1-unique portion is hidden within type 2 capsids and appears on the surface of the capsids during the infectious pathway in cells (19). The VP1-unique portion is well conserved among different AAVs. Comparison of the portion among serotypes 1 through 4 and 6 revealed that one serotype is more than 80% identical to another. The type 5 VP1-unique portion is 70 to 75% identical to that of other serotypes, while the sequence alignment of VP2 or VP3 of AAV1 through AAV6 showed that type 5 is approximately 55% identical to other serotypes. The initial trial mutation of the start codon for type 5 VP1 gene to ACG failed to produce infectious rAAV5 particles due to low synthesis of VP1 polypeptide (Fig. 2C). However, the successful generation of rAAV2 particles in insect cells and the notion that the VP1-specific region is well conserved among different serotypes led us to construct a chimeric type 5 VP1 polypeptide whose N-terminal portion was partially replaced by the equivalent portion of type 2. The transduction of COS cells and mouse muscles with rAAV5 produced in

insect cells clearly indicated that the chimeric VP1, VP254, could confer infectivity to it (Fig. 7 and 8).

The strategy of producing AAV "pseudotyped" vectors, typically consisting of AAV2 ITR and non-AAV2 capsids, such as AAV4 and AAV5, has been reported previously (2, 3, 26, 34). We first tested similar pseudopackaging of rAAV DNA type 2 ITRs into type 5 capsids with type 2 RepBac in insect cells. However, the yields of vector particles produced were four times lower than those reached by packaging type 5 DNA into type 5 capsids (data not shown). We also examined the production of rAAV5 by packaging type 2 AAV DNA with type 2 Rep78 and type 5 Rep52 into type 5 capsids, which also resulted in low yields of rAAV5 (data not shown). The production of type 5 vector in 293 cells by transfection with a type 5 vector plasmid and a type 5 *rep cap* plasmid usually yields more than 10^4 particles per HEK293 cell, and the production of pseudotyped type 5 vectors by using a type 2 AAV vector plasmid and type 2 *rep* and type 5 *cap* plasmid recovers 3×10^5 particles per cell (unpublished observation), an observation consistent with the production in Sf9 cells.

Using Sf9 cells, we found that Rep52 proteins of other serotypes were capable of packaging DNA with type 5 ITRs into type 5 capsids more efficiently than type 5 Rep52. Type 2 small Rep protein has been shown to package the AAV2 genome into preformed capsid with its helicase activity in collaboration with large Rep protein (7, 17). The small Rep protein associates with Rep78/68 (24) and probably specifically interacts with large Rep protein during encapsidation (7). The basis for the improved AAV packaging with non-type 5 Rep52 remains to be elucidated. To exclude the possibility that cellular proteins and/or baculovirus proteins played a major role in packaging type 5 DNA, we used a RepBac that expressed only type 5 Rep78 for production of type 5 rAAV. No rAAV5 particles were recovered from the recombinant baculovirus-infected Sf9 cells (data not shown), suggesting that the small Rep protein is absolutely required for generating rAAV5 in insect cells. As shown in Fig. 5F, the fact that the partial replacement of the VP1-unique portion with the corresponding portion of type 2, the strategy we took to generate infectious type 5 particles in insect cells, inhibited type 5 Rep52-mediated introduction of type 5 ITR genomes into type 5 capsids may only indicate the role of the type 2 VP1-unique portion as a physical barrier during packaging of rAAV genomes into capsids. We believe that under a special circumstance, such as in invertebrate cells, heteroserotypic Rep52 is superior to type 5 Rep52 in packaging rAAV DNA with type 5 ITR into type 5 capsids. It is interesting to examine whether other serotypes of Rep52 can package type 5 rAAV DNA into type 5 capsid in mammalian cells. We are currently investigating the packaging of type 5 genome into type 5 capsids with different serotypes of Rep52 in HEK293 cells.

The majority of the vector genome of rAAV5 produced in HEK293 cells in the present study is in single-stranded monomeric form, irrespective of the size of the vector genome (Fig. 7C and D and 8B). However, when the size of vector DNA is shorter than the size of the wild-type AAV genome, insect cells tend to package longer, 4.7-kb DNA into type 5 capsids. The 4.7-kb longer virion DNA in Sf9-produced rAAV5 appears to be a cleavage product of multimers of replicated vector genomes. If the size of a multimer is within the packaging limit,

it is efficiently introduced into AAV capsids. If a multimer is larger than 4.8 kb in size, a partially truncated multimer is packaged into AAV capsids in insect cells (6). Sequencing of 4.7-kb DNA packaged into virions will be a key to disclosing the difference between packaging of vector DNA into capsids in HEK293 cells and insect cells. The difference in packaged virion DNA between rAAV5 produced in human cells and in insect cells provides important information on designing vector DNA for production of rAAV5 in insect cells.

In summary, we developed a new method for production of rAAV5 in insect cells, which offers a better alternative to the existing production methods of rAAV5, although the vector genomes packaged into capsids differ in size from rAAV5 produced in HEK293 cells. The robust generation in suspension culture will facilitate the use of type 5 rAAV not only for basic studies but also for clinical studies.

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Specific and Efficient Transduction of Cochlear Inner Hair Cells with Recombinant Adeno-associated Virus Type 3 Vector

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Recombinant adeno-associated virus (AAV) vectors are of interest for cochlear gene therapy because of their ability to mediate the efficient transfer and long-term stable expression of therapeutic genes in a wide variety of postmitotic tissues with minimal vector-related cytotoxicity. In the present study, seven AAV serotypes (AAV1–5, 7, 8) were used to construct vectors. The expression of EGFP by the chicken β -actin promoter associated with the cytomegalovirus immediate-early enhancer in cochlear cells showed that each of these serotypes successfully targets distinct cochlear cell types. In contrast to the other serotypes, the AAV3 vector specifically transduced cochlear inner hair cells with high efficiency *in vivo*, while the AAV1, 2, 5, 7, and 8 vectors also transduced these and other cell types, including spiral ganglion and spiral ligament cells. There was no loss of cochlear function with respect to evoked auditory brain-stem responses over the range of frequencies tested after the injection of AAV vectors. These findings are of value for further molecular studies of cochlear inner hair cells and for gene replacement strategies to correct recessive genetic hearing loss due to monogenic mutations in these cells.

Key Words: adeno-associated virus, serotype, gene transfer, cochlea, hair cells

INTRODUCTION

The total number of hair cells in the cochlea is finite. They are not renewed and there is very little (if any) redundancy in this population. The irreversible loss of cochlear hair cells is presumed to be a fundamental cause of permanent sensorineural hearing loss. Gene transfer into hair cells presents numerous opportunities for protecting these cells. There is considerable interest in the development of viral vectors to deliver genes to the cochlea to counteract hearing impairment, and recent studies have focused on vectors based on adenovirus [1–3], herpes simplex virus [4–6], lentivirus [7], and adeno-associated virus (AAV) [8,9]. The patterns of vector-encoded transgene expression have been found to differ significantly among vectors. Cochlear hair cells can be efficiently transduced with adenovirus vectors [10–12].

However, these vectors were found to provoke a strong immune response that could damage recipient cells and compromise cochlear function [10,13,14]; they are also incapable of mediating prolonged transgene expression [15,16]. Although AAV vectors might overcome these problems, the transduction of hair cells by AAV2-derived vectors is controversial [8,10,17]. To our knowledge, other AAV serotypes have not yet been tested as cochlear gene transfer vectors *in vitro* or *in vivo*. AAV vectors are of interest in the context of gene therapy because they mediate efficient transfer and long-term stable expression of therapeutic genes in a wide variety of postmitotic tissues with minimal vector-related cytotoxicity.

In this study, we assessed the utility of seven AAV serotypes as vectors with the chicken β -actin promoter associated with a cytomegalovirus immediate-early

enhancer (CAG)-driven enhanced green fluorescent protein (EGFP) gene [18] in the murine cochlea. Vectors were introduced by microinjection through the round window membrane [19]. As a result, we determined that the specific and efficient gene transduction of inner hair cells could be achieved by using AAV type 3 vectors.

RESULTS

Expression Profile of EGFP in the Cochlea

Several cell types line the cochlear duct and support the hair cells (Fig. 1A). We carefully made a small opening in the tympanic bulla and injected vectors derived from the AAV1–4, 7, and 8 pseudotypes into the cochlea of two strains of mice (C57BL/6) and ICR) through the round window membrane (Fig. 1B). The mode of EGFP expression in various murine cochlear hair cells had a close similarity and was essentially equal for both strains. We determined the distribution of AAV vector-mediated EGFP expression throughout the cochlea for all serotypes tested (Table 1). A principal finding is that the inner hair cells in the organ of Corti showed clear evidence of EGFP expression with all of the AAV serotype-derived vectors except for the AAV4 vector (Fig. 2). This result indicates that most of the vectors (AAV1–5, 7, and 8) could efficiently transduce cochlear inner hair cells *in vivo* when slowly infused into the scala tympani. The AAV3-based vector was the most efficient and specific of the serotypes in transducing cochlear inner hair cells (Fig. 3). Transduction with 5×10^{10} genome copies (gc)/cochlea of the AAV3 vector resulted in robust transgene expres-

sion in the inner hair cells. The spiral ganglion cells showed significantly higher levels of fluorescence per unit area with the AAV5-based vector (Fig. 2n), and the spiral ligament cells were transduced prominently with the AAV1 and AAV7 vectors (Figs. 2d and 2r). Histological sections of cochleae injected with the AAV4 vector identified EGFP-positive cells predominantly in connective tissue within the mesothelial cells beneath the organ of Corti and in mesenchymal cells lining the perilymphatic fluid spaces (Figs. 2j and 2l). Furthermore, we detected intense expression with the AAV5- and AAV8-based vectors in the inner sulcus cells and in Claudius' cells (Figs. 2p and 2x). We did not detect notable levels of gene expression in the outer hair cells, supporting pillar cells, or stria vascularis cells for any serotype.

Long-term Expression of EGFP

We examined cochlear expression of the EGFP transgene in animals sacrificed at 1–12 weeks. Expression persisted in cochlear tissues for up to 3 months after infusion, while the extent of expression peaked at 2 weeks.

Transgene Activity

We determined the percentage of inner hair cells transduced with the AAV3 vector. The mid- to high-frequency regions of the cochlea were efficiently transduced, as shown in Fig. 3. Almost all of the inner hair cells in the basal and middle cochlear regions were transduced with the AAV3 vector (Fig. 4). Transgene expression was not detected in the hair cells of the apical turn of the cochlea. The predominant expression in the middle and basal cochlear turns is reasonable, as the virus

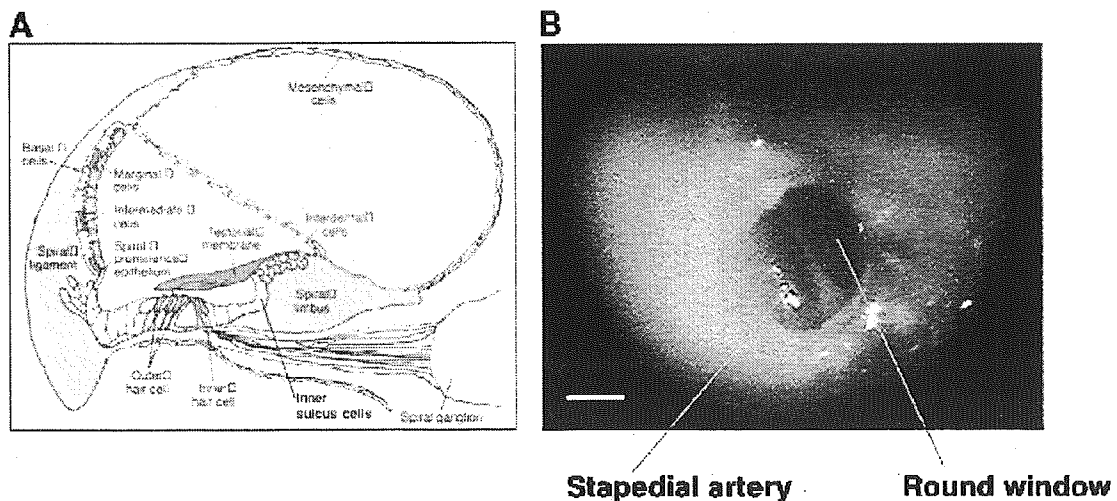


FIG. 1. (A) Schematic diagram of a cross section of the cochlea, demonstrating the scala vestibuli, scala tympani, and scala media or cochlear duct. The organ of Corti rests on the basilar membrane, with the hair cell cilia embedded in the gelatinous tectorial membrane. The outer margin of the cochlear duct contains the stria vascularis. Reproduced, by permission of the publisher, from [44]. (B) Direct visualization of the round window membrane in the right ear. The upper side of the picture is the back of the mouse and the right side is the head of the animal. The stapedial artery, a branch of the internal carotid artery, transverses an open bony semicanal within the round window niche. Bar denotes 500 μ m, 15 \times original magnification.

TABLE 1: Expression of transgene in the mouse cochlea with vectors derived from the AAV1-4, 7, and 8 pseudotypes

| Vector | Inner hair cells | Outer hair cells | Spiral ganglion | Stria vascularis | Spiral ligament | Spiral limbus | Spiral membrane | Reissner's membrane | Inner and outer pillar cells | Inner sulcus cells | Deiters' cells | Claudius' cells | Hensen's cells | Mesenchymal cells |
|--------|------------------|------------------|-----------------|------------------|-----------------|---------------|-----------------|---------------------|------------------------------|--------------------|----------------|-----------------|----------------|-------------------|
| AAV1 | +++ | - | ++ | - | ++ | ++ | ++ | - | - | + | - | - | - | ++ |
| AAV2 | ++ | - | + | - | + | + | - | - | - | - | - | - | - | - |
| AAV3 | ++++ | - | - | - | - | - | - | - | - | - | - | - | - | - |
| AAV4 | - | - | - | - | - | - | - | - | - | - | - | - | - | + |
| AAV5 | +++ | - | +++ | - | + | ++ | + | + | ++ | ++ | - | + | - | - |
| AAV7 | +++ | - | + | - | +++ | ++ | - | - | - | - | - | - | - | ++ |
| AAV8 | ++++ | - | - | - | + | + | - | - | - | ++ | - | + | - | + |

The level of expression was graded by fluorescence intensity on a four-level scale (+, ++, +++, +++) depending on the pixel/unit area count. +++++ means the strongest intensity of EGFP expression, + means the weakest intensity of EGFP expression, while - means no fluorescence.

was slowly infused into the scala tympani adjacent to the most basal turn of the cochlea. The percentage of transduced inner hair cells from the basal (high frequencies) to the apical (low frequencies) cochlear regions is shown in Fig. 4.

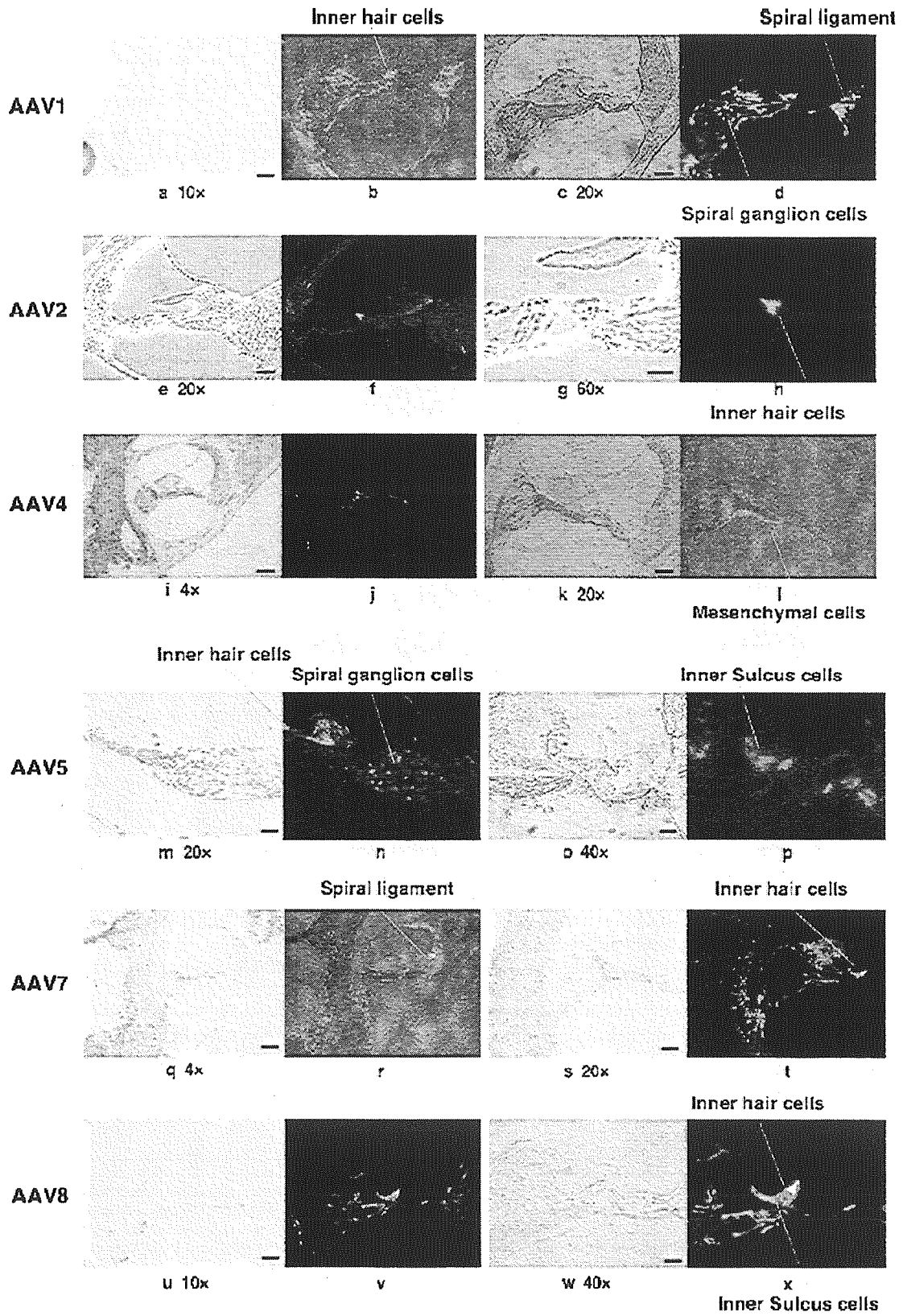
Cytotoxicity

We detected no deleterious effects on the viability of transduced cells. We compared evoked auditory brainstem response (ABR) threshold levels before and after injection, using a two-way repeated measure of the analysis of variance. There was no significant loss in ABR and hence no change in cochlear function for up to 10 days following vector infusion (Figs. 5A and 5B). In addition, the cellular and tissue architecture of experimental cochleae remained intact. There was no evidence of endolymphatic hydrops after AAV vector injection in any of the animals. We observed no significant destruction of the inner or outer hair cells (Fig. 5C).

DISCUSSION

In the present study, we assessed the utility of vectors derived from seven AAV serotypes for gene delivery into the cochlea. Our results showed that the AAV3 vector was the most efficient and specific in transducing cochlear inner hair cells, although these cells could also be transduced with AAV1, 2, 5, 7, and 8 vectors. The transduction efficiency of the spiral ganglion by the AAV5 vector was particularly high, followed by that of the AAV1, AAV2, and AAV7 vectors. The efficient and specific transduction of inner hair cells with the AAV3 vector suggests that it recognizes a unique host range with a distinct cellular receptor. Transduction efficiency is dependent on initial viral binding (a property of the viral capsid), entry, and various postentry processes such as intracellular trafficking and second-strand synthesis [20-22]. The genome size of AAV vectors has also been demonstrated to affect transduction efficiency [23]. Comparisons of the serotypes have indicated that heterogeneity in the capsid-encoding regions and a differential ability to transduce cells may be associated with different receptor and co-receptor requirements for cell entry [24]. However, the receptors and co-receptors of AAV3 have not yet been clearly identified.

In the current study, we found that cochlear inner hair cells could be transduced with six AAV serotypes, although Lalwini *et al.* [8] reported that outer hair cells could be transduced with a low titer (1×10^6 viral particles/ml) of AAV2 *in vivo*. After injecting the AAV2 vector, we found that the spiral ganglion neurons, the inner hair cells, and the cells in the spiral ligament were all transduced. This transduction pattern differs from that reported in previous studies [8,10,17], and this discrepancy might be due to the different delivery methods and dissimilar promoters. Although the CAG promoter directs



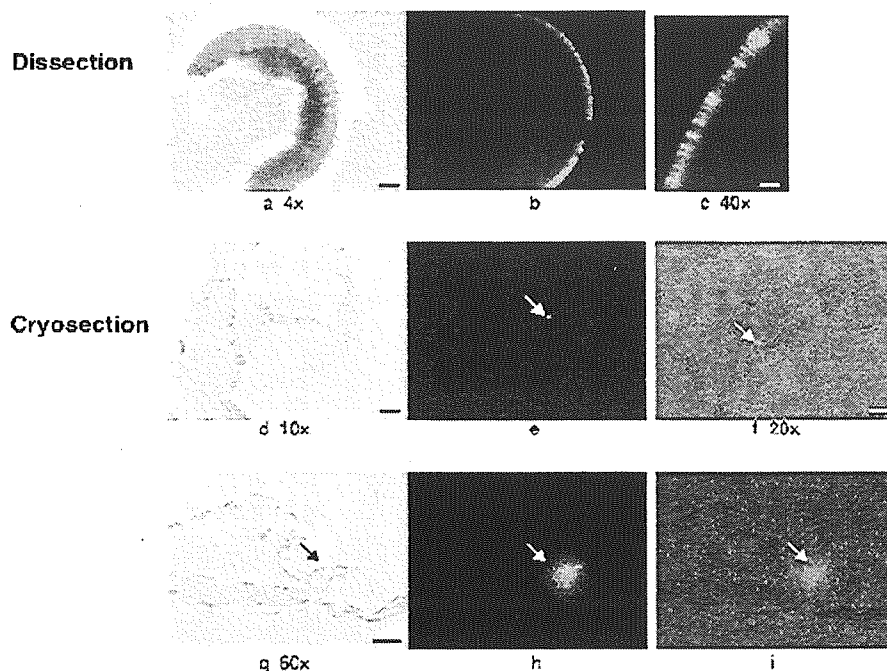


FIG. 3. Cochlear transduction with AAV3-CAG-EGFP. Dissected cochleae and cryosections show transgene expression in inner hair cells. (a) A light photomicrograph of the basal turn of the cochlea is shown, illustrating its laminar structure. (b) A fluorescence photomicrograph of this dissection. (c) A higher magnification view of the dissection shown in (b), illustrating a row of inner hair cells in the organ of Corti expressing EGFP. (d–i) Representative photomicrographs from three magnifications of a radial cochlear cryosection. (d) Light photomicrograph of an intact cochlear duct. Fluorescence photomicrograph of this duct is shown in (e). (h and i) A higher magnification of (e), illustrating EGFP expression within inner hair cells. Cryosections show transgene expression in the inner hair cells (arrows). Scale bars: 4 ×, 250 μm; 10 ×, 100 μm; 20 ×, 50 μm; 40 ×, 25 μm; 60 ×, 25 μm.

higher expression than do the cytomegalovirus (CMV) and EF-1 α promoters [25], each promoter drives reporter gene expression in different cell types [26,27].

Cell-specific or -selective infectivity of the viral vectors suggests the presence of various factors to introduce the distinct expression patterns of the transgenes. Spiral ganglion neurons and glial cells can be transduced with a lentivirus-GFP construct *in vitro* but not *in vivo* [7]. The differential transducibility under *in vivo* and *in vitro* conditions reflects a high degree of structural isolation of the spiral ganglion and other cell types—such as the cells on the periphery of the endolymph—from the perilymph into which the viral vector was introduced. The strict separation of the endolymph from the perilymph is maintained by tight junctions that line the boundary between these fluid chambers. The size of the viral particle may contribute to the observed variability in transgene expression promoted by different vectors. The diameters of adenovirus and retrovirus (including lentivirus) particles are approximately 75 nm and greater than 100 nm, respectively, while the diameters of AAV vectors are typically 11–22 nm [28,29]. Thus, the larger size of lentiviruses and adenoviruses may limit their subsequent

dissemination from the perilymph into the endolymph. The variable patterns of adenovirus- and lentivirus-mediated gene expression seen with different methods of inoculation may be due to the inoculation route, the volume and number of viral particles, differences in viral preparation, or differences in the method of transgene detection. The introduction of adenovirus vectors by cochleostomy or with an osmotic pump via the round window leads to a more efficient transduction of cochlear hair cells [30–32]. The apical domain (apical membrane and stereocilia) of cells in the sensory epithelium (hair cells and supporting cells) is bathed in endolymph, while the basal-lateral domain is immersed in perilymph. Access of the viral vectors to the endolymphatic space by cochleostomy may facilitate the transduction of hair cells and supporting cells. However, although the cochleostomy procedure has been tested, inoculation into the membranous labyrinth could not be confirmed [32]. In the present study, AAV vectors were found to infect cochlear hair cells easily *in vivo*, via round window injection.

Gene transfer into the cochlea through the round window membrane is ideal, because this procedure

FIG. 2. Transduction of the cochleae by AAV1-, AAV2-, AAV4-, AAV5-, AAV7-, and AAV8-based vectors. (a, c, e, g, i, k, m, o, q, s, u, and w) Light photomicrographs of cochlear cryosections. (b, d, f, h, j, l, n, p, r, t, v, and x) Fluorescence photomicrographs (green fluorescence from transgene). The spiral ligament cells were transduced prominently with the AAV1 and AAV7 vectors (d and r). Transgene expression in inner hair cells was detected with AAV1-, AAV2-, AAV5-, AAV7-, and AAV8-based vectors (b, h, n, l, and x). AAV4-based vector faintly transduced mesenchymal cells (j and l). The spiral ganglion cells showed significant levels of fluorescence with the AAV5-based vector (n). Intense fluorescence was detected with the AAV5- and AAV8-based vectors in the inner sulcus cells (p and x). Scale bars: 10 ×, 100 μm; 20 ×, 50 μm; 40 ×, 25 μm; 60 ×, 25 μm.

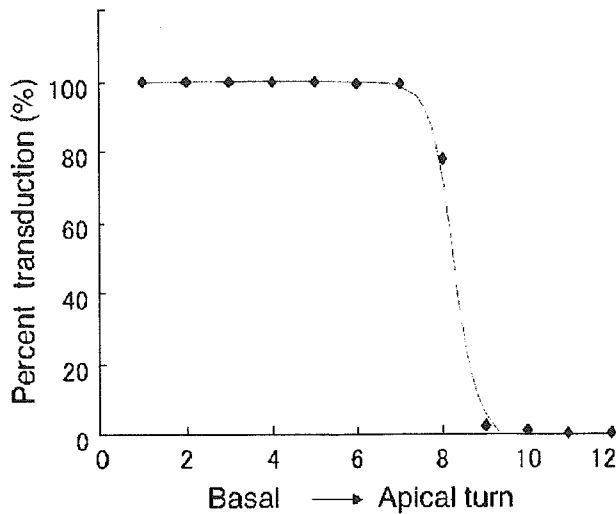


FIG. 4. EGFP expression profile of inner hair cells transduced with AAV3, as shown for a cross section subdivided into 12 segments ranging from the basal (high frequencies) to the apical (low frequencies) cochlear regions.

requires simple surgery without cochlear trauma [19]. Another critical factor in assessing the utility of a gene transfer vector is safety. Factors determining safety include the toxicity of the gene transfer agent itself, the provocation of immune responses, the generation of replication-competent virus, and the risk of creating genetically modified cells by insertional mutagenesis. The cells and tissues within the AAV-EGFP-perfused cochleae were free from inflammation and were generally intact. No pathological changes were observed in the organ of Corti, stria vascularis, or spiral ganglion cells. The long-term expression of EGFP within the cochlear tissues is consistent with data obtained from other animal models and different organ systems [9,33]. Since EGFP is known to introduce cellular toxicity, vectors expressing physiologically therapeutic proteins would achieve longer transduction periods than EGFP. Gene transfer into the inner hair cells presents numerous opportunities for auditory neuroscience. Potential applications include the localization of proteins by expression of tagged constructs, the generation of dominant-negative or antisense knockouts of endogenous proteins, the rescue of mutant phenotypes to identify disease genes, and perhaps even the treatment of auditory disorders. Advances in the molecular basis of auditory diseases have allowed the identification of a number of genetic disorders such as presbycusis, acoustic trauma, and ototoxicity. The development of gene therapy now allows us to evaluate the effects of transferring therapeutic genes into the inner ear by several different strategies. The expression of marker genes in the inner ear tissue has been demonstrated. Further studies will improve our understanding of cochlear function as well as provide

for the development of novel therapies for a wide variety of inner ear diseases. Intracochlear gene transfer using AAV vectors has been established as a viable experimental proposition. Future study will include the transfer of functioning genes *in vivo* and the development of alternative vectors. While clinical application may be some way off, it is vital that gene delivery techniques are optimized in anticipation of future need.

In conclusion, the data presented in this paper demonstrate successful gene transfer into several types of cochlear cells *in vivo* with AAV-based vectors. Interestingly, the AAV3 vector promoted inner hair cell-specific transduction. These findings are of value for further molecular studies of the cochlear inner hair cells and for gene replacement strategies to correct hereditary hearing loss due to specific monogenic mutations affecting cochlear inner hair cells.

MATERIALS AND METHODS

Construction and preparation of proviral plasmids. The AAV vector proviral plasmid pAAV2-*LacZ* harbors an *Escherichia coli* β -galactosidase expression cassette with the CMV promoter, the first intron of the human growth hormone gene, and the SV40 early polyadenylation sequence, which are flanked by inverted terminal repeats (ITRs) [34]. The *LacZ* expression cassette of pAAV2-*LacZ* was ligated to *NotI*-excised pAAV5-RNL [35] to form the proviral plasmid pAAV5-*LacZ*. The pAAV2-CAG-EGFP-WPRE construct consists of the EGFP gene under the control of the CAG promoter (the chicken β -actin promoter associated with the cytomegalovirus immediate-early enhancer) and WPRE (woodchuck hepatitis virus posttranscriptional regulatory element) flanked by ITRs. The WPRE cassette augments the stability of transgene mRNA [36] and increases EGFP expression levels, thereby ensuring long-term transgene expression. A *Bam*HI-*Xba*I fragment containing the EGFP cDNA excised from pEGFP-1 and a *Hind*III fragment containing the WPRE sequence excised from pBS II SK⁺WPRE-B11 (a gift from Dr. J. Donello) was ligated to *Xho*I linkers and cloned into an *Xho*I site of pCAGGS (a gift from Dr. J.-I. Miyazaki) to create pCAG-EGFP-WPRE. The EGFP expression cassette from pCAG-EGFP-WPRE was ligated to the *NotI*-excised pAAV2-*LacZ* and pAAV5-RNL [35] to form the proviral plasmids pAAV2-CAG-EGFP-WPRE and pAAV5-CAG-EGFP-WPRE, respectively. The AAV-helper plasmid harbors Rep and Cap. The adenovirus helper plasmid pAdeno5 (identical to pVAE2AE4-5) encodes the entire E2A and E4 regions and the VA RNA I and II genes [37]. Plasmids were purified with the Qiagen plasmid purification kits (Qiagen K.K., Tokyo, Japan).

Recombinant AAV vector production. Vectors derived from the AAV1-4, 7, and 8 pseudotypes were produced with the AAV packaging plasmid pAAV1RepCap (for AAV1) [38], pHLP19 (for AAV2), pAAV3RepCap (for AAV3) [39], pAAV4RepCap (for AAV4) [40], pAAV7RepCap (for AAV7) [41], or pAAV8RepCap (for AAV8) [41] and the AAV proviral plasmid pAAV2-*LacZ* or pAAV2-CAG-EGFP-WPRE. The plasmids pAAV5RepCap [35] and pAAV5-*LacZ*, or pAAV5-CAG-EGFP-WPRE, were used to produce vector with the AAV5 pseudotype [42]. Seven AAV serotype vectors were produced as previously described by the three-plasmid transfection adenovirus-free protocol [37]. Briefly, three days before transfection, 293 cells were plated onto a 10-tray Cell Factory (Nalge Nunc International, Rochester, NY, USA; 6×10^7 cells/10-tray). The cells were cotransfected with 650 μ g each of the proviral plasmid, the AAV vector packaging plasmid, and the adenovirus helper plasmid pAdeno5 [34] by the calcium phosphate coprecipitation method. The medium was changed following incubation for 6-8 h at 37°C. Recombinant AAV was harvested 72 h after transfection by three freeze/thaw cycles. The crude viral lysate was purified twice on a cesium chloride two-tier centrifugation

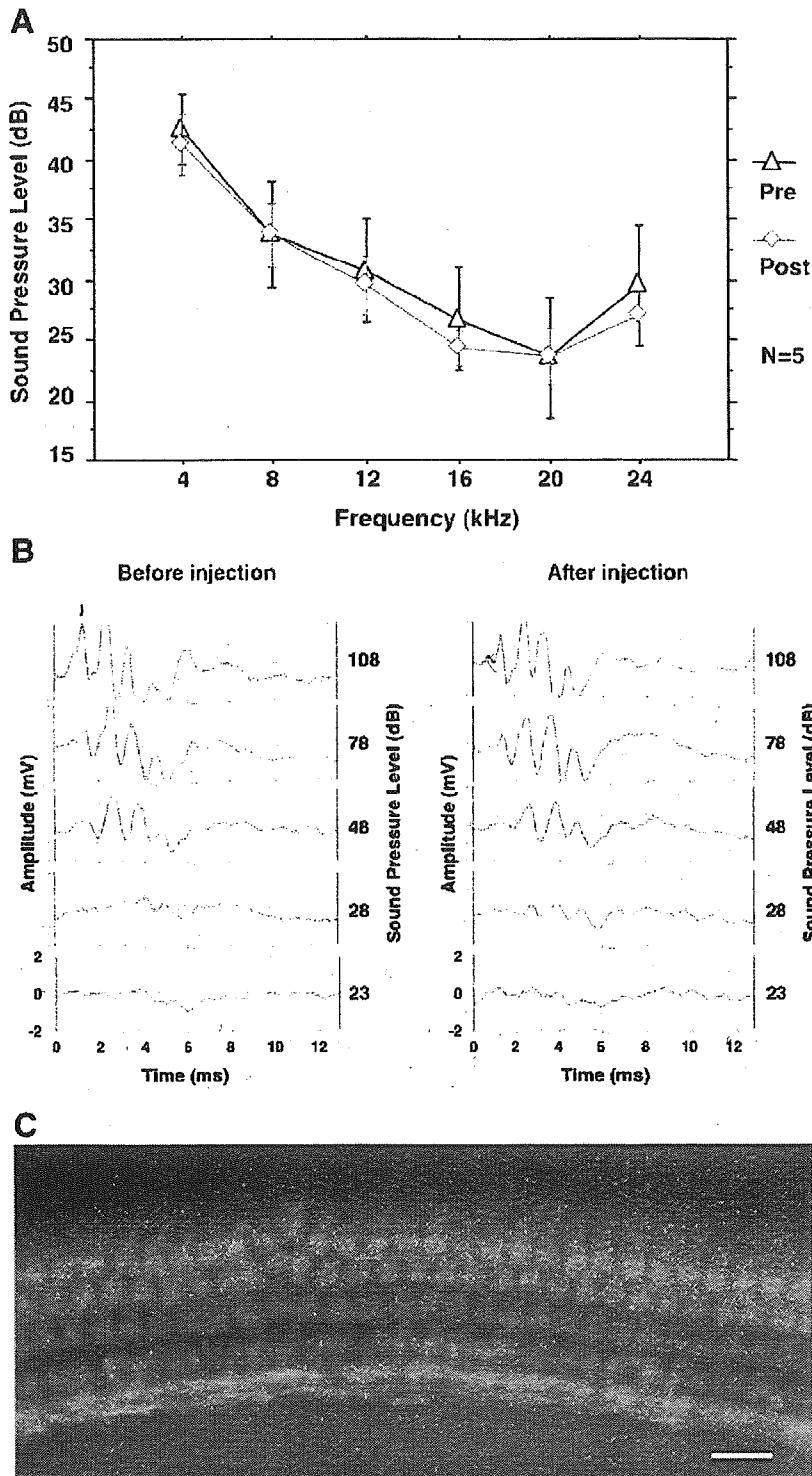


FIG. 5. (A) ABR threshold (mean \pm SD) at each frequency tested preoperatively (pre) versus postoperatively (post). (B) Example of ABR waveforms in C57BL/6J at various stimuli (16 kHz; 108 dB, 78 dB, 48 dB, 28 dB, and 23 dB). ABR were tested in the transduced ear prior to viral injection and 10 days after injection. Wave I was measured to analyze the activity of the cochlea. (C) F-actin staining showing that no outer hair cells were lost from inoculated cochleae. Original magnification 40 \times ; scale bar, 25 μ m.

gradient as described previously [24]. The viral stock was treated with DNase and titrated by quantitative real-time PCR with plasmid standards [43].

Surgical procedures and cochlear perfusions. All animal studies were performed in accordance with the guidelines issued by the committee on animal research of Jichi Medical School and approved by its ethics



committee. Sixty female C57BL/6J mice (4 weeks of age; CLEA Japan, Tokyo, Japan) and 40 male ICR mice (2 months of age; Japan SLC, Shizuoka, Japan) were utilized. The mice were initially anesthetized with ketamine (50 mg/kg) and the analgesic xylazine (5 mg/kg). A postauricular approach was used to expose the tympanic bony bulla. A small opening (2 mm) in the tympanic bulla was carefully made to allow access to the round window membrane. In the tested groups, 5 μ l AAV vector solution (5×10^{10} gc) was microinjected into the cochlea through the round window over 10 min with a glass micropipette (40 μ m in diameter) fitted on a Univentor 801 syringe pump (Serial No. 170182, High Precision Instruments, Univentor Ltd., Malta) [19]. A small plug of muscle was used to seal the cochlea and the surgical wound was closed in layers and dressed with antibiotic ointment. Five mice of each strain received control cochlear perfusions with artificial perilymph (145 mM NaCl, 2.7 mM KCl, 2 mM MgSO₄, 1.2 mM CaCl₂, 5 mM Hepes) alone. Each AAV-EGFP serotype was injected into five mice of each strain. Another 20 C57BL/6J mice were injected with the AAV3 vector to study long-term expression.

Cochlear function assessment using ABR. To assess the physiological status of experimental ears, auditory thresholds were determined with multiple frequency and intensity tone bursts by performing ABR audiometry with Tucker-Davis Technologies and Scope v3.6.9 software (Power Lab/200; ADInstruments, Castle Hill, Australia). Tone pipes were introduced into the operated ears of the anesthetized mice, and evoked potentials were recorded using needle electrodes inserted through the skin. ABR were elicited and measured 256 times at 4, 8, 12, 16, 20, and 24 kHz frequencies with tone bursts in systematic 5-dB steps. The rise/fall times for the tone bursts were 0.1 ms rise/ms flat (cosine gate). Free-field system was used as a calibration procedure. Wave I was measured to analyze the activity from the cochlea. The lowest stimulus level that yielded a detectable ABR waveform was defined as the threshold. ABR were tested in the infused ear prior to surgery and 10 days postsurgery. Data were statistically analyzed using repeated-measures analysis of variance followed by paired Student's *t* test performed with StatView 5.0 software (SAS Institute Inc., Cary, NC, USA). Values of *P* < 0.05 were considered significant.

Histology. Cochlear transgene expression patterns were determined for all AAV serotypes by visualizing EGFP expression. The animals were sacrificed 10 days after injection, and intracardiac perfusion was performed with 4% paraformaldehyde (PFA) in 0.1 M phosphate buffer, pH 7.4. The cochleae were harvested and the stapes footplates were removed. For AAV3-mediated transduction, the animals (five mice for each time point) were sacrificed 1, 2, 4, 8, or 12 weeks after inoculation. Postfixation was carried out in 4% PFA for 4 h at 4°C, and decalcification was performed in 10% EDTA for 12 days at room temperature. The cochlear half-turns were microdissected and processed and the other half-turns were prepared by cryosection (10 μ m) to detect EGFP expression by using an Olympus IX70 (Olympus Corp., Tokyo, Japan) fluorescence microscope with a standard fluorescein isothiocyanate filter set and Studio Lite software (Olympus Corp.). Cells that exhibited fluorescence were considered positive for transgene expression. The level of expression was graded by fluorescence intensity on a four-point scale (+, ++, +++, ++++) depending on the pixel/unit area count. Hair cell counts were carried out with dissected cochlea.

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Technical Report

Large-Scale Production of Recombinant Viruses by Use of a Large Culture Vessel with Active Gassing

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ABSTRACT

Adenovirus and adeno-associated virus (AAV) vectors are increasingly used for gene transduction experiments. However, to produce a sufficient amount of these vectors for *in vivo* experiments requires large-capacity tissue culture facilities, which may not be practical in limited laboratory space. We describe here a large-scale method to produce adenovirus and AAV vectors with an active gassing system that uses large culture vessels to process labor- and cost-effective infection or transfection in a closed system. Development of this system was based on the infection or transfection of 293 cells on a large scale, using a large culture vessel with a surface area of 6320 cm². A minipump was connected to the gas inlet of the large vessel, which was placed inside the incubator, so that the incubator atmosphere was circulated through the vessel. When active gassing was employed, the productivity of the adenovirus and AAV vectors significantly increased. This vector production system was achieved by improved CO₂ and air exchange and maintenance of pH in the culture medium. Viral production with active gassing is particularly promising, as it can be used with existing incubators and the large culture vessel can readily be converted for use with the active gassing system.

OVERVIEW SUMMARY

Large-scale production of recombinant viruses, using a large culture vessel with active gassing, is superior to protocols using standard tissue culture plates or flasks because of the higher capacity for cell growth. Although a previous protocol for recombinant virus production in a large culture vessel had the problem of insufficient transduction efficiency resulting from inadequate gas exchange, a method to use active gassing successfully improved productivity of recombinant viruses. Development of a vector production system on a large scale, using commercially available large culture vessels, allows us to process labor- and cost-effective manipulation in a closed system.

INTRODUCTION

ADENOVIRUS AND ADENO-ASSOCIATED VIRUS (AAV) VECTORS are highly efficient for transduction in many gene therapy studies (Okada *et al.*, 2002b, 2004; Ito *et al.*, 2003; Nomoto *et al.*, 2003; Yamaguchi *et al.*, 2003; Mochizuki *et al.*, 2004; Yoshioka *et al.*, 2004; Liu *et al.*, 2005). However, current production methods rely on the manipulation of many individual flasks and are not generally considered appropriate for scaling-up of production because it would be a time-consuming and labor-intensive process. Therefore, alternative tissue culture vessels with higher capacity for cell growth, such as a 10-tray Cell Factory (CF10; Nalge Nunc International, Rochester, NY) with a surface area of 6320 cm², could be suitable for scaling-up of

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vector production (Okada *et al.*, 2002a). This device is easy to handle and can be used for efficient cell culture on a large scale in a closed system requiring only an air filter (Berger *et al.*, 2002; Tuyaerts *et al.*, 2002). Nevertheless, a previous protocol for recombinant virus production in the CF10 had the problem of insufficient scaling-up of vector production (Liu *et al.*, 2003). In that protocol, inadequate gas exchange between the culture vessel and the incubator might have been the cause of the inefficient yield.

We consequently adapted an active gassing system to generate large numbers of recombinant viruses in the CF10. The purpose of this active gassing is to control and maintain CO₂ tension and pH in the growth medium by passing a gas mixture through the CF10. For many types of cells, pH is an important parameter for controlling cell growth. This can be achieved by gassing with CO₂ in atmospheric air in the incubator. Enhanced gas exchange in a large culture vessel should improve both viral infectivity and plasmid transfection efficiency. In combination with the previously described method of using the CF10 (Okada *et al.*, 2002a), we have now created a simple and highly efficient system of producing vector stock on a large scale. Presented here is a labor- and cost-effective method for large-scale production of adenovirus and AAV vectors with an active gassing system that uses a large culture vessel to achieve transfection or infection in a closed system.

MATERIALS AND METHODS

Cell culture with active gassing

Propagation of vectors was based on the infection or transfection of human embryonic kidney-derived 293B cells (Yamaguchi *et al.*, 2003) by using either a flask with a surface area of 225 cm² (Falcon, T-225; BD Biosciences Discovery Labware, Bedford, MA) or the CF10, as described previously (Okada *et al.*, 2002a). Cells were cultured in Dulbecco's modified Eagle's medium and nutrient mixture F12 (DMEM-F12: Invitrogen, Grand Island, NY) with 10% fetal bovine serum (FBS; Sigma-Aldrich, St. Louis, MO), penicillin (100 units/ml), and streptomycin (100 µg/ml) at 37°C in a 5% CO₂ incubator. First, cells were plated at 2.3×10^6 cells per T-225 or at 6.5×10^7 cells per CF10 to achieve a monolayer at 20 to 40% confluency when cells initially attach to the surface of the flask. The volume of medium used per flask was 40 ml per T-225 or 1120 ml per CF10. Subsequently, cells were grown to a confluency of 70–90% over the next 48 to 72 hr for adenovirus infection or plasmid transfection. An aquarium pump (NISSO, Tokyo, Japan) was used to circulate air through the CF10 with 5% CO₂ and humidity control by an incubator. The CF10 was mounted with a bacterial air filter (bacterial air vents; Pall Gelman Sciences, Ann Arbor, MI) to connect the aquarium pump. The pump was connected to the gas inlet of the CF10 and the CF10 was placed inside the incubator, so that the incubator atmosphere was circulated through the CF10. The flow through the CF10 was maintained at 500 ml/min. Culture medium was sampled periodically, and the CO₂ concentrations and pH were estimated with a blood gas analyzer (Nova PHOX; Diamond Diagnostics, Holliston, MA). Glucose levels of the culture medium were also estimated with a glucose meter (Glutest Sensor, Glutest Ace GT-1640; Sanwa Kagaku Kenkyusho, Nagoya, Japan).

Construction and propagation of adenoviral vectors

A recombinant adenoviral vector, Ad-EGFP, was constructed using an adenoviral DNA–protein complex without a transgene insert (AVC2.null) (Okada *et al.*, 1998); it carried the cytomegalovirus (CMV) promoter, cloning sites, a simian virus 40 (SV40) intron, and the SV40 polyadenylation signal. To generate Ad-EGFP encoding enhanced green fluorescent protein (EGFP), a *SpeI*–*ClaI* fragment containing the *EGFP* cDNA excised from pEGFP-1 (BD Biosciences Clontech, Palo Alto, CA) was inserted into the *XbaI* and *NspV* sites in the DNA–protein complex, AVC2.null, using the direct *in vitro* ligation technique (Okada *et al.*, 1998). The ligated DNA–protein complex was introduced into 293 cells by the calcium phosphate transfection method. Viral plaques on 293 cells were isolated, amplified, and titrated by standard techniques. To amplify the vector in 293 cells, half the medium in the tissue culture flasks was exchanged with fresh DMEM-F12 containing 10% FBS 1 hr before infection. Cells were infected with the virus at 10 multiplicities of infection (MOI) per cell. Cells were incubated to reach full cytopathic effect, and crude viral lysate was purified by two rounds of CsCl two-tier centrifugation. The average number of plaque-forming units (PFU) was assessed on the basis of the 50% tissue culture infective dose. The number of vector particles was estimated by dot-blot hybridization of DNase I-treated stocks with plasmid standards.

Construction and propagation of AAV vectors

AAV1-EGFP, a recombinant AAV type 1 expressing the *EGFP* gene under the control of the CAG promoter (modified chicken β -actin promoter) with the CMV-IE enhancer, was generated by the following procedure. A *Bam*HI–*XbaI* fragment containing EGFP cDNA excised from pEGFP-1 and a *Hind*III fragment containing the woodchuck hepatitis virus posttranscriptional regulatory element (WPRE) sequence excised from pBluescript II SK(+)/WPRE-B11 (a gift from T. Hope, University of Illinois at Chicago, Chicago, IL) was cloned into an *XhoI* site of pCAGGS (a gift from J.-i. Miyazaki, Osaka University Graduate School of Medicine, Japan) to create pCAG-EGFP-WPRE, using an *XhoI* linker. The EGFP expression cassette in pCAG-EGFP-WPRE was ligated to *NotI*-excised pAAV-LacZ to form the proviral vector plasmid pAAV2-CAG-EGFP-WPRE. AAV viral stocks were prepared according to a previously described protocol (Okada *et al.*, 2002a) with minor modifications. Half the medium in tissue culture flasks was exchanged with fresh DMEM-F12 containing 10% FBS 1 hr before plasmid transfection. Subsequently, cells were cotransfected with 23 µg (per T-225) or 650 µg (per CF10) of each of the following plasmids: a proviral vector plasmid, AAV-1 chimeric helper plasmid p1RepCap (Mochizuki *et al.*, 2004), and adenoviral helper plasmid pAdeno, by a calcium phosphate coprecipitation method. Each of the vector and helper plasmids was added to 4 ml (per T-225) or 112 ml (per CF10) of 300 mM CaCl₂. This solution was gently added to an equal volume of 2× HEPES-buffered saline (HBS: 290 mM NaCl, 50 mM HEPES buffer, 1.5 mM Na₂HPO₄, pH 7.0) and immediately mixed by gentle inversion three times to form a uniform solution. This solution was immediately mixed with fresh DMEM-F12 containing 10% FBS outside the flasks to produce a homogeneous plasmid solution mixture. Subsequently, medium in the

culture flasks was entirely replaced with this plasmid solution mixture. At the end of incubation for 6 hr, the plasmid solution mixture in the culture flasks was replaced with pre-warmed fresh DMEM-F12 containing 2% FBS. Cell suspensions were collected 72 hr after transfection and centrifuged at $300 \times g$ for 10 min. Each cell pellet was resuspended in 2 ml (per T-225) or 56 ml (per CF10) of Tris-buffered saline (TBS: 100 mM Tris-HCl [pH 8.0], 150 mM NaCl). Recombinant AAV was harvested by three cycles of freeze-thawing of each resuspended pellet. Crude viral lysate was then purified twice by passage through a CsCl two-tier centrifugation gradient, as described previously (Okada *et al.*, 2002b). The viral stock was titrated by dot-blot hybridization of DNase I-treated stocks with plasmid standards. To confirm transgene expression with the propagated vector *in vivo*, 5-week-old male Sprague-Dawley rats were injected via the anterior tibial muscle with AAV1-EGFP (1×10^{11} genome copies per rat). Fifteen weeks after injection, the rats were sacrificed and expression was confirmed by fluorescence microscopy.

Statistical analysis

Statistical significance was determined on the basis of an unpaired, two-tailed *p* value and Student *t* test, and a *p* value less than 0.05 was considered significant.

RESULTS

Improved gas exchange and maintenance of pH in medium after recombinant adenovirus infection

Propagation of vectors was based on infection or transfection of 293 cells on a large scale. A minipump was connected to the gas inlet of the CF10 and placed inside the incubator, so that the atmosphere in the incubator, containing 5% CO₂, was circulated through the CF10. The gas flow for circulation through the CF10 was maintained at 500 ml/min. An appropriate gas flow rate was important to give a uniform distribution of the gas in the individual trays of the CF10. A flow less than 200 ml/min gave uneven distribution of the gas, and significantly influenced cell growth. Gas flow that was too high also disturbed the uniformity of cell density. Appropriate cell density and uniform distribution of cells are critical to achieve successful gene transduction. Application of active gassing significantly increased cell growth in the CF10 (Table 1). CO₂ concentrations in the media stayed at their initial levels when using either a T-225 or CF10 with active

gassing (Fig. 1A). In contrast, the CO₂ concentration inside the CF10 increased subsequent to adenovirus infection in the absence of active gassing. The pH of culture medium in the CF10 with active gassing was close to that in the T-225 and significantly higher than that in the CF10 without active gassing (Fig. 1B).

Monitoring of cell numbers and time point for harvest

The glucose level was monitored as an index for tracing cell growth and cytopathic effect in the CF10 to avoid the necessity for a specialized microscope to monitor cells in the large culture vessel. The glucose level decreased with increasing cell confluency and progression of cytopathic effect (CPE) (Fig. 2). When 80% CPE was reached, the glucose level was reduced to about 50 mg/ml. When glucose levels were less than 25%, the cells showed full CPE and this was regarded as the appropriate time for harvest.

Improved adenovirus vector production in a large culture vessel with active gassing

We estimated the adenovirus vector yield propagated by using 28 T-225 flasks with a surface area of 225 cm², a CF10 with a surface area of 6320 cm², or a CF10 in the presence of active gassing. When active gassing was used with the CF10, the productivity of the adenovirus vectors was dramatically increased, by 53.4 times compared with that in the CF10 without active gassing (Fig. 3). The vector yield per producer cell in the CF10 was also significantly improved in the presence of active gassing (Table 1). The PFU-to-particle ratios for vectors produced in the T-225, CF10, and CF10 with active gassing were 1:7, 1:15, and 1:10, respectively.

Efficient AAV vector production in a large culture vessel with active gassing

Enhanced gas exchange in a large culture vessel should also improve vector production through plasmid transfection. AAV vectors were produced in a large vessel by a three-plasmid transfection adenovirus-free protocol (Okada *et al.*, 2002b). Three days after plasmid transfection, the CO₂ concentrations in medium from the CF10 in the presence of active gassing were significantly less than those without active gassing (Table 2). The pH of the culture medium in the CF10 with active gassing was also improved. The CF-10 with active gassing was compatible with the three-plasmid transfection protocol for recombinant AAV production. When we used active gassing, the vec-

TABLE 1. INCREASED CELL GROWTH AND VECTOR YIELD WITH ACTIVE CO₂ AND AIR EXCHANGE^a

| Flask | Number of cells harvested | Vector yield per cell (PFU/cell) |
|---------------------------|---|----------------------------------|
| 225-cm ² flask | $(1.4 \pm 0.2) \times 10^9$ (per 28 flasks) | 7.9×10^3 |
| CF10 | $(4.9 \pm 1.6) \times 10^8$ | 4.1×10^2 |
| CF10 + AG | $(1.3 \pm 0.3) \times 10^9$ | 8.2×10^3 |

^aAt the time of cell harvest after adenovirus infection, cell growth and vector yield per cell in a CF10 with a surface area of 6320 cm² in the presence or absence of active gassing (AG) were compared with that in 28 flasks with a surface area of 225 cm² each.