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# Self-complementary adeno-associated viral vectors for gene therapy of hemophilia B: progress and challenges

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### Abstract

Therapies currently used for hemophilia involve injection of protein concentrates that are expensive, invasive and associated with side effects such as development of neutralizing antibodies (inhibitors) that diminish therapeutic efficacy. Gene transfer is an attractive alternative to circumvent these issues. However, until now, clinical trials using gene therapy to treat hemophilia have failed to demonstrate sustained efficacy, although a vector based on a self-complementary adeno-associated virus has recently shown promise. This article will briefly outline a novel gene-transfer approach using self-complementary adeno-associated viral vectors using hemophilia B as a target disorder. This approach is currently being evaluated in the clinic. We will provide an overview of the development of self-complementary adeno-associated virus vectors as well as preclinical and clinical data with this vector system.

#### Keywords

factor IX; hemophilia B; inhibitors; inverted terminal repeats; neutralizing antibodies; recombinant adeno-associated virus; self-complementary adeno-associated virus; single-stranded adeno-associated virus; terminal resolution sites

The term 'hemophilia' is used to refer to a group of disorders that are characterized by impairment of blood coagulation. Each of these disorders is caused by mutations in a single gene, resulting in deficiency of a single clotting factor in the coagulation cascade. The most common manifestation of these disorders is spontaneous hemorrhage in the absence of injury, predominantly in weight-bearing joints, resulting in painful chronic arthropathy. Spontaneous bleeding into a closed space such as the skull can result in death.

Hemophilia B is an X-linked recessive condition caused by deficiency of factor IX (FIX), and is the second most common of the hemophilias, affecting approximately one in 25,000 males. A number of mutations can occur in the gene encoding FIX, each compromising

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function of the clotting factor to a different degree. This disorder therefore occurs along a spectrum of severity that varies inversely with the levels of functional FIX in the plasma. Generally, individuals with <1% function of clotting factor are classified as severe hemophiliacs, those with 1-5% function as moderate hemophiliacs and those with >5% function as mild hemophiliacs [1].

Current therapy of hemophilia B involves administration of clotting factor concentrates delivered as injections, which have the disadvantages of being invasive and inconvenient to the patient, besides being prohibitively expensive for many patients around the world. Another concern with this mode of treatment is the development of circulating antibodies, or 'inhibitors' that neutralize the clotting factor administered as concentrate, thus rendering these ineffective. Acquisition of blood-borne diseases such as HIV/AIDS or hepatitis B through contaminated blood products was a large problem in the past but is less of a problem now, due to the availability of recombinant factors and more effective screening techniques on blood products [2].

The hemophilias are ideal candidates for gene therapy as they are caused by diminished function of a single protein, which is in turn caused by alteration (by mutation or deletion) of a single gene; restoring a functional copy of the affected gene could thereby completely ameliorate the clinical manifestations of the disease. It has also been shown that restoration of circulating clotting factor to 1-2% of its physiological levels greatly improves the bleeding diathesis, and so the therapeutic goals are modest. Clinical approaches for gene and cell therapy of the related disorder hemophilia A (summarized in Table 1) have so far not yielded sustained correction of the bleeding diathesis [3,4], but have on one occasion been associated with an inflammatory response triggered by the vector [5]. These studies have revealed a need for a vector that is safer and more efficacious than the vectors that have been used thus far.

#### Adeno-associated viral vectors for gene therapy of hemophilia B

Adeno-associated viruses are single-stranded DNA viruses that infect human cells but are naturally replication deficient, and are not associated with human disease. Recombinant vectors derived from these viruses consist of two 145 nucleotide inverted terminal repeats flanking an expression cassette encoding a therapeutic transgene, with deletion of all the viral open reading frames (Figure 1) [6,7]. Recombinant adeno-associated viral (rAAV) vectors are less immunogenic than several other viral vectors [8], but an immune response may still occur against the vector capsid proteins as well as the transgene-encoded protein. rAAV vectors have demonstrated promise due to their efficacy as a vector for gene transfer in nondividing tissues *in vivo*, such as neurons [9,10], photoreceptor cells [11,12], hepatocytes [13,14] and muscle cells [15,16]. In clinical settings, rAAV vectors have been shown to be efficacious in persistent transgene expression upon transduction of skeletal muscle [17,18] or retinal pigment epithelium [19,20].

The rAAV genome is maintained in a predominantly episomal state within transduced cells [21], although a small degree of genomic integration has been observed in some studies [22]. Administration of rAAV vector in a rodent model of mucopolysaccharoidosis VII resulted in genomic vector integration and hepatocellular carcinoma development, raising concerns that rAAV vectors may not be completely free of genotoxic effects [23]. Nevertheless, other studies have failed to demonstrate genotoxic effects following rAAV vector transfer [24–26]. These studies raise hopes that rAAV vectors may be safer than vector systems such as oncoretroviruses that have been associated with genotoxicity and tumorigenesis in clinical trials [27,28].

Two clinical trials have attempted to use rAAV vectors in hemophilia B patients (summarized in Table 1). In the first trial, patients were administered rAAV to express human factor IX (hFIX) by intramuscular injection. Although all patients showed effective gene transfer at the site of injection, the clinical response was modest, with only one patient (out of eight) showing an increase in hFIX levels above 1% [29-31]. The second trial involved liver-targeted delivery of rAAV by direct infusion of vector into the hepatic artery, for several reasons: the liver is the natural site of hFIX synthesis; liver-directed expression (by intraportal administration of vector) leads to 1.5–4-fold higher levels of hFIX in the circulation when compared with intramuscular or tail vein administration in mice [32]; and expression from the liver has been shown to mediate immune tolerance, thereby reducing the risk of developing neutralizing antibodies [32,33]. Therapeutic levels of hFIX were obtained in two patients treated at the highest dose  $(2 \times 10^{12} \text{ vector genomes } [vg]/kg)$ , although this effect was transient, lasting only 2-4 weeks after vector administration. In one of these patients, the decline in hFIX levels was associated with a transient rise in liver transaminases, which may have been caused by a cellular immune response to the adenoassociated virus (AAV)2 capsid that destroyed the transduced hepatocytes, thus abrogating clinical benefits. Patients administered low or intermediate doses of vector ( $8 \times 10^{10}$  vg/kg or  $4 \times 10^{11}$  vg/kg) did not develop therapeutic levels of hFIX, although one patient treated at the intermediate dose developed an immune response targeting transduced hepatocytes [34].

High vector doses necessary for a therapeutic response may therefore be associated with liver toxicity due to an immune response to the vector capsid, which appears to be dose dependent, occurring at higher vector doses than are required for a clinical response, and abrogating the possibility of stable expression of the transgene product at therapeutic levels. High vector doses are shown to be associated with nonspecific biodistribution of vector in animal models [32,35]. Production of clinical-grade vector at 10<sup>14</sup> vg per patient can also be cost-prohibitive [36]. It may be possible to abrogate the immune response to vector by using immunosuppressive therapy [37], but this has not yet been demonstrated in human subjects. An attractive alternative would therefore be the development of more potent vectors that can stably express the transgene at a lower vector dose without triggering an immune response.

#### Self-complementary AAV vectors

The AAV genome is single stranded, but upon transduction of target cells the viral genome must first be converted into a transcriptionally active double-stranded form, which is a crucial rate-limiting step in transgene expression from an rAAV vector [38]. This step may be facilitated by coadministration of adenovirus [39], expression of the adenoviral genes *E1* and *E4* [40], or coadministration of DNA-damaging agents [41]. However, these methods are associated with toxicity to the target cells that are to be transduced.

A different strategy towards overcoming this limiting step was the development of selfcomplementary AAV (scAAV) vectors (Figure 1). Here, the vector genome is designed as a single-stranded inverted repeat, which folds back upon itself to form a double-stranded genome when entering into infected cells. A genome <2.5 kilobases (kb; approximately half the size of the wild-type AAV genome) can therefore be packaged as a dimer, with the two inverted repeats pairing along their length, closed covalently by a hairpin at the terminal repeat. There is therefore no need for complementary strand synthesis, and this rate-limiting step is bypassed [42]. Dimerization of the genome in this manner can be stabilized by mutation or deletion of one of the two terminal repeat [43]), which prevents cleavage by AAV Rep proteins to form monomers [44]. The replication fork initiates at the wild-type trs and proceeds through the genome and through the mutant trs, which is unable to facilitate resolution, causing the replication fork to proceed back across the genome where it

terminates at the wild-type trs. The resultant self-complementary molecule is thus flanked by wild-type trs and has a mutant trs in the middle, and dimerizes along its length when packaged into the AAV [45].

Upon transduction of target cells, the scAAV genome exists either as circular genomes or concatemers, the former being much more effective in transgene expression [46]. scAAV vector genomes are more stable and more prone to circularization upon transduction of *in vivo* tissues than single-stranded AAV (ssAAV) vector [44]. The ends of the scAAV vector resemble a double-stranded DNA break, and circularization of the scAAV genome depends on the action of a number of double-stranded break repair proteins, such as Mre11, Nbs1, BLM, WRN, DNA-PK(CS) and ATM, the latter being required for circularizing of the scAAV genome in nondividing cells *in vivo*. This suggests that DNA repair via both homologous recombination and nonhomologous end repair pathways plays a role in scAAV genome circularization in target cells [47,48].

Unlike ssAAV, transduction by scAAV is not affected by coinfection with adenovirus or treatment with hydroxyurea or proteosome inhibitors, as the limiting step of second-strand synthesis is circumvented [49,50]. When used to transduce cells in culture, scAAV vectors were shown to be 5–140-fold more effective than the corresponding ssAAV [49]. A number of cell types have been effectively transduced by scAAV vectors in a therapeutically relevant context. Mesenchymal stem cells transduced with scAAV showed gene expression for up to 3 months when transplanted into rat brain [51]. scAAV vectors have been used to deliver CD40 ligand to lung carcinoma cells, and intratumoral administration of this vector can inhibit tumor growth by CD40 ligand-mediated activation of the immune response [52]. A therapeutic transgene can be targeted to hepatoblastoma cells and hepatocellular carcinoma cells by scAAV vectors, effecting apoptosis of the target cells *in vitro* [53]. Immature human dendritic cells can be transduced by scAAV without detectable change to their surface marker profiles, their functional properties or their ability to differentiate into mature dendritic cells [54].

The versatility of scAAV vectors is also demonstrated by their ability to transduce several different kinds of tissues in an in vivo or ex vivo setting. scAAV and ssAAV vectors were used to express erythropoietin upon injection into mouse muscle in vivo; the scAAV vectors expressed higher levels of erythropoietin, also leading to an increased hematocrit level due to expansion of red blood cell numbers by the transgenically expressed erythropoietin. An scAAV vector expressing green fluorescent protein of a liver-specific promoter transduced hepatocytes stably for 3 months in vivo. Transduction of cells in the CNS was also more widespread using scAAV as compared with ssAAV [45]. Dispersion of scAAV vector across various cell types in the CNS is accomplished by either intravenous or intracisternal administration [55]. scAAV vectors have been used to successfully transduce human and murine pancreatic islets without affecting their function [56]. scAAV vectors transduced the trabecular meshwork of the eye in rats and monkeys effectively with stable transgene expression lasting for months (in rats) to years (in monkeys) [57]. Transduction of an intact trabecular meshwork in the human eye [58] or the retinal pigment epithelium in the mouse [59,60] is more effective with scAAV than ssAAV vectors. scAAV vectors effectively transduce retinal ganglion cells of the primate retina [61], and can be used to deliver a therapeutic transgene into mouse retinal ganglion cells in vivo [62]. Transduction of the mouse myocardium is accomplished effectively with either vector, but earlier expression of transgene is observed with scAAV compared with ssAAV [63]. Percutaneous transendocardial delivery of scAAV vector results in effective transduction of approximately 60% of the canine myocardium [64,65]. Transduction of spinal cord motor neurons in rats was accomplished by retrograde axonal transport of scAAV vector following either intramuscular or intranerve injection [66]. scAAV vectors also enabled successful

transduction of the thymus in murine and primate models [67], and the joint spaces of rabbit models of arthritis [68], as well as chondrocytes and synoviocytes from equine joint tissue [69].

In addition to an increased level of transgene expression by scAAV vectors as compared with ssAAV, subtle changes in patterns of expression have also been noted. An equivalent dose of scAAV was shown to transduce up to 90% of hepatocytes *in vivo*, compared with 2% of hepatocytes transduced by ssAAV [44]. At a slightly lower dose of vector, scAAV transduced 25–50% of mouse hepatocytes, compared with less than 5% by ssAAV [45]. Expression of hFIX from an ssAAV vector led to intense expression of transgene from a small proportion (5%) of hepatocytes, while expression from an scAAV vector led to more homogenous but moderate expression from a large population of hepatocytes. Expression of the scAAV vector thus paralleled more closely the natural expression of hFIX, which is normally expressed in a homogenous manner throughout the liver [70].

#### Limitations of scAAV & attempts to optimize scAAV-mediated gene transfer

Several natural serotypes of AAV have been identified [71,72], primarily based on the surface capsid protein, which determines viral tropism and transduction efficiency, as well as immunogenicity in a human host. The most common serotype used thus far has been AAV2, which effectively transduces a variety of cell types. As humans are the natural host for AAV2, seropositivity to AAV2 is highly prevalent in the general population, which may diminish the utility of AAV2 as a therapeutic vector; results obtained in animal studies using AAV2 may not be entirely predictive of clinical responses in human subjects [73]. Other serotypes such as AAV8, which was first noted for its efficacy in tranducing liver cells [71], are lower in prevalence in the general population. The AAV8 capsid protein is less immunogenic than AAV2 [74] and is also uncoated earlier following transduction [75], further reducing the likelihood of development of a neutralizing immune response. In studies performed on nonhuman primates, the AAV8 vector was found to be more effective than AAV2 in liver cell transduction [76]. Pre-existing immunity to AAV8, though rare, does considerably diminish efficacy of gene transfer in these primate models, redirecting the vector to the spleen and away from the liver [77,78]. AAV8 also mediates effective gene transfer to the thymus [67]. Other AAV serotypes have been shown to mediate effective gene transfer to other tissues, such as AAV6 for cardiac cells [64,79], AAV1 for hematopoietic stem cells [26,80] and AAV3 for liver tumor cells [53]. Nevertheless, the AAV2 capsid is the most effective in transduction of the majority of cell types tested thus far [52,79,81,82]. Pre-existing immunity to specific AAV serotypes, as well as the need for these serotypes to transduce a target cell type, may therefore limit the potential for AAVmediated gene transfer.

A significant limitation with the scAAV vector system is its small packaging capacity; as the AAV genome is 4.7 kb in size, the maximum packaging capacity of the scAAV vector is approximately half of this length, or about 2.3 kb. The packaging capacity can be extended to 3.3 kb genomes, but the proportion of single-stranded genomes increases linearly with genome length. Genomes larger than 3.5 kb are packaged almost solely as single-stranded forms. AAV Rep proteins encoded by helper plasmids during vector production have been shown to influence the generation of single-stranded genomes, and the use of helper plasmids that resulted in lower levels of Rep enabled the generation of scAAV vectors that were capable of packaging up to 6.6 kb genomes in their double-stranded forms [83]. The use of these modified helper plasmids may therefore enable packaging of larger therapeutic transgenes into scAAV vectors. Another study demonstrated that packaging capacity may be capsid dependent, with the AAV5 capsid enabling packaging of up to 8.9 kb of single-

stranded genomes [84], although other studies find no such variation between serotypes [85].

The small size of the scAAV genome does not preclude delivery of siRNA, and therefore scAAV vectors may be ideal for this mode of therapy. The *PDHA1* gene was successfully targeted by scAAV-delivered siRNAs *in vitro*, leading to a decline in enzyme activity levels [86,87]. *PDHA1* siRNAs have also been delivered directly to rat striatum, resulting in decreased levels and activity of target protein in the striatum [88]. The *MDR1* gene, which confers a drug resistant phenotype to tumor cells, has been targeted *in vitro* with scAAV-delivered siRNA [89]. scAAV vectors have also been used to deliver an anti-hepatitis C virus miRNA cluster that showed efficacy both *in vitro* and *in vivo* in liver cells [90].

A gender-specific imbalance in therapeutic efficacy following scAAV vector administration has been noted in some animal models, with benefits more pronounced in males than in females. Administration of an scAAV–*LP1–apo3* vector in a mouse model of atherosclerosis showed retardation of aortic atherosclerotic lesions of up to 58% in male animals, but only 33% in female animals [91]. scAAV-mediated hFIX expression was also shown to be higher in male mice than in female mice [70]. Pretreatment of the mice with bortezomib before scAAV transfer or exposure of the mice to adenovirus 10–20 weeks after scAAV transfer were both successful in raising transgene expression in female animals [92]. The cause of this gender imbalance in these studies, both of which involved liver-mediated transgene expression, is unclear.

Transgene expression and transduction via AAV vectors has been shown to be inhibited by the cellular EGF receptor–protein tyrosine kinase signaling pathway [93]. The EGF receptor–protein tyrosine kinase pathway was shown to directly phosphorylate surface tyrosine residues on AAV2 capsid proteins, which inhibited transduction in a manner that was independent of cell entry and second-strand DNA synthesis [94]. The authors hypothesized that tyrosine phosphorylation of the AAV2 capsid proteins led to ubiquitination and proteosomal degradation of the AAV2 vector. They therefore substituted the surface tyrosine residues on the AAV2 capsid, and demonstrated decreased ubiquitination of the capsid, which was associated with greater transduction, more efficient intracellular trafficking of the vector and higher levels of transgenic protein [95]. These tyrosine-substituted AAV2 capsids were shown to effectively transduce mouse retina, skeletal muscle and liver *in vivo* [96–99].

The cellular factor FKBP52 has also been shown to phosphorylate AAV capsids and mark them for degradation. FKBP52 is itself dephosphorylated and inactivated by the cellular Tcell protein tyrosine phosphatase (TC-PTP). scAAV–TC-PTP vectors, when coadministered with ssAAV vectors, therefore act as helper viruses by preventing degradation of the ssAAV capsid [80,100]. FKBP52 can also be dephosphorylated and inactivated by protein phosphatase 5 (PP5). Coadministration of scAAV–PP5 and scAAV–TC-PTP with an ssAAV2 vector increased the transduction efficiency of the ssAAV2 vector by several fold following tail vein injection in mice, without having any adverse effects on transduced hepatocytes [101,102]. As ssAAV vectors have a higher packaging capacity than scAAV vectors, and as there are larger transgenes for which ssAAV cannot be substituted for scAAV, the aforementioned strategy is a novel use of scAAV vectors to deliver cellular factors that can increase transgene expression from the ssAAV vector.

Adeno-associated viral vectors have been shown to trigger the immune system by directly modulating cellular signaling pathways. AAV transduction was shown to activate both the canonical and noncanonical NF- $\kappa$ B pathways, leading to the expression of proinflammatory molecules. This effect could be blocked by the use of inhibitors to the NF- $\kappa$ B pathway

[103]. In a comparative study, scAAV vectors were found to have a more profound stimulatory effect on local innate immune responses than ssAAV vectors. Several factors such as Toll-like receptor (TLR)-9, TLR-2, MyD88, TNF- $\alpha$ , IFN- $\alpha$  and IFN- $\beta$  were upregulated. scAAV vectors also increased infiltration of neutrophils, macrophages and

natural killer cells into the liver. These effects were found to be mediated by TLR-9 signaling that was stimulated by scAAV, and could be partially blocked by TLR-9 inhibition [104]. It will be important to identify the molecular mechanisms by which scAAV modulates the immune system and to design therapies that can circumvent these effects.

Owing to the small packaging capacity of the AAV genome, it is important to determine the ideal regulatory elements that could optimally express a transgene from an scAAV vector. Among 14 different small promoters that were tested, the liver-specific transthyretin (*TTR*) enhancer promoter produced optimal hFIX expression upon vector administration directly into the portal vein in C57BL/6 mice. Addition of the minute virus of mice (*MVM*) small intron and the bovine growth hormone polyA (*BGHpA*) was also shown to facilitate transgene expression. Codon optimization of hFIX led to a fourfold increase in expression *in vivo* compared with endogenous hFIX, with the increase in expression shown to be independent of vector copy number. Interestingly, the scAAV vector was distributed evenly in liver cells, thereby mimicking endogenous hFIX synthesis, whereas an ssAAV vector administered in a parallel study was localized to a small subsection of liver cells [70]. These principles (using smaller promoters that may also be tissue-specific, gene-regulatory elements such as introns and codon-optimized coding sequences) can be established for a number of gene transfer conditions [105].

#### Manufacture of clinical-grade scAAV vector for therapy

Several protocols have been described for the generation of AAV vector on a large scale for gene therapy [106–108]. One such report describes the generation of good manufacturing practice-grade scAAV vector at a large scale for hemophilia B clinical trials [109]. A calcium phosphate-based method was used to transiently transfect two plasmids (one expressing the transgene of interest in the scAAV backbone along with adenoviral helper sequences and the other expressing AAV helper sequences) into 293T cells expanded into ten-stack culture chambers. After 40–48 h the cells were harvested, lysed, benzonase-treated, filtered and subjected to size-exclusion and ion-exchange chromatography. The material obtained was pooled, subjected to another round of size-exclusion chromatography and eluted in a lower volume. Eluates from several such preparations were pooled to obtain the final vialed product. A total of 432 ten-stack culture chambers yielded  $2 \times 10^{15}$  vector genomes.

Transfection efficiency was assessed by Western blotting, and viral titer assessed by quantitative PCR (qPCR) at each stage of the manufacturing process. Standard assays were used to test for sterility, presence of DNA contaminants, mycoplasma, adventitious viruses, retroviruses, bovine viruses and porcine viruses. sodium dodecyl sulfate polyacrylamide gel electrophoresis confirmed the presence of the three viral capsid proteins, normal and alkaline agarose gel electrophoresis confirmed the presence of AAV-like particles. Binchoninic acid and UV were used to measure the total protein and the capsid protein, respectively, and qPCR and UV to measure viral titer. An replication-competent AAV assay was used to test for recombinants with replicative potential. The product was also tested for potency at various doses in mice, and transgenic protein (hFIX) levels assessed at various time points by ELISA. The final vialed product was tested for sterility, pH and general appearance, titer, general safety and seal integrity, and endotoxins.

Of importance, it was found that qPCR underestimates the viral titer by as much as tenfold as compared with UV due to the fact that re-annealing of genome templates interferes with primer binding, whereas dot-blot hybridization yields titers comparable to UV [109]. This is consistent with a report showing that restriction endonuclease treatment of the viral genome to cleave the hairpins and facilitate primer binding yields higher titers consistent with UV [110]. An underestimation of viral titer used in clinical trials may influence the interpretation of the efficacy of scAAV-based therapy. Also of importance is the fact that the final product contained a proportion of empty capsids along with viral vector; the A260 nm/ A280 nm reading suggests 5.3 capsid equivalents per vg, while binchoninic acid suggests a twofold increase in capsid protein [109]. The clinical relevance of this is significant, as capsid proteins may trigger a T-cell-based immune response against the vector and host cells transduced by the vector [34,73].

#### scAAV vectors in preclinical & clinical studies for hemophilia B

A mini-hFIX cassette was developed to facilitate its packaging into scAAV vectors. The hFIX transgene was expressed off the synthetic liver promoter 1 (LP1), which consists of liver-specific elements from the human apolipoprotein E/C-I gene locus control region and the human  $\alpha$ 1-antitrypsin promoter. The LP1 promoter has the advantages of being compact and highly liver-specific. Also included in this construct are an SV40-modified small T antigen intron and an SV40 late polyA. The hFIX cDNA is a codon-optimized 1.6 kb fragment. This construct was inserted into a modified AAV2 back-bone with an intact 5' trs and a deleted 3' trs. Relatively low doses of vector could deliver therapeutic levels of hFIX in FIX-knockout mice, leading to correction of the bleeding diathesis. The scAAV vector successfully transduced up to 90% of hepatocytes in 6 weeks, and had a several fold increased expression of hFIX than the corresponding ssAAV vector. Analysis of the livers of these mice showed persistence of AAV genomes as concatamers or monomeric circles. Although scAAV genomes were detected in all tissues 6 weeks after administration, transgene expression was only detected from the liver. Upon administration to nonhuman primates, scAAV vectors were shown to express therapeutic levels of hFIX that were stably maintained when administered at the low dose of  $4 \times 10^{11}$  vg/kg [111].

Following these promising results, the scAAV–LP1–hFIX vector was further tested for therapeutic efficacy in nonhuman primates. Upon pseudotyping with the hepatotropic AAV8 capsid, administration of the vector via peripheral vein led to stable therapeutic levels of hFIX over a period of 9 months. These levels were similar to those obtained upon administration via portal vein; an important point as patients with a bleeding diathesis may not tolerate portal vein injection, and the efficacy of peripheral vein administration of AAV8-coated vector may significantly improve therapeutic ease and safety. hFIX protein expressed off the transgene was found to be post-transcriptionally processed appropriately and was biologically active *in vivo*. In animals with pre-existing immunity to AAV8, showing that pre-existing immunity to different strains of AAV can be circumvented by switching between different capsids [14].

A follow-up on these animals showed that even the highest dose of vector  $(2 \times 10^{12} \text{ vg/kg})$  produced no abnormalities detected by clinical or biochemical tests, while leading to the highest levels of hFIX expression with nearly 100% of hepatocytes transduced. A lower dose of vector  $(2 \times 10^{11} \text{ vg/kg})$  led to expression of hFIX at >10% of physiological levels, which was maintained for 5 years of observation, although some decline in transgene copy number and the proportion of transduced hepatocytes was observed, and all animals did develop antibodies to vector capsid proteins. Clinical, biochemical and histological assays as

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well as ultrasound analysis during this period revealed no toxicity, demonstrating the safety and efficacy of this vector system [35].

A clinical trial is underway to evaluate efficacy of scAAV2/8–LP1–hFIXco in hemophilia B patients, and preliminary results have been reported at the 2010 Annual Meeting of the American Society of Hematology [112] and the 2011 Annual Meeting of the American Society of Gene and Cell Therapy [113]. Encouragingly, patients treated with the lowest vector dose ( $2 \times 10^{10}$  vg/kg) showed sustained low-level hFIX expression (1–2%) with no side effects or immune response against the vector. At a slightly higher dose of vector, hFIX expression was again detected (2–4%) but, in addition, an expansion of capsid-specific T cells was detected; however, this was not associated with elevation in liver enzymes or a decline in hFIX expression. Further data from this trial will shed more light on the efficacy of the scAAV vector system.

#### Expert commentary

The development of scAAV vectors has greatly improved the prospects of successful gene transfer in the clinic. However, this approach is limited to a few diseases where the therapeutic transgene is approximately 1.5 kb, and thus small enough to be packaged into scAAV vectors. Possible target disorders include Gaucher's disease [114], Krabbe's disease, phenylketonurea [115] and ornithine transcarbamylase deficiency [116]. Promising results have been obtained for hemophilia B, a disease where liver-directed expression of hFIX to 1–2% of physiological levels can result in a marked improvement in clinical outcome.

#### Five-year view

Within the next few years, we will have mature data from the scAAV clinical trial for hemophilia B. This will inform us whether scAAV vectors are associated with improved clinical outcomes when compared with ssAAV vectors, as preliminary data from the first patients seem to suggest, although there are other novel aspects of the vector design used in the current clinical trial. This clinical trial will also shed light on whether higher vector doses are associated with significantly higher transgene expression, and whether this is associated with significant toxicity. The knowledge gained from these studies will directly impact therapy for other monogenic disorders.

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#### Key issues

- Adeno-associated virus (AAV)-based vectors show promise as these viruses are nonpathogenic, efficacious in transduction of nondividing tissues *in vivo* and lead to long-term expression of transgenes.
- AAV-based vectors have thus far shown the greatest promise for gene therapy of hemophilia B, but progress has been hampered by the need for high vector doses for efficient gene transfer and to compensate for a concomitant immune response to the vector capsid.
- Self-complementary AAV (scAAV) vectors in particular are characterized by greater expression of transgene, more efficient transduction of target cells and more favorable patterns of expression.
- These vectors can be encapsidated within capsid proteins from different serotypes in order to target them to particular tissues such as the liver, or to overcome pre-existing immunity to other serotypes.
- A limitation of the AAV-based vector system is its small packaging capacity (which is further halved in scAAV vectors).
- Manufacturing protocols have been described for production of large-scale quantities of good manufacturing practice-grade vectors for clinical use.
- An scAAV2/8 vector expressing human factor IX from a liver-specific *LP1* promoter has shown promise in animal models, and a clinical trial using this vector for hemophilia B is underway.
- The ongoing clinical trial using scAAV vectors will be crucial to determine the direction of future clinical trials for hemophilia B, as well as other disorders.



#### Figure 1. Adeno-associated virus variant genomes

(A) Wild-type adeno-associated virus (AAV) genome, which consists of two inverted terminal repeats flanking coding sequences for four Rep proteins (Rep 78, Rep 68, Rep 52 and Rep 40) involved in genome replication, and four capsid proteins (VP1, VP2, AAP and VP3). (B) Recombinant AAV genome, where the coding sequences of the AAV genome are substituted with the transgenic cassette, which is flanked by the 3' and 5' inverted terminal repeats. Upon entry into the transduced cell (blue arrow), the single-stranded genome is converted by host factors into a transcriptionally active double-stranded form (dashed green arrow). (C) scAAV genome, in which the transgenic cassette is expressed as an inverted repeat with a deleted/mutated 3' terminal repeats in the middle, and flanked by two intact 5' terminal repeats. Upon entry into the transduced cell, the inverted repeats pair along their length, creating a double-stranded transcriptionally active genome, bypassing the need for second strand synthesis by host cell factors. Deletion/mutation of the 3' terminal repeats prevents the generation of monomeric forms by the Rep endonucleases, thereby stabilizing the self-complementary AAV genome in its dimeric form.

# Table 1

Clinical studies for gene and cell therapy of hemophilia A and B.

		administration		Kel.
Hemophilia A 7 h d	Iransient transfection of hFVIII into autologous lermal fibroblasts	Transplantation of fibroblasts into omentum	Transient elevation of FVIII levels in three out of six patients	[3]
Hemophilia A F E	Retroviral vector expressing 3-domain deleted hFVIII	Intravenous: peripheral vein	Transient elevation of FVIII levels in nine out of 13 patients	[4]
Hemophilia A C	Jutless adenoviral vector xpressing full-length hFVIII	Intravenous	Toxic reaction in one patient, resulting in discontinuation of treatment	[5]
Hemophilia B r	AAV2 vector expressing hFIX	Intramuscular: skeletal muscle	Effective gene transfer into muscle tissue. Therapeutic levels of FIX were not seen	[29–31]
Hemophilia B r	AAV2 vector expressing hFIX	Right hepatic artery	Therapeutic levels of FIX attained for 2–4 weeks at the highest dose of administered vector. This dose was also associated with an immune response against transduced hepatocytes, abrogating clinical benefits	[34]

FIX: Factor IX; FVIII: Factor VIII; hFIX: Human factor IX; hFVIII: Human factor VIII; rAAV: Recombinant adeno-associated viral.