

# Direct Intracranial Injection of AAVrh8 Encoding Monkey $\beta$ -N-Acetylhexosaminidase Causes Neurotoxicity in the Primate Brain

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GM2 gangliosidoses, including Tay–Sachs disease and Sandhoff disease, are lysosomal storage disorders caused by deficiencies in  $\beta$ -N-acetylhexosaminidase (Hex). Patients are afflicted primarily with progressive central nervous system (CNS) dysfunction. Studies in mice, cats, and sheep have indicated safety and widespread distribution of Hex in the CNS after intracranial vector infusion of AAVrh8 vectors encoding species-specific Hex  $\alpha$ - or  $\beta$ -subunits at a 1:1 ratio. Here, a safety study was conducted in cynomolgus macaques (cm), modeling previous animal studies, with bilateral infusion in the thalamus as well as in left lateral ventricle of AAVrh8 vectors encoding cm Hex  $\alpha$ - and  $\beta$ -subunits. Three doses ( $3.2 \times 10^{12}$  vg [ $n=3$ ];  $3.2 \times 10^{11}$  vg [ $n=2$ ]; or  $1.1 \times 10^{11}$  vg [ $n=2$ ]) were tested, with controls infused with vehicle ( $n=1$ ) or transgene empty AAVrh8 vector at the highest dose ( $n=2$ ). Most monkeys receiving AAVrh8-cmHex $\alpha/\beta$  developed dyskinesias, ataxia, and loss of dexterity, with higher dose animals eventually becoming apathetic. Time to onset of symptoms was dose dependent, with the highest-dose cohort producing symptoms within a month of infusion. One monkey in the lowest-dose cohort was behaviorally asymptomatic but had magnetic resonance imaging abnormalities in the thalami. Histopathology was similar in all monkeys injected with AAVrh8-cmHex $\alpha/\beta$ , showing severe white and gray matter necrosis along the injection track, reactive vasculature, and the presence of neurons with granular eosinophilic material. Lesions were minimal to absent in both control cohorts. Despite cellular loss, a dramatic increase in Hex activity was measured in the thalamus, and none of the animals presented with antibody titers against Hex. The high overexpression of Hex protein is likely to blame for this negative outcome, and this study demonstrates the variations in safety profiles of AAVrh8-Hex $\alpha/\beta$  intracranial injection among different species, despite encoding for self-proteins.

**Keywords:** Tay-Sachs disease, hexosaminidase, intracranial delivery, adeno-associated virus, gene therapy, AAV

## INTRODUCTION

TAY–SACHS DISEASE (TSD) and Sandhoff disease (SD) belong to a family of lysosomal storage disorders (LSDs) known as GM2 gangliosidoses. Both TSD and SD result from a deficiency in the lysosomal enzyme  $\beta$ -N-acetylhexosaminidase (Hex, hexosa-

minidase), which breaks down the cellular metabolite GM2 ganglioside into GM3 ganglioside by hydrolyzing the terminal N-acetyl galactosamine residue in the lysosome. HexA is a heterodimer of  $\alpha$ - and  $\beta$ -subunits, which are encoded by the *HEXA* and *HEXB* genes, respectively. TSD

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results from mutations in *HEXA*, while SD results from mutations in *HEXB*. Other isozymes of Hex include HexB, a  $\beta/\beta$  dimer, and HexS, an  $\alpha/\alpha$  dimer, neither of which can cleave GM2 ganglioside in humans. HexA deficiency leads to accumulation of GM2 ganglioside in the lysosome, causing dysfunction and eventual cell death. GM2 gangliosidoses primarily affect the central nervous system (CNS), leading to progressive neuronal degeneration.<sup>1</sup>

The three clinical forms of GM2 gangliosidoses are infantile, juvenile, and late-onset. The most severe infantile form is characterized by premature death within the first 5 years of life. Infants develop normally for the first 6 months, and patients are usually diagnosed at approximately 1 year of age due to regression of or failure to reach developmental milestones. Symptoms include seizures, deficits in motor functions, and dysphagia. Patients eventually decline to a vegetative state before succumbing to the disease. Presently, the care for these patients consists of palliative measures because there is no effective treatment.<sup>2</sup>

The current treatment for many LSDs is enzyme replacement therapy (ERT), where recombinant enzyme is delivered repeatedly, usually through intravenous or intramuscular injections.<sup>3</sup> However, due to the inability of lysosomal enzymes to cross the blood–brain barrier (BBB), this strategy is ineffective to treat the majority of LSDs, which have a CNS component, including TSD and SD.<sup>4,5</sup>

Gene therapy has proven to be one of the most promising approaches for treating many LSDs, with the potential to be a permanent, one-time treatment that can circumvent the issues associated with the BBB and recombinant enzyme delivery. Both ERT and gene therapy utilize the cross-correction mechanism where lysosomal enzymes secreted by a cell, especially when overexpressed, can be taken up by neighboring cells through the mannose-6-phosphate receptor and targeted correctly to the lysosome. Therefore, stable gene delivery to a relatively small subset of cells may be sufficient to supply the entire body with therapeutic levels of the missing enzyme. In the brain, distribution of lysosomal enzymes occurs through diffusion,<sup>6</sup> cerebral spinal fluid (CSF) flow,<sup>7–9</sup> and axonal transport.<sup>10–15</sup>

Recombinant adeno-associated virus (AAV) vectors have become the leading platform for development of *in vivo* gene therapies for LSDs with neurological involvement, as they transduce dividing and non-dividing cells in the CNS at high efficiency, and mediate long-term gene expression with minimal or no toxicity.<sup>16</sup> The most effective gene therapy approaches based on intracranial infusion

of AAV vectors have targeted structures highly interconnected with other regions of the brain such as the striatum,<sup>17–19</sup> ventral tegmental area,<sup>20</sup> deep cerebellar nuclei (DCN),<sup>21</sup> thalamus,<sup>22</sup> or CSF.<sup>8,23,24</sup> These intracranial delivery strategies reduce the number of injections necessary to achieve widespread distribution of lysosomal enzymes in the brain and have demonstrated efficacy in large animal models of different LSDs.<sup>25–32</sup>

SD mice and cats, and TSD sheep are the three available animal models of GM2 gangliosidoses whose phenotypes mimic the symptoms seen in human patients. Traditionally, both *HEXA* and *HEXB* genes are delivered to both SD and TSD animal models in order to produce ideal HexA isozyme overexpression.<sup>33–35</sup> SD mice (*Hexb*<sup>−/−</sup>) were generated by targeted disruption of *Hexb*. SD mice have been treated by intracranial injections of a combination of two AAV vectors, encoding human Hex  $\alpha$ - and Hex  $\beta$ -subunits. Treated SD mice live as long as 2 years, or the normal life-span of wild-type mice.<sup>17,25,26</sup> Both the SD cat and TSD sheep are naturally occurring large animal models that accumulate GM2 ganglioside and show neurological and motor dysfunction as well as a reduction in life-span, living approximately 4.5 and 9 months, respectively.<sup>25,36,37</sup> SD cats have a 25-base pair inversion in the 3' end of *Hexb* that results in <3% residual enzyme activity in brain.<sup>38</sup> TSD sheep carry a mutation in the *Hexa* gene that leads to skipping of exon 11, and thus HexA activity to the artificial substrate 4-methylumbelliferyl 6-sulfo-2-acetamido-2-deoxy- $\beta$ -D-glucopyranoside potassium salt (MUGS) is reduced to 6% and 29% of normal in liver and brain, respectively.<sup>36</sup> Simultaneous delivery of two AAVrh8 vectors encoding feline Hex  $\alpha$ - and Hex  $\beta$ -subunits has been successful in treating feline SD CNS pathology and extending life-span through bilateral injection into the thalamus alone or in combination with the DCN.<sup>25,26</sup> Unpublished data from the authors' lab has demonstrated similar efficacy in treating SD mice using AAVrh8 vectors encoding mouse Hex subunits, as shown previously for other AAV constructs.<sup>17,39</sup> Furthermore, a single injection of AAV vectors into the cerebral lateral ventricle (ICV) is as effective as DCN injection in complementing the therapeutic effect of bilateral thalamic injections in mice and cats.<sup>27</sup> Because ICV injection is considered safer than DCN injection, preclinical safety studies were conducted using bilateral thalamic injections in combination with ICV delivery of an equimolar formulation of two AAVrh8 vectors encoding Hex  $\alpha$ - and Hex  $\beta$ -subunits.<sup>16</sup>

Many preclinical safety studies have been conducted in nonhuman primates (NHPs) due to phy-

logenetic similarities, allowing testing of the feasibility of scale-up from rodent models to humans. Numerous studies have been conducted in NHPs to assess safety of AAV gene therapy for metabolic and neurological diseases using systemic, cerebrospinal fluid, and intracranial delivery routes.<sup>40–47</sup> However, such safety studies differ widely in delivery route, AAV vector design/dosing, therapeutic gene, and volume/rate of injection. Thus, extrapolation of safety to this paradigm is not advisable, as a novel capsid (AAVrh8) and target (thalamus) is being used, which have never been tested in humans. Previous studies using direct intracranial injections of AAVrh8 vectors encoding species-specific Hex  $\alpha$ - or  $\beta$ -subunits at 1:1 ratio in mice, cats, and sheep have all indicated safety, widespread enzyme distribution, and therapeutic efficacy.<sup>25,27</sup> In the present study, the gene therapy strategy used in SD cats was tested for preclinical safety and feasibility using normal monkeys.

## MATERIALS AND METHODS

### AAVrh8-CBA-Hex $\alpha$ / $\beta$ -WPRE<sup>mut6 $\Delta$ ATG</sup>

#### vector production

A single-stranded rAAV vector plasmid described previously<sup>48</sup> was used to generate the AAVrh8-CBA-cmHex $\alpha$ -WPRE<sup>mut6 $\Delta$ ATG</sup> and AAVrh8-CBA-cmHex $\beta$ -WPRE<sup>mut6 $\Delta$ ATG</sup> plasmids. The woodchuck hepatitis virus post-transcriptional regulatory element was modified to include the previously described mut6 mutations in the putative promoter region of protein X,<sup>49</sup> as well as replace all putative ATG codons with TTG. This modified WPRE<sup>mut6 $\Delta$ ATG</sup> was synthesized by overlapping polymerase chain reaction (PCR). Modified WPRE sequence: GATAATCAA CCTCTGGATTACAAAATTTGTGAAAGATTGAC TGGTATTCTTAACTTTGTTGCTCCTTTTACGC TTTGTGGATACGCTGCTTATTGCCTTTGTAT CTTGCTATTGCTTCCCCTTTGGCTTTCATTTT CTCCTCCTTGTATAAATCCTGGTTGCTGTCTC TTTTGGAGGAGTTGTGGCCCGTTGTCAGGCA ACGTGGCGTGGTGTGCACTGTGTTTGCTGAC GCAACCCCACTGGTTGGGGCATTGCCACCA CCTGTCAGCTCCTTTCCGGGACTTTCGCTTTC CCCCTCCCTATTGCCACGGCGGAATCATCG CCGCTGCCTTGCCCGCTGCTGGACAGGGGC TCGGCTGTTGGGCACTGACAATTCCTGTTG TGTGCGGGAAATCATCGTCCCTTTCCTTGGC TGCTCGCTGTGTTGCCACCTGGATTCTGCG CGGGACGTCTTCTGCTACGTCCCTTCGGCC CTCAATCCAGCGGACCTTCCCTTCCCGCGGCC TGCTGCCGGCTCTGCGGCCTTTCGCGTCT TCGCTTCCGCCCTCAGACGAGTCGGATCTCCC TTTGGGCCGCTCCCGCATCGGACTAG.

AAVrh8 vectors were produced by triple transient transfection of 293T cells followed by purification using iodixanol gradient and fast protein liquid chromatography, as previously described.<sup>50</sup> Vectors were then concentrated using Acrodisc<sup>®</sup> Units with Mustang<sup>®</sup> Q Membranes (Pall Corporation), buffer exchanged to phosphate-buffered saline (PBS) using 10 kDa Slide-A-Lyzer<sup>™</sup> Dialysis Cassettes (Thermo Fisher Scientific), and filtered using Millex-GV Syringe Filter Unit, 0.22  $\mu$ m, PVDF, 4 mm (EMD Millipore). The titers of the vectors were measured by real-time quantitative PCR with primers and probe to the bovine growth hormone polyadenylation signal, as described previously.<sup>50</sup>

## ANIMALS

Male and female cynomolgus macaques (cm; 1.5–2.5 years old) were purchased from Worldwide Primates, Inc. and selected for this study based on the absence of AAVrh8 neutralizing antibodies in serum (<1:10), measured as previously described.<sup>51</sup> All experiments were reviewed and approved by the Institutional Animal Care and Use Committee at the University of Massachusetts Medical School, and performed in compliance with the National Institutes of Health Guide for Care and Use of Laboratory Animals, 8th edition.

## INTRACEREBRAL INJECTION OF AAVRH8 VECTORS OR PBS AND IMAGING IN ANIMALS

Three doses ( $3.2 \times 10^{12}$  vg [1  $\times$  cohort,  $n = 3$ ],  $3.2 \times 10^{11}$  vg [1/10th cohort,  $n = 2$ ], or  $1.1 \times 10^{11}$  vg [1/30th cohort,  $n = 2$ ]) containing a 1:1 ratio of AAVrh8 vectors encoding cmHex $\alpha$ - and cmHex $\beta$ -subunits, along with control groups injected with vehicle (PBS,  $n = 1$ ) or  $3.2 \times 10^{12}$  vg of an AAVrh8 vector without a transgene (TGE,  $n = 2$ ), were tested. Neurosurgical procedure, injection, and imaging were performed as previously described.<sup>16,52,53</sup> Targeting and trajectory planning occurred through fusion of live operative fluoroscopy and cone-beam computed tomography scans with presurgical magnetic resonance imaging (MRI). Bilateral injections into the thalamus occurred simultaneously (150  $\mu$ L per side) with an infusion rate of 2  $\mu$ L/min using a 16G SmartFlow<sup>™</sup> neuroventricular cannula (MRI Interventions, Inc.). Approximately 10 min after infusion, the cannulas were withdrawn from the brain and followed by injection into the left cerebral lateral ventricle (300  $\mu$ L) at a rate of 50  $\mu$ L/min using a 16G SmartFlow<sup>™</sup> neuroventricular cannula (MRI Interventions, Inc.). In all cohorts, each thalamus received 1/4th of the total AAV vector dose, and the left cerebral lateral

ventricle received half. Injection volumes were kept constant across cohorts.

### **Clinical monitoring and neurological assessment**

Monkeys were assessed in their home cage before surgery and weekly as described.<sup>54</sup> Behavior of monkeys in their home cage was also evaluated using a camcorder to document behavior before surgery and at 7, 15, 30, 60, and 90 days post surgery. Spontaneous behavior (recording the monkey for 3.5 min while absent from the room), food presentation and response (to handing animal treats or placing treats in the food hopper for 1 min), and response to threat (approaching animal's cage while looking directly at the monkey for 1 min) were all documented.

### **Measurement of antibody titers in serum**

Serum antibody titers toward AAVrh8 vector or purified human Hex  $\alpha$ -subunit (Sigma–Aldrich) were measured using an enzyme-linked immunosorbent assay (ELISA), as previously described.<sup>25</sup> Titers  $\leq$ 1:8 were considered not detectable.

### **Tissue collection, processing, histopathology, histochemistry, and immunohistochemistry**

Fresh, unperfused tissue was collected from animals in the highest-dose cohort ( $3.2 \times 10^{12}$  vg) as well from the PBS control group. Animals from the  $3.2 \times 10^{11}$  vg dose cohort,  $1.1 \times 10^{11}$  vg dose cohort, and control animals receiving AAVrh8 vector without a transgene were perfused with cold PBS prior to tissue collection. Following death, a complete necropsy was performed. All tissues collected were fixed in 10% neutral buffered formalin and were paraffin embedded according to standard procedures. Five micrometers of histologic sections were routinely processed and stained with hematoxylin and eosin (H&E). Luxol fast blue histochemical staining was performed based on standard procedures. To characterize the immune cell infiltrate of the lesions, immunohistochemistry for CD3, CD20, and CD68 was performed. Briefly, all staining protocols used an antigen retrieval protocol of microwaving in 0.1 M of sodium citrate buffer for 20 min followed by 20 min of cooling at room temperature (RT). CD3 (Dako; catalog #A0452) was applied for 30 min at room temperature at a dilution of 1:600. CD20 (Dako; catalog #M0755) and CD68 (Dako; catalog #M0814) were applied overnight at 4°C at a dilution of 1:179 and 1:410, respectively. Secondary antibodies were biotinylated horse anti-mouse (Vector Laboratories) or biotinylated goat anti-rabbit (Vector Laboratories) as appropriate,

diluted 1:200, and incubated for 30 min at room temperature. Tertiary reagent in each case was Vectastain ABC Elite reagent (CD3) or Vectastain ABC Standard reagent (CD20, CD68) incubated for 30 min at RT (Vector Laboratories). All slides were developed with DAB chromogen (Dako) and counterstained with Mayer's hematoxylin. Negative controls consisted of identical brain sections stained with irrelevant species-, isotype-, and concentration-matched antibodies. A lymph node served as the control tissue for all antibodies.

### **Western blotting**

Tissues were lysed from 3 mm of biopsy punches of the thalamus in T-PER™ Tissue Protein Extraction Reagent (Thermo Scientific) containing cOmplete Protease Inhibitor Cocktail (Roche). Total protein was measured using QuickStart™ Bradford Protein Assay with serial dilutions of bovine serum albumin as protein standard (Bio-Rad). Total protein lysates (20  $\mu$ g) were resolved using SDS-polyacrylamide gel electrophoresis with 4–20% Mini-PROTEAN® TGX™ (Bio-Rad) precast gel in 25 mM Tris 190 mM Glycine 0.1% sodium dodecyl sulfate followed by wet transfer to a 0.22  $\mu$ m nitrocellulose membrane (Maine Manufacturing, LLC) in 25 mM Tris 190 mM Glycine. Membranes were blocked for 1 h in 5% nonfat dry milk in Tris-buffered saline with 0.05% Tween™ 20 (TBST) and probed with rabbit anti-Hex  $\alpha$ -subunit antibody (LifeSpan BioSciences, Inc.) or mouse anti-Hex  $\beta$ -subunit antibody<sup>25</sup> diluted 1:1,000 in 5% nonfat dry milk in TBST and incubated overnight at 4°C. Horseradish peroxidase (HRP)-conjugated secondary antibodies (anti-mouse or anti-rabbit; GE Healthcare) were diluted 1:5,000 in 5% nonfat dry milk TBST and incubated for 1 h at room temperature. Signal was detected using Pierce ECL Western Blotting Substrate (Thermo Fisher Scientific) followed by exposure to Amersham Hyperfilm ECL (GE Healthcare). Membranes were stripped using 100 mM glycine (pH 4.0) for 25 min, washed in TBST, and then blocked and probed again using HRP-conjugated mouse anti- $\beta$ -actin antibody (GenScript) diluted 1:1,000 in 5% nonfat dry milk in TBST for 1 h and detected as before.

### **Hexosaminidase activity**

Hexosaminidase isozyme activities were measured from fresh frozen tissue either from 3 mm biopsy punches from the thalamus of 4 mm coronal blocks or from 2 mm slices from thoracic spinal cord coronal blocks using 4-methylumbelliferyl N-acetyl- $\beta$ -D-glucosaminide (MUG; Sigma) and MUGS (Toronto Research Chemicals), and normalized to total protein measured by QuickStart™ Bradford Protein

Assay with serial dilutions of bovine serum albumin as protein standard (Bio-Rad), as previously described.<sup>25</sup> Enzyme activity is represented as nmol 4MU/h/mg protein.

### Fluorescent immunostaining

Formalin-fixed paraffin sections (5  $\mu$ m thick) were used for neuropathological evaluation. Paraffin was removed from tissue-mounted slides with two 8 min incubations in xylenes and then rehydrated with serial 4 min incubations in 100%, 95%, and 80% ethanol followed by 1 $\times$  PBS. For antigen retrieval, slides were boiled in 10 mM Tris 1 mM EDTA (pH 9.0) for 5 min and then cooled for 20 min. Blocking with permeabilization was done for 1 h in 5% normal goat serum (SouthernBiotech) in PBS with 0.05% Tween-20 (Acros). Sections were incubated with primary antibodies diluted in 5% normal goat serum in PBS with 0.05% Tween-20 for 1 h at room temperature at the following concentrations: rabbit anti-HEXA, 1:1,000 (LifeSpan BioSciences, Inc.), mouse anti-HEXB, 1:1,000,<sup>25</sup> or NeuN, 1:1,000 (EMD Millipore). Sections were washed three times in TBST, and secondary antibodies against the appropriate species and conjugated to the different fluorophores (Alexa-488 or Alexa-555; Thermo Fisher Scientific) were diluted 1:1,000 in 5% normal goat serum in PBS with 0.05% Tween-20 and incubated for 30 min followed by three washes in PBS. Sections were stained for 1 min in Thermo Fisher Scientific Pierce™ DAPI Nuclear Counterstain diluted in PBS. Coverslips were mounted on slides using PermaFluor aqueous mounting medium (Thermo Fisher Scientific).

### In situ hybridization

RNAscope fluorescent multiplex kit (Advanced Cell Diagnostics) was used for *in situ* analysis of vector derived mRNA in tissue sections using a WPRE<sup>mut6</sup> $\Delta$ ATG specific probe set (Advanced Cell Diagnostics). Formalin-fixed 4  $\mu$ m paraffin brain sections were used according to the manufacturer's instructions.

### Silver stain of AAVrh8 vector and densitometry

AAVrh8 vector purity was assessed by gel electrophoresis using 4–20% Mini-PROTEAN® TGX™ (Bio-Rad) precast gels followed by silver staining (Invitrogen) and densitometry.

### Electron microscopy of AAVrh8 vector preparations

AAVrh8 vector preparations were analyzed by the Core Electron Microscopy Facility at the University of Massachusetts Medical School (Worcester, MA) using negative staining.

### Endotoxin detection

Endotoxin levels were measured on AAV vector preparations using the Endpoint Chromogenic LAL Assays (QCL-1000™) from Lonza, according to the manufacturer's instructions.

## RESULTS

### Experimental design

The goal of this study was to assess the safety of a formulation of two AAVrh8 vectors (1:1 ratio) encoding Hex  $\alpha$ - and  $\beta$ -subunits injected bilaterally into the thalamus and one cerebral lateral ventricle in cm (Supplementary Fig. S1; Supplementary Data are available online at [www.liebertpub.com/hum](http://www.liebertpub.com/hum)). Vector purity, endotoxin levels, and empty capsid ratios were measured (Supplementary Fig. S1). These vectors encoded cm Hex proteins to reduce the likelihood of a confounding immune response triggered by delivery of Hex proteins from other species. All subjects were selected for the absence of AAVrh8 neutralizing antibody titers (<1:10). The study included three dosing cohorts:  $3.2 \times 10^{12}$  vg (1 $\times$  cohort,  $n = 3$ ),  $3.2 \times 10^{11}$  vg (1/10th cohort,  $n = 2$ ), and  $1.1 \times 10^{11}$  vg (1/30th cohort,  $n = 2$ ) of AAVrh8-cmHex vectors. Control groups were injected with vehicle (PBS,  $n = 1$ ) or  $3.2 \times 10^{12}$  vg of a TGE AAVrh8 vector ( $n = 2$ ). The 1 $\times$  dose was based on the efficacious dose in SD cats by vg/kg brain weight.<sup>27</sup> Safety was determined by neurological behavioral assessments, MRI data obtained at the study endpoint (90 days), and pathological examination of tissues.

### Clinical observations

All monkeys tolerated and recovered well from the surgical procedure. However, monkeys in the highest-dose cohort (1 $\times$ ) started to show neurological symptoms at 2–3 weeks after the injection. Symptoms included general weakness of limbs, ataxia, inability to perch, and reduced food intake. The neurological function of these animals declined rapidly, and they became apathetic, which led to premature euthanasia between 20 and 28 days. In the 1/10th dosing cohort, animals appeared normal until 33 and 48 days after the injection when they began to develop chorea and dyskinesia during voluntary movement, such as when reaching out for food. From this cohort, the animal that developed symptoms at 33 days required euthanasia at 57 days due to lethargy and unresponsiveness. However, the other animal in this cohort survived to the 90-day study endpoint. In the 1/30th dosing cohort, one animal lost voluntary use of its right arm at 59 days and required euthanasia due to

**Table 1.** Summary of neurological symptoms observed in study animals

Animal ID	Vector dose	Symptom onset (days)	Survival (days)	Clinical signs
0908068	1 $\times$	14	20	Lethargy, unresponsive
0908087	1 $\times$	17	23	Ataxia, unable to perch
0909589	1 $\times$	27	28	Generalized weakness especially of right leg, reduced food intake
8722431552	1/10th	33	57	Dyskinesias, chorea
09087133	1/10th	48	$\geq$ Endpoint	Dyskinesias, chorea
0909587	1/30th	59	66	Loss of voluntary use of right arm, eventual lethargy
0908997	1/30th	n/a	$\geq$ Endpoint	None
8963934563	PBS	n/a	$\geq$ Endpoint	None
5347096796	TGE (1 $\times$ )	n/a	$\geq$ Endpoint	None
7667353701	TGE (1 $\times$ )	n/a	$\geq$ Endpoint	None

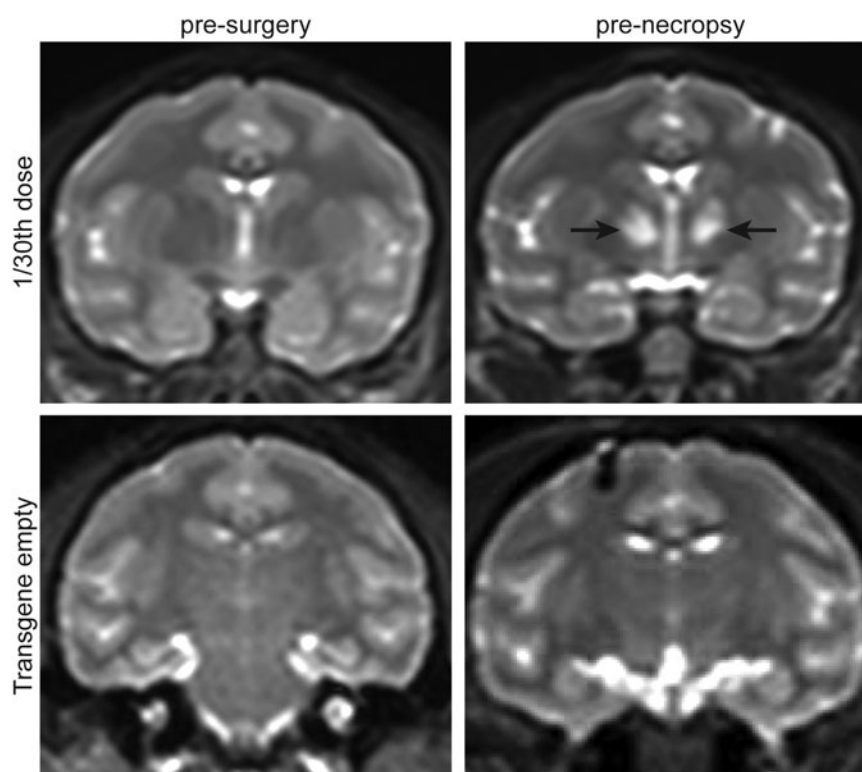
For each animal, vector dose, day of symptom onset, length of survival, and observed clinical signs are indicated. The predetermined endpoint of the study was 90 days.

TGE, transgene empty; PBS, phosphate-buffered saline.

lethargy at 66 days. The other animal in this cohort did not develop any major clinical signs and survived to the 90-day study endpoint (Table 1). Despite no outward clinical symptoms in this animal, a brain MRI before necropsy indicated alterations in the thalamus (Fig. 1). No clinical signs were observed in the control cohorts, which consisted of a PBS-injected animal, as well as two animals injected with a TGE AAVrh8 vector at the 1 $\times$  dose (Table 1 and Fig. 1).

### Pathology

Serum antibody titers to human recombinant HexA and AAVrh8 capsid were measured both before the injection and at necropsy. There were no detectable titers to the HexA protein before the injection or at necropsy in all animals. At necropsy, animals receiving vector (including the TGE control) had AAVrh8 capsid serum antibody titers ranging from 1:128 to 1:49,152. The PBS-injected animal did not have any AAVrh8 capsid antibody titers (Table 2).



**Figure 1.** Corresponding coronal sections of brain T2-weighted magnetic resonance imaging before surgery and before necropsy indicating signal alterations at the study endpoint in the treated animal. Hyper-intensities (*arrows*) detected in the thalami of an animal injected with  $1.1 \times 10^{11}$  vg (1/30th dose) of AAVrh8 vector formulation encoding cmHex $\alpha$  or cmHex $\beta$  subunits. No hyper-intensities were noted at any time point in the thalami of animals treated with a transgene empty vector.

**Table 2.** Serum titers against recombinant human Hex $\alpha$ -subunit and AAVrh8 capsid

Animal ID	Titer to hHex $\alpha$ -subunit		Titer to rh8 capsid	
	Before surgery	At necropsy	Before surgery <sup>a</sup>	At necropsy
<i>1<math>\times</math></i>				
0908068	ND 1:8	ND 1:8	ND	1:3,072
0908087	ND 1:8	ND 1:8	ND	1:2,048
0909589	ND 1:8	ND 1:8	ND	1:1,024
<i>1/10th</i>				
8722431552	ND 1:8	ND 1:8	ND	1:49,152
09087133	ND 1:8	ND 1:8	ND	1:49,152
<i>1/30th</i>				
0909587	ND 1:8	ND 1:8	ND	1:128
0908997	ND 1:8	ND 1:8	ND	1:384
<i>PBS</i>				
8963934563	ND 1:8	ND 1:8	ND	ND 1:8
<i>TGE</i>				
5347096796	ND 1:8	ND 1:8	ND	1:49,152
7667353701	ND 1:8	ND 1:8	ND	1:1,024

The titers before surgery and at necropsy measured by enzyme-linked immunosorbent assay against Hex $\alpha$  and AAVrh8 capsid for each animal and their dosing cohorts are indicated.

<sup>a</sup>Presurgery AAVrh8 titers were measured by neutralizing antibody screen to select animals.

ND, not detected.

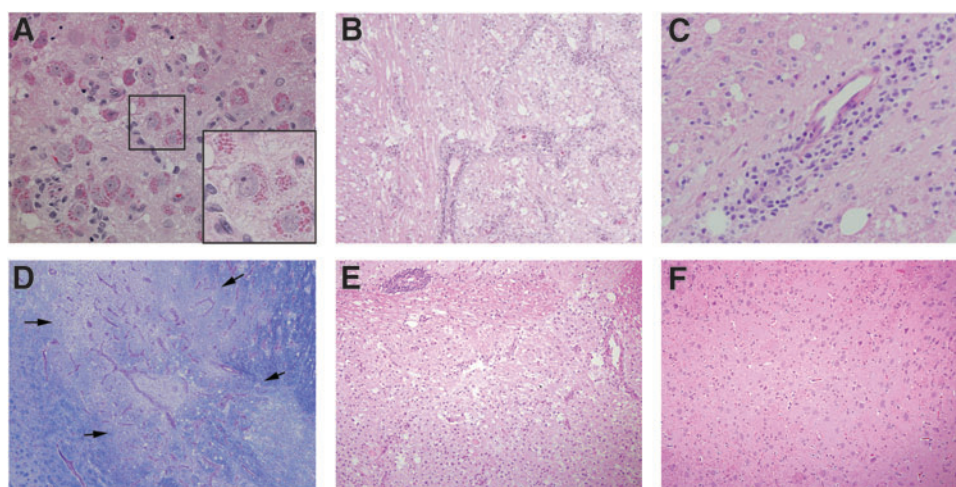
Histological examination of the brain revealed widespread neuropathology in the thalamus of animals injected with AAVrh8 encoding Hex. Alterations included the presence of granular eosinophilic inclusions in neurons (Fig. 2A), necrotic areas with cellular loss and vacuolation (Fig. 2B), perivascular cuffing (Fig. 2C), and white-matter loss indicated by lack of luxol fast blue stain (Fig. 2D). Neurons with eosinophilic inclusions

were also observed in the thalamus of the other NHPs injected with lower doses of AAVrh8-cmHex vectors (Fig. 2E). There was no evidence of neuropathology in other regions of the brain, with the exception of that associated with needle tracks. All control cohorts had mostly normal histological appearance in the thalamus (Fig. 2F), with only small lesions present near the injection site or needle tracks.

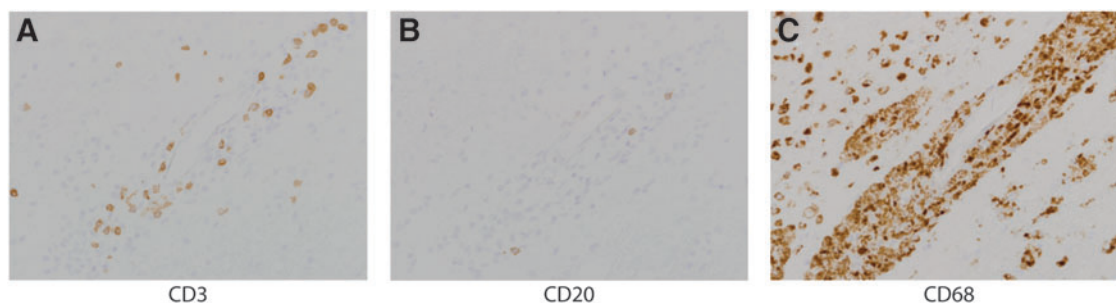
Analysis of the types of immune cells by immunohistochemistry revealed the presence of T cells (CD3<sup>+</sup>) and B cells (CD20<sup>+</sup>) in perivascular cuffs, but little to none elsewhere in the thalamus (Fig. 3A and B). However, there was a large macrophage/microglial presence, as shown in CD68 immunostaining in the thalamus (Fig. 3C). These findings were common to all subjects treated with AAV-cmHex vectors, but not in NHPs treated with either PBS or TGE AAV vector.

#### Hexosaminidase expression and localization

Western blot analysis of Hex  $\alpha$ - and Hex  $\beta$ -subunit protein levels revealed increased levels in the thalamus of animals injected with AAVrh8 vectors encoding cmHex $\alpha/\beta$  (Fig. 4A). Accordingly, hexosaminidase activity in the thalamus and spinal cord of animals injected with AAVrh8 vectors encoding cmHex $\alpha/\beta$  was considerably higher than in control animals, in some instances by >100-fold (Fig. 4B). This large increase in hexosaminidase expression was still present, despite the cellular loss observed in the thalamus (Fig. 2).



**Figure 2.** Histological findings in the thalamus. (A) Neurons with granular eosinophilic inclusions were present in the thalamus of AAVrh8-Hex injected animals. Box indicates location of magnified region. 40 $\times$ . (B) Necrotic area in the thalamus with cell loss and vacuolation. 10 $\times$ . (C) Example of perivascular cuffing in the thalamus. 20 $\times$ . (D) Luxol fast blue staining revealed a large area (delineated by arrows) of necrosis, vascular proliferation, and white matter loss (note pallor compared to adjacent dark blue color). 10 $\times$ . (E) Neuropathological findings were identical in the lowest dosing cohort. 10 $\times$ . (F) Normal histology of the thalamus in phosphate-buffered saline (PBS)-injected animal; 10 $\times$  (1 $\times$  cohort, A–D; 1/30th cohort, E; PBS-injected, F). Tissue sections for histological assessment shown in (A), (B), (C), (E), and (F) were stained with hematoxylin and eosin.



**Figure 3.** Immunological profile of perivascular cuffs in AAVrh8-cmHex $\alpha/\beta$  injected monkey thalamus. **(A)** CD3 immunostaining for T cells. **(B)** CD20 immunostaining for B cells. **(C)** CD68 immunostaining for macrophages/microglia. **(A–C)**, 1/10th cohort animal). 20 $\times$ .

Despite cellular loss (Fig. 2), there were still numerous cells in the thalamus containing supra-physiological levels of both Hex subunits, as shown by double immunofluorescence staining for Hex  $\alpha$ - and Hex  $\beta$ -subunits (Fig. 5A). Many of these cells

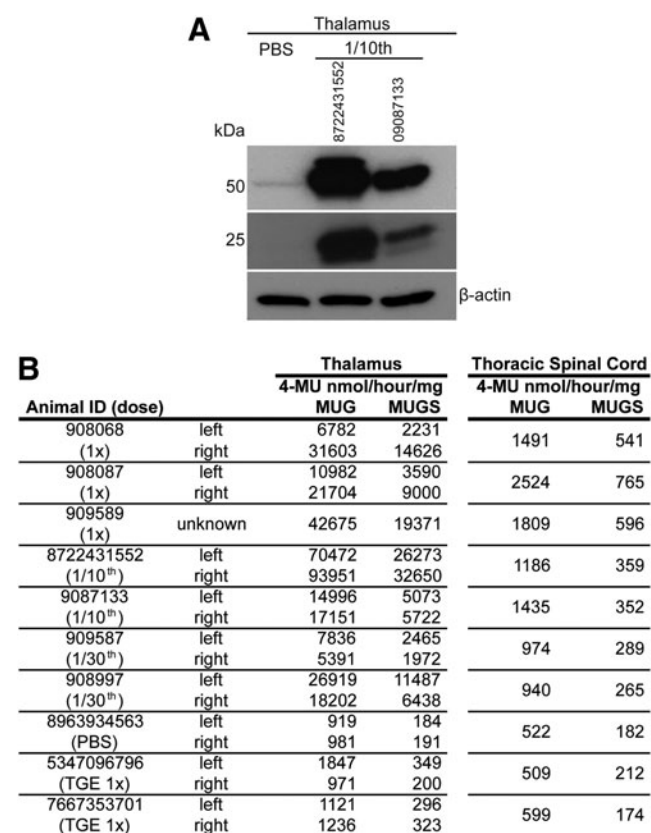
were neurons given the co-localization of Hex  $\alpha$ -subunit with NeuN (Fig. 5B). *In situ* hybridization in parallel sections using a WPRE probe indicated many AAV vector transduced cells (Fig. 5C).

## DISCUSSION

Production of Hex through direct thalamic delivery of AAVrh8 vectors encoding species-specific subunits proved to be toxic in normal NHPs, probably due to overexpression. The severe neurological symptoms that manifested over time were surprising, given the demonstrated efficacy and safety of bilateral injections of AAV-Hex vectors (AAV1, AAV2, and AAVrh8) in normal and GM2 affected disease mice, cats, and sheep.<sup>17,25,27,39</sup> The absence of neuropathology elsewhere in the CNS suggests that delivery of AAVrh8-Hex vectors into CSF through the cerebral lateral ventricles is safe.

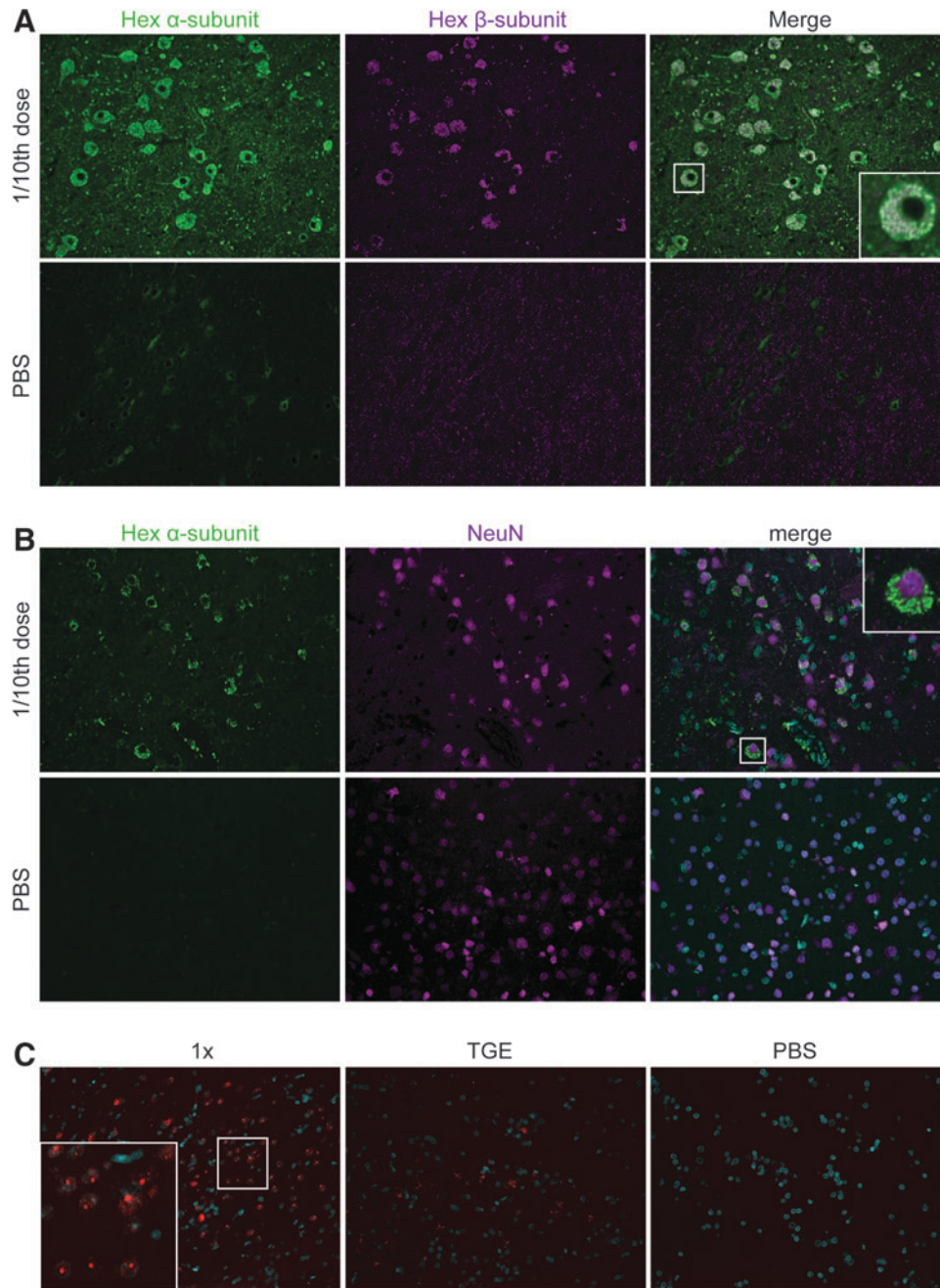
The most likely explanation for the neurologic deterioration is the extensive cellular loss in the thalamus, revealed by neuropathological examination of the NHP brains injected with AAVrh8-cmHex $\alpha/\beta$ . Unlike the severe histological findings documented here, a prior study in NHPs injected with an AAV2 vector expressing human acid sphingomyelinase in the thalamus and brain stem also developed severe neurological symptoms but with no apparent cellular loss. Instead, there was robust infiltration of injected brain structures with inflammatory cells.<sup>55</sup>

Several lines of evidence in the present study suggest the degenerative process triggered by thalamic injection of AAVrh8-cmHex $\alpha/\beta$  vectors in normal NHPs is unlikely to have been caused by an immunological response. The human and cm hexosaminidase  $\alpha$ - and  $\beta$ -subunits differ by 11 and 23 amino acids, respectively (Supplementary Figs. S2 and S3). Intracranial injection of AAV vectors encoding non-self-proteins in NHPs and other species has been shown to trigger strong immune responses, resulting in extensive cellular loss. In addition to the



**Figure 4.** Increased Hex expression and activity in the central nervous system. **(A)** Western blot analysis of Hex  $\alpha$ - or  $\beta$ -subunit protein levels in thalamus lysates from the PBS-injected control and two animals from the 1/10th cohort. **(B)** Hex activity was measured in the thalamus and thoracic spinal cord on all cohorts. 4-Methylumbelliferyl N-acetyl- $\beta$ -D-glucosaminide substrate is cleaved by HexA, HexB, and HexS isozymes. 4-Methylumbelliferyl 6-sulfo-2-acetamido-2-deoxy- $\beta$ -D-glucopyranoside potassium salt substrate is preferentially cleaved by HexA and HexS isozymes. Note: Due to an emergency euthanasia and necropsy, the identity of the thalamus (left or right) in animal is 909589 is unknown.





**Figure 5.** Hex $\alpha$ - or Hex $\beta$ -subunit localization in monkey thalamus. **(A)** Immunofluorescence staining for Hex $\alpha$  (green) and Hex $\beta$  (magenta) subunits and their co-localization (white). Box indicates location of magnified region. 40 $\times$ . **(B)** Immunostaining for Hex $\alpha$ -subunit (green) and NeuN (magenta). 40 $\times$ . **(C)** RNA *in situ* hybridization with probe for WPRE (red) expressed by AAV transduced cells. DAPI stain (cyan). 20 $\times$ .

antigenic nature of the transgene, the AAV capsid tropism to antigen presenting cells in the brain may also be a contributing factor.<sup>56,57</sup> Moreover, as patients afflicted with lysosomal storage diseases treated by enzyme replacement therapy often develop antibodies to recombinant human proteins,<sup>58,59</sup> the differences in Hex subunits between species may be a significant factor that could confound the interpretation of the results. As such, the AAVrh8-Hex vectors encoded cm hexosaminidase  $\alpha$ -

and  $\beta$ -subunits, which would make it unlikely that cellular loss was caused by an adaptive immune response to the transgenes. The presence of large numbers of transduced neurons in the thalamus near areas of heavy cellular loss further supports the notion of a non-immune mechanism. In addition, inflammatory cells in the thalamus were mostly of the macrophage lineage (CD68<sup>+</sup>), with very few CD3<sup>+</sup> T cells only found near some blood vessels. These findings suggest that the inflammatory cell

infiltration was likely secondary to a primary neurodegenerative insult. Also, no evidence was found of antibodies to HexA at the endpoint, although this finding has a caveat that the ELISA assay was based on commercially available human Hex, and thus it is possible that it failed to detect antibodies to NHP specific epitopes.

Despite having low or absent neutralizing antibodies to AAVrh8 (titer <1:10) prior to injection, all NHPs injected with AAVrh8 vectors had high anti-capsid antibody titers at the endpoint. However, a humoral response to the capsid is unlikely to have a significant contribution to the cellular loss, as control NHPs injected with the transgene-empty AAVrh8 vector showed no evidence of neuropathology beyond that associated with the needle track. Moreover, previous studies have reported the presence of high anti-AAV antibody titers after intracranial delivery in NHPs with no apparent adverse effects.<sup>46,47</sup>

Despite extensive cellular loss, Hex activity in the thalamus of NHP injected with AAVrh8-cmHex $\alpha/\beta$  vector was >100-fold above normal. These high Hex expression levels appear to be driven by the remaining transduced neurons laden with eosinophilic granules filled with Hex, as indicated by co-immunofluorescence staining with antibodies to Hex  $\alpha$ - and  $\beta$ -subunits. Neurons with identical intracellular features were also found in the thalamus of long-lived cats treated with AAVrh8 vectors encoding feline Hex subunits, but with no evidence of cellular loss or neurological deficits.<sup>25</sup> Transgenic mice overexpressing lysosomal beta-glucuronidase (GUSB) also displayed eosinophilic granules in different cell types, and brain expression levels were ~100–700-fold above normal, with no apparent neurological deficits. This indicates that massive overexpression of at least GUSB in neurons is well tolerated.<sup>60</sup> However, there are fundamental differences between transgenesis and AAV-mediated *de novo* overexpression, namely developmental adaptation of cellular pathways to high-level expression of a lysosomal enzyme. Also, tissue level activity in transgenic mice is likely the result of relatively uniform enzyme activity in most cells in the tissue. On the other hand, the efficiency of AAV gene delivery to the brain by intraparenchymal injection is far from achieving transduction of all cells in a particular structure, and thus the measured activities are the result of enzyme expression in a subset of cells. The >100-fold above normal HexA levels in the thalamus of AAVrh8 injected NHPs at the endpoint appeared to be produced by a relatively small number of neurons, and thus it is likely

that activities in those cells are considerably higher than the average measured in tissue punches but are still compatible with survival. It seems reasonable to postulate that HexA expression crossed an unknown threshold in some cells that triggered a neurodegenerative process. A possible mechanism is that continuous high-level expression of HexA overtaxes the protein folding capacity in some AAV transduced neurons, leading to a chronic unfolded protein response that ultimately results in cell death.<sup>61</sup> The exact limit of overexpression tolerated by neurons is unknown, but the notion that lysosomal enzymes can be expressed at any level with no consequences to genetically modified cells should be reconsidered, although the possibility that these adverse events are HexA specific cannot be excluded. An example of overexpression-associated toxicity of a lysosomal enzyme in a particular target cell type is that described for *ex vivo* lentivirus gene therapy in Krabbe disease, where galactocerebrosidase proved to be toxic to hematopoietic stem cells but not to their progeny.<sup>62</sup> This has not been documented for other lysosomal enzymes where this *ex vivo* strategy has been successful for storage diseases such as metachromatic leukodystrophy.<sup>63</sup> Presently, the possibility that toxicity is the result of unique properties of cm hexosaminidase subunits cannot be excluded. Further studies with hexosaminidases from different species will be necessary to address this possibility.

The reasons for the difference in outcomes between NHPs and normal cats injected with AAVrh8 vectors carrying the same expression elements at comparable doses are unknown. It is possible the CBA promoter and/or WPRE in the transgene expression cassette have different potencies across species, and also that AAVrh8 transduces NHP cells more efficiently, given that it was cloned from rhesus macaque tissues.<sup>64</sup> The  $3 \times 10^{12}$  vg dose tested in NHP was based on a prior study<sup>27</sup> where GM2 cats were treated with  $1.1 \times 10^{12}$  vg of AAVrh8 vector delivered bilaterally to the thalamus and CSF via the lateral ventricle. The dose/brain weight is comparable at  $5.3 \times 10^{13}$  vg/kg, considering average brain weights of  $60.4 \pm 5.5$  g and  $20.9 \pm 1.6$  g in young cm and cats, respectively. An important aspect to consider is that the thalamic dose in cats was delivered in a columnar fashion, where the needle was raised 0.15 cm after each 10–20  $\mu$ L bolus, while in NHP convection-enhanced delivery in a single site in the thalamus was used. The two infusion techniques likely generate different patterns of AAV vector distribution in the thalamus and possibly trans-

duction efficiency that might be reflected in differences in AAV genome copy number per cell. Nonetheless, a 30-fold reduction in AAV vector dose in NHP was insufficient to prevent toxicity in NHP. This suggests that differences in infusion technique and ultimately AAV distribution in thalamus are unlikely to explain the surprising difference in outcomes in monkeys and cats. The degree of neuronal loss documented in monkeys prevents a meaningful comparison of AAV vector genome content in the thalamus between the two species and others, as it is impossible to know how much of the initial dose has been lost in monkeys due to toxicity. Since dose reduction proved ineffective in preventing thalamic toxicity in monkeys, an alternative approach is to decrease transgene expression strength from the AAV vector. This may be accomplished through removal of the WPRE element and using weaker promoters. However, the optimization process will be mostly empirical because the available data on safe levels of hexosaminidase overexpression in the brain of other species injected with AAV-Hex<sup>17,26,27</sup> are at the tissue level comprised of AAV transduced and non-transduced cells. Presently, there is no information on the degree of enzyme overexpression in individual AAV transduced cells (neurons) in the brain that is compatible with long-term safety and efficacy. A recent study using systemic delivery of AAV9 encoding another lysosomal enzyme,  $\alpha$ -N-acetylglucosaminidase, to treat mucopolysaccharidosis (MPS) IIIB resulted in liver toxicity associated with transgene overexpression in wild-type but not MPS IIIB mice.<sup>65</sup> The difference in safety outcomes across and within species raises concerns about the predictive value of preclinical dose ranging and safety studies when translating AAV gene therapies from animals to humans.

Prior studies assessing the safety of multisite intracranial delivery of AAVrh10 vectors in NHPs reported no serious adverse events.<sup>46,66</sup> However,

neuropathological examination revealed cellular loss at the injection sites, but because the studies were carried out with AAVrh10 vectors encoding human enzymes, it is not possible to distinguish between an immunologically driven effect and toxicity due to overexpression as documented here. The absence of neurological side effects in those studies may simply be due to the small injection volumes per site (10  $\mu$ L) in non-motor cortical regions where the effect of cellular loss in small numbers may not be easily detectable. The mechanistic basis for the cellular loss/spongiosis observed in those studies has different implications for the success in translation of these multisite approaches to patients. Clinical trials have been initiated with AAVrh10 vectors tested in NHPs and the outcomes in patients will ultimately reveal the safety and efficacy of the multisite delivery approach (NCT01801709).

Overexpression and secretion of lysosomal enzymes from genetically modified cells is the fundamental tenet of the current *ex vivo* and *in vivo* gene therapy approaches for lysosomal storage diseases. Though the exact tolerability limits for lysosomal enzyme overexpression in cells is unknown and may vary across enzymes, the notion that these proteins can be safely expressed at any level with no consequences should be evaluated carefully.

## ACKNOWLEDGMENTS

We would like to thank Dr. Richard P. Moser, MD, for use of his MRI compatible stereotaxic frame. This work was supported by grants from Cure Tay–Sachs Foundation, National Tay–Sachs and Allied Diseases Association, and the National Institutes of Health (U01NS064096).

## AUTHOR DISCLOSURE

No competing financial interests exist.

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Received for publication July 13, 2016;  
accepted after revision January 23, 2017.

Published online: January 26, 2017.