

Presentation #OR54

Vector genome loss and epigenetic modifications impact long-term transgene expression of AAV5 vectors produced in mammalian HEK293 and insect *Sf* cells

Ashrafali Mohamed Ismail, Britta Handyside, Lening Zhang, **Bridget Yates**, Lin Xie, Choong-Ryoul Sihn, Ryan Murphy, Taren Bouwman, Chan Kyu Kim, Sherry Bullens, Stuart Bunting, Sylvia Fong

BioMarin Pharmaceutical Inc., Novato, CA, USA

Disclosures

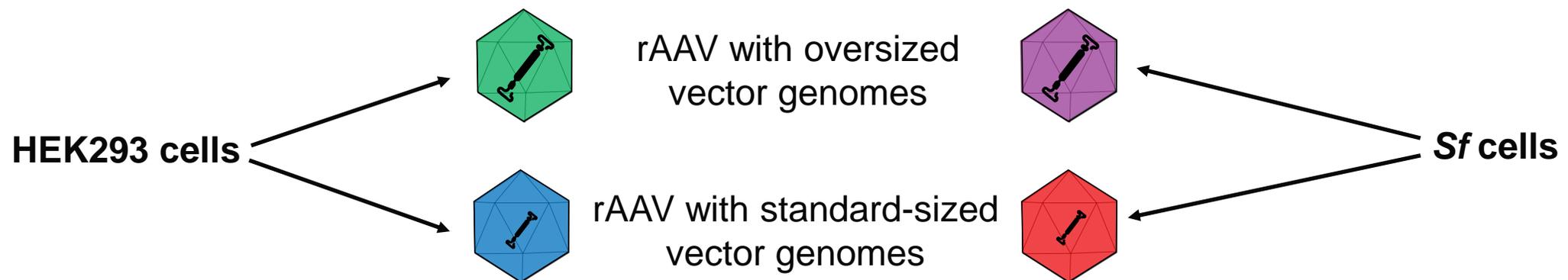
- Funding for this study was provided by BioMarin Pharmaceutical Inc.
- All authors are employees and stockholders of BioMarin Pharmaceutical Inc.

Introduction

- rAAV can be produced in human HEK293 cells or *Sf* insect-cell systems^{1–4}
- Valoctocogene roxaparvovec (AAV5-hFVIII-SQ) is a gene therapy for the treatment of severe haemophilia A produced with *Sf* cells⁵
 - The vector genome slightly exceeds the optimal packaging capacity of AAV vectors⁶

Objective:

- To compare the long-term durability of expression in mice treated with “oversized” (~5000 nt) and “standard-sized” (4600 nt) rAAV5 vectors produced in HEK293 or *Sf* cells
- To investigate the mechanistic factors affecting long-term transgene expression



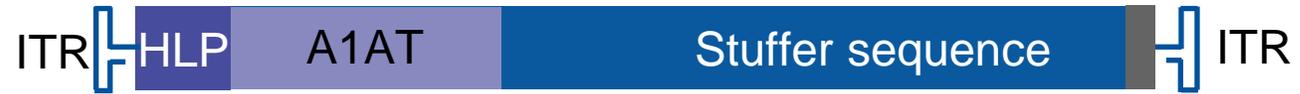
AAV, adeno-associated virus; AAV5, AAV serotype 5; HEK293, HEK293 cells; hFVIII-SQ, human factor VIII, SQ variant; rAAV, recombinant AAV; *Sf*, *Spodoptera frugiperda* cells.

1. Chahal PS, et al. *J Virol Methods*. 2014;196:163–73. 2. Kotin RM, et al. *Hum Gene Ther*. 2017;28(4):350–60. 3. Kondratov O, et al. *Mol Ther*. 2017;25(12):2661–75. 4. Kurasawa JH, et al. *Mol Ther Methods Clin Dev*. 2020;19:330–40. 5. Rangarajan S, et al. *N Engl J Med*. 2017;377(26):2519–30. 6. Bunting S, et al. *Mol Ther*. 2018;26(2):496–509.

Study design

- C57BL/6 WT mice were administered an AAV5-hA1AT vector IV at a dose of 6×10^{13} vg/kg
 - Serial bleed cohort: blood was collected weekly during the first 4 weeks after dosing, then monthly through week 57
 - Takedown cohort: livers were collected for oversized vector genome analysis at weeks 1, 3, 12, 24, and 57

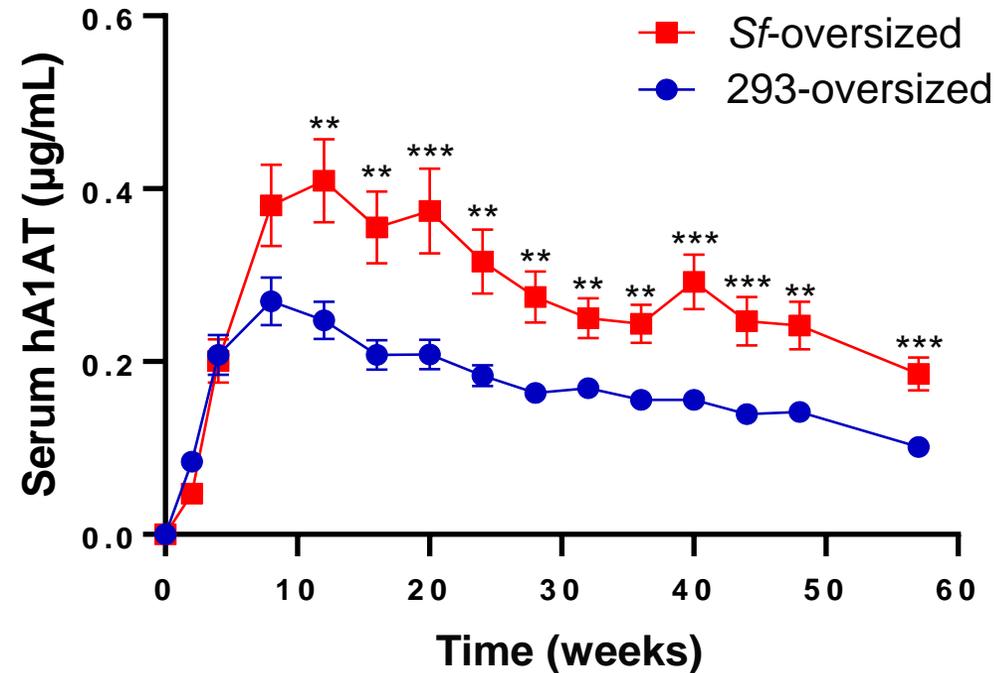
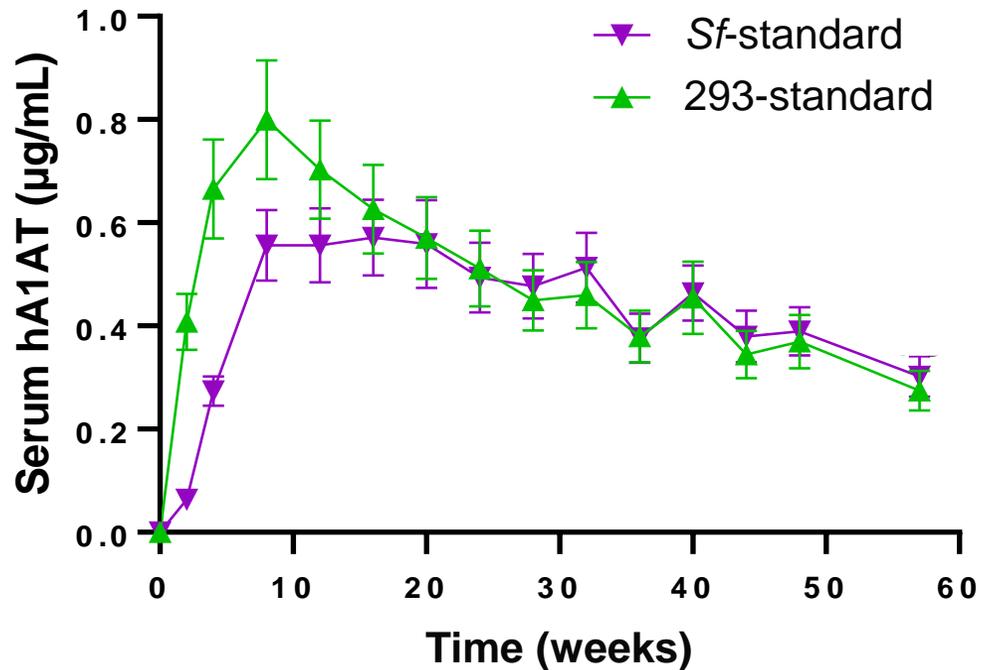
Oversized vector
(~5000 nt)



Standard-sized vector
(4600 nt)

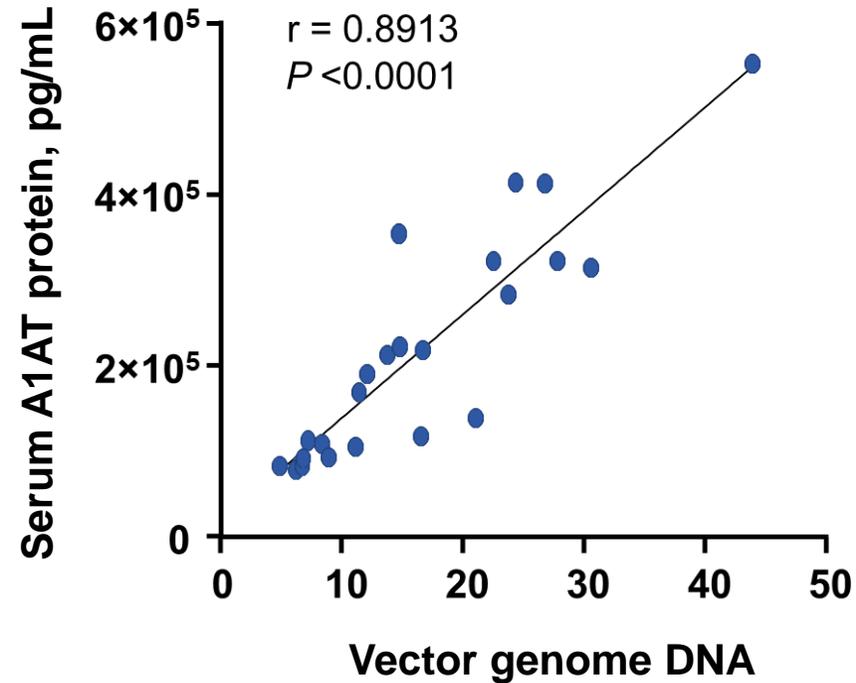
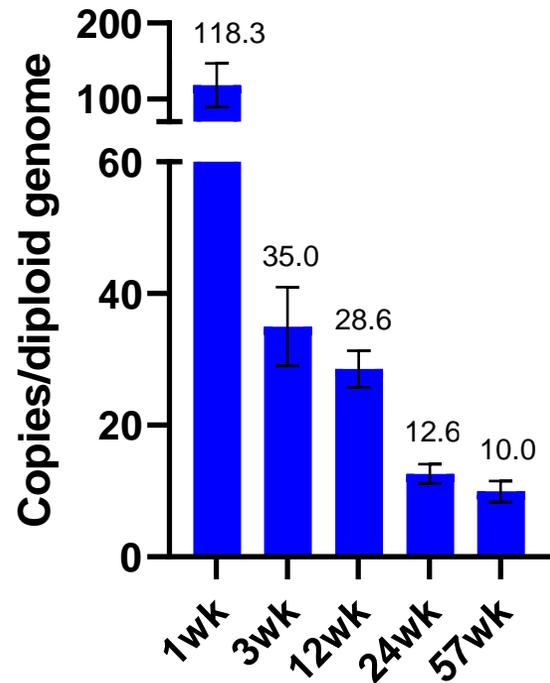


Oversized vectors produced in *Sf* cells demonstrated higher transgene expression long-term compared with 293



** $P < 0.01$; *** $P < 0.001$ using a repeated measure analysis on log transformed serum hA1AT. Data are mean \pm SE. 293, human HEK293 cells; hA1AT, human alpha-1 antitrypsin; SE, standard error; *Sf*, *Spodoptera frugiperda* cells.

Degradation of the vector genome in the liver is the primary mechanism mediating decline in transgene expression with 293-produced vector



- Peak protein expression occurred at 12 weeks

293-produced vectors elicit a heightened immune response compared with *Sf*-produced vectors

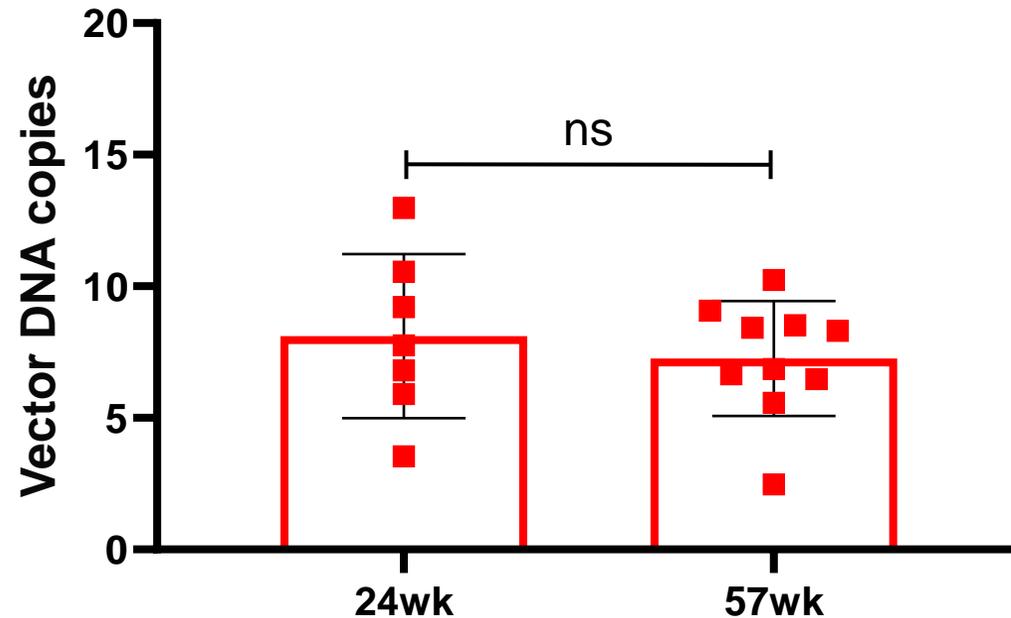
- RNA-seq analysis of liver tissue from mice at 12 weeks post-dose
 - Significantly upregulated pathways in 293-treated mice were associated with innate immune response
 - 293-treated mice also showed higher innate immune responses compared with the vehicle control

Top upregulated pathways	293- vs <i>Sf</i> -produced vector	
	<i>P</i> adj	NES
TNF α signaling via NF κ B	3.02E-08	2.09
G2m checkpoint	0.0416	1.37
Myc targets v2	0.0409	1.69
Myc targets v1	0.0043	1.60
Inflammatory response	0.0409	1.42
INF γ response	0.0071	1.57
TGF β signaling	0.0409	1.58
Allograft rejection	0.0409	1.39
Apoptosis	0.0409	1.44
INF α response	0.0071	1.69
P53 pathway	0.0409	1.37

The Benjamini-Hochberg method was used to adjust *P*-values for differential expression and gene set enrichment analyses. NES determines whether a gene set is moving up (indicating that it is positively regulated) or down (indicating that it is negatively regulated) in the gene rankings for 293 vs *Sf* vector-treated mice (*n* = 7 per group). 293, human HEK293 cells; INF α , interferon- α ; INF γ , interferon- γ ; Myc targets v, Myc targets version; NES, normalised enrichment score; NF κ B, nuclear factor kappa B; *P* adj, adjusted *P*-value; TGF β , transforming growth factor- β ; TNF α , tumour necrosis factor- α ; *Sf*, *Spodoptera frugiperda* cells.

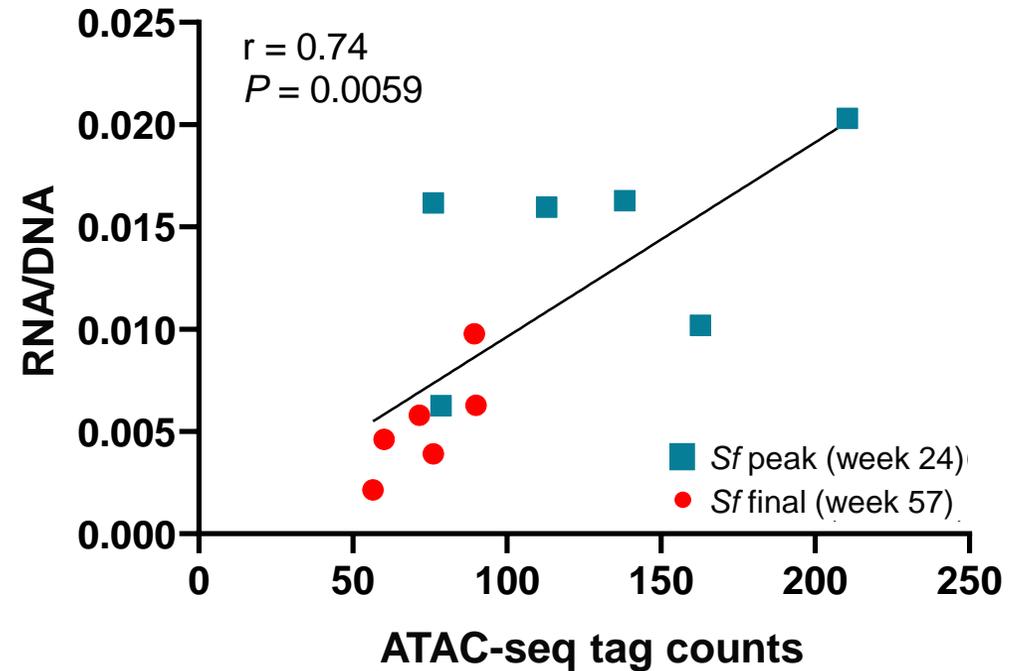
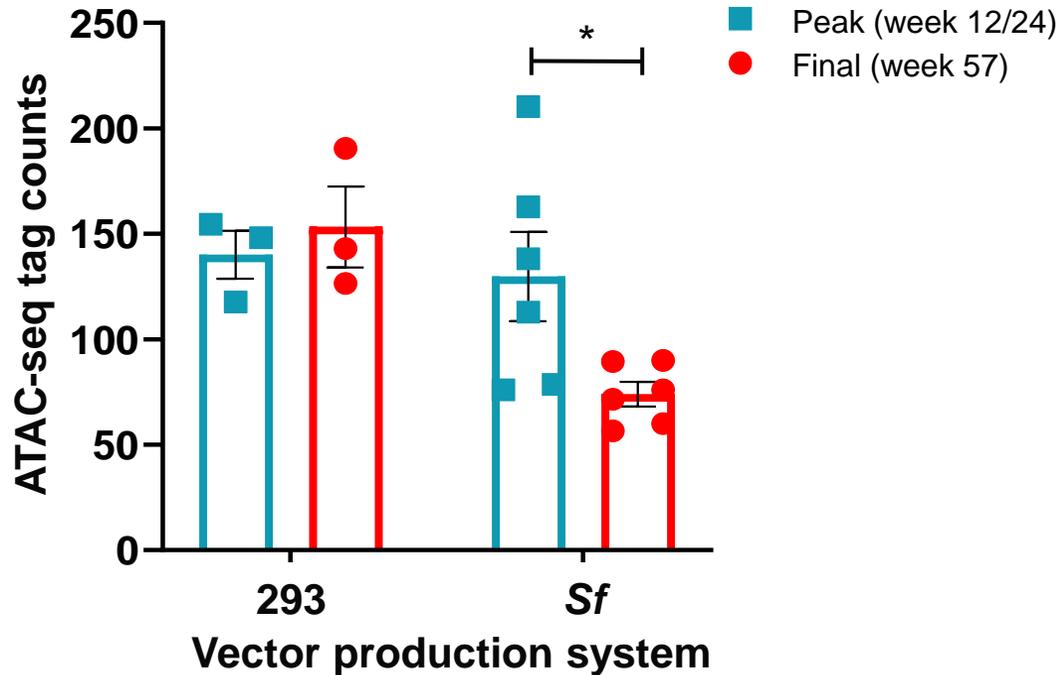
Copy numbers of *Sf*-produced vector genomes did not decrease from peak expression to the final time point

Sf-oversized overall genomes



Ns, not significant using a Welch's t-test. Data are mean \pm SE. Vector DNA copies are per diploid genome. SE, standard error; *Sf*, *Spodoptera frugiperda* cells; wk, week.

A decrease in genome accessibility may impact the decline in transgene expression with *Sf*-produced vector

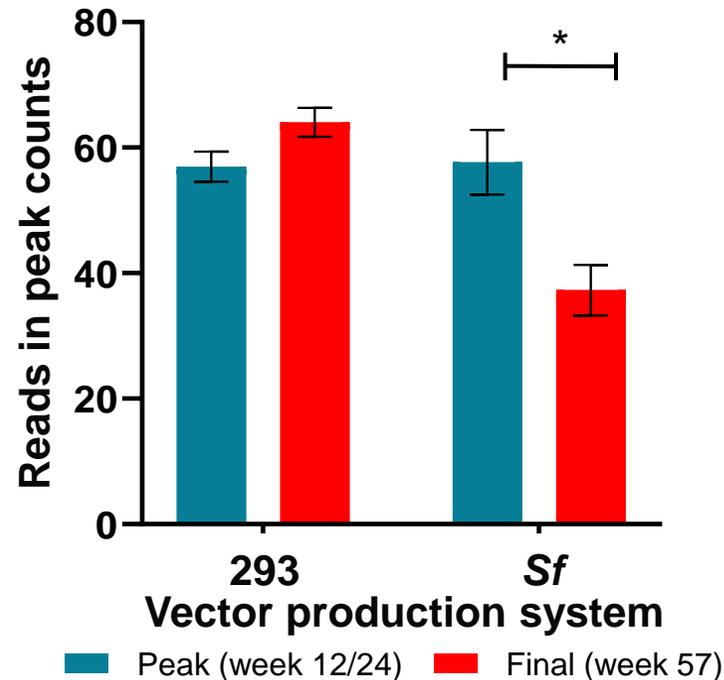
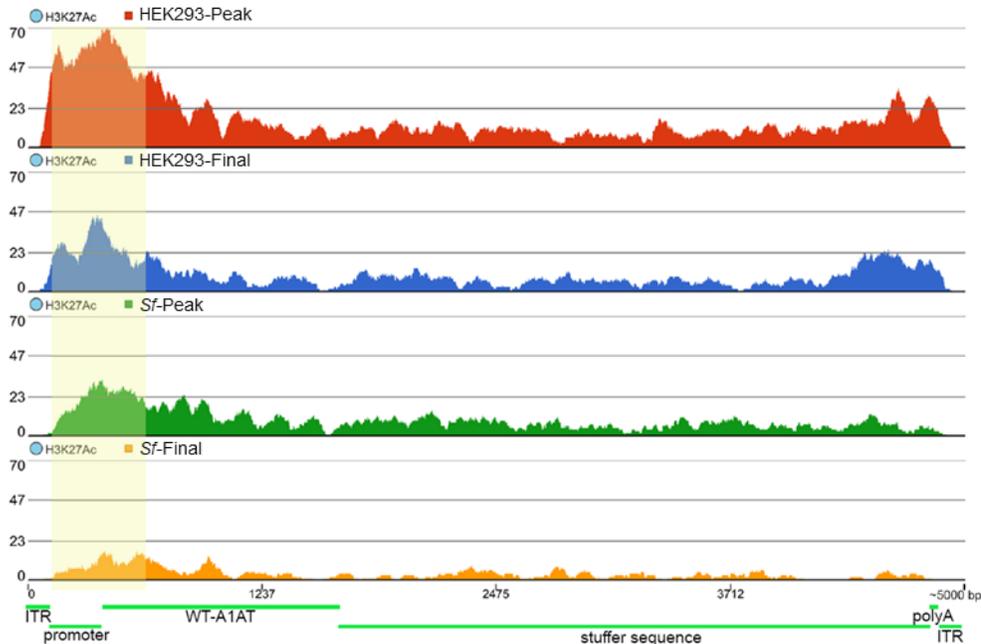


* $P < 0.05$ using a Welch's t-test on the log transformed ATAC-seq tag counts. Data are mean \pm SE. Tag counts are ATAC-seq peak regions in the AAV5 promoter (1–660 bp) corrected for sequencing depth and normalised to vector genome copies.

293, human HEK293 cells; ATAC-seq, assay for transposase-accessible chromatin sequencing; SE, standard error; *Sf*, *Spodoptera frugiperda* cells.

Changes in chromatin and histone modifications mediate decrease in transgene expression for *Sf*-produced vectors

H3K27ac-based epigenetic modifications



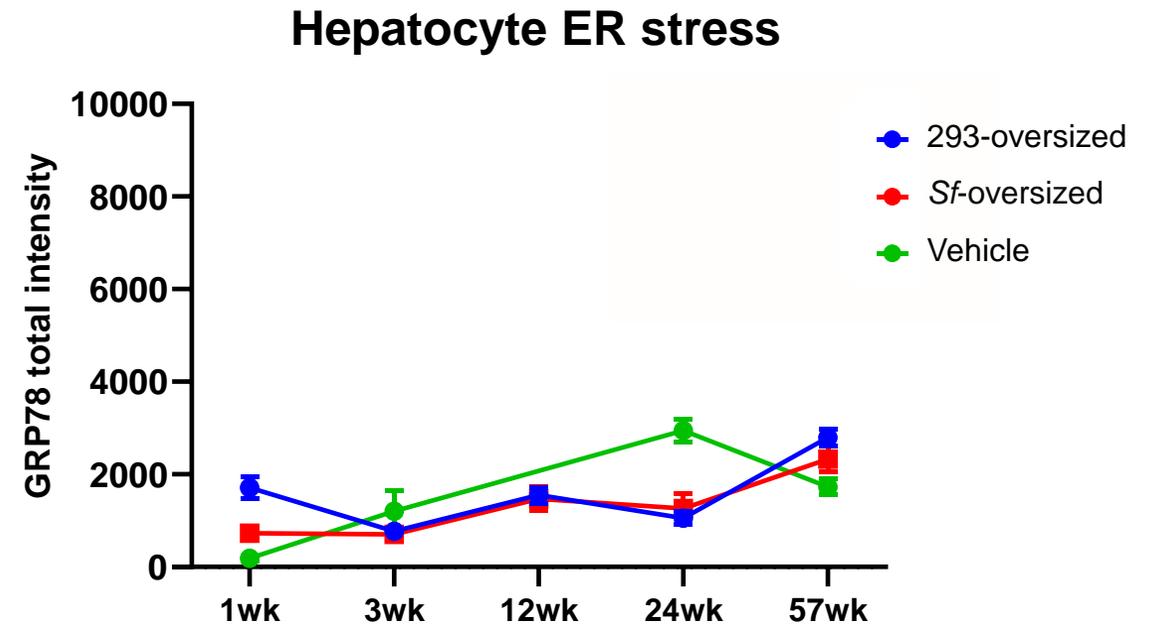
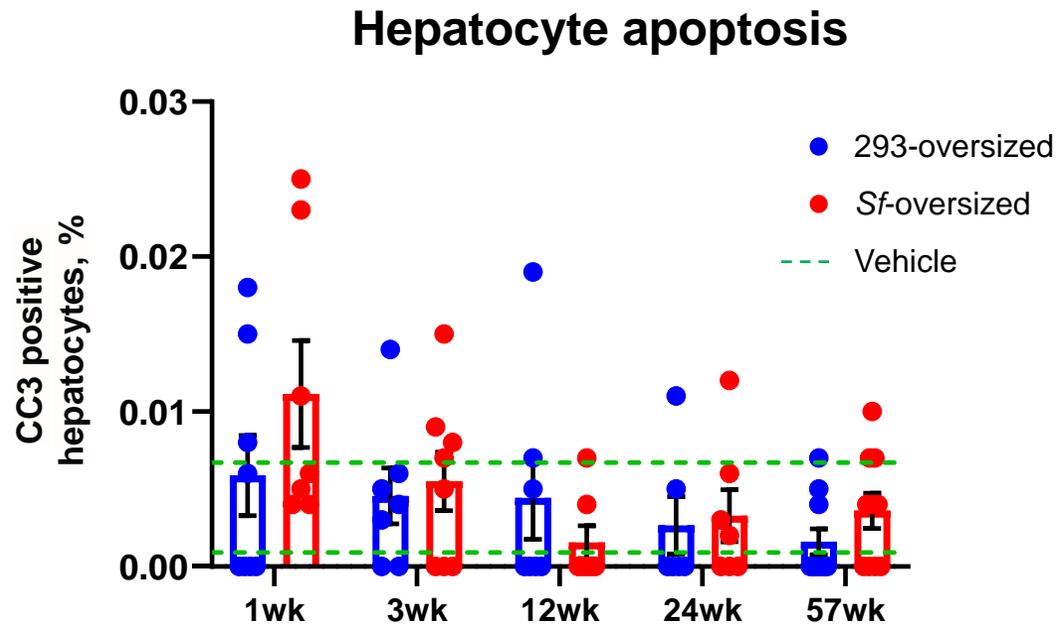
- H3K4me3, another active histone marker, had similar results
- No associations observed with the repressive histone marker H3K9me3

The highlighted yellow portion corresponds to the region where mean and SE peak counts were assessed as shown in the bar graph.

* $P < 0.05$ using a Student's t-test within each cell line. Data are mean \pm SE.

293, human HEK293 cells; H3K27ac, acetylation of lysine 27 on histone H3; ITR, inverted terminal repeat; polyA, polyadenylation signal; SE, standard error; *Sf*, *Spodoptera frugiperda* cells; WT-A1AT, wild-type alpha-1 antitrypsin.

Liver histology evaluations showed minimal changes in hepatocyte apoptosis or ER stress



Data are mean ± SE.

293, human HEK293 cells; CC3, cleaved-caspase 3; ER, endoplasmic reticulum; GRP78, glucose regulatory protein 78; SE, standard error; *Sf*, *Spodoptera frugiperda* cells; wk, week.

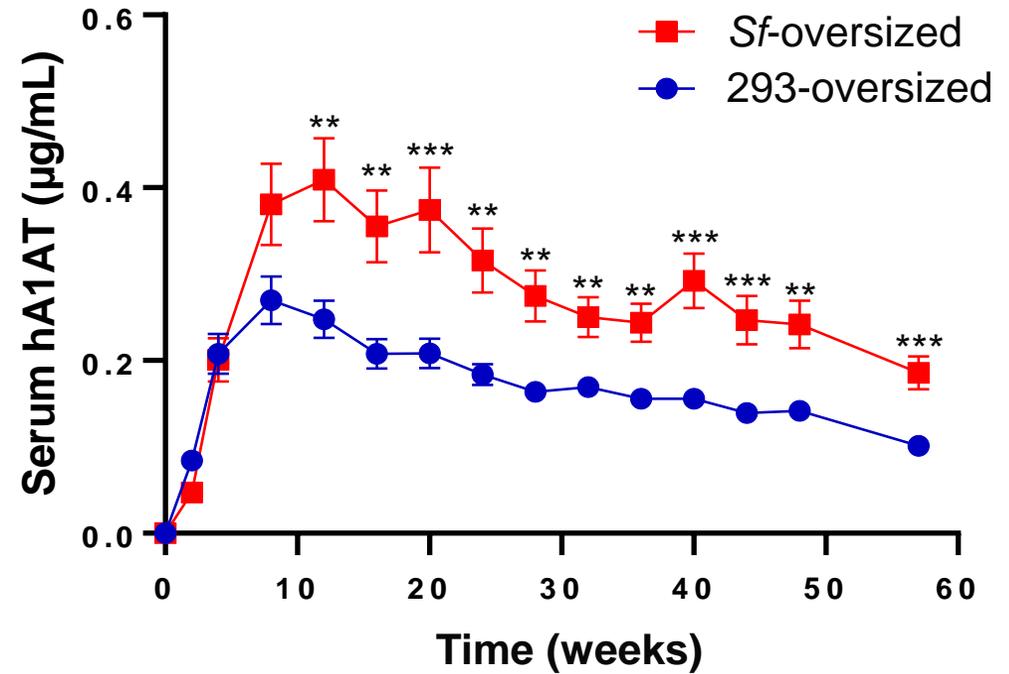
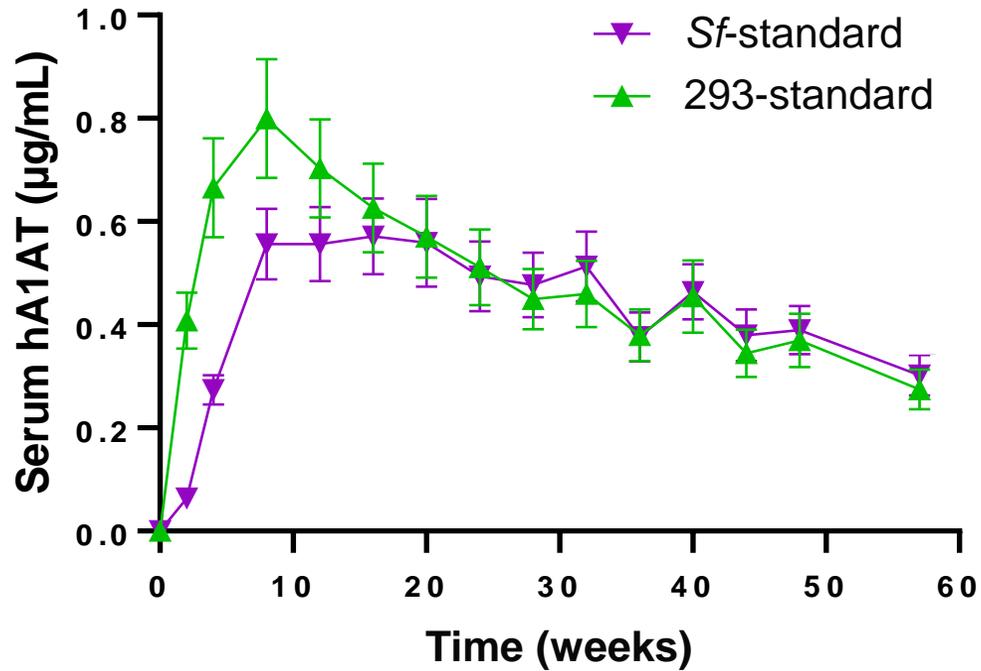
Conclusions

- rAAV5 vectors produced in HEK293 and *Sf* cells showed similar long-term durability of expression in mice, despite distinct mechanisms contributing to the decline in transgene expression over time
- Oversized vectors produced in *Sf* cells may represent a viable alternative to vectors produced in HEK293 cells
- Genome metabolism and more pronounced innate immune responses may mediate the decline in transgene expression by 293-produced vector
- Dynamics of genome accessibility including transcription factors or histone binding and other epigenetic modifications, may distinctly impact durable transgene expression of *Sf*-produced vector
 - Additional analysis of epigenetic regulation of rAAV vectors would deepen our understanding of AAV-mediated transgene expression
- No liver tumours were observed after more than 1 year of follow-up in mice dosed with AAV5-hA1AT vectors produced in either manufacturing system

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Question and answer session



Please direct additional inquiries to Dr. Sylvia Fong via email at sfong@bmrn.com