#### Cellular, Tissue, and Gene Therapies Advisory Committee September 2-3, 2021 Meeting Presentation

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# rAAV Integration: In Vitro & Mice

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### Disclosures

Co-Founder – JAYA Biosciences

Consultant – BioMarin Pharmaceutical

Consultant – Pfizer

Consultant – Taysha Gene Therapies

Scientific Advisory Board – M6P Therapeutics

# rAAV Integration: In Vitro & Mice

### Chronological:

- A) 1997-2001 Prior to the observation of hepatocellular carcinoma (HCC)
- B) 2001-2016 rAAV integration and associated HCC in mice
- C) Conditions affecting rAAV-associated HCC

(1997-2001)

JOURNAL OF VIROLOGY, Nov. 1997, p. 8429–8436 0022-538X/97/\$04.00+0 Copyright © 1997, American Society for Microbiology

Vol. 71, No. 11

### Adeno-Associated Virus Vector Integration Junctions

ELIZABETH A. RUTLEDGE AND DAVID W. RUSSELL\*

- 1) In vitro (HeLa cells) integration analysis
- 2) Random integration
- 3) None of the integrated vectors were fully intact

(1997-2001)

JOURNAL OF VIROLOGY, Dec. 1997, p. 9231–9247 0022-538X/97/\$04.00+0 Copyright © 1997, American Society for Microbiology Vol. 71, No. 12

#### Cellular Recombination Pathways and Viral Terminal Repeat Hairpin Structures Are Sufficient for Adeno-Associated Virus Integration In Vivo and In Vitro

C. C. YANG,<sup>1</sup> X. XIAO,<sup>1</sup> X. ZHU,<sup>1</sup>† D. C. ANSARDI,<sup>1</sup>‡ N. D. EPSTEIN,<sup>2</sup> M. R. FREY,<sup>3</sup> A. G. MATERA,<sup>3</sup> AND R. J. SAMULSKI<sup>1</sup>\*

- 1) Integration analysis in cell lines and a cell-free system
- 2) No obvious site preference for integration.
- 3) No intact ITRs were identified
- 4) Favor actively transcribed regions

(1997-2001)

JOURNAL OF VIROLOGY, July 1999, p. 5438–5447 0022-538X/99/\$04.00+0 Copyright © 1999, American Society for Microbiology. All Rights Reserved. Vol. 73, No. 7

Isolation of Recombinant Adeno-Associated Virus Vector-Cellular DNA Junctions from Mouse Liver

HIROYUKI NAKAI, 1,2,3\* YUICHI IWAKI, MARK A. KAY, AND LINDA B. COUTO1

- 1) In vivo integration analysis (mouse liver)
- 2) Integrated vectors were rearranged (ITRs and vector sequences)
- 3) Two integrants were identified in genes (\alpha1 collagen, rRNA)

(1997-2001)

JOURNAL OF VIROLOGY, Aug. 2001, p. 6969–6976 0022-538X/01/\$04.00+0 DOI: 10.1128/JVI.75.15.6969–6976.2001 Copyright © 2001, American Society for Microbiology. All Rights Reserved. Vol. 75, No. 15

#### Extrachromosomal Recombinant Adeno-Associated Virus Vector Genomes Are Primarily Responsible for Stable Liver Transduction In Vivo

HIROYUKI NAKAI, STEPHEN R. YANT, THERESA A. STORM, SALLY FUESS, LEONARD MEUSE, AND MARK A. KAY\*

- 1) Extrachromosomal rAAV is the primary source of expression
- 2) Only 5-10% of rAAV vector are integrated into the host genome
- 3) Low level integration, increased safety profile

# Summary 1 (1997-2001)

- 1) Relatively low level of integration
- 2) Chromosome/vector junctions are within or near ITRs
- 3) ITRs are rearranged
- 4) No or minimal homology between cellular and vector sequences
- 5) No integration 'hot spots' were identified
- 6) Integration mechanism is unknown
- 7) No toxicity observed up to this point

(2001-2016)

Gene Therapy (2001) 8, 1343–1346 © 2001 Nature Publishing Group All rights reserved 0969-7128/01 \$15.00

www.nature.com/gt

#### **BRIEF COMMUNICATION**

## Observed incidence of tumorigenesis in long-term rodent studies of rAAV vectors

A Donsante<sup>1</sup>, C Vogler<sup>2</sup>, N Muzyczka<sup>3</sup>, JM Crawford<sup>4</sup>, J Barker<sup>5</sup>, T Flotte<sup>3</sup>, M Campbell-Thompson<sup>4</sup>, T Daly<sup>1,6</sup> and MS Sands<sup>1</sup>

- 1) Long-term (18mo) study in MPSVII mouse
- 2) IV administration at birth (~1x10<sup>14</sup> vg/kg)
- 3) Persistent expression, dramatic clinical/behavioral improvements
- 4) 3/5 rAAV-treated animals had HCC at 18 months of age
- 5) Impossible to determine if rAAV caused HCC

(2001-2016)

# Adeno-associated virus vectors integrate at chromosome breakage sites

Daniel G Miller<sup>1</sup>, Lisa M Petek<sup>2</sup> & David W Russell<sup>2,3</sup>

NATURE GENETICS VOLUME 36 | NUMBER 7 | JULY 2004

- 1) rAAV does not increase mutation rate
- 2) rAAV integrates at spontaneous or induced double strand breaks.

(2001-2016)

JOURNAL OF VIROLOGY, Mar. 2005, p. 3606–3614 0022-538X/05/\$08.00+0 doi:10.1128/JVI.79.6.3606–3614.2005 Copyright © 2005, American Society for Microbiology. All Rights Reserved. Vol. 79, No. 6

### Large-Scale Molecular Characterization of Adeno-Associated Virus Vector Integration in Mouse Liver

Hiroyuki Nakai,<sup>1</sup>\* Xiaolin Wu,<sup>2</sup> Sally Fuess,<sup>1</sup> Theresa A. Storm,<sup>1,3</sup> David Munroe,<sup>2</sup> Eugenio Montini,<sup>4</sup>† Shawn M. Burgess,<sup>5</sup> Markus Grompe,<sup>4,6</sup> and Mark A. Kay<sup>1,3</sup>

- 1) Analyzed 347 rAAV integration sites in mouse liver
- 2) Integration 'hot spot' was found in rRNA gene repeats
- 3)>50% of integrations occurred near transcription start sites or CpG islands

(2001-2016)

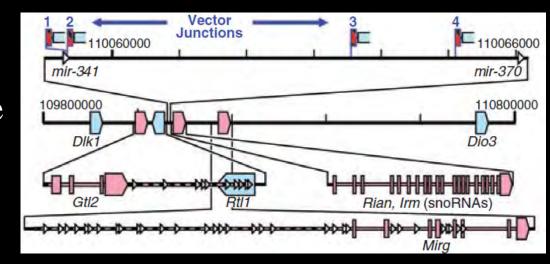
# AAV Vector Integration Sites in Mouse Hepatocellular Carcinoma

Anthony Donsante, \*\* Daniel G. Miller, \*\* Yi Li, \*\* Carole Vogler, \*\* Elizabeth M. Brunt, \*\*

David W. Russell, \*\*, \*\* Mark S. Sands\*\*, \*\*

SCIENCE VOL 317 27 JULY 2007

- 1) 15/34 rAAV-treated mice developed HCC (6/18 MPSVII, 9/16 WT)
- 2) Integrated rAAV sequences were isolated from HCC samples from 4 mice
- 3) In each case, rAAV integrated in the mouse *Rian* locus
- 4) Several downstream genes and µRNAs were dysregulated



(2001-2016)

# Long-Term Correction of Sandhoff Disease Following Intravenous Delivery of rAAV9 to Mouse Neonates

Jagdeep S Walia<sup>1,2,3,4</sup>, Naderah Altaleb<sup>2</sup>, Alexander Bello<sup>5,6</sup>, Christa Kruck<sup>2</sup>, Matthew C LaFave<sup>7</sup>, Gaurav K Varshney<sup>7</sup>, Shawn M Burgess<sup>7</sup>, Biswajit Chowdhury<sup>2</sup>, David Hurlbut<sup>8</sup>, Richard Hemming<sup>2</sup>, Gary P Kobinger<sup>5,6</sup> and Barbara Triggs-Raine<sup>2,3,4</sup>

- 1) 8/10 rAAV-treated Sandhoff mice developed HCC
- 2) IV injection of rAAV in neonatal mice (2.5 x 10<sup>14</sup> vg/kg)
- 3) Several tumors had rAAV integrations in the Rian locus

(2001-2016)

# Vector design influences hepatic genotoxicity after adeno-associated virus gene therapy

Randy J. Chandler,<sup>1</sup> Matthew C. LaFave,<sup>2</sup> Gaurav K. Varshney,<sup>2</sup> Niraj S. Trivedi,<sup>3</sup> Nuria Carrillo-Carrasco,<sup>4</sup> Julien S. Senac,<sup>1</sup> Weiwei Wu,<sup>5</sup> Victoria Hoffmann,<sup>6</sup> Abdel G. Elkahloun,<sup>5</sup> Shawn M. Burgess,<sup>2</sup> and Charles P. Venditti<sup>1</sup>

J Clin Invest (2015) 125:870-880

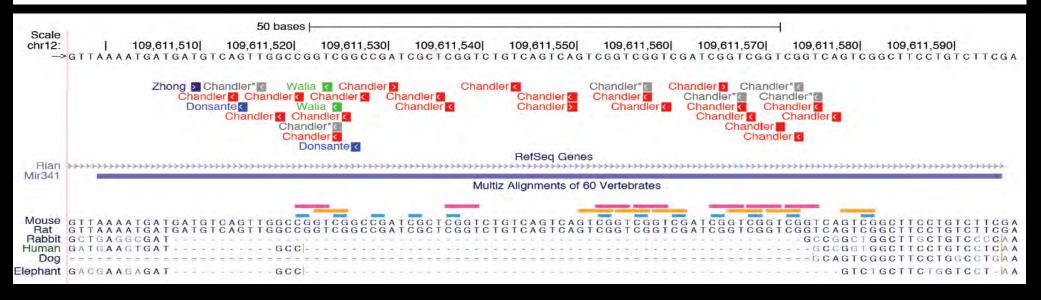
- 1) 64/95 mice treated with rAAV (1x10<sup>14</sup> vg/kg) at birth developed HCC
- 2) Confirmed the *Rian* locus as a 'hot spot' for rAAV integration
- 3) HCC appears to be dose-dependent
- 4) Strong promoter/enhancer combinations increased incidence of HCC
- 5) HCC independent of transgene
- 6) HCC independent of genotype

(2001-2016)

# Genotoxicity in Mice Following AAV Gene Delivery: A Safety Concern for Human Gene Therapy?

Randy J Chandler<sup>1</sup>, Matthew C LaFave<sup>2</sup>, Gaurav K Varshney<sup>2</sup>, Shawn M Burgess<sup>2</sup> and Charles P Venditti<sup>1</sup>

Molecular Therapy vol. 24 no. 2 february 2016



- 1) Compilation of rAAV integration sites across several studies
- 2) Many *Rian* integrations occur in a region (~60bp) unique to rodents

# Summary 2 (2001-2016)

- 1) Systemic delivery of a rAAV vector can cause HCC in mice
- 2)rAAV integration into the mouse *Rian* locus is associated with HCC
- 3) Many rAAV integrants are located in *Rian* sequences unique to rodents
- 4) Highest frequency of HCC if administered during the newborn period
- 5) Strong promoter/enhancer combinations increase frequency
- 6) rAAV-associated HCC appears to be dose-dependent
- 7) Low frequency if administered in adult animals
- 8) No HCC following CNS-directed rAAV-mediated gene therapy

(recent findings)

# Liver Injury Increases the Incidence of HCC following AAV Gene Therapy in Mice

Dhwanil A. Dalwadi,<sup>1,2</sup> Laura Torrens,<sup>3</sup> Jordi Abril-Fornaguera,<sup>3</sup> Roser Pinyol,<sup>3</sup> Catherine Willoughby,<sup>3</sup> Jeffrey Posey,<sup>2</sup> Josep M. Llovet,<sup>3,4,5</sup> Christian Lanciault,<sup>6</sup> David W. Russell,<sup>7,8</sup> Markus Grompe,<sup>2</sup> and Willscott E. Naugler<sup>1</sup> Molecular Therapy Vol. 29 No 2 February 2021

- 1) Increased incidence of HCC in <u>adult</u> mice with non-alcoholic fatty liver
- 2) Rian-targeted construct 100% penetrance
- 3) Non-targeted construct 50% penetrance

(recent findings)

### Enhanced Efficacy and Increased Long-Term Toxicity of CNS-Directed, AAV-Based Combination Therapy for Krabbe Disease

Yedda Li,<sup>1</sup> Christopher A. Miller,<sup>1</sup> Lauren K. Shea,<sup>1</sup> Xuntian Jiang,<sup>1</sup> Miguel A. Guzman,<sup>2</sup> Randy J. Chandler,<sup>3</sup> Sai M. Ramakrishnan,<sup>1</sup> Stephanie N. Smith,<sup>3</sup> Charles P. Venditti,<sup>3</sup> Carole A. Vogler,<sup>2</sup> Daniel S. Ory,<sup>1</sup> Timothy J. Ley,<sup>1,4</sup> and Mark S. Sands<sup>1,4</sup>

Molecular Therapy Vol. 29 No 2 February 2021 @

- 1) HCC development following CNS-directed rAAV-mediated gene therapy
- 2) Combination therapy (HSCT transplantation + small molecule drug)
- 3)~95% penetrance in combination-treated Krabbe & WT mice

### Targeted Sequence Capture

Table 1. AAV Integration Sites in or near Cancer- and Cell-Growth/Death-Associated Genes							
Animal ID	Agea (months)	Chr	Integration <sup>b</sup> Start Site (bp)	Read Counts	Gene	Gene Description	Reference
6657	13.4	12	109643597°	857	Rian	microRNA cluster	33
6675	7	12	109618074 <sup>c</sup>	1,390	Rian	microRNA cluster	33
	14.7	6	99150006 <sup>c</sup>	4,883	Foxp1	Forkhead protein, tumor suppressor	28,29
		8	39104763°	199	Tusc3	endoplasmic reticulum (ER) protein, candidate tumor suppressor	30
6722	17.6	12	109631801 <sup>c</sup>	3,060	Rian	microRNA cluster	33
		7	75627402°	1,682	Akap 13	A-kinase anchor protein, double oncogene homology, breast cancer	25,34
6815	14.7	12	109609803°	1,684	Rian	microRNA cluster	33
6824		12	109631309 <sup>c</sup>	213	Rian	microRNA cluster	33
	15.0	1	$192215900^{c}$	55	Kcnh1	K <sup>+</sup> channel, increased expression confers growth advantage	26,35
		6	81973046 <sup>d</sup>	87	Evala	regulator of programmed cell death	27,36
6828	13.1	12	109625075°	3,737	Rian	microRNA cluster	33
6902		12	109613953 <sup>c</sup>	85	Rian	microRNA cluster	33
	14.9	12	109671965 <sup>ε</sup>	825	Rian	microRNA cluster	33
	****	6	94142817 <sup>c</sup>	65	Magil	membrane-associated guanylate kinase, candidate tumor suppressor	31,32
7025	15.5	12	109615046	1,430	Rian	microRNA cluster	33
7045	16.8	12	109606198 <sup>c</sup>	3,519	Rian	microRNA cluster	33
7046	14.2	12	109611352 <sup>c</sup>	387	Rian	microRNA duster	33

- 1) All tumors had rAAV integrations in the Rian locus
- 2) 4/10 tumors had rAAV integrations in candidate tumor suppressors, cancer-associated genes, or genes that regulate cell growth or death

### Conclusions

- 1) rAAV vectors can stably integrate into the mouse genome ( $\leq 10\%$ )
- 2) The vast majority of integrated vectors appear to be grossly rearranged
- 3) The integration mechanism is unknown
- 4) Integration into the mouse *Rian* locus is associated with HCC
- 5) HCC formation appears to be dose dependent
- 6) Strong promoter/enhancer combinations increase HCC frequency
- 7) Low frequency of HCC in adult animals
- 8) Low frequency of HCC following CNS-directed gene therapy
- 9) Liver injury (eg. NAFL) can exacerbate HCC phenotype
- 10) Adjunct therapies with mild oncogenic potential exacerbate HCC phenotype

## Acknowledgements

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### <u>NIH</u>

Randy Chandler Charles Venditti

### University of Washington

Daniel Miller David Russell

### Additional Information

Recent round table discussion on rAAV integration

Sponsored by the American Society of Gene and Cell Therapy (ASGCT)

August 18, 2021

https://www.youtube.com/watch?v=L\_4luK3fNU0