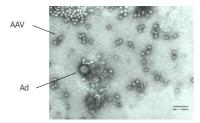
Adeno-associated virus (AAV)

Taxonomy

Family: Parvovirus Subfamily: Parvovirinae Genus: Dependovirus Type: AAV 1-12



Morphology

Particles are icosahedral, non-enveloped, 18-26 nm diameter, 50% protein (VP1-3) 50% DNA. Resistant to inactivation by solvents, pH and heat.

Genome

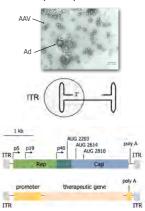
Linear, non-segmented, ssDNA \sim 5 kb. AAVs package equal amounts of (+) and (-) strands.

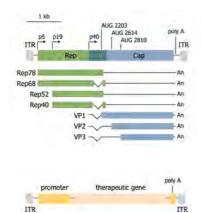


Xie et al 2002

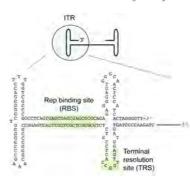
Gene transfer using Adeno-Associated Virus (AAV) vectors

- 1. Based on a broadly diffuse, non pathogenic virus
- 3. Vectors do not express viral proteins (not inflammatory and not immunogenic); long term persistence
- 4. Expression of the therapeutic gene can be driven by any desirable promoter
- 5. High titer vector preparations can be obtained by virion purification
- 6. Cells are transduced at high multiplicity of infection; mixing of different rAAV preparations results the simultaneous expression of gene combinations in vivo





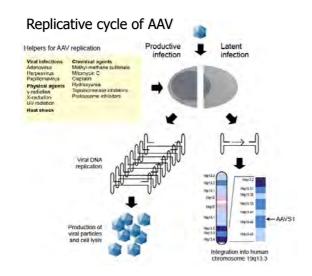
AAV Inverted Terminal Repeat (ITR)

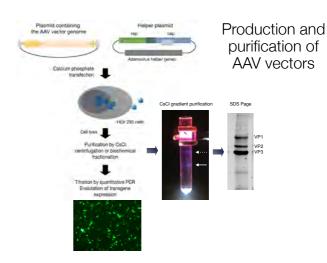


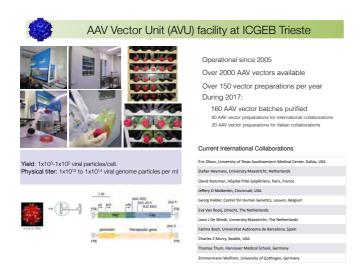
The AAV coding region is flanked by two ~145-nt-long ITRs, having an internal complementarity stretch in their first 125 nt and thus forming a T-shaped halipin structure, identical at the two viral ends.

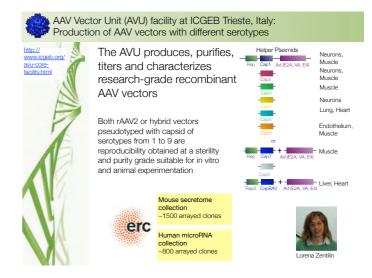
This palindromic sequence is the only cis-acting genetic element necessary for all AAV functions, including:
- viral DNA replication
- site-specific integration into the host cell DNA packaging of wirions

The first two activities (replication and integration) require the presence of Rep68 or Rep78 proteins, which specifically bind a sequence within the ITR, the Rep binding site (RBS), and cleave in a site- and strand-specific manner at the terminal resolution site (TRS) located 13 nucleotides (nt) upstream of the RBS. An almost identical sequence in human chromosome 19q.13.4 represents the minimal sequence necessary and sufficient for AAV site-specific integration



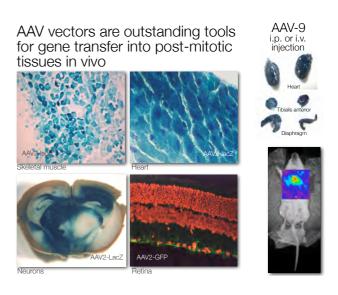


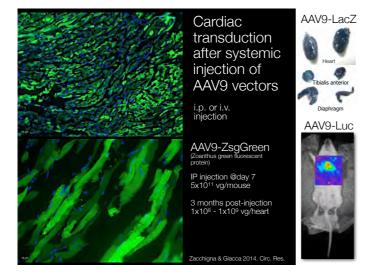




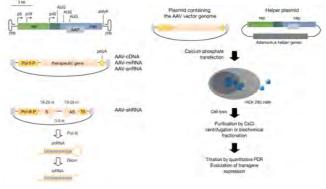
The origin of common AAV isolates, their receptors and tissue tropism







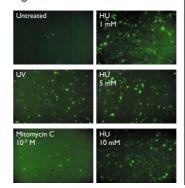




Tropism of AAV for postmitotic cells in vivo.

Why?

Induction of rAAV-GFP transduction by genotoxic agents in wt CHO cells



Helpers for AAV replication

Viruses Adenovirus (E4-ORF 6) HSV-1

Physical agents
γ-radiation
X-ray
UV
Heat shock

Chemical agents

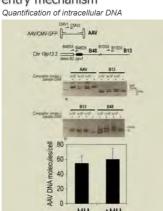
nemical agents
Methyl methan sulfonate
Mitomycin C
Cisplatinum
Hydroxyurea
Topoisomerase inhibitors
(novobiocin, etoposide, campthotecin)
Protease inhibitors



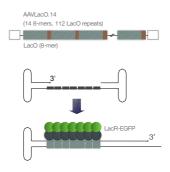
Zentilin et al. J. Virol. 2001

Hydroxyurea induces permissivity to rAAV transduction through a post-entry mechanism

Efficiency of transduction MRC + 1 mM HU + AAV/GFP MRC + AAV/GFP 40 30 20 10 -HU

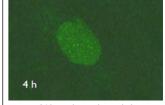


An intracellular reporter to visualize nuclear dsAAV DNA



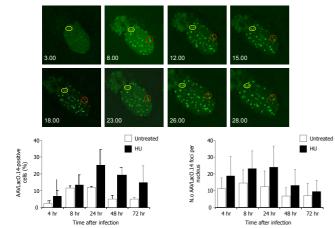
Formation of nuclear dsDNA AAV foci after transduction of poorly permissive cells

MRC/GFP-LacR cells

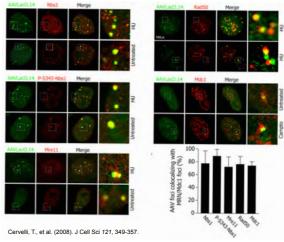


24 hour observation period

Kinetics of formation of AAVLacO.14 foci

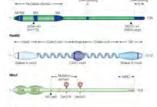


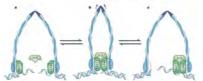
AAVlacO.14 foci co-localize with DDR foci



MRN complex

- Multisubunit nuclease composed of Mre11, Rad50 and Nbs-1
- Binds both ss and ds DNA and has pivotal role in sensing damaged or hairpin structured DNA

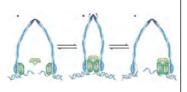




D'Amours and Jackson, Nat Rev Mol Cell Biol, 3, 317-327, 2002

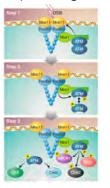
MRN complex

- Multisubunit nuclease composed of Mre11, Rad50 and Nbs-1
- Binds both ss and ds DNA and has a pivotal role in sensing damaged or hairpin structured DNA and process it



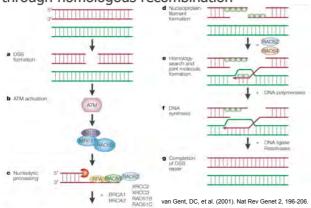
D'Amours and Jackson, Nat Rev Mol Cell Biol, 3, 317, 2002

MRN complex activates ATM and checkpoint signalling

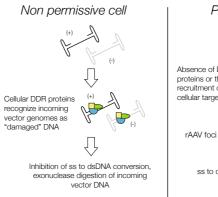


Lee JH, Paull TT, Science 308, 551, 2005

Double-stranded DNA break repair through homologous recombination



Cellular DNA Damage Response (DDR) proteins restrict AAV transduction



"The DDR inhibition model"

Permissive cell



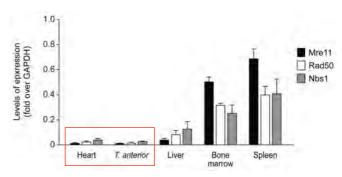
ss to dsDNA conversion and efficient transduction

Zentilin L et al. 2001. J Virol 75, 12279 Cervelli T et al. 2008. J Cell Sci 121, 349 Schwartz RA et al. 2007. J Virol 81, 1293 Lovric J et al. 2012. Mol Ther 20, 2087

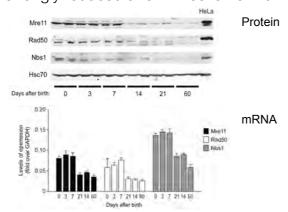
Why are post-mitotic cells permissive in vivo?

Post-mitotic tissues express low levels of MRN proteins in vivo

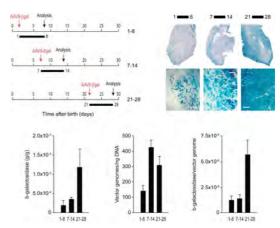




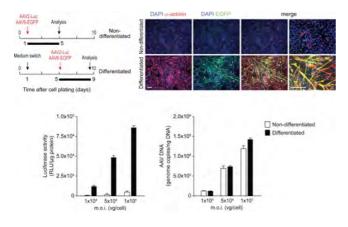
Cardiac MRN complex expression is strongly reduced after 2 weeks from birth

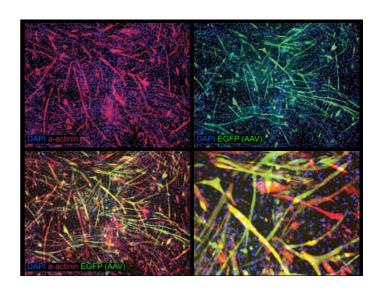


Cardiomyocyte permissivity to AAV transduction increases after birth



C2C12 differentiation increases permissivity to AAV transduction



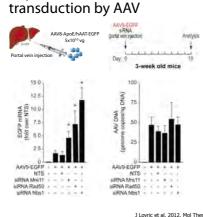


AAVB-EGFP
Mre11 siRNA

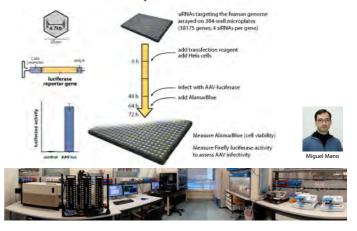
AAVB-EGFP
RadSo siRNA

AAVB-EGFP
Nbs1 siRNA

Transient knockdown of MRN improves in vivo liver transduction by AAV

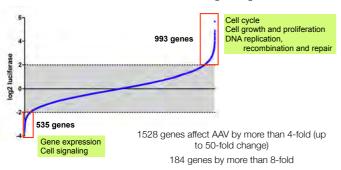


High-throughput screening of a whole genome siRNA library for AAV transduction

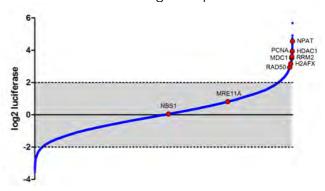


Molecular determinants of AAV transduction

GENOME-WIDE SCREENING - 18175 gene targets

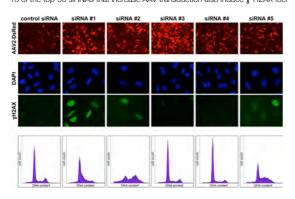


Molecular determinants of AAV transduction: enrichment for genes involved in the DNA Damage Response

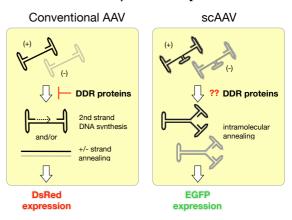


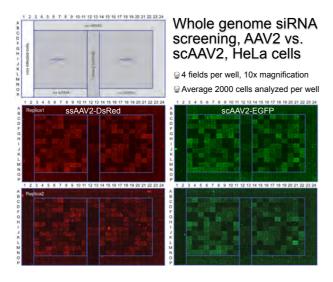
Screening for the induction of yH2AX foci by siRNAs modulating AAV permissivity

5 of the top-10 siRNAs that increase AAV transduction also induce γ -H2AX foci 13 of the top-50 siRNAs that increase AAV transduction also induce γ -H2AX foci



Self-complementary AAV





Summary

- Cellular DNA Damage Reponse (DDR) proteins restrict AAV transduction
- Lack of MRN correlates with AAV permissitivity of post-mitotic tissues in vivo
- Results of whole genome, siRNA screening for AAV permissivity: 900+ genes improving AAV transduction; several genes involved in DNA repair or required for maintenance of DNA integrity
- No significant difference between ssAAV and scAAV to siRNAs inducing AAV permissivity