AAV Genome Sequencing





GENEWIZ[®] Multiomics & Synthesis Solutions from Azenta Life Sciences offers a suite of in-depth next generation sequencing solutions that can be leveraged for adeno-associated virus (AAV) applications to provide a streamlined approach to AAV-based gene therapy research. Using Illumina[®] short-read and PacBio[®] long-read (>10kb) sequencing, these solutions enable high-throughput monitoring of rAAV quality throughout the AAV gene therapy workflow – from initial construct assembly to host transfection.



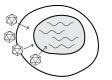
Transgene expression cassettes are cloned into AAV vectors.



AAV Gene Therapy Work Flow



packaged with the transgene.



rAAV is delivered to host cells.



Transfected cells express the transgenic protein in the host.



Confirm Plasmid Sequences

Perform highly-accurate confirmation of AAV plasmids and inserts using:

- Illumina short-read DNA sequencing
- PacBio long-read DNA sequencing

PCR + Sanger sequencing is also available

Validate Packaged Material

Confirm rAAV vector sequences and identify DNA contaminants in AAV preparation with:

Short-read or long-read whole genome sequencing

Now offering single-stranded to double-stranded conversion of AAV genomes

Quantify AAV Expression

Measure AAV expression of the target gene post-infection using:

- Targeted sequencing
- RNA sequencing at the RUO and CLIA levels

qPCR is also available

Monitor Host Response

Stratify and characterize host immune response during and after infection via:

- RNA sequencing
- Single-cell sequencing
- Spatial genomics (i.e., digital spatial profiling)

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AAV Genome Sequencing Data Analysis

Variant Calling Distribution of variant allele frequency as compared to the reference AAV genome



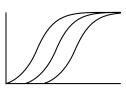
Contamination Testing Mapping distribution of on- and off-target packaged DNA sequences

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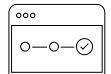
ITR Truncation Distribution of truncation hot spots across the AAV genome

Features & Benefits

- **Trusted partner** supporting over 150 global pharmaceutical and biotech customers in their cell and gene therapy research
- **RUO- and CLIA-grade**, high-capacity, and highthroughput sequencing for projects of virtually any size and sample type
- Optimized, automated, and scalable workflows with stringent quality control producing high-quality, consistent results
- **Customized and extensive** bioinformatics solutions with interactive, hands-on analysis



Superior Data Quality Exceeds manufacturer's benchmarks



Real-Time Project Updates
Through our online system



Ph.D.-Level Support At every step

NGS Platforms

GENEWIZ Multiomics & Synthesis Solutions is a certified service provider of Illumina NGS platforms and the PacBio Sequel[®].







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