

PCR Methods for the Characterization of AAV-based Drug Product from Titer to Potency

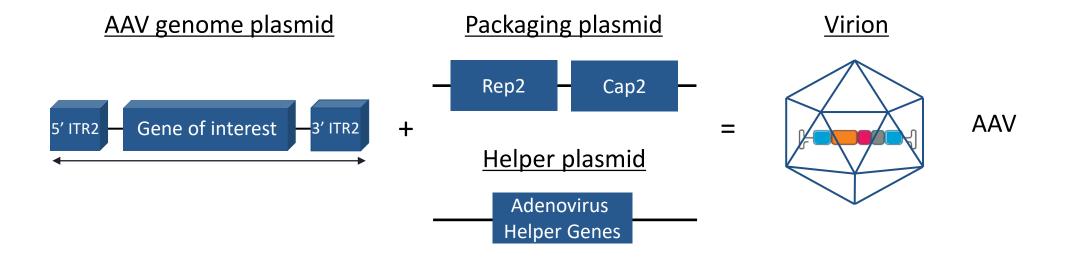
Pavlina Wolf



Gene Therapy Analytical Development 2022

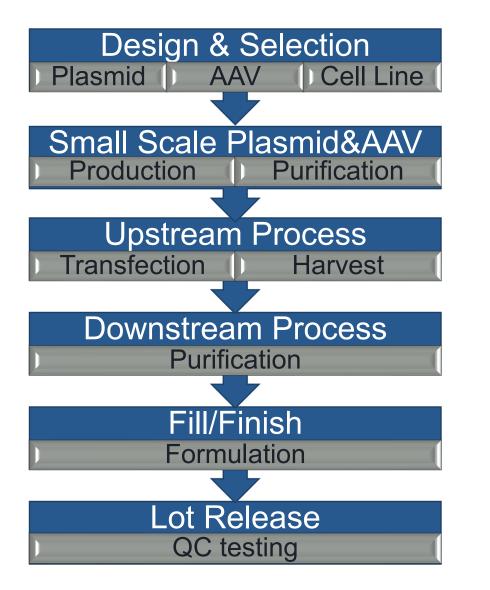
Adeno-Associated Virus (AAV) is an Ideal Vector for Gene Transfer

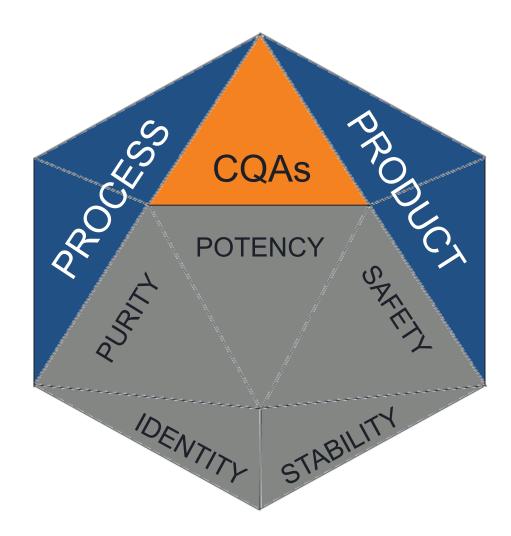
- AAV is a small (4.7 Kb), single stranded, non-pathogenic DNA virus
- Transduces both dividing and non-dividing cells
- Long-term transduction following a single administration
- Different AAVs exhibit distinct tropism for various tissues and organs in different species





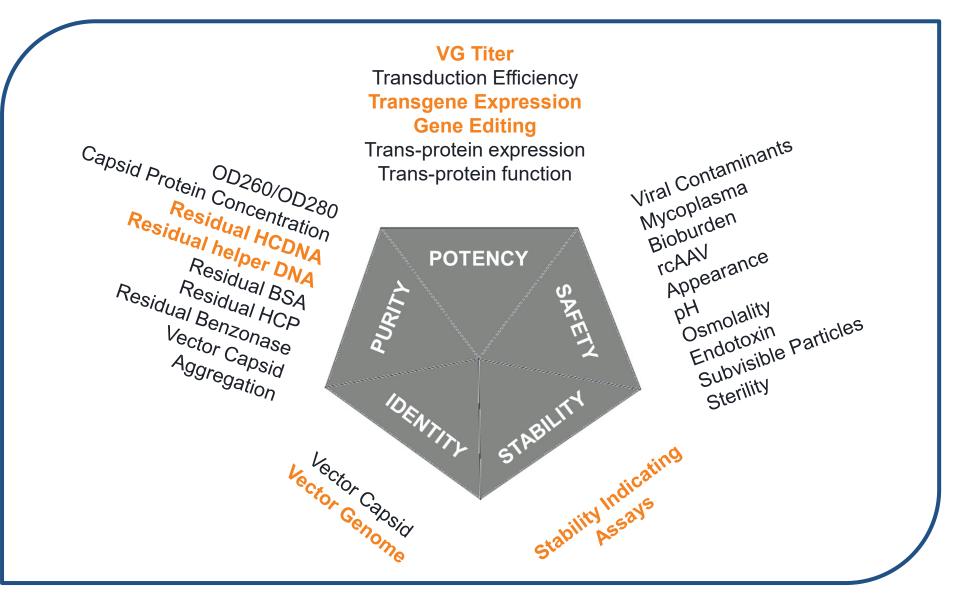
rAAV CQAs Emerge as a Function of Vector and Process Design





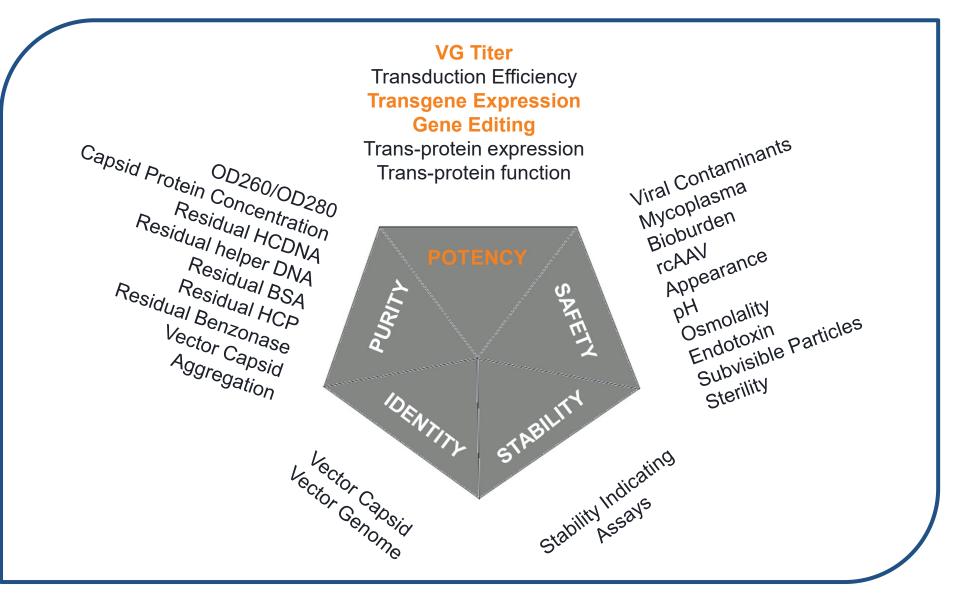


PCR Methods in a Typical QC Testing Panel for AAV Lot Release



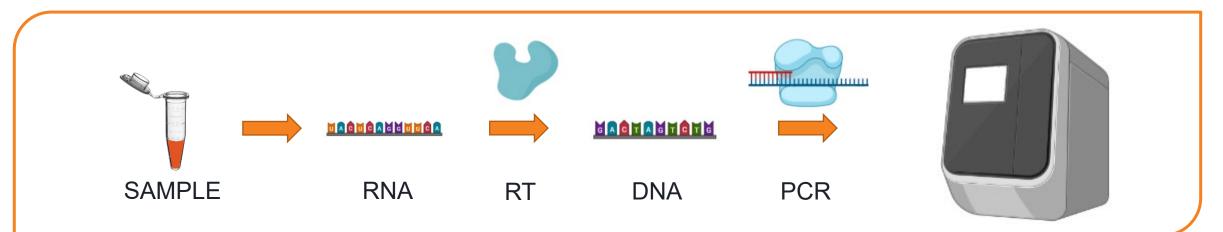


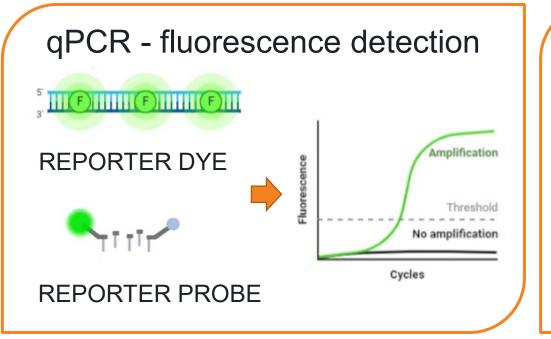
PCR Methods for Potency Evaluation



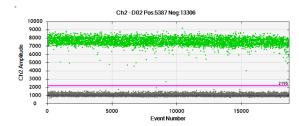


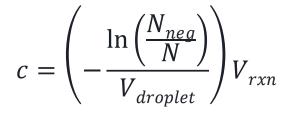
Two Main Technologies for PCR Quantitation





ddPCR - direct counting of positive partitions adjusted for probability of cooccupancy with random distribution



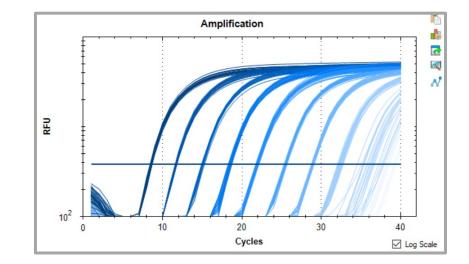


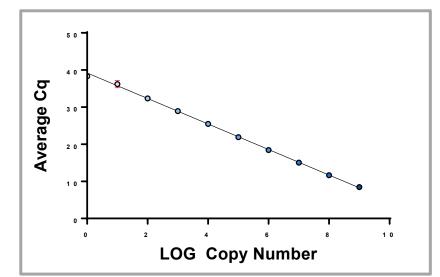


Real Time PCR Quantitation

qPCR¹

- 1. Estimates target abundance against a calibration curve
 - Requires standards of known concentration
- 2. Precision depends on multiple variables from instrument to inhibitors
 - Measure 1.25-1.5-fold differences at best²
 - Assay precision deteriorates at low copy number (CN)
 - Sensitive to effect of inhibitors and differences in amplification efficiency
- 3. Dynamic range of ~9 logs
 - > ddPCR
- 4. Rare targets may not amplify
- 5. Multiplexing for \geq 4 targets

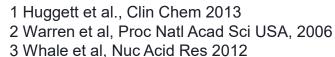




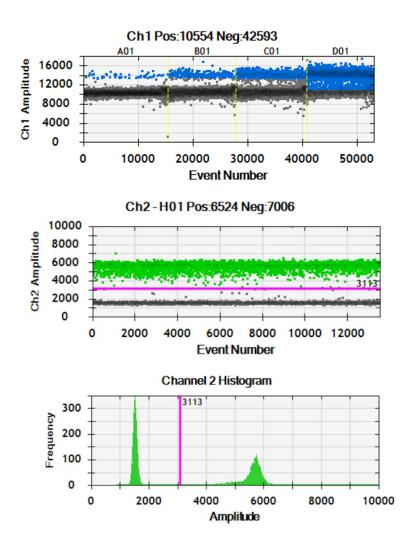
Droplet Digital PCR Quantitation

ddPCR¹

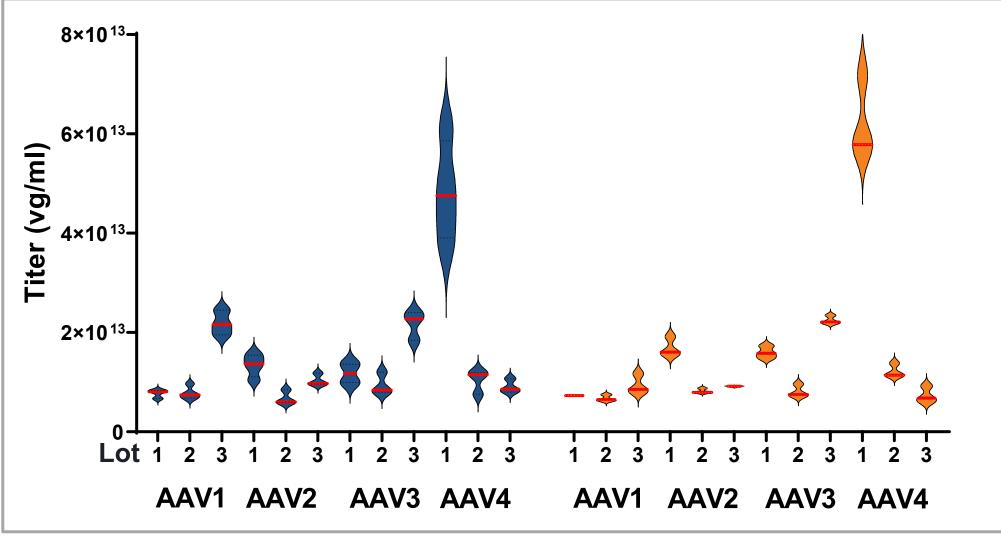
- 1. Estimates target abundance from positive to total partition ratio²
 - Requires only known volume. Prior estimate of titer helpful but not essential
- 2. Precision \approx # of partitions counts
 - Measure differences as low as <1.2-fold³
 - Assay precision drops at both ends
 - Tolerant of inhibitors
- 3. Dynamic range \approx # of partitions
 - In practice < qPCR (~5 logs)</p>
- 4. Sensitivity ≈ # of partitions
 - Frequency as low as 1/100K has been measured⁴
- 5. Only recently >2 channels for multiplexing



4 Heyries et al Nat Meth 2011; Pekin et al, Lab chip 2011



Case Study: Comparison of VG Titers by qPCR vs ddPCR



Average %CV by qPCR = 17%

Average %CV by ddPCR = 11%



qPCR

ddPCR

Analytical Criteria for PCR based Method Development

SENSITIVITY	SPECIFICITY	ACCURACY	PRECISION	LINEARITY
 # of copies accurately measured in the assay LOD ≤ LOQ The concentration of template at 95%Cl ≥ 3 due to poisson distribution limits Methods vary across industry LOQ The lowest quantifiable concentration with acceptable precision and accuracy Best determined empirically 	 Detects target sequence Does not detect any other sequences also present in the sample matrix 	 Differences between measured and known titers presented as fold change or copy # estimates 	 Repeatability Precision and robustness of the assay with the same samples analyzed repeatedly in the same assay Intermediate Precision Variation in results between assay runs and/or laboratories Test multiple reagent lots, instruments, operators 	 Dilutional 3-5 non-zero points Cal Curve (qPCR) ≥6 non-zero points Run in duplicate Identical matrix as samples Linear regression on log transformed data



Acceptance Criteria for PCR Based Titer

SENSITIVITY	SPECIFICITY	ACCURACY	PRECISION	LINEARITY
 Acceptance criteria depend on assay For vg titer at LOQ %CV<25% and accuracy at 70-130% 	 NTC < LOD no interference in extracted gDNA Matrix effect evaluated on QC samples 	 QCs spiked in relevant matrix with established acceptance criteria Ideally QCs spanning the range of the assay 	 CV ≤ 25% for intermediate precision Tighter for repeatability 	 qPCR amplification efficiency between 90-110% R^2 ≥ 0.980

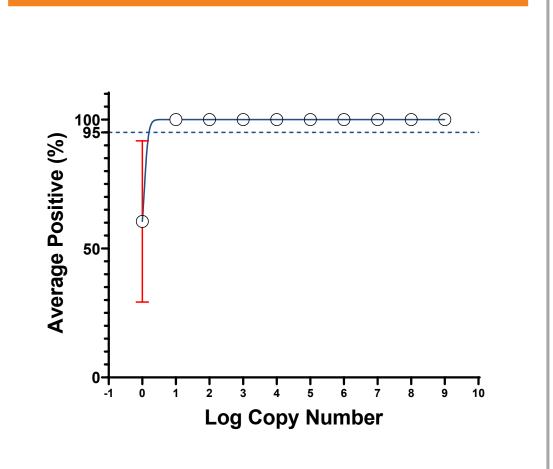


LoB/LoD and LOQ in qPCR

Methods to Establish LoD and LoQ

- LoB/LoD Calculation per CLSI guideline EP17
 - $LoB = mean \ blank + 1.645 \times \sigma_{blank}$
 - $LoD = LoB + 1.645 \times \sigma_{low concentration sample}$
- Problem: above equations apply to linear response
 - qPCR response is logarithmic and does not cross threshold for negative sample
- Alternative: establish LoD based on % positive response in replicate standard curves¹
- For LOQ run the same 20 replicates and select the point with %CV<25% and accuracy 70-130%

LoD Based on 20 Replicate Std Curves





1 Forootan et al., Biomolecular Detection and Quantitation, 2017

Other AAV VG Titer Considerations

qPCR vs ddPCR?

- Is ddPCR preferred over qPCR for regulatory filings?
 - No specific guidance but ddPCR has greater precision
 - Either assay should be acceptable if fitfor-purpose and properly validated
 - If feasible start with ddPCR in preclinical assay development to facilitate easy transfer to CMC

Titer Considerations

- Research material typically uses vendor titer with probe against ITR region
 - Overestimated concentration
- Confirm vendor titer with target gene specific assay
- Accurate titer assures equivalent dosing for reliable evaluation of AAV potency

Accurate vg Titer Critical for Accurate Potency Measurement



Regulatory Guidance for Potency Testing of GT Products

What is Potency?

 Potency is defined as "the specific ability or capacity of the product, as indicated by appropriate laboratory tests or by adequately controlled clinical data obtained through the administration of the product in the manner intended, to effect a given result."

(21 CFR 600.3(s))

Required for Lot Release, Stability, and Comparability

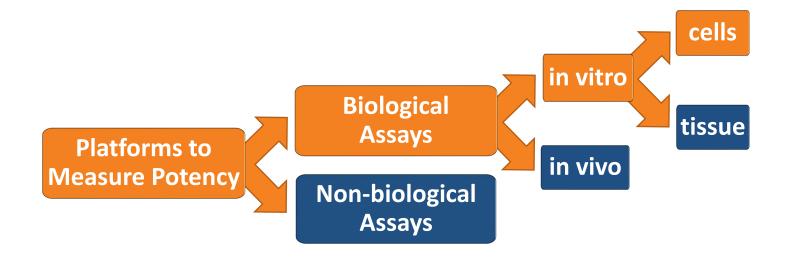
Guidance Documents

- FDA-2008-D-0520: Guidance for Industry: Potency Tests for Cellular and Gene Therapy Products
- USP-NF<1032>: Design and Development of Biological Assays
- USP-NF<1034>: Analysis of Biological Assays
- ICH Q14: Analytical Procedure
 Development
- ICH Q2(R2): Validation of Analytical Procedure



Potency of AAV Drug Product

- A gene therapy vector relies on 2 main biological activities for its potency
 - 1. the ability to transfer a genetic sequence to a cell
 - 2. the biological effect of the expressed genetic sequence





EDIT-103 for Autosomal Dominant Retinitis Pigmentosa (adRP)

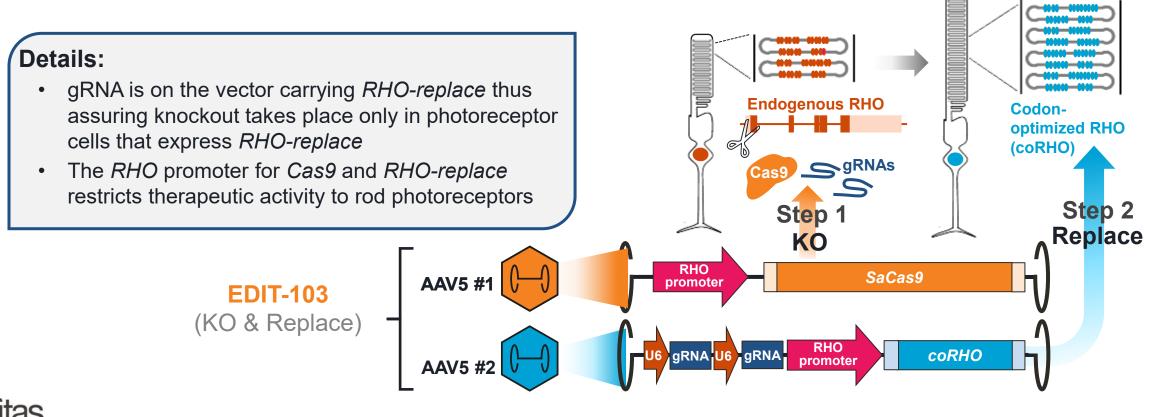
- An inherited autosomal dominant retinal disease leading to blindness in later life
- Symptoms:
 - Decreased night vision (nyctalopia)
 - Loss of peripheral vision (tunnel vision), and eventually significant decline in central vision
- No approved treatments





EDIT-103: Dual AAV-Based "Knockout and Replace" Therapeutic Strategy

- Agnostic to any RHO mutation thus will knockout any dominant gain-of-function rhodopsin mutant
- Step 1: Both mutant and normal endogenous RHO will be knocked out in the treated area
- Step 2: Exogenous normal RHO (resistant to editing) will replace endogenous RHO
- One-time subretinal administration aimed to restore/prevent vision loss

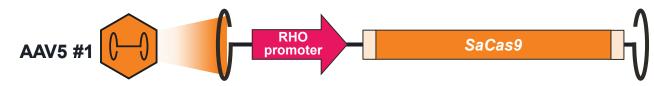


Dual Vector Strategy Adds Complexity to Potency Assay Design

• EDIT-103 delivers two transgenes

Vector 1

• Expression of Cas9 endonuclease



Vector 2

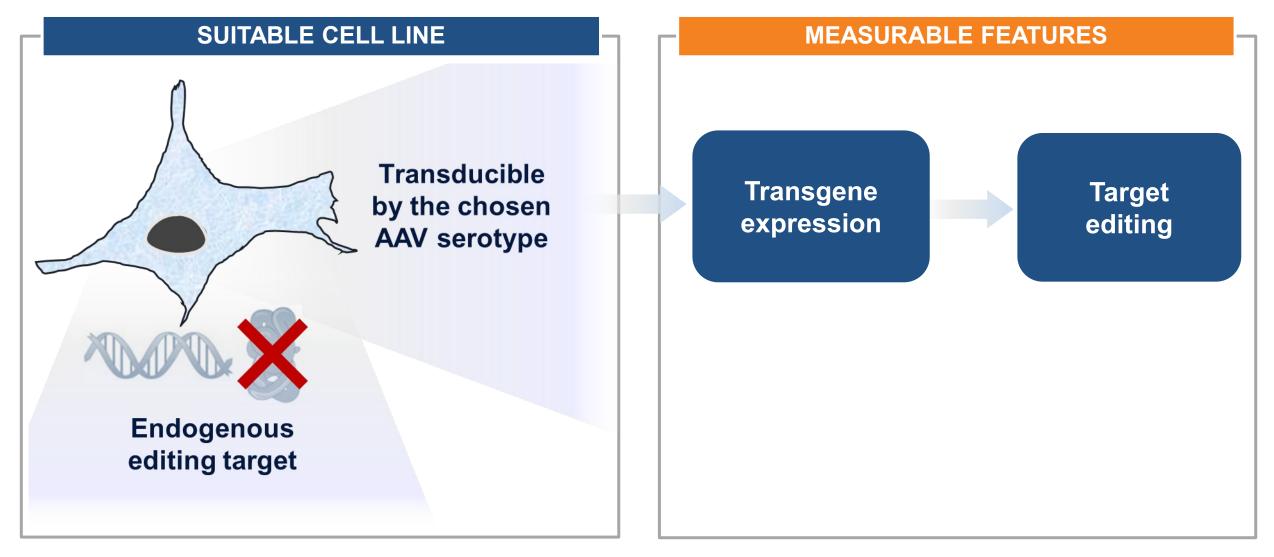
- Expression of gRNA
- Expression of coRHO replacement at mRNA → RT-ddPCR readout
- coRHO expression-driven restoration to the structure/function of the photoreceptor

Vector 1 & 2

Editing of both endogenous RHO alleles by the Cas9 endonuclease → ddPCR readout



In-Vitro Potency Assay Platform Strategy

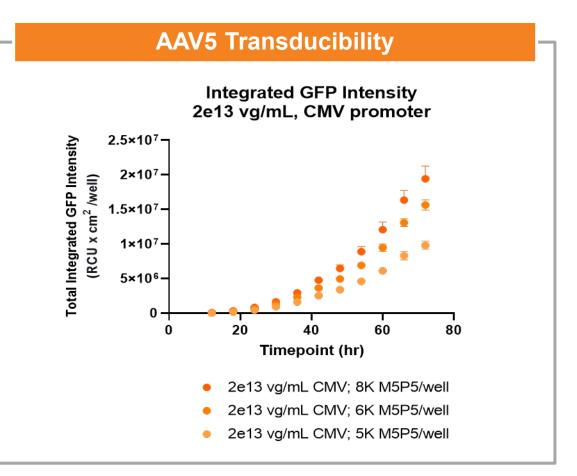




In-Vitro Potency Assay Platform Selection

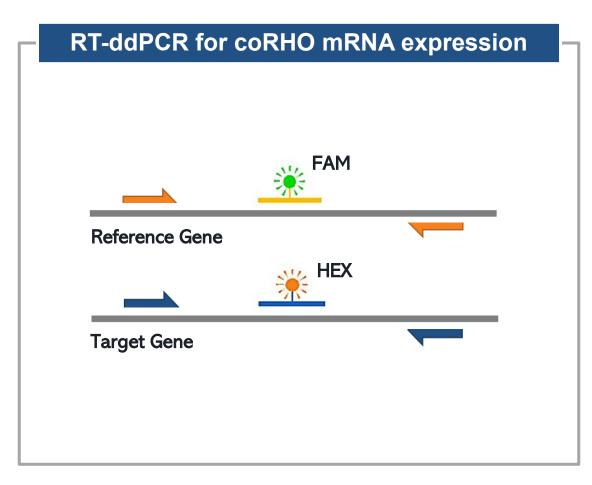
In-Vitro Platform

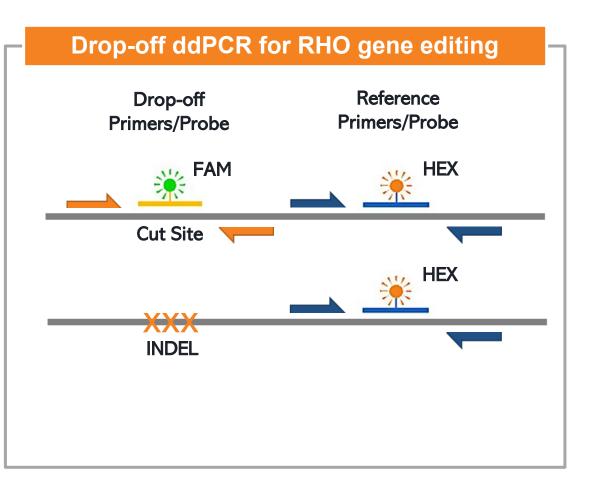
- HEK293T cells
 - contain but don't express endogenous RHO target
 - AAV5 transducible
- Additional cell line engineering required to enable the expression of transgenes under RHO promoter
- Cell culture conditions optimized for efficient transduction





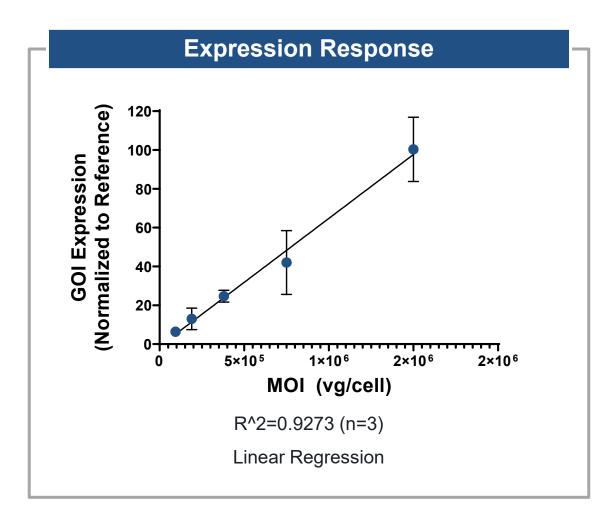
ddPCR Readouts for mRNA Expression and Target Editing







Protein Replacement Potency Assay

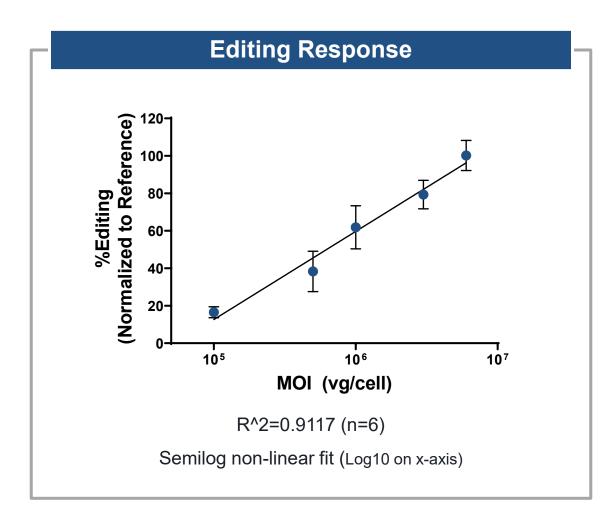


Next Steps in Development

- Assay will be transferred to AD/QC for method optimization / validation
 - Establish Relative Potency
 - Optimize key assay steps
 - Establish system suitability criteria
 - Assay acceptance criteria
 - Sample acceptance criteria
 - Method qualification and transfer to QC for method validation



Editing Potency Assay



Next Steps in Development

- Assay will be transferred to AD/QC for method optimization / validation
 - Establish Relative Potency
 - Optimize key assay steps
 - Establish system suitability criteria
 - Assay acceptance criteria
 - Sample acceptance criteria
 - Method qualification and transfer to QC for method validation



Summary

VG Titer Assay Development

- Don't rely on ITR based vendor titers
- Develop target gene specific assays
- Test primers and probes *in silico*
- Design against cross-reactivity to matrix components and AAV cargo
- Select a platform that will easily transfer to regulated environment

Potency Assay Development

- Start potency assay development early!
 - Technically challenging
 - Slow TAT on assays
 - PCR based assays to prove editing and expression efficacy at IND
 - Functional potency for release of Phase 3 clinical material

Let the Full Lifecycle of the Program Guide Decisions in Assay Development



Acknowledgements

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* Alphabetical order according to last names

