

# Development of a triplex one-step RT-ddPCR method as a quantitative potency assay for the vMiX™ platform



R. Lyth, J. Kenth, L. Reilly, R. Joubert, L. Li, I. Blount, J. Isaac, A. Bloom, A. Martorana, H. Cumar, M. MacDonald, P. Boyce

AviadoBio Ltd, London, UK

## OBJECTIVE

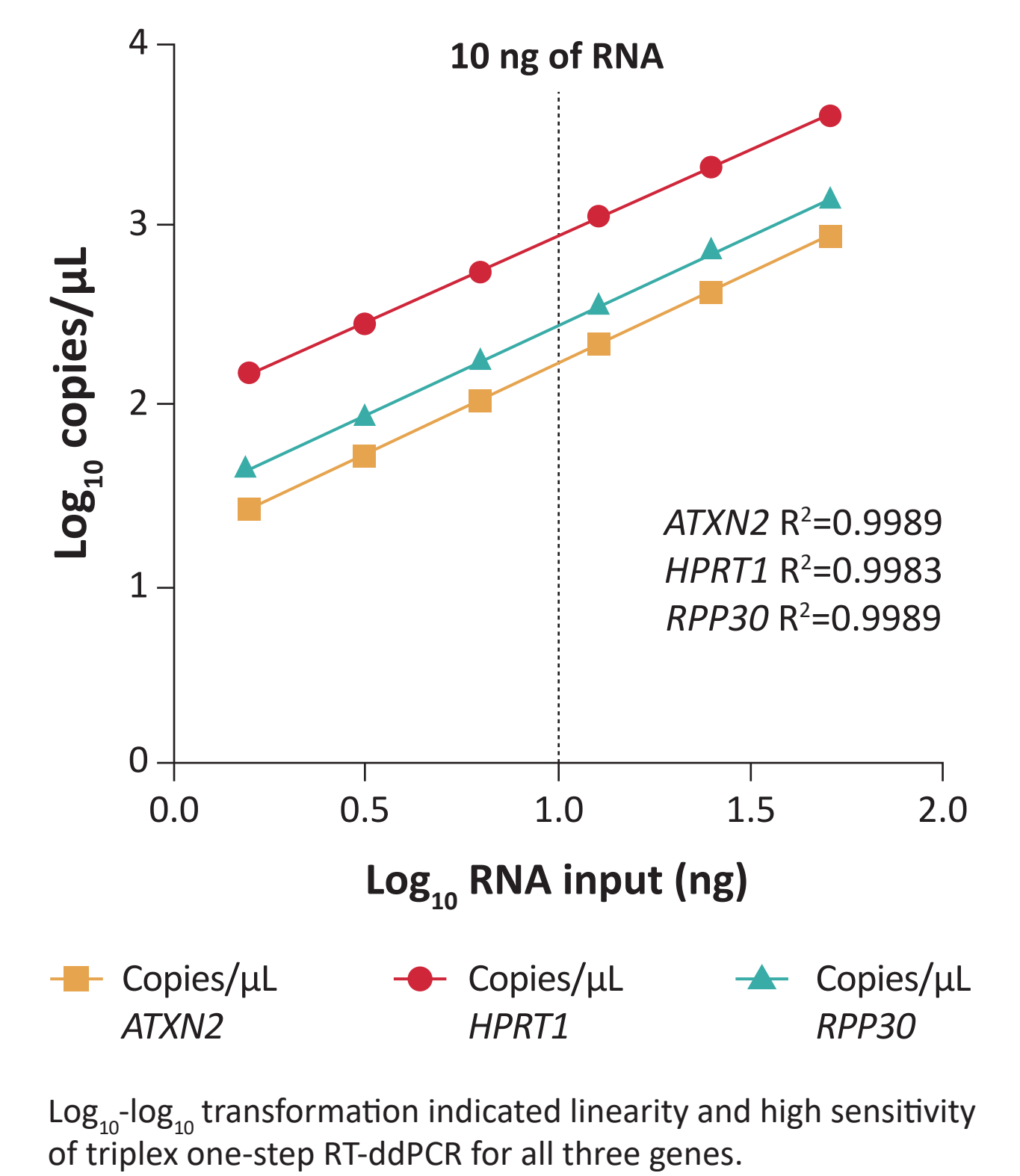
To develop a triplex one-step reverse transcription droplet digital polymerase chain reaction (RT-ddPCR) method to quantify the extent of ataxin 2 (*ATXN2*) mRNA knockdown (KD) by AVB-205. This approach can be applied across the vMiX™ gene silencing platform for rapid gene silencing potency assay development.

## RESULTS

### Determination of total RNA load, linearity and sensitivity

- To determine total RNA load in the linear range of all three target genes, a multiplexing experiment was performed where 10 ng of RNA was identified to be an optimum RNA load (Figure 3).
- RNA load was assayed in both simplex and multiplex (data not shown) to ensure the multiplex reaction had no impact on gene expression quantification.
- All three genes demonstrated high sensitivity and good precision between replicates with a linear detection range of 1.56–50 ng of RNA input.

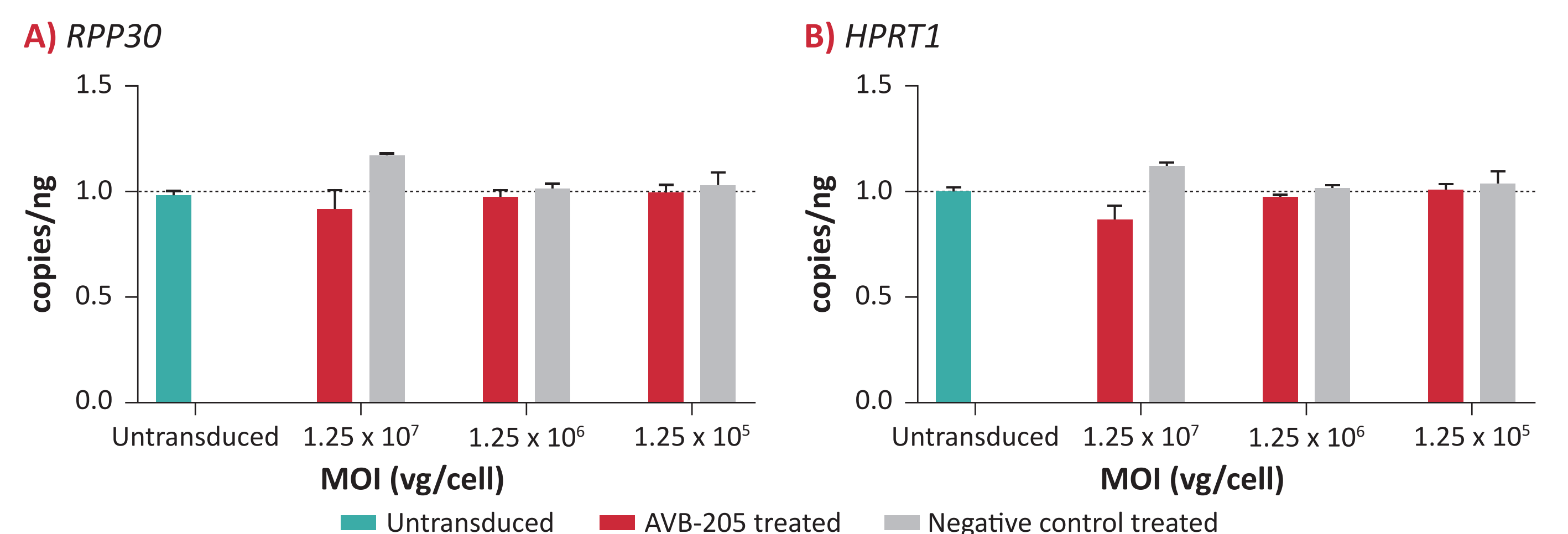
**Figure 3:** *ATXN2*, *HPRT1* and *RPP30* multiplex of total RNA load and linearity



### Assessing the stability of chosen HKGs across a range of MOIs

- HeLa cells were transduced with either AVB-205 or a negative control with non-targeting mRNA across a 3 log range.
- Both HKG expressions were stable across the range of MOIs regardless of AAV9 infection and subsequent *ATXN2* mRNA KD (Figure 4).
- A ratio of *RPP30*:*HPRT1* expression was used as an assay control to monitor variability in cell culture within the assay.

**Figure 4:** HKG stability in AVB-205-treated HeLa cells



A) The gene expression profile of *RPP30* (copies/ng) and of B) *HPRT1* was stable in HeLa samples treated with AVB-205 (red) against the negative control with non-targeting mRNA (in grey) across three MOIs and the untransduced sample (in teal). All samples were normalized to untransduced samples. *RPP30* and *HPRT1* gene expression was stable.

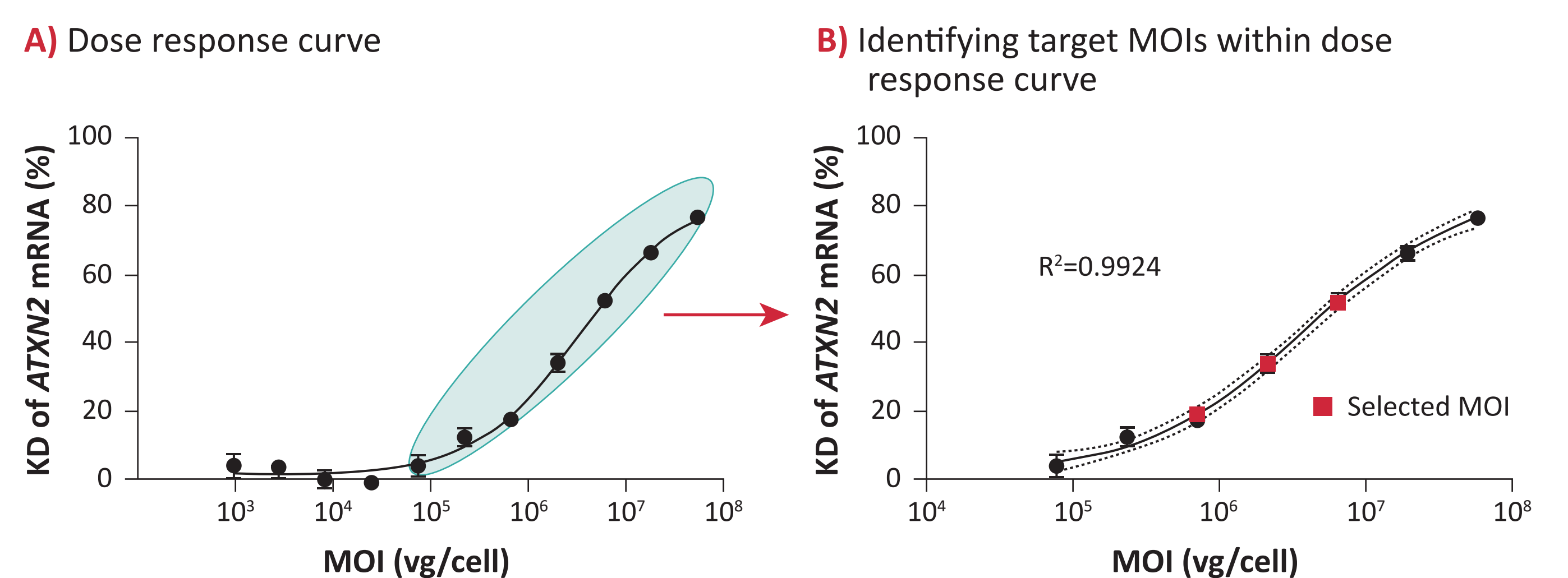
### Dose response curve of *ATXN2* mRNA percentage knockdown

- Three MOIs were selected for testing *ATXN2* mRNA percentage KD routinely:  $6.95 \times 10^5$ ,  $2.06 \times 10^6$  and  $6.25 \times 10^6$  (Figure 5).

### Inter-assay precision

- The assay was tested three times between two operators evaluate inter-assay precision. All values were below a 15% coefficient of variation (CV; Table 1).

**Figure 5:** Dose response curve of *ATXN2* mRNA percentage knockdown



A) HeLa cells were transduced with MOIs covering a 6-log range. MOIs between  $5.6 \times 10^7$  and  $9.48 \times 10^3$  were fitted with a four-parameter logistic sigmoidal curve. B) The three MOIs highlighted in red were selected for further assay development.

**Table 1:** Inter-assay precision knockdown of *ATXN2*

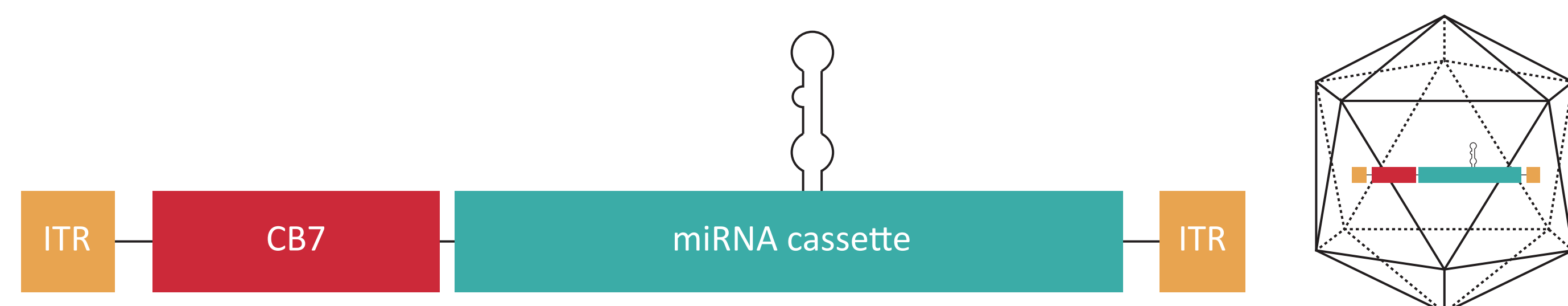
MOI	Run 1, Operator 1 KD (%)	Run 2, Operator 2 KD (%)	Run 3, Operator 1 KD (%)	Average KD (%)	CV (%)
$6.25 \times 10^6$	53.3	50.5	58.7	54.2	7.1
$6.95 \times 10^5$	20.3	21.5	23.7	21.9	10.6

Inter-assay precision shown by KD (%) achieved at two MOIs ( $6.25 \times 10^6$  and  $6.95 \times 10^5$ ) across three runs by two operators. The CV (%) was calculated using independent triplicate values from each run at the respective MOIs. CV values less than 15% indicate high precision of the assay.

## INTRODUCTION

- AVB-205 is an investigational gene therapy product encoding a vMiX™ gene silencing microRNA (miRNA) that targets and degrades *ATXN2* mRNA transcripts (Figure 1).

**Figure 1:** AVB-205 miRNA payload design



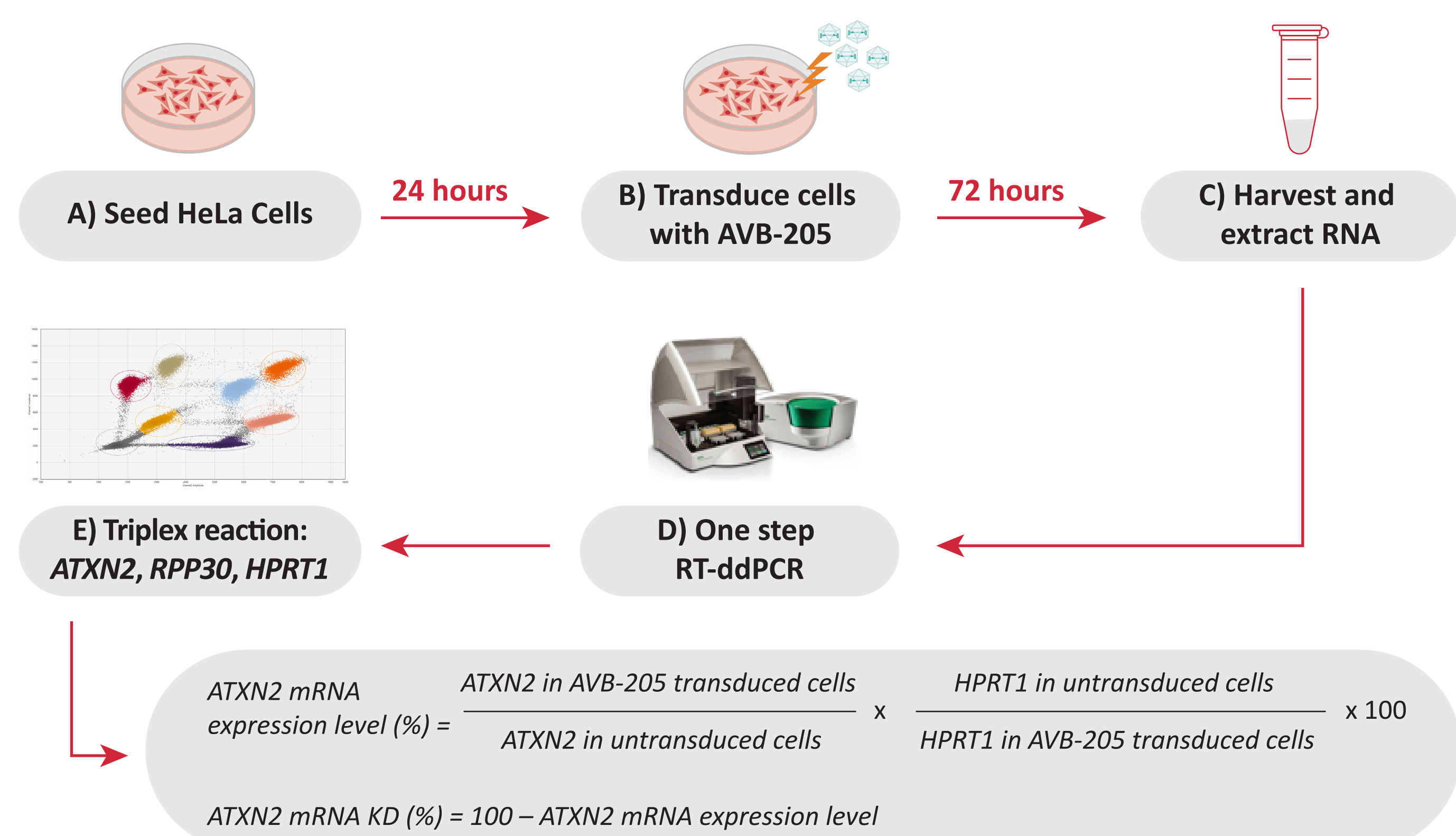
AVB-205 is composed of a recombinant adeno-associated virus serotype 9 (AAV9) vector containing DNA encoding an miRNA construct targeting *ATXN2* mRNA, under control of the ubiquitous chicken  $\beta$ -actin promoter (CB7). The position of inverted terminal repeats (ITRs) and miRNA optimizes miRNA processing, and the single-stranded genome optimizes packaging and manufacturability.

- ATXN2* mRNA degradation significantly slows disease progression in TAR DNA binding protein 43 (TDP-43)-overexpressing mouse models of amyotrophic lateral sclerosis (ALS), so has emerged as a potential therapeutic strategy.<sup>1</sup>
- Here, we describe the development of a triplex one-step RT-ddPCR *in vitro* potency assay that can quantify the KD of endogenous *ATXN2* mRNA caused by the transduction of AVB-205.
- For such gene silencing approaches, measuring the KD of endogenous genes in relation to multiple housekeeping genes (HKGs) in a single reaction can be used to assess mechanism of action, and potency for routine testing of batch release and stability.

## METHODS

- The RT-ddPCR method was developed using the following steps (Figure 2):
  - Determine total RNA load, linearity, and sensitivity of RT-ddPCR multiplexing
  - Evaluation of HKG stability in response to an AAV9 infection and *ATXN2* KD
  - Determination of dose-response curve to assess linearity and range
  - Evaluation of assay precision

**Figure 2:** Method for measuring *ATXN2* knockdown (%) in cells treated with AVB-205



A) HeLa cells are seeded at  $8 \times 10^4$  cells/well in 500  $\mu$ L of complete media then incubated at 37°C for 24 hours. B) Cells are transduced with AVB-205 at selected multiplicities of infection (MOIs) in the linear range of the dose response curve. C) Cells are further incubated at 37°C for 72 hours before automated RNA extraction using the Qiacube by Qiagen. D) RNA samples are loaded at a concentration of 10 ng/well in the one-step ddPCR Biorad machine. E) The triplex gene clusters are selected and analyzed, and the KD (%) is calculated from the formula above.

## CONCLUSIONS

- This one-step RT-ddPCR *in vitro* potency assay shows excellent sensitivity and repeatability, working in triplex to detect two HKGs alongside the gene of interest and confidently reporting knockdown of *ATXN2*.
- The one-step RT-ddPCR method provides a platform approach for vMiX™ potency assay development.