# Dyno-86m: Optimizing Intravitreal Delivery to the Non-Human Primate Retina with Machine-Guided AAV Capsid Design //

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### Abstract

There is an unmet need for safer and more efficient payload delivery methods to enable gene therapies with improved clinical outcomes in the eye. Dyno's platform enables machine-guided design of optimized capsid variants, in vivo characterization of capsid libraries, and in vivo validation of capsid transduction properties. Through this approach we designed and validated an AAV capsid variant, dyno-86m, with remarkably improved efficiency in transducing the neural retina following intravitreal (IVT) delivery into cynomolgus macaque nonhuman primate (NHP) eyes. Single-cell sequencing and histological analyses confirmed that dyno-86m dramatically outperforms AAV2 and is consistently better than external engineered IVT capsid variants in transducing photoreceptors, at rates with clinically meaningful potential.

### Summary

- Single-nuclei RNA sequencing demonstrates transduction of all retinal cell types by dyno-86m in cynomolgus macaque
- > Quantification of transduction by histology confirmed snRNAseq and NGS based measurements
- > As shown by histology and snRNAseq, dyno-86m transduced NHP retina cells up to 3x more efficiently than an external engineered capsid, and by NGS 80x more efficiently than AAV2, with no reduction in AAV production efficiency
- dyno-86m capsid is available for immediate licensing. Contact ► The bd@dynotx.com to learn more about licensing and emerging validated capsids from more recent NHP studies



Figure 1. Dyno's capsid discovery platform. (A) Dyno utilizes machine guided design and high throughput DNA synthesis to engineer variants with improved functional properties in large, high quality libraries. (B) Optimization of variants is iterative and builds on previous data. Variants are designed in silico and characterized in vivo by NGS of variant-associated barcodes from NHP tissues. The data are used to train machine learning models to enable design of next-generation libraries.



Figure 2. Readouts for capsid discovery and validation studies. Initial characterization of variants employs complex (>1e5 variants) capsid libraries and utilizes NGS readouts to measure biodistribution and transduction in ocular and non-ocular tissues. In vivo validation is performed in smaller scale studies and provides bulk and single-cell NGS (<100 variant/experiment scale) as well as histology based readouts (<10 variant/experiment scale).





retinal cell types in 8 separate 10X runs. (C-D) UMAP projection shaded by transduction events by the external engineered capsid and dyno-86m, respectively. (E) Transduction rates (fraction of transduced nuclei normalized by dosage of test articles) in retinal cell types. (F) Fold improvement of dyno-86m transduction rate as compared to the external engineered capsid. RGC=retinal ganglion cells; AC=amacrine cells; BC=bipolar cells; MG=Müller glia.

Figure 8. The dyno-86m capsid transduces cells in all retinal layers in the NHP macula. Transduction by dyno-86m shown in green and by the external engineered capsid in magenta.



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## Figure 9. The dyno-86m capsid transduces cells in all NHP retinal layers up to 3-fold more efficiently than the

external engineered capsid. (A) Example images showing automated quantification of eGFP and mCherry expressing cells. (B) Quantification of dyno-86m and the external engineered capsid transduction in different retinal layers and regions or (C) across the whole retinal cross-section normalized to the dose per vector (n=4 sections/eye averaged across 2 injected eyes).