

# Identification of Dose-Dependent Immune Landscape Signatures Following Administration of Ixo-vec in Non-Human Primates



Julian N. Ramos

Principal Scientist, Adverum Biotechnologies

May 18, 2023



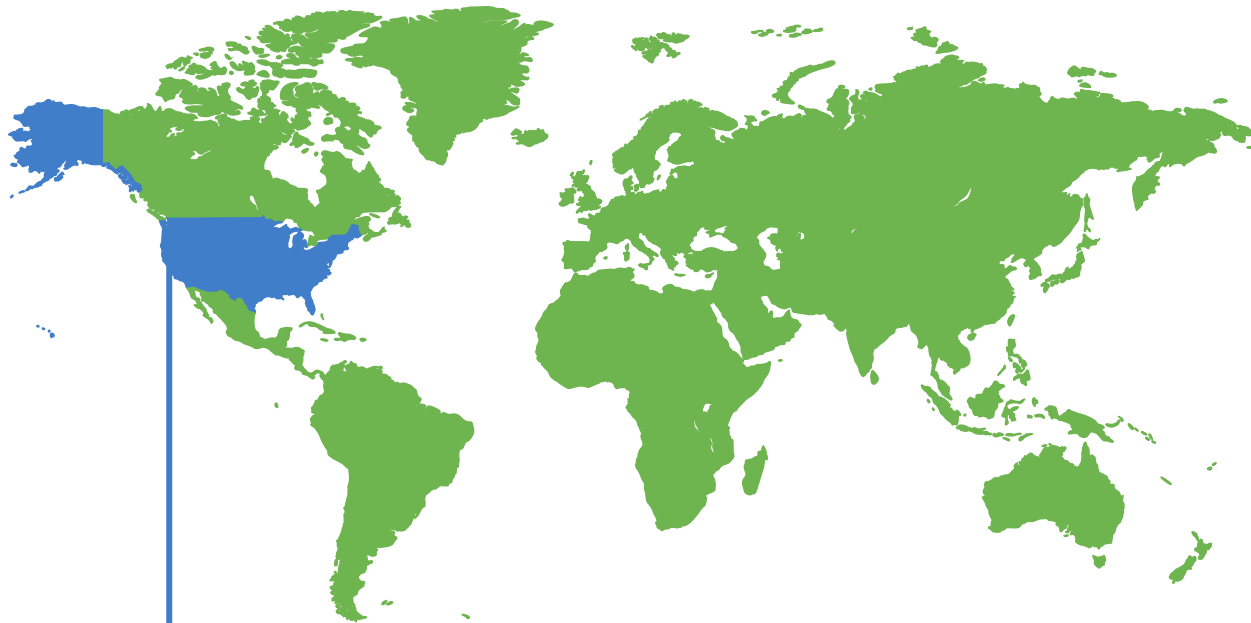
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- **Adverum Biotechnologies:** employee, stockholder.
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# Wet AMD: Leading Cause of Blindness in Patients Over 65

**20M** people living with wet AMD worldwide<sup>1,2</sup>



**1.5M** people impacted by wet AMD in the U.S.<sup>1,2</sup>

## Opportunity for New Wet AMD Treatment

### Current Standard of Care Drives Suboptimal Outcomes

Frequent bolus anti-VEGF IVT injections required leading to poor compliance<sup>3,4</sup>

Lifetime need for frequent injections overburdens patients, caregivers, and healthcare providers

\$139 billion in U.S. annual economic burden of vision loss, including falls, cognitive decline and depression<sup>5</sup>

### Gene Therapy: Overcomes Wet AMD Treatment Burden

Single, easily administered in-office IVT treatment

Durable, disease modifying therapy, potentially “one and done” functional cure

Readily deployable in the U.S. and around the world addressing direct and indirect costs

<sup>1</sup>Bright Focus Foundation. Age-Related Macular Degeneration: Facts & Figures.

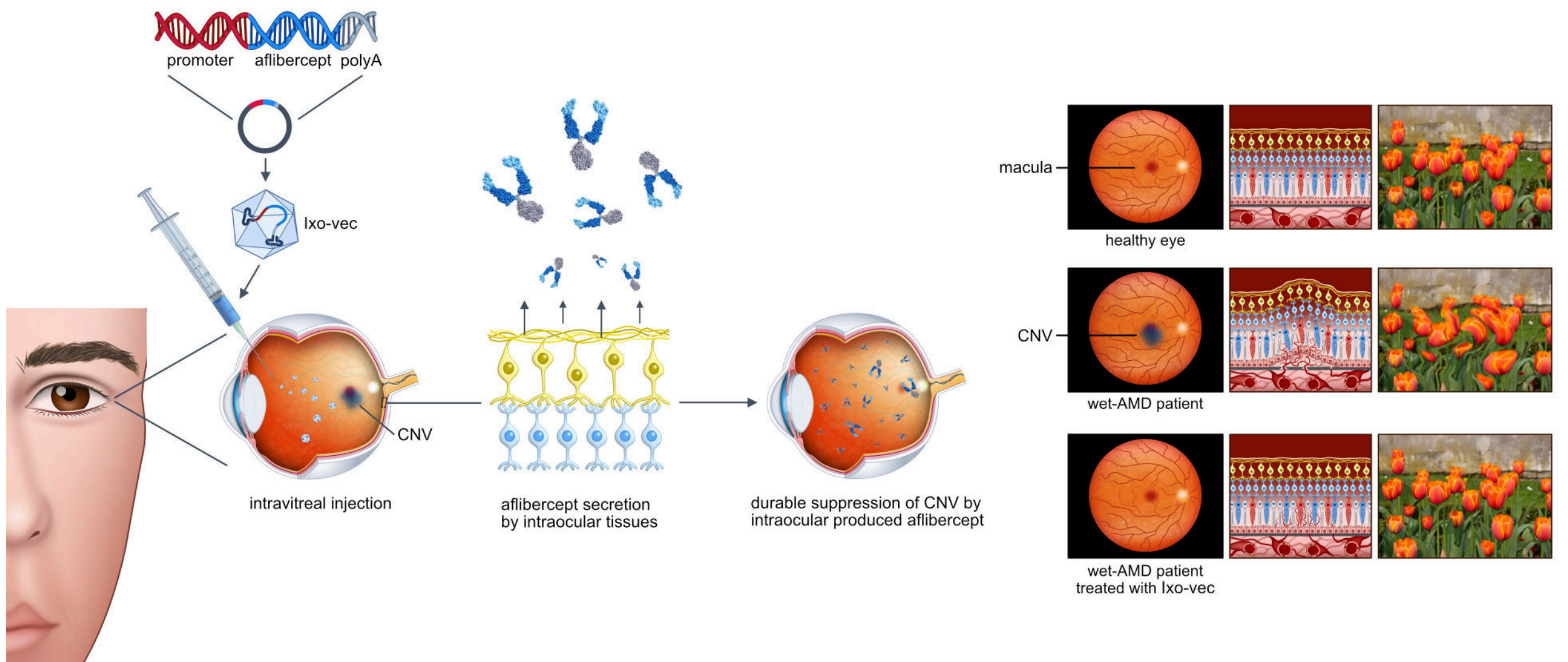
<sup>2</sup>Wong WL, et al. Global prevalence of age-related macular degeneration and disease burden projection for 2020 and 2040: a systematic review and meta-analysis. *Lancet Glob Health*. 2014;2:106–16.

<sup>3</sup>Holz FG et al. *Br J Ophthalmol* 2015; 99 (2): 220–226.

<sup>4</sup>Khanani A, et al. *Ophthalmol. Retina* 2020 Feb; 4(2):122-123.

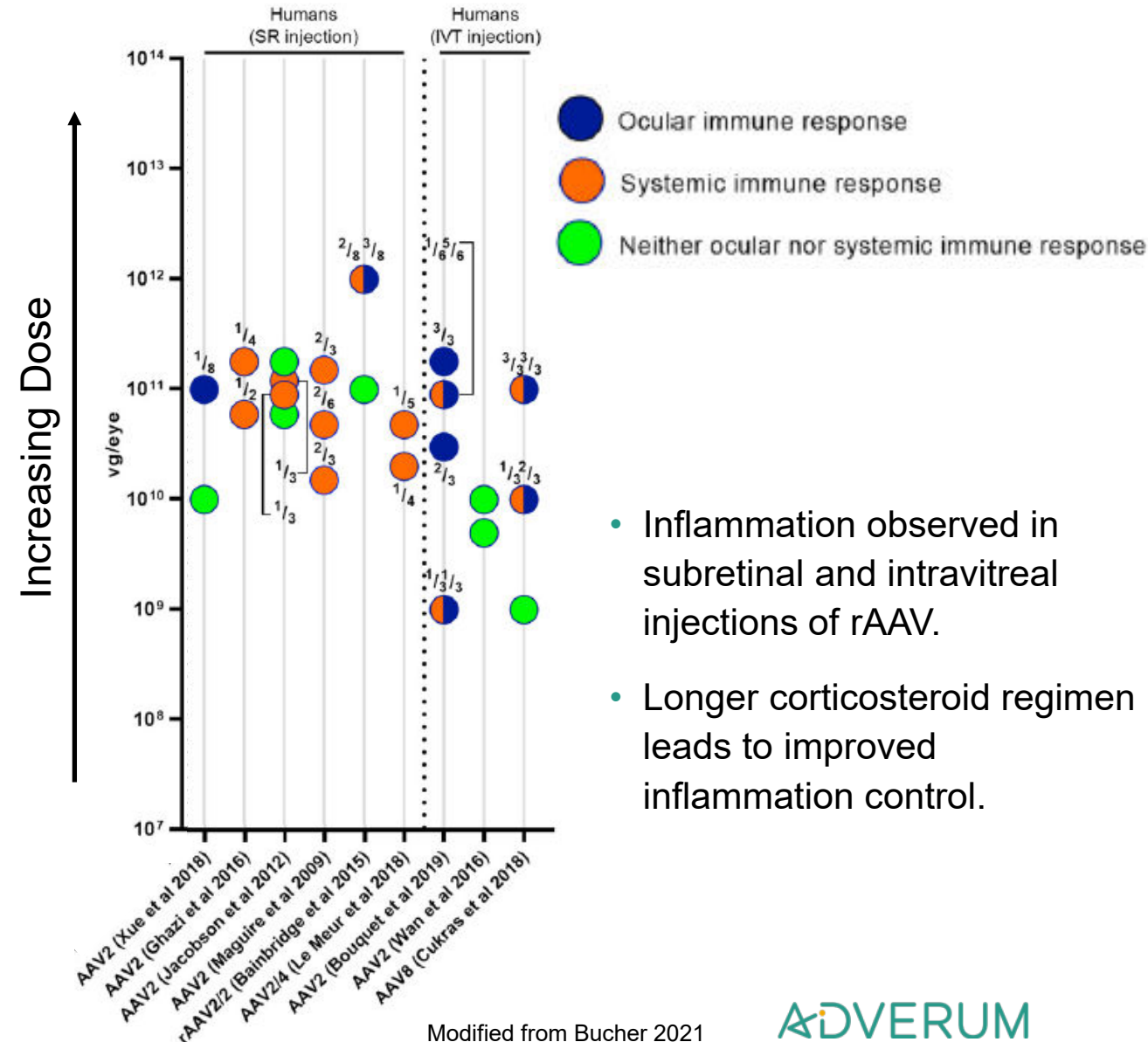
<sup>5</sup>Angiogenesis Foundation: Patient-centered Outcomes in Wet Age-related Macular Degeneration, Boston, MA, October 2017

# Ixo-vec is a gene therapy biofactory approach designed for continuous delivery of aflibercept (anti-VEGF) by single intravitreal injection



# Clinical implementation of gene therapy products is influenced by rAAV immune mediated inflammatory response

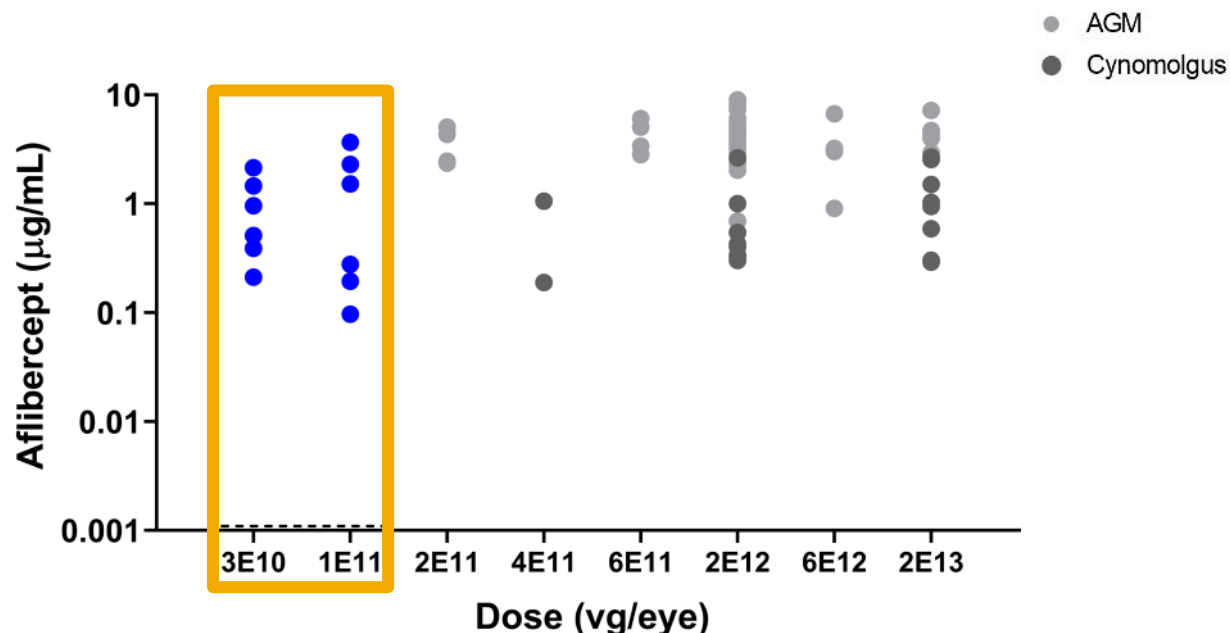
- Dose-dependent inflammation is observed across the gene therapy field, irrespective of route of administration and therapeutic modality.
- rAAV immunogenicity can be mitigated by dose reduction and improved corticosteroid regimen.
- Systems-wide analyses of nonclinical models could help identify molecular signatures reflective of the dose-dependent inflammation commonly associated with gene therapy.



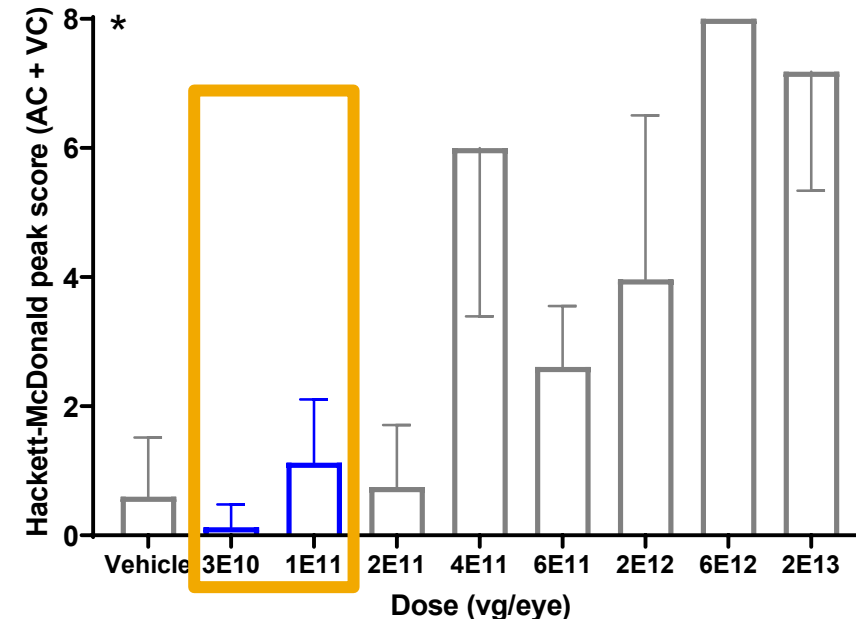
# Ixo-vec potency enables ability to dose down to improve inflammation profile

Non-dose proportional aflibercept levels enable dose reduction to improve Ixo-vec inflammation profile while maintaining efficacy.

## Vitreous Humor



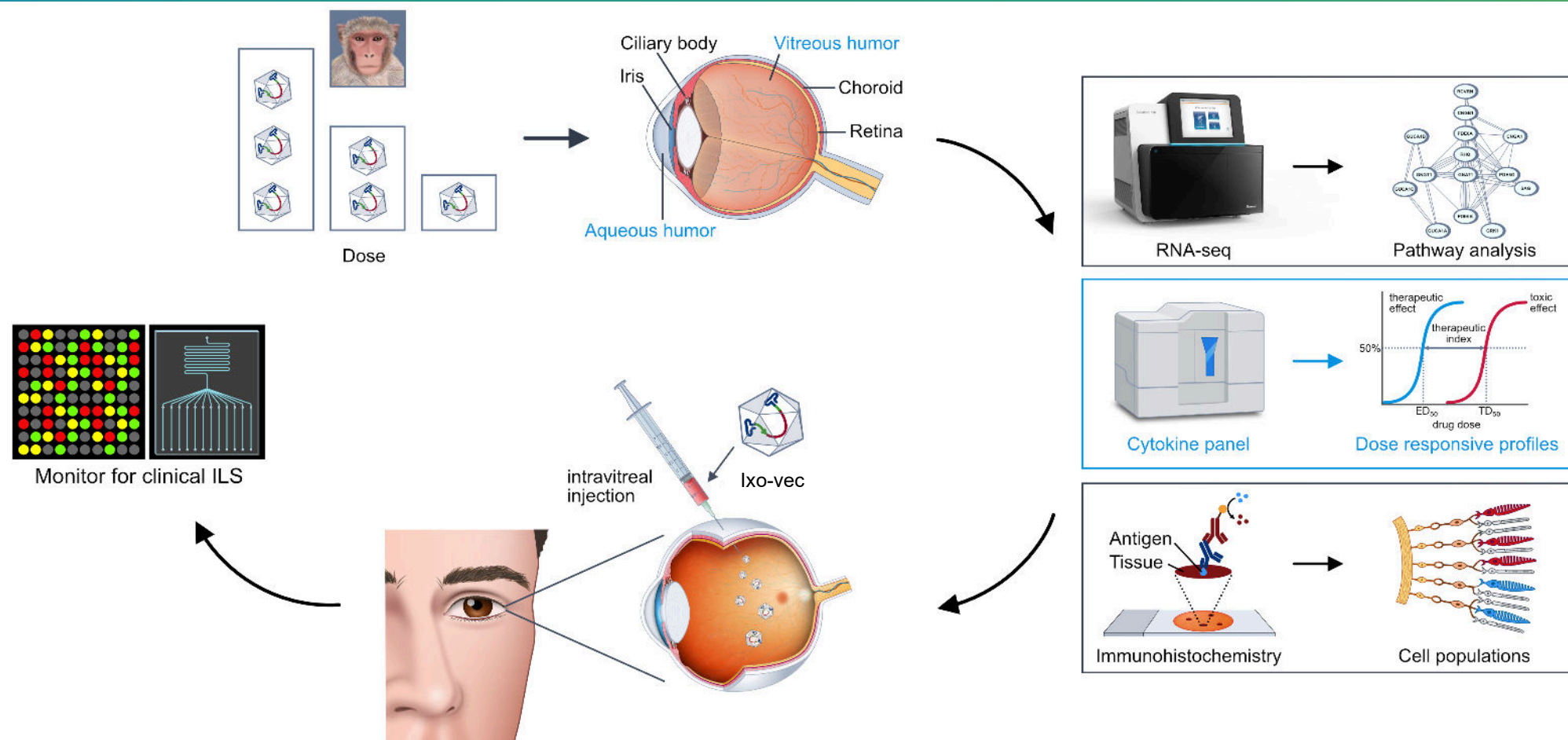
## Inflammation Scores



- NOAEL established at NHP dose of 1E11 vg/eye = 2E11 vg/eye human equivalent dose (HED).
- NHP dose of 3E10 vg/eye = 6E10 vg/eye HED.
- NHP dose of 1E11 vg/eye = 2E11 vg/eye HED.
- Doses in current LUNA clinical study outlined in orange boxes.

\*Scale is cumulative of two parameters for maximum score of 8.

# Defining immune landscape signatures of Ixo-vec at supra-clinical doses



- Doses spanning 3-log fold range above NOAEL (1E11 vg/eye):
  - 4E11 vg/eye (low), 2E12 vg/eye (mid), 2E13 vg/eye (high)
- Single eye injected for each animal.
- Endpoint: 3 months post-dose.

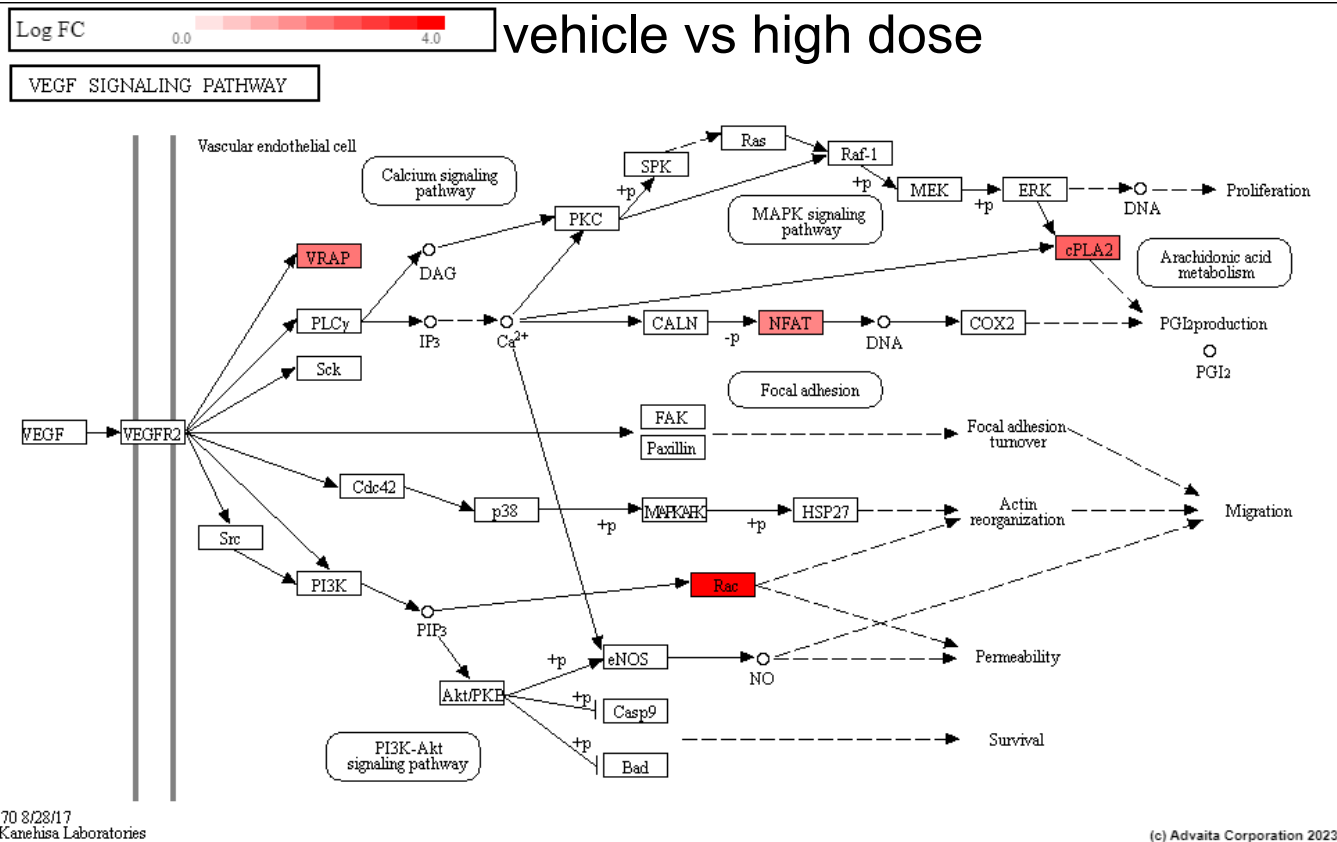
- Multiple regions of retinal anatomy submitted for bulk RNA-seq.
- Pathway analysis was performed on RNA-seq samples.
- Parallel effort characterized inflammation by histology.

Illustration: Gardenia Gonzalez Gil

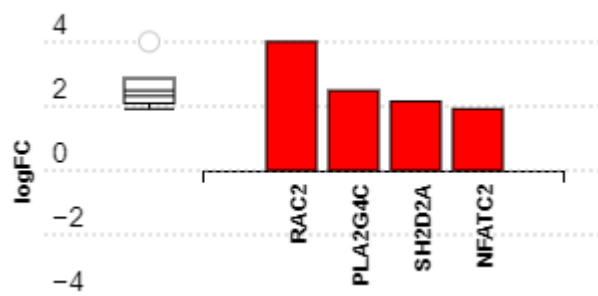
## Converging dataset strongly supports dose-dependent inflammatory response

- Pathway analysis was performed using multiple methods (topology and non-topology based).
  - Modulation is determined by the number of differentially expressed genes within a set as well as their position and magnitude of role within a pathway.
- Transcriptomic analysis did not support unfolded protein response (UPR), oxidative or endoplasmic reticulum (ER) stress, neovascularization, nor ciliary body dysfunction.
- Ixo-vec expression detected only in the dosed eye.
- Robust and dose-dependent activation of immune responses, consistent in nature across ocular tissues.
  - Severity in line with anticipated vector biodistribution.
- No evidence that ciliary body architecture was directly affected by Ixo-vec.
  - Only indirectly as dose-dependent inflammation increases.
  - Histology showed no disruption of ciliary body fold formation at any dose.
- RNA-seq and histopathology indicate activation of innate and adaptive immune systems consistent with dose-dependent rAAV-associated inflammation.

# VEGF signaling is not activated in ocular tissues



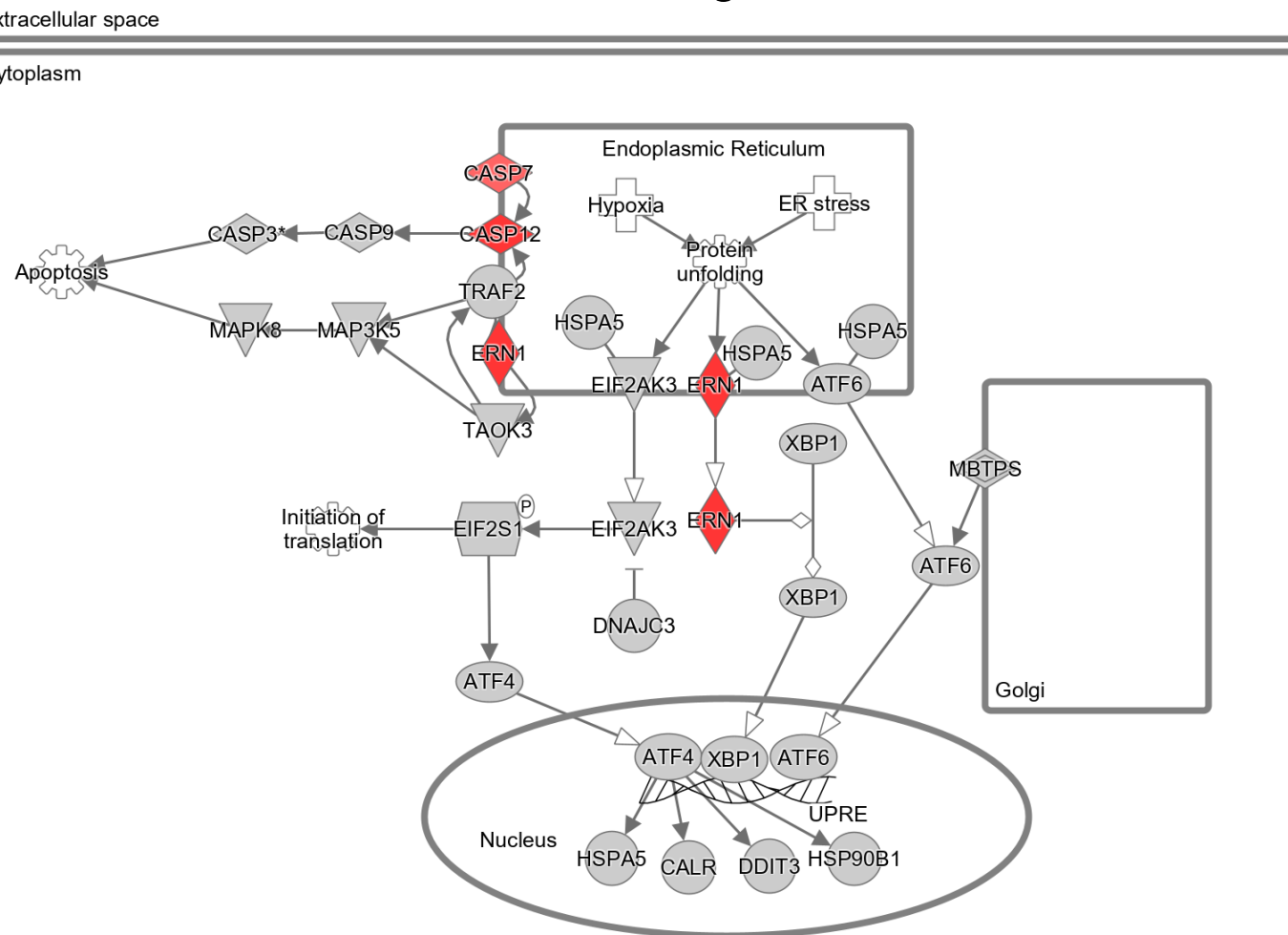
- Observation was similar across doses for all ocular tissues examined.
- Ixo-vec unlikely to exacerbate VEGF-mediated pathology of wet AMD.



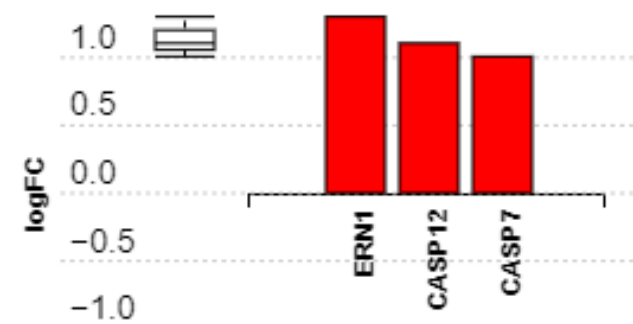
# No evidence of UPR or ER stress by Ixo-vec dosing

## Endoplasmic Reticulum Stress Pathway

### vehicle vs high dose

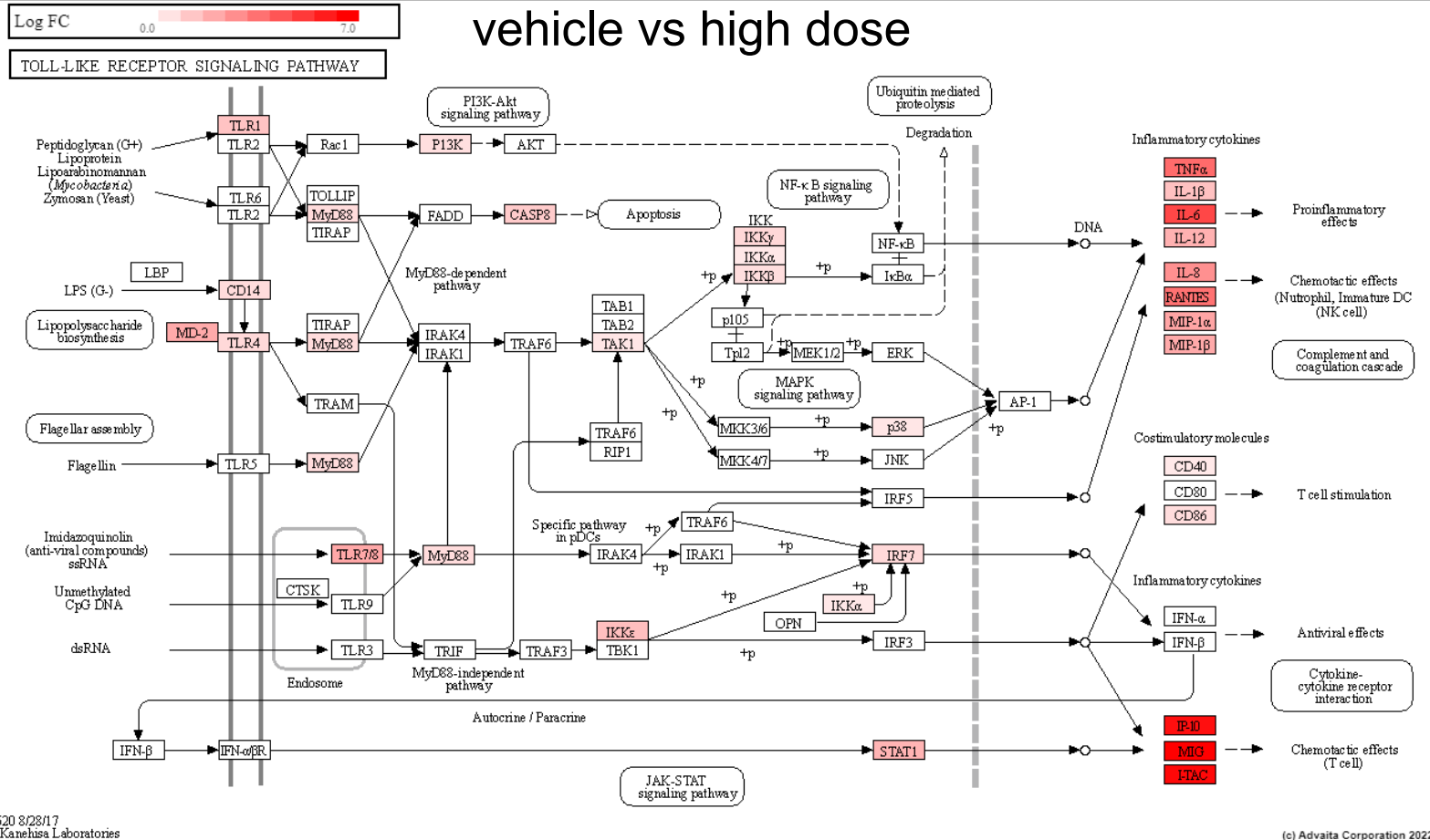


- Cellular toxicity has been attributed to inefficiencies in transgene translation, folding, and post-transcriptional modification.<sup>1, 2</sup>
- Ixo-vec expression unlikely induces cellular stress.
- UPR reportedly upregulated by rAAV transduction.<sup>3, 4</sup>
- Transduction-mediated UPR unlikely to be observed at study endpoint.

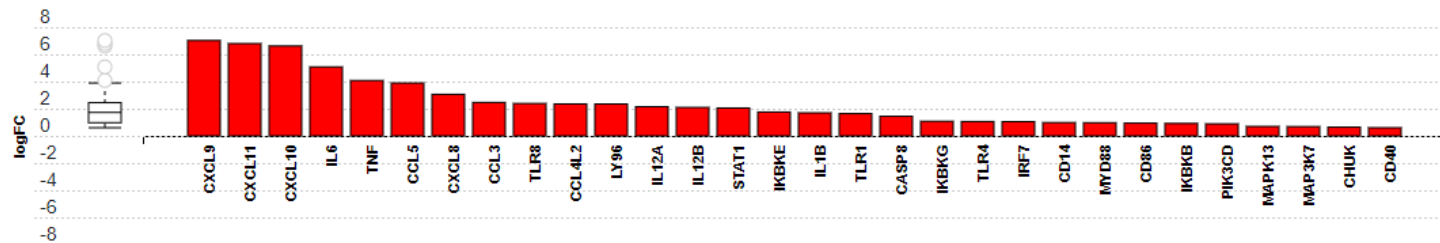


- mRNA expression level exhibited little dose-dependence in ocular tissues. No alternative splicing observed.

# Toll-like receptor signaling

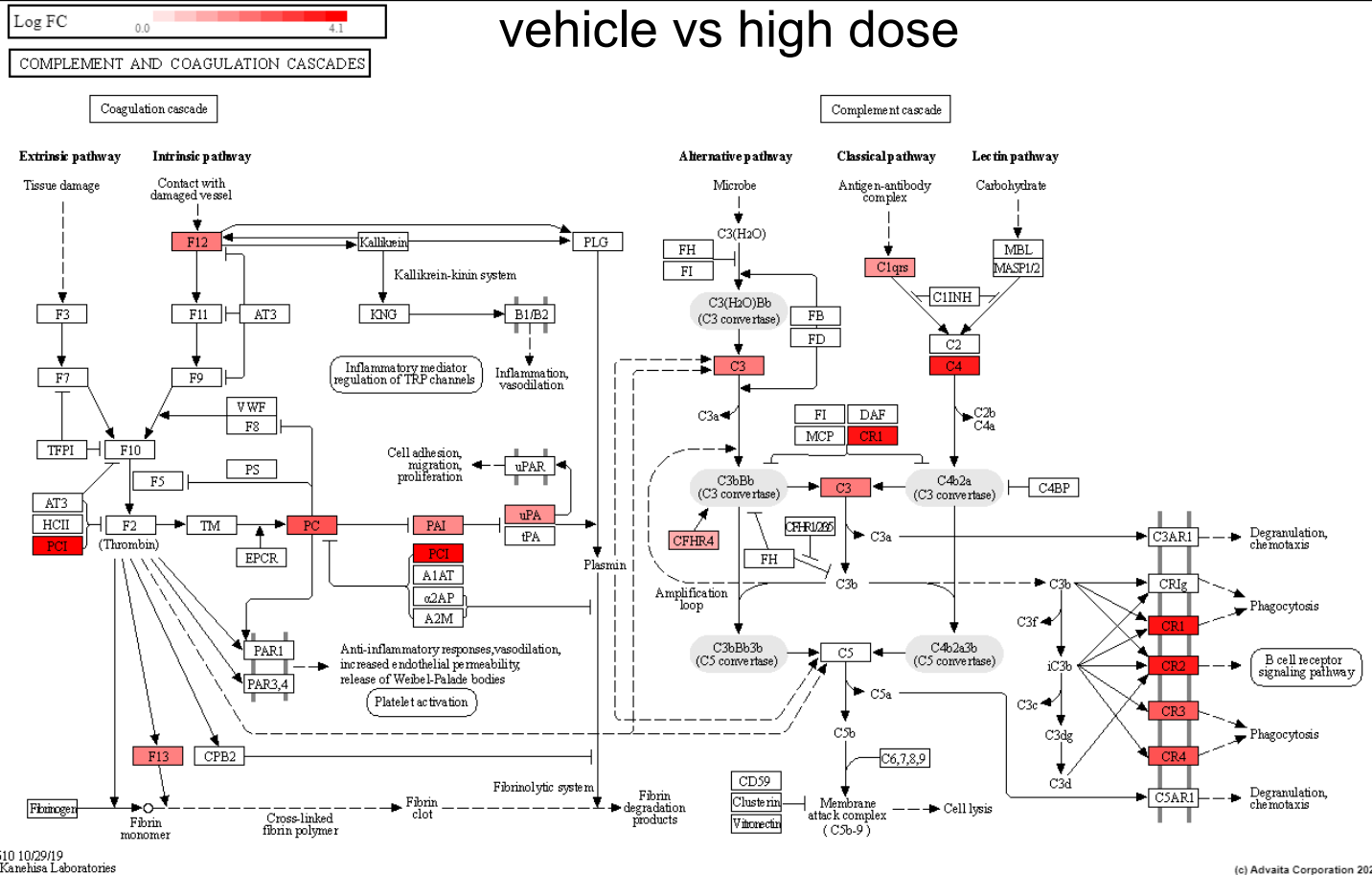


- Unmethylated CpG DNA – TLR9 – MYD88 axis not activated.
- TLR8 suggests response to exogenous mRNA load.<sup>1</sup>
- TLR1/2/4 have reported activation by viral proteins.<sup>2, 3</sup>

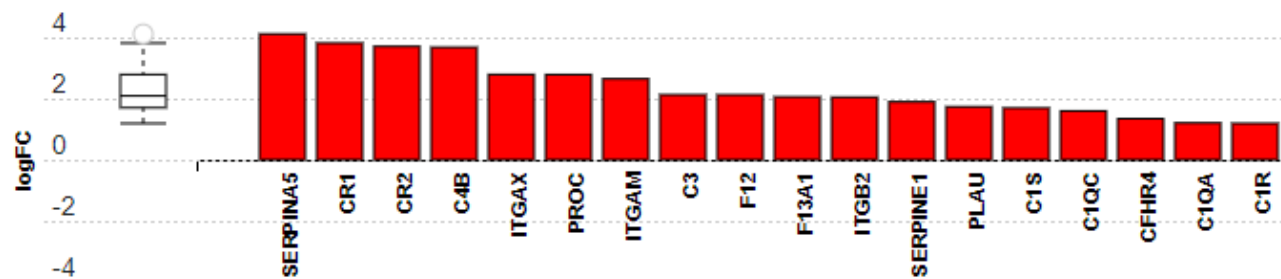


<sup>1</sup>Lester 2014 *JMB* ; <sup>2</sup>Hosel 2012 *Hepatology* ; <sup>3</sup>Zhou 2021 *J Med Virol* ;

# Complement-mediated response observed in dose-responsive manner

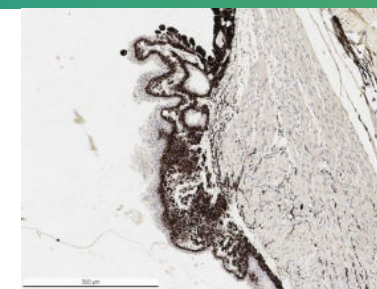
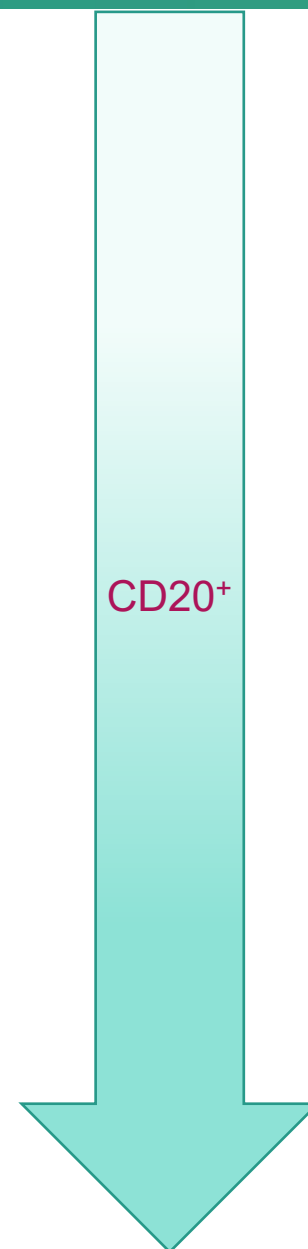
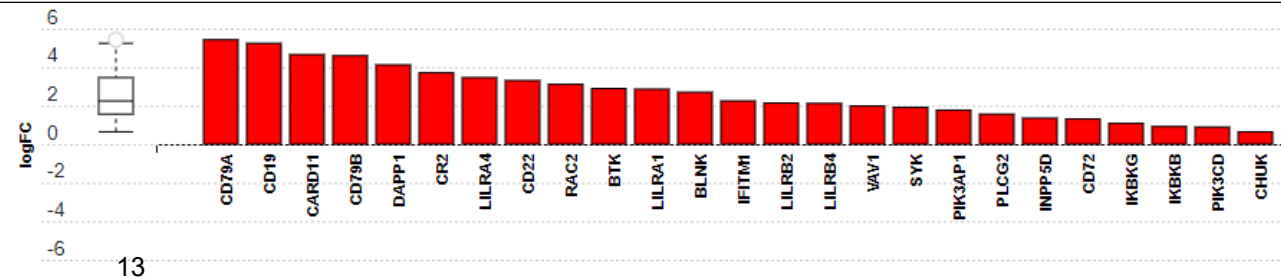
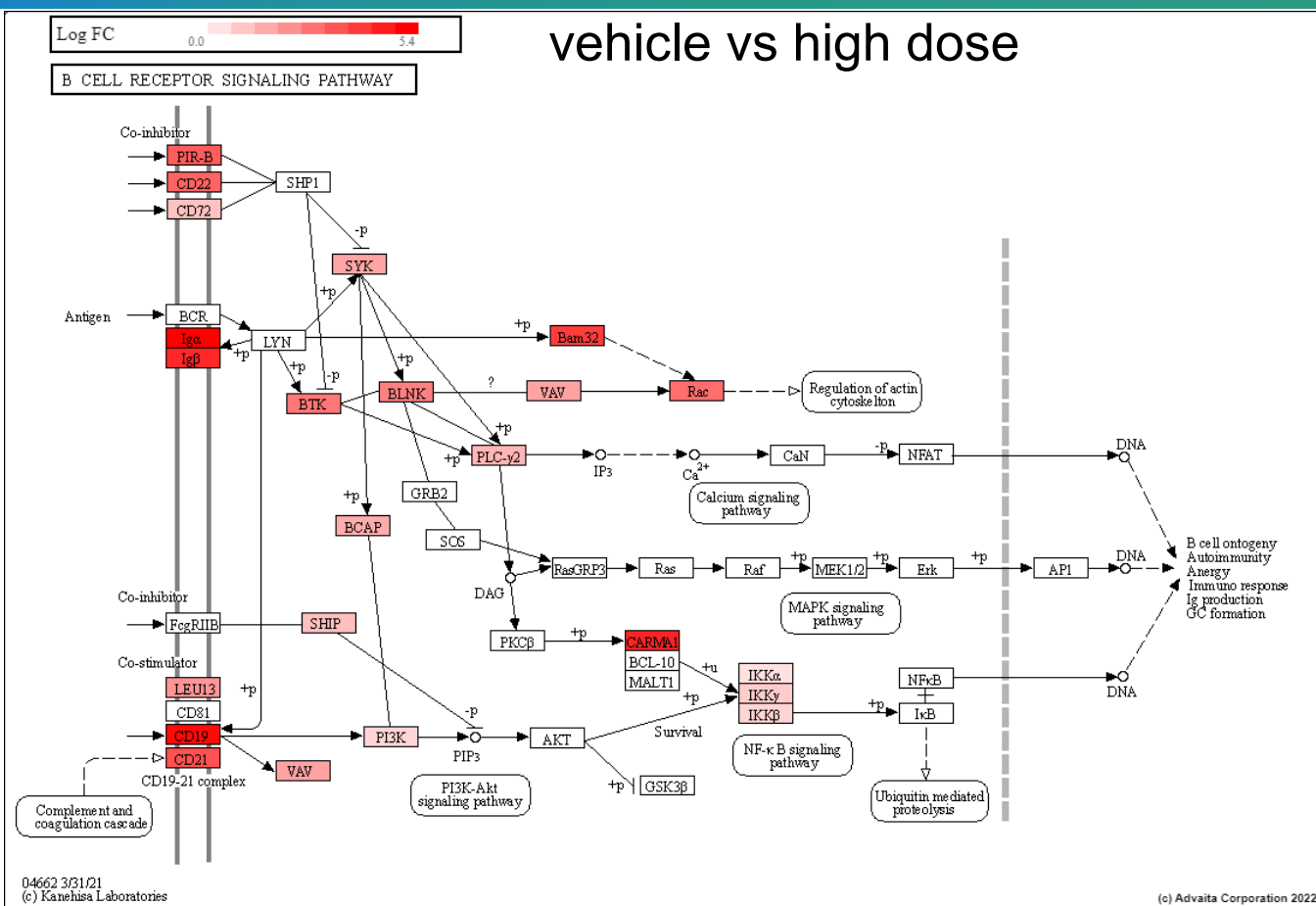


- Could represent antibody complexes bound to rAAV capsid/protein.
- Complement activation previously reported as inflammatory response to high systemic dose of rAAV in clinical trials.<sup>1, 2</sup>
- Complement activation has been known to activate B cells.

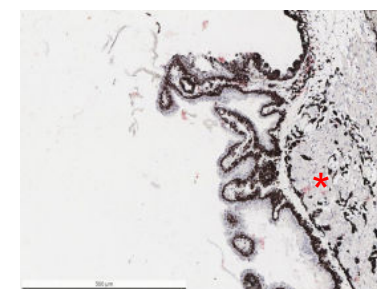


<sup>1</sup>Pfizer 2019 *Press release* ; <sup>2</sup>Solid Biosciences 2019 *Press release*

# B lymphocyte activation and presence increased in dose-responsive manner



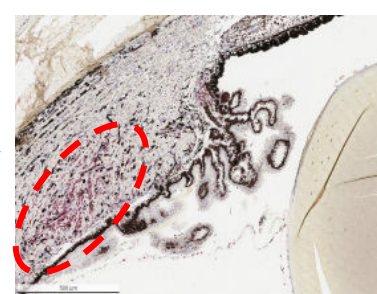
Vehicle



Low dose



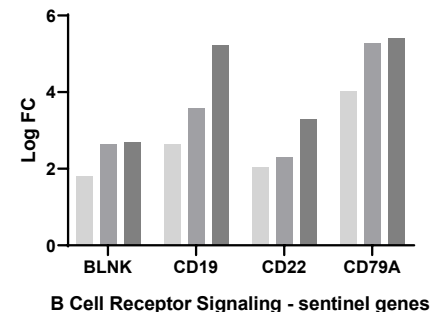
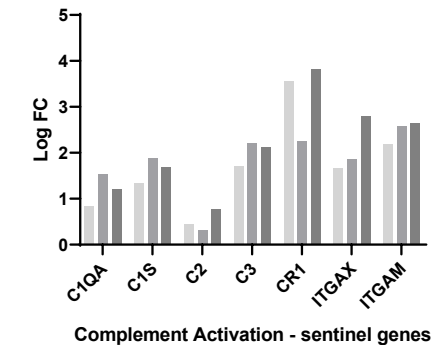
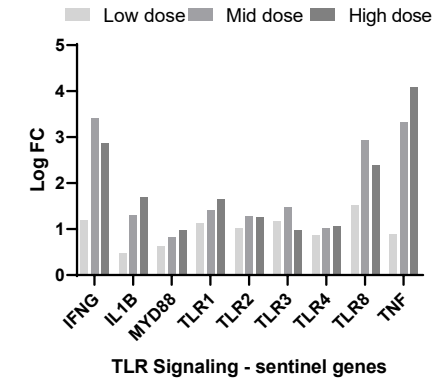
Mid dose



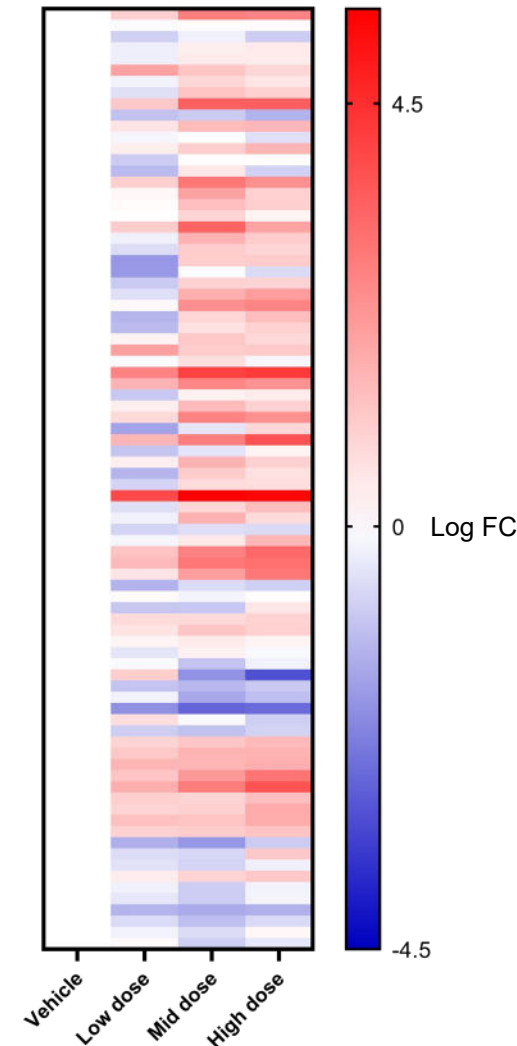
High dose

# Creation of immune landscape signatures

- Sentinel genes were chosen to represent pathways identified by RNA-seq analysis.
- Concordance found between RNA-seq and RT-qPCR.
- Immune landscape signature (ILS) panel could enable:
  - Nonclinical evaluation to interrogate balance between efficacy/tolerability axis
  - Predict patient outcomes
  - Identify high risk patients (e.g., with microvascular comorbidities)
  - Stratify patients that may not need corticosteroids from those that are more susceptible to rAAV-immunogenicity and require them



## RT-qPCR Panel



# Conclusions

**Pathway analysis and IHC converged on dose-dependent activation of innate and adaptive immune response pathways, consistent with rAAV-associated inflammation.**

- Aflibercept expression levels plateaued across doses, unlike dose-dependent adaptive immune response.
- Immune landscape signatures of supra-clinical doses are consistent in nature across ocular tissues – predominately in the anterior tissues of the eye and retina.
- Assessed 3-log fold range above NOAEL.
- No evidence that harnessing ocular cells as biofactories to produce aflibercept leads to expression-associated toxicity/inflammation.
- No evidence that ciliary body architecture was directly affected by Ixo-vec – only indirectly and at supra-clinical doses secondary to dose-dependent increase in inflammation.
- Use of lower doses and an improved steroid prophylaxis, implemented in the LUNA trial, are expected to improve Ixo-vec inflammation profile while preserving therapeutic levels of aflibercept.

# Acknowledgements

- *Brahim Belbellaa\**
- *Kristina Bender\**
- Ruslan N. Grishanin
- Kelly Hanna
- Jie Ma
- Julio Nieves
- Brigit E. Riley
- *Alex Tai\**
- *Maryam Tarazkar\**
- Jenny Vo



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