Exploratory Safety Profile of EDIT-101, a First-in-Human in vivo CRISPR Gene Editing Therapy for CEP290-Related Retinal Degeneration

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Introduction

- CEP290 protein localizes to connecting cilium of photoreceptors and is essential for ciliary assembly and protein trafficking¹.
- CEP290 mutations frequently cause Leber congenital amaurosis type 10 (LCA10), a retinal degenerative condition resulting in severe visual impairment².
- The most common CEP290 mutation causing LCA10 is c.2991+1655A>G within intron 26 (CEP290-IVS26), which results in a nonfunctional CEP290 protein (Figure 1A)².
- Currently, there are no approved LCA10 treatments. To address this medical need, we developed EDIT-101, a Cas9-based gene editing medicine that specifically removes the CEP290-IVS26 mutation to restore photoreceptor function.
- The safety, tolerability, and efficacy of EDIT-101 are being assessed in the BRILLIANCE clinical trial (Figure 2). Viral shedding is a key exploratory safety end-point and here we present viral shedding data from EDIT-101 treated patients.

Figure 1. EDIT-101, a first-in-human *in vivo* CRISPR gene editing medicine for CEP290related retinal degeneration



Local delivery to key subretinal space

CEP290-Related Retinal Degeneration, EDIT-101, and Gene Editing Therapeutic Concept overview. A. The most common CEP290 mutation is c.2991+1655A>G within intron 26 (CEP290-IVS26), which generates a non-functional CEP290 and progressive retinal degeneration that results in LCA10^{1,2}. **B.** EDIT-101 utilizes the photoreceptor-tropic adeno-associated virus type 5 (AAV5) to deliver DNA encoding Staphylococcus aureus Cas9 (SaCas9) expressed under the photoreceptor-specific GRK1 promoter and two gRNAs under U6 promoters. Both gRNAs are highly specific to the CEP290 locus, with no off-targets identified³. C. The two gRNAs direct SaCas9-mediated DNA cleavage up- and downstream of the CEP290-IVS26 mutation, excising it, and restoring CEP290 splicing and function³.



BRILLIANCE study overview: The BRILLIANCE study (NCT03872479) is a Phase 1/2, open-label, single ascending dose clinical trial designed to evaluate safety, tolerability, and efficacy of EDIT-101 in patients with CEP290-related retinal degeneration. Up to 34 patients will be enrolled in up to 5 cohorts to evaluate up to 3 dose levels of EDIT-101. Patients receive a single subretinal injection of EDIT-101 into their eye with the worst visual acuity. Study follow up occurs for 3 years, after which patients are asked to consent to participate in a long-term follow-up study for an additional 12 years.

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Methods

Table 1. Assay sensitivity and linear range

Assay sensitivity and linear range										
Matrix	Linear range (copy number)	Inter-assay precision (% Ct CV)	E (%)	R ²						
Blood	25 to 10 ⁷	0.8–1.8	94–104	≥0.998						
Tears	25 to 10 ⁷	0.6–2.0	92–103	≥0.998						
Nasal mucosa	25 to 10 ⁷	0.3–1.2	94–98	≥0.999						

Sensitivity and linear range assessment: Assay sensitivity in all blood, tears, and nasal mucosa was determined over 5 runs using a linearized plasmid encoding SaCas9 (PLA323) as a surrogate for EDIT-101. Linear range is reported in copies of linearized PLA323 or double stranded EDIT-101 viral vector DNA. Limit of detection (LoD) and limit of quantification (LoQ) were determined to be 10 and 25 copies of either PLA323 or EDIT-101 DNA, respectively. Inter-assay precision indicates the coefficient of variation (CV) between different runs. PCR efficiency and goodness of fit are represented by the E and R² values, respectively.

Table 2. Matrix effect

Matrix effect of sample DNA on qPCR						
Matrix	% of nominal concentration					
Blood	78–81					
Tears	109–115					
Nasal mucosa	110-113					

Matrix effect assessment: Matrix effect in blood, tears, and nasal mucosa was determined by spiking PLA323 with matrix DNA and dividing the observed PLA323 copy number by the input copy number. Matrix DNA alone was not amplified above the LoD, indicating the assay is specific for the SaCas9 DNA sequence (data not shown). Nominal value is equal to 1.28e¹² EDIT-101 viral genomes/mL for blood and tears and 5.49e¹¹ EDIT-101 viral genomes/mL for nasal mucosa.

Table 3. Stability of EDIT-101 in matrices at different temperatures and timepoints

Stability of EDIT-101 in matrices (% of EDIT-101 at T = 0)													
Storage temperature (°C)	18-	-25	2-	2–8 –10 to –30		-60 to -90							
Storage time	4 ł	4 hrs		24 hrs		1 m		Freeze/thaw (n=3)		1 m		6 m	
Viral spike	Low	High	Low	High	Low	High	Low	High	Low	High	Low	High	
Blood	101	102	93	91	90	93	89	82	76	77	95	92	
Tears	103	93	110	101	121	104	121	110	87	90	91	84	
Nasal mucosa	99	97	93	107	109	112	95	104	109	127	109	126	

Stability of EDIT-101 viral vector in matrices mimicking anticipated clinical sample storage conditions: Stability was determined by spiking EDIT-101 into blood, tears, or nasal mucosa samples, subjecting the samples to conditions listed above, and then processing them for DNA extraction and subsequent qPCR analysis. Values within the table represent the observed value divided by the T = 0 value. Values at T = 0 were determined by spiking EDIT-101 into blood, tears, or nasal mucosa samples and immediately processing them for DNA extraction and qPCR analysis. High spike of EDIT-101 at T = 0 were $1.0e^5$, $1.0e^4$, and $5.0e^5$ for blood, tears, and nasal mucosa, respectively. Low spike of EDIT-101 at T = 0 were $1.0e^4$, $1.0e^3$, and $5.0e^4$ for blood, tears, and nasal mucosa, respectively.

Conclusions

- immunogenicity profile for EDIT-101.





We developed a SaCas9-specific qPCR assay to evaluate EDIT-101 viral shedding in patient samples from the BRILLIANCE trial. Detected EDIT-101 viral genome levels in patient samples were significantly lower than administered doses (<1%). Viral genomes were detected more frequently, and at a higher concentration, in tear samples from the treated eye than any other matrix analyzed.

and P30 EY010572. EDIT-101 shedding was transient, with viral clearance below the LoQ observed for most patients by Day 7 and one HD patient by Acknowledgments: We would like to acknowledge and thank all patients in the BRILLIANCE clinical trial, their families, clinical and translational Week 4 in tears and Day 7 in blood and nasal mucosa, suggesting very little risk of systemic viral persistence following treatment. operations, data management, Tricia Gooljarsingh, external PIs, clinical sites, Foundation Fighting Blindness, and partner CROs. No correlation between EDIT-101 doses and viral shedding levels was observed. Taken together, these data suggest a favorable





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