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Gene Therapy Updates for Inherited Retinal Dystrophies

Introduction

Inherited retinal dystrophies (IRDs) encompass a group of genetically diverse disorders, each uniquely influencing distinct retinal cell pathways and retinal areas. IRDs currently affect an estimated 5.5 million individuals worldwide, exerting a profound impact on the quality of life of those affected.¹ Depending on the mutated gene, typical presentations often manifest as colour or night blindness, or peripheral vision blindness progressing to complete blindness.² Consequently, patients grappling with IRDs face not only the physical challenges of their condition, but also endure significant psychosocial and economic repercussions.3

Historically, IRDs were diagnosed and classified based solely on clinical characterization, with no available treatment options. However, advances in genetic characterization have led to the identification of over 270 causative genes, enabling the development of more targeted therapies aiming to restore the function of these mutated genes.2 It is therefore not surprising that this remains an active field of research, aiming to find treatments that can potentially slow down, halt, or even reverse vision loss.

This review aims to provide an updated summary of the current state of IRD treatments, and to discuss recent advancements and emerging therapeutic strategies. The main classifications that will be explored are macular dystrophies; stationary cone dystrophies; rod-cone dystrophies; Leber congenital amaurosis (LCA); and chorioretinal dystrophies. In this review,

Table 1. Comprehensive overview of clinical trials for targeted therapies in common inherited retinal diseases (as of January 1, 2024). Table 1. Comprehensive overview of clinical trials for targeted therapies in common inherited retinal diseases (as of January 1, 2024).

Phase III clinical trials (registered on clinicaltrials. gov) were selected for literature analysis. In cases where no Phase III trials were available, Phase I or II clinical trials with results were analyzed. Stationary rod dystrophies as well as progressive cone and cone-rod dystrophies will not be discussed, as available studies for these disorders are limited to the preclinical phase (Table 1).

Gene Therapy

Gene therapy involves the introduction or modification of genetic material within cells to replace the function of mutated genes. The eye is an ideal target for gene therapy because of its tight blood-ocular barrier, making it relatively immune privileged. In addition, the retina is readily accessible, and a patient's response to therapy can easily be monitored through clinical examinations and imaging. Managing monogenic autosomal-recessive and X-linked mutations is facilitated by the loss of function of these abnormal proteins.4 Conversely, dominant mutations are less amenable to genetic therapies, as the abnormal gain-of-function proteins impede the action of the correctly synthesized ones post-treatment.4

In gene therapy, three main approaches are used to address mutations. The first, and most prevalently used, is gene augmentation.4 This technique is uniquely employed for monogenic recessive or X-linked inherited diseases, and introduces a wild type copy of the pathogenic gene into target retinal cells, thereby augmenting the production of a functional protein.⁴ Second, gene editing may be used for dominant mutations.4 This involves the Clustered Regularly Interspaced Short Palindromic Repeats (CRISPR‑Cas9) technology, where a gene-specific guide RNA is linked to a Cas9 endonuclease and identifies, cuts and removes specific portions of DNA to be replaced.4 The downside to this technique, however, is the potential of creating novel mutations.4 Last, gene inactivation can also be used for dominant mutations.4 Here, small interfering RNAs (siRNAs) or antisense oligonucleotides (AONs) can correct or block the production of the mRNA transcribed from the mutated DNA gene.4

 Subretinal and intravitreal injections are the most common modes of delivery. Subretinal injections are used for outer retinal targets and require smaller amounts to achieve a therapeutic effect.5 They are locally administered

between photoreceptors and the RPE layer, and complications resemble those of pars plana vitrectomy.⁵ Intravitreal injections are used for inner retinal targets, but are more immunogenic and it is harder to transduce photoreceptors and RPE cells because of the barrier effect from the inner limiting membrane.⁵

Apart from AONs, genetic therapy is delivered to target retinal cells via viral or non-viral vectors. Within viral vectors, adeno-associated vectors (AAV) have a smaller gene size carrying capacity (4.5 kb to 4.9 kb) and do not integrate into the host's genome.⁶ Lentiviruses (LV) carry genes up to 8 kb but integrate the host's genome, causing a small risk of insertional mutagenesis.⁶ On the other hand, non-viral vectors have a lower risk of genotoxicity and immunogenicity, but have a lower specificity and are less stable than viral vectors.⁶

Macular Dystrophies

Macular dystrophies include Stargardt disease, Best vitelliform macular dystrophy (BVMD), X-linked retinoschisis (XLRS) and pattern dystrophies. In this review, Stargardt disease, BVMD and XLRS will be discussed.

Stargardt disease

Inherited in an autosomal recessive manner, Stargardt affects 1:8000 to 1:10,000 individuals, making it a leading cause of juvenile macular degeneration.7 Patients affected by this dystrophy often have a mutated ATP-binding cassette sub-family A gene (ABCA4).⁷ Lack of this protein causes toxic accumulation of bisretinoid compounds in the RPE, leading to RPE dysfunction and causing visual impairment.⁷ The main obstacle in developing therapies for this gene is its large size (6.8 kb).7 Preclinical studies are currently exploring the use of non-viral delivery systems such as covalently closed and circular DNA (C3DNA).7 Studies on porcine and non-human primate retinas have provided evidence of sustained ABCA4 protein expression up to six months post-treatment, showing promising results for possible human applications in the future.7

Best Vitelliform Macular Dystrophy

Best disease is usually inherited in an autosomal dominant pattern and affects 1:10,000 individuals. 8 It is caused by a mutation in the BEST1 gene, responsible for the expression of the transmembrane protein Bestrophin 1, which is

greatly implicated in the calcium homeostasis of the RPE.⁹ Best disease evolves through six stages, from the subclinical/previtelliform phase where the fundus appears normal, to the vitelliform stage with classic egg yolk lesions on the macula, gradually evolving toward the atrophic stage.9 The most commonly reported symptoms are vision dimness, metamorphopsia and scotoma, but these symptoms vary largely between individuals.9 Although genetic testing is needed for definitive diagnosis, there are no available pre-clinical or clinical gene therapy studies.

X-linked retinoschisis

X-linked retinoschisis causes predominant central vision loss in 1:5,000 to 1:25,000 males and is associated with a mutation in the retinoschisin 1 gene (RS1).¹⁰ This retinoschisin membrane protein is involved in retinal cell layer organization and cell adhesion, explaining why patients develop macular schisis that may even extend to the peripheral retina.10 As opposed to subretinal injections, intravitreal gene therapy delivery is the preferred approach, as patients have a higher predisposition to retinal detachments.10 Although preclinical studies have shown effective gene augmentation therapies in non-human models, the results of two Phase I/II trials introducing AAVs intravitreally (AAV8-RS1) showed inflammation in almost all patients with no improvement in visual function.10

Stationary Cone Dystrophies

Stationary cone dystrophies such as achromatopsia and blue cone monochromatism represent a group of IRDs characterized by a stable and non-progressive impairment of cone photoreceptors. As achromatopsia is the more classic example, it will be discussed in this review.

Achromatopsia

Achromatopsia (ACHM) is an autosomal recessive, inherited disorder affecting approximately 1:30,000 people.11 It is characterized by the early, insidious loss of photoreceptor cones, leading to vision loss; colour blindness; hemeralopia; photophobia; and nystagmus.¹¹ Approximately 90% of patients with ACHM carry mutations in the cyclic nucleotide-gated channel alpha 3 (CNGA3) or beta 3 (CNGB3) gene, which encodes essential components of the phototransduction cascade.11 Although there are currently no approved therapies, ongoing Phase I

and II AAV gene augmentation clinical studies, including two that have shown gains in visual acuity and contrast sensitivity in all nine treated patients with CNGA3 mutations, are underway.12

Rod-cone Dystrophies

Rod-cone dystrophies include retinitis pigmentosa (RP), Usher syndrome, enhanced S-cone syndrome, and Bietti crystalline dystrophy. In this review, RP and Usher syndrome will be discussed.

Retinitis pigmentosa

Retinitis pigmentosa is the most common IRD, affecting 1:4,000 people.13 It can be inherited through autosomal dominant, autosomal recessive, or X-linked genetic patterns, reflecting its complex genetic etiology. It is characterized by primary rod and secondary cone Degeneration.¹³ Symptoms include nyctalopia, followed by progressive peripheral visual field loss.13

Accounting for up to 15% of all RP cases, X-linked RP is the most severe form of the condition.13 Seventy to seventy-five percent of these patients have mutations in the GTPase regulator gene (RPGR), a protein involved in ciliary transport and critical in maintaining photoreceptor integrity.13 In a Phase I/II dose escalation gene augmentation trial (AAV2/5‑RPGR), six out of seven patients treated with low or intermediate doses showed stability or improvement in retinal sensitivity at 12 months.¹⁴ In the higher dose cohort however, two out of three treated patients showed signs of inflammation and no signs of visual improvement.¹⁴

The USH2A gene codes for usherin, a protein necessary for basement membrane and photoreceptor integrity.15 In patients with Usher syndrome Type 2a and some non-syndromic forms of RP, mutations have been found in the USH2A exon 13, leading to clinical studies using an AON designed to skip this exon $(QR-421a)$.¹⁵ In the STELLAR study, all 20 treated patients had visual acuity improvement, objectified by an average gain of six letters or improvement in total retinal sensitivity at 48 weeks post-treatment.¹⁵ There were no reported serious adverse events.¹⁵

Leber Congenital Amaurosis

Commonly inherited in an autosomal recessive manner and affecting 1:50,000 to 1:100,000 people, LCA is one of the most severe forms of retinal dystrophy.16 In addition to severe vision loss, patients often have accompanying sensory nystagmus, near-absent pupillary response, and a non-detectable electroretinogram response.17 Voretigene neparvovec‑rzyl (AAV2‑hRPE65v2) is currently the only FDA approved gene therapy for IRDs. Recent surgical technique enhancements avoid bleeding and inadvertent macular hole formation.18 This subretinal gene augmentation therapy targets biallelic RPE65 mutations frequently found in LCA type 2 patients, accounting for up to 16% of all LCA cases.¹⁶ The RPE65 gene is responsible for converting trans-retinyl esters to 11-cis-retinols, and its dysfunction leads to an inability to regenerate pigments in photoreceptors.16 Recent clinical studies have demonstrated sustained partial rescue of photoreceptor function for up to four years post-treatment.16 This was objectified via multi-luminance mobility tests, visual field testing, and full-field stimulus tests.16 The most commonly reported adverse event post-treatment was central retinal thinning.¹⁹

In type 10 LCA, the gene encoding the centrosomal protein 290 (CEP290) is frequently mutated, leading to faulty photoreceptor cilia function.20 In Phase 1b and II clinical trials, AONs (QR‑110) have been used to correct the faulty mRNA before protein translation.²⁰ Five out of 11 treated patients showed a -0.3 logMAR improvement in visual acuity one year post-treatment; the most common adverse event was the development of cataracts.²⁰ A Phase III trial has been completed, although its results are yet to be released. 21 In addition, there are ongoing Phase I/II clinical trials using the CRISPR-Cas9 system to eliminate the IVS26 mutation in this same CEP290 gene.²¹

Chorioretinal Dystrophies

Chorioretinal dystrophies are a distinct subgroup of IRD characterized by progressive degeneration in both the choroid and retina. The most prevalent example, choroideremia (CHM), will be discussed in this review.

Choroideremia

CHM is an X-linked recessive dystrophy affecting 1:100,000 to 1:200,000 males.²² It is characterized by the centripetal loss of photoreceptors, RPE cells and the choriocapillaris, even reaching the fovea in severe cases.²³ Patients begin to report nyctalopia and peripheral vision loss in late childhood, progressing to near-complete vision loss by the age of 40.23 Choroideremia is caused by a mutation in the CHM gene, which encodes the Rab escort protein 1 (REP1), an enzyme essential for intracellular trafficking of vesicles.23 A recent Phase III STAR study treating one eye per patient with an either low or high dose of timrepigene emparvovec (BIIB111/AAV2‑REP1) allowed patients to gain in visual acuity when compared to the control eye.²² However, the number of treated patients meeting this three-line improvement did not reach statistical significance, which is why there has been no regulatory approval.22 In Phase I and II clinical trials, there were rare cases of adverse events, with two cases of retinal holes over a non-functional retina and one case of intraretinal immune response.²²

Conclusion

 In summary, ongoing research endeavours are focused on the development of sustainable gene therapies for IRDs, previously considered untreatable. The primary challenges include the development of delivery methods with reduced immunogenicity, ensuring enduring treatment effects, and establishing therapies that minimize host mutagenesis. Optimal treatment candidates appear to be patients with early-stage diagnoses and gradual disease progression, as these factors provide a broader window for treatment before the degeneration of target cells. This article provides an overview of a select number of ongoing clinical studies, indicating a cautious yet hopeful outlook for the future of IRD treatments.

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