

24 January 2023 EMADOC-628903358-34665 European Medicines Agency

# Statement on the amended policy on orphan designations for inherited retinal dystrophies

The Committee on Orphan Medicinal Products (COMP) has adopted a statement explaining its amended policy on orphan designations (OD) for inherited retinal dystrophies (IRD), the scientific and regulatory rationale for this change, and potential next steps for developers of orphan medicinal products.

# Amended policy on orphan designations for inherited retinal dystrophies

Based on a thorough review, supported by a consultation of IRD clinical experts and patients, the COMP has adopted a new approach for designating conditions in IRDs. The COMP has decided that three options will be available for orphan conditions in an IRD OD application.

The three options are:

- For therapies that are relatively broadly applicable in IRDs, terms (e.g. *Rod-dominant phenotype*) can be selected from table 1 for orphan designated conditions. If a particular broad therapy could target more than one group, multiple orphan designations may be needed.
- For targeted gene therapies, the OD condition can be constructed from the term "inherited retinal dystrophy due to dysfunction in the target-gene.
- Finally, for some IRDs which may not fit the table 1 scheme, an occasional singular orphan designation outside the table 1 structure may still be necessary for non-gene therapy product(s).

**Table 1** Grouping for inherited retinal diseases for the purpose of orphan designation

- 1. Non-syndromic IRD
  - 1.1. Cone-dominant phenotype\*
  - 1.2. Rod-dominant phenotype
  - 1.3. Macular dystrophy
- 2. Syndromic IRD
  - 2.1. Cone-dominant phenotype
  - 2.2. Rod-dominant phenotype
  - 2.3. Macular dystrophy
- 3. Inherited choroidal dystrophies
- 4. Hereditary vitreoretinopathies
- \* Phenotypes include inherited pathological dysfunction as well as inherited progressive degenerations

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# Scientific and regulatory rationale

Inherited retinal dystrophies are a group of diseases affecting retinal structure and function. The classical names of IRDs (e.g. retinitis pigmentosa, Leber congenital amaurosis) were developed when the underlying genetics were not known or not well understood, and were based on clinical appearance, and other signs and symptoms. The genetics of IRDs are very complex.<sup>1,2,3,4,5,6,7,8</sup> It is now understood that one abnormal gene can have different clinical appearances (phenotypes) e.g. abnormalities in the *RDH12* gene has been associated with early-onset severe retinal dystrophy/Leber congenital amaurosis (EOSRD/LCA), (mild) retinitis pigmentosa, cone-rod dystrophy, and macular dystrophy. It is also understood that one phenotype (e.g. autosomal recessive retinitis pigmentosa) can be associated with variants in more than 60 different genes such as the *ABCA4* gene, *AGBL5* gene, *AHR* gene and others.

Upon request by sponsors, the COMP has the mandate to consider the submitted application and designate an orphan condition if all criteria are satisfied. *The <u>Guideline on the format and content</u> of applications for designation as orphan medicinal products and on the transfer of designations from one sponsor to another*, and <u>Commission notice 2016/C 424/03</u> provides information on what constitutes a *valid condition*, and what may be an acceptable subset of a disease for the purpose of the orphan designation.

Previously, COMP has used classical IRD terms as OD conditions. Given the complexity of IRDs, defining the OD condition in IRD is not straight forward. In addition, using classically derived IRD names for the orphan condition may mean that some otherwise treatable patients may be out of scope of an approved treatment if these patients show different signs and symptoms to the classical group. According to scientific literature, the identification of the genetic cause of the disease represents a hallmark for the patients IRD.<sup>9,10,11</sup> With the approval a targeted gene-therapy, it is also timely to consider the OD policy in the light of such a licensed indication.

Therefore, the COMP undertook a review to ascertain the *state of the art* in IRDs to assess what would be the best set of terms for orphan designation in this therapeutic setting. The review included a literature review, overview of 64 active OD in IRDs in the EU including the nature of the product, possible alternative grouping systems of IRDs <sup>12,13,14,15,1</sup>, and a consultation of IRD clinical experts and patients. This consultation included perspectives from patients and clinicians on several aspects such as prevalence, <sup>16,9,17</sup> on current clinical practice, the most up to date sources of information, and optimal clinical groupings. Finally, COMP considered of the pros and cons of different possible approaches to setting the OD for IRDs.

Based on these in-depth considerations, supported by the outcome of the consultation, the COMP has adopted a new approach for designating conditions in IRDS. The COMP has decided that three options as identified above will be available for orphan designated condition in an IRD OD application.

## **Impact on sponsors**

Sponsors with existing ODs in IRD may consider amending their OD before filing a marketing authorisation application (MAA) or protocol assistance in the event that the existing designation would not cover the intended target patient population, and hence a potential therapeutic indication in the case of MAA.

For new OD submissions, sponsors should specify the orphan condition applied for and fully justify the chosen approach in line with the recommendations in this document and the orphan legislation and guidance as these will be considered by the COMP when deciding on the OD application.

# Information sources

Information on genetics of IRDs:

https://web.sph.uth.edu/RetNet/disease.htm

https://www.omim.org/

### Information on orphan designations in the EU:

- Active orphan designations in the EU: the Community Registry of Orphan medicinal Products https://ec.europa.eu/health/documents/community-register/html/reg\_od\_act.htm?sort=a
- Guidance

Orphan designation: Overview | European Medicines Agency (europa.eu)

### References

- Leroy B. Brave new world; gene therpay for inherited retinal diseases. EyeNet. July 1, 2018. Accessed Dec 13, 2022. <u>https://www.aao.org/assets/bebfbaef-a092-45b0-9883-</u> <u>c563331546ae/636649294795430000/july-2018-eyenet-supplement-pdf?inline=1</u>
- Cremers FPM, Boon CJF, Bujakowska K, Zeitz C. Special Issue Introduction: Inherited Retinal Disease: Novel Candidate Genes, Genotype-Phenotype Correlations, and Inheritance Models. *Genes (Basel)*. Apr 16 2018;9(4)doi:10.3390/genes9040215
- Tsang SH, Sharma T. Leber Congenital Amaurosis. *Adv Exp Med Biol*. 2018;1085:131-137. doi:10.1007/978-3-319-95046-4\_26
- 4. Tatour Y, Ben-Yosef T. Syndromic Inherited Retinal Diseases: Genetic, Clinical and Diagnostic Aspects. *Diagnostics (Basel)*. Oct 2 2020;10(10)doi:10.3390/diagnostics10100779
- 5. Fuster-Garcia C, Garcia-Bohorquez B, Rodriguez-Munoz A, et al. Usher Syndrome: Genetics of a Human Ciliopathy. *Int J Mol Sci*. Jun 23 2021;22(13)doi:10.3390/ijms22136723
- Talib M, Boon CJF. Retinal Dystrophies and the Road to Treatment: Clinical Requirements and Considerations. *Asia Pac J Ophthalmol (Phila)*. May-Jun 2020;9(3):159-179. doi:10.1097/APO.00000000000290
- Schneider N, Sundaresan Y, Gopalakrishnan P, et al. Inherited retinal diseases: Linking genes, diseasecausing variants, and relevant therapeutic modalities. *Prog Retin Eye Res.* Jul 2022;89:101029. doi:10.1016/j.preteyeres.2021.101029
- 8. Verbakel SK, van Huet RAC, Boon CJF, et al. Non-syndromic retinitis pigmentosa. *Prog Retin Eye Res*. Sep 2018;66:157-186. doi:10.1016/j.preteyeres.2018.03.005
- Perea-Romero I, Gordo G, Iancu IF, et al. Genetic landscape of 6089 inherited retinal dystrophies affected cases in Spain and their therapeutic and extended epidemiological implications. *Sci Rep.* Jan 15 2021;11(1):1526. doi:10.1038/s41598-021-81093-y
- Kellner U, Jansen S, Bucher F, Stingl K. [Diagnosis of inherited retinal dystrophies. Relevance of molecular genetic testing from the patient's perspective]. *Ophthalmologie*. Aug 2022;119(8):820-826. Diagnostik erblicher Netzhautdystrophien. Stellenwert molekulargenetischer Diagnostik aus Patientenperspektive. doi:10.1007/s00347-022-01602-w
- Lorenz B, Tavares J, van den Born LI, Marques JP, Scholl HPN, Group EVn. Current Management of Inherited Retinal Degeneration Patients in Europe: Results of a Multinational Survey by the European Vision Institute Clinical Research Network. *Ophthalmic Res.* 2021;64(4):622-638. doi:10.1159/000514540

- 12. Fenner BJ, Tan TE, Barathi AV, et al. Gene-Based Therapeutics for Inherited Retinal Diseases. *Front Genet*. 2021;12:794805. doi:10.3389/fgene.2021.794805
- 13. Kohler S, Gargano M, Matentzoglu N, et al. The Human Phenotype Ontology in 2021. *Nucleic Acids Res*. Jan 8 2021;49(D1):D1207-D1217. doi:10.1093/nar/gkaa1043
- 14. Sergouniotis PI, Maxime E, Leroux D, et al. An ontological foundation for ocular phenotypes and rare eye diseases. *Orphanet J Rare Dis.* Jan 9 2019;14(1):8. doi:10.1186/s13023-018-0980-6
- 15. Georgiou M, Fujinami K, Michaelides M. Inherited retinal diseases: Therapeutics, clinical trials and end points-A review. *Clin Exp Ophthalmol*. Apr 2021;49(3):270-288. doi:10.1111/ceo.13917
- Gong J, Cheung S, Fasso-Opie A, et al. The Impact of Inherited Retinal Diseases in the United States of America (US) and Canada from a Cost-of-Illness Perspective. *Clin Ophthalmol*. 2021;15:2855-2866. doi:10.2147/OPTH.S313719
- Hanany M, Sharon D. Allele frequency analysis of variants reported to cause autosomal dominant inherited retinal diseases question the involvement of 19% of genes and 10% of reported pathogenic variants. *J Med Genet.* Aug 2019;56(8):536-542. doi:10.1136/jmedgenet-2018-105971