

Resource Summary Report

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Hereditary Hearing Loss Homepage

RRID:SCR_006469

Type: Tool

Proper Citation

Hereditary Hearing Loss Homepage (RRID:SCR_006469)

Resource Information

URL: <http://hereditaryhearingloss.org/>

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Description: Overview of the genetics of hereditary hearing impairment for researchers and clinicians. The site lists data and references for all known gene localizations and identifications for nonsyndromic hearing impairment, and several for syndromic hearing loss. For syndromic hearing impairment, only a few of the most frequent forms are covered. An atlas of cochlea with genes listed can be accessed from this site.

Abbreviations: Hereditary Hearing Loss

Resource Type: atlas, database, topical portal, portal, data or information resource

Keywords: cochlea, syndromic, nonsyndromic, gene, genetics, hearing impairment, hearing, ear, FASEB list

Related Condition: Hereditary hearing impairment, Hearing impairment

Resource Name: Hereditary Hearing Loss Homepage

Resource ID: SCR_006469

Alternate IDs: nif-0000-00075, OMICS_01542

Record Creation Time: 20220129T080236+0000

Record Last Update: 20260728T043233+0000

Ratings and Alerts

No rating or validation information has been found for Hereditary Hearing Loss Homepage.

No alerts have been found for Hereditary Hearing Loss Homepage.

Data and Source Information

Source: [SciCrunch Registry](#)

Usage and Citation Metrics

We found 455 mentions in open access literature.

Listed below are recent publications. The full list is available at [nidm-terms](#) .

Li Y, et al. (2025) SLC26A4 C.317C > A Variant: Functional Analysis and Patient-Derived Induced Pluripotent Stem Line Development. Molecular genetics & genomic medicine, 13(4), e70098.

Jagannath K, et al. (2025) Identification of known and novel genetic variants in sensorineural hearing loss: insights from whole exome sequencing in Indian families. Molecular biology reports, 52(1), 883.

Ahmed S, et al. (2025) Large-scale audiometric phenotyping identifies distinct genes and pathways involved in hearing loss subtypes. Communications biology, 8(1), 1515.

Kim JA, et al. (2025) Systematic genetic assessment of hearing loss using whole-genome sequencing identifies pathogenic variants. Experimental & molecular medicine, 57(4), 775.

Jang SH, et al. (2025) Recent preclinical and clinical advances in gene therapy for hereditary hearing loss. Molecules and cells, 48(12), 100285.

Wang W, et al. (2025) Clinical application of preimplantation genetic testing based on low-coverage next-generation sequencing with linkage analyses in hereditary hearing loss families. Journal of assisted reproduction and genetics, 42(6), 1989.

Kim JA, et al. (2025) In vivo consequences of varying degrees of OTOA alteration elucidated using knock-in mouse models and pseudogene contamination-free long-read sequencing. Genes & diseases, 12(3), 101533.

Swetha J, et al. (2025) A 250-kb Microdeletion Identified in Chromosome 16 Is Associated With Non-Syndromic Sensorineural Hearing Loss in a South Indian Consanguineous Family. Journal of audiology & otology, 29(1), 31.

Bazazzadegan N, et al. (2025) A Novel Candidate Gene MACF1 is Associated with Autosomal Dominant Non-syndromic Hearing Loss in an Iranian Family. Archives of Iranian medicine, 28(1), 63.

Yalcouyé A, et al. (2025) Whole-exome sequencing reveals known and candidate genes for hearing impairment in Mali. HGG advances, 6(1), 100391.

Yan D, et al. (2025) Generation and Auditory Phenotypic Characterization of Prps1 p.Ala87Thr Mouse Knock-In Model for Human DFNX1 Deafness. Clinical genetics, 108(5), 566.

Arai Y, et al. (2025) Novel OTOG Variants and Clinical Features of Hearing Loss in a Large Japanese Cohort. *Genes*, 16(1).

Ukaji T, et al. (2025) AAV-mediated base editing restores cochlear gap junction in GJB2 dominant-negative mutation-associated syndromic hearing loss model. *JCI insight*, 10(5).

Morovvati S, et al. (2025) The clinical and genetic spectrum of twenty-six individuals with hearing loss affected by MYO15A variants. *Scientific reports*, 15(1), 14320.

Xia R, et al. (2025) Therapeutic restoration of miR-96 prevents hearing loss in mice through modulation of noise-induced and genetic pathways. *iScience*, 28(5), 112355.

Lee SY, et al. (2025) Comprehensive genetic profiling of sensorineural hearing loss using an integrative diagnostic approach. *Cell reports. Medicine*, 6(7), 102206.

Zafeer MF, et al. (2025) Human organoids for rapid validation of gene variants linked to cochlear malformations. *Human genetics*, 144(4), 375.

Qi M, et al. (2025) NOX2 Contributes to High-Frequency Outer Hair Cell Vulnerability in the Cochlea. *Advanced science (Weinheim, Baden-Wurttemberg, Germany)*, 12(34), e08830.

Shadab M, et al. (2025) Recurrent and Novel Pathogenic Variants in Genes Involved with Hearing Loss in the Pakistani Population. *Molecular diagnosis & therapy*, 29(4), 519.

Vitry S, et al. (2025) Advancing precision ear medicine: leveraging animal models for disease insights and therapeutic innovations. *Mammalian genome : official journal of the International Mammalian Genome Society*, 36(2), 417.