



CLIA: 10D2337214

Hearing Impairment Panel

Requisition & Consent Form

Patient Information

Last Name	First Name	DOB	Sex	MRN
Address		City	State	Zip
Phone Number				
Email				
Ethnicity ☐ African American ☐ Ashkenazi Jewish ☐ East Asian ☐ Hispanic ☐ Middle Eastern ☐ White ☐ Other				

Insurance/Billing Information

Insurance Name	Member ID	Group ID
Primary Name	Primary DOB	Relationship
☐ Self ☐ Spouse ☐ Child		

Ordering Provider/ Facility

Provider Name	NPI	Facility Name	Email
Address		City	State
Zip		Phone Number	

Specimen Information

Specimen Type		
Collector Name	Collection Date/ Time	Collection Site
☐ Office ☐ Lab ☐ Home ☐ Other		

Confirmation of Medical Necessity for Genetic Testing

I certify that I am a licensed healthcare provider (or authorized representative/genetic counselor) ordering genetic testing. I confirm that the patient has been informed of the purpose, scope, limitations, and potential risks of this testing, has had the opportunity to ask questions and seek genetic counseling, and has provided informed consent, which is on file. I attest that this testing is medically necessary and will be used in the clinical management of the patient. I further confirm that the patient authorizes **Revive Genetics** to release relevant information to third-party payors for reimbursement purposes, to receive direct payment for services rendered, and, where applicable, to act as the patient's designated representative for insurance appeals. I acknowledge that the patient understands and accepts financial responsibility for any amounts not covered by insurance, including payments made directly to the patient.

Medical Professional Signature: _____ Date: _____

Print Name:

Website www.revivegenetics.us Email info@revivegenetics.us	Address 9000 SW 152 nd St Suite107, Palmetto Bay, FL 33157	Lab Director Uzma Farooq, MD	Phone Number 786-250-3419
--	---	--	-------------------------------------



CLIA: 10D2337214

Hearing Impairment Panel

Requisition & Consent Form

Test Selection

Select	Panel	Genes	Brief Description
	Express Gene™ Hearing Impairment Genetic Testing Panel	254 Genes	Identifies inherited genetic variants associated with hearing impairment by evaluating genes involved in auditory function, cochlear development, inner ear structure, sensory hair cell function, and auditory signal transmission.

Indications for Testing (Check All That Apply)

☐ Diagnostics
 ☐ Family History
 ☐ Recurrent or Unexplained Infections
 ☐ Patient Management

Clinical Information

Genetic Testing Done Previously: ☐ Yes ☐ No	If Yes what test:	Date of Previous Test: ____/____/____
---	-------------------	---------------------------------------

The panel includes all genes listed below. If only specific gene(s) are requested, please select them below.
For additional information about this test, please visit our company website.

ABHD12, ACTB, ACTG1, ADCY1, ADGRV1, AIFM1, ALMS1, AMMECR1, ANKH, ANLN, ARSG, ASAH1, ATP2B2,
 ATP6V1B1, BCS1L, BDP1, BLOC1S6, BRAF, BSND, BTBD, CABP2, CACNA1D, CATSPER2, CCDC50, CD151, CD164,
 CDC14A, CDC42, CDH23, CDKN1C, CEACAM16, CEP250, CEP290, CEP78, CHD7, CHSY1, CIB2, CISD2, CLDN14,
 CLIC5, CLPP, CLRN1, COCH, COL11A1, COL11A2, COL17A1, COL1A1, COL2A1, COL3A1, COL4A3, COL4A4,
 COL4A5, COL4A6, COL5A1, COL5A2, COL9A1, COL9A2, COL9A3, CRYM, CYBA, DCAF17, DCDC2, DIABLO, DIAPH1,
 DIAPH3, DLX5, DMXL2, DNMT1, DSPP, EDN3, EDNRA, EDNRB, EFTUD2, EIF3F, ELMOD3, EPS8, EPS8L2, ERCC2,
 ERCC3, ESPN, ESRRB, EYA1, EYA4, FAM136A, FAM65B, FDXR, FGF3, FGFR1, FGFR2, FGFR3, FITM2, FOXC1, FOXI1,
 GATA3, GDF6, GFER, GIPC3, GJA1, GJB2, GJB3, GJB6, GLI3, GPSM2, GREB1L, GRHL2, GRXCR1, GRXCR2, GSDME,
 HARS, HARS2, HGF, HOMER2, HOXA1, HOXA2, HOXB1, HSD17B4, IGF1, ILDR1, KARS, KCNE1, KCNJ10, KCNQ1,
 KCNQ4, KIT, KITLG, KMT2D, LARS2, LHFPL5, LHX3, LMX1A, LOXHD1, LRP2, LRTOMT, MAN2B1, MANBA,
 MARVELD2, MASP1, MET, MGP, MIR96, MITF, MPZL2, MSRB3, MT-ATP8, MT-CO1, MT-CO2, MT-CYB, MT-ND1,
 MT-ND4, MT-ND6, MT-RNR1, MT-TC, MT-TE, MT-TF, MT-TH, MT-TK, MT-TL1, MT-TS1, MT-TS2, MT-TV, MT-TW,
 MYH14, MYH9, MYO15A, MYO1A, MYO3A, MYO6, MYO7A, NARS2, NDP, NDRG1, NEFL, NF2, NIPBL, NLRP3,
 NOG, OPA1, OSBPL2, OTOA, OTOF, OTOG, OTOGL, P2RX2, PAX1, PAX2, PAX3, PCDH15, PDSS1, PDZD7, PEX1,
 PEX6, PHEX, PJK, PMP22, PNPT1, POLR1C, POLR1D, POU3F4, POU4F3, PRPS1, PRRX1, PTPRQ, RDX, REST,
 RIPOR2, ROR1, S1PR2, SEMA3E, SERAC1, SERPINB6, SIX1, SIX5, SLC12A2, SLC17A8, SLC19A2, SLC26A4, SLC26A5,
 SLC4A11, SLC5A2, SLC5A3, SLITRK6, SMAD4, SMPX, SNAI2, SOBP, SOX10, SOX9, SPATA5, SPTBN4, STRC,
 SYNE4, SYT2, TBC1D24, TBL1X, TCOF1, TECTA, TIMM8A, TJP2, TMC1, TMEM126A, TMEM132E, TMIE, TMPRSS3,
 TNC, TNFRSF11B, TPRN, TRIOBP, TSPEAR, TWNK, USH1C, USH1G, USH2A, WFS1, WHRN

List relevant family history of disease (affected family members and relationship to patient):

Website www.revivegenetics.us
Email info@revivegenetics.us

Address
 9000 SW 152nd St Suite107, Palmetto Bay, FL
 33157

Lab Director
 Uzma Farooq, MD

Phone Number
 786-250-3419



**Revive
Genetics
Diagnostics
Laboratory**

CLIA: 10D2337214

Informed Consent

Consent to Testing and Use of Results

I confirm that the submitted specimen belongs to the identified patient and has been voluntarily provided for analysis by **Revive Genetics**. I authorize the laboratory and my healthcare provider to analyze my clinical and genetic information, perform the ordered Hearing Impairment genetic testing, release test results to the ordering healthcare provider, and disclose information necessary for billing and reimbursement to my insurance carrier, when applicable.

Consent for Hearing Impairment Genetic Testing

Your healthcare provider has ordered a **Revive Genetics Hearing Impairment Genetic Testing Panel** to evaluate genes associated with inherited hearing disorders. This test analyzes genes involved in auditory function, cochlear development, inner ear structure, sensory hair cell function, auditory signal transmission, and other biological pathways essential for normal hearing and balance. Depending on your personal and family history, this evaluation may help identify inherited genetic variants associated with nonsyndromic and syndromic hearing loss, congenital deafness, progressive hearing impairment, auditory neuropathy, vestibular disorders, and other inherited auditory conditions.

Genetic testing may identify disease-causing (pathogenic) variants, likely pathogenic variants, variants of uncertain significance (VUS), or no clinically significant findings. A positive result may help confirm a diagnosis, guide medical management, assist with treatment decisions, identify at-risk family members, support early intervention and hearing rehabilitation strategies, determine eligibility for emerging targeted therapies or clinical trials, and provide information regarding inherited hearing loss risk. However, a positive result does not predict disease severity, age of onset, rate of hearing loss progression, or response to treatment. Likewise, a negative result does not eliminate the possibility of an inherited hearing disorder.

Please be aware of the following limitations:

This test analyzes only the genes included in the selected panel.

Some genetic changes, including deep intronic variants, balanced chromosomal rearrangements, epigenetic alterations, repeat expansions (unless specifically included), low-level mosaicism, structural variants, mitochondrial variants (unless specifically included), or variants in genes not included on the panel, may not be detected.

Not all inherited hearing disorders have known genetic causes, and some disease-causing variants may not be detectable with current technology.

Variants of Uncertain Significance (VUS) may be identified. These findings currently have insufficient evidence to determine whether they are disease-causing and should not be used alone for medical decision-making.

Genetic test results should always be interpreted together with your personal medical history, family history, physical examination, audiologic evaluation, hearing assessments, vestibular testing (when indicated), imaging studies, laboratory findings, and other clinical information.

Additional testing, including comprehensive audiologic evaluations, vestibular function testing, imaging studies, deletion/duplication analysis, mitochondrial testing, or broader genomic testing such as whole exome or whole genome sequencing, may be recommended based on your clinical presentation and family history.

Genetic counseling is strongly recommended before and/or after testing to discuss the potential benefits, limitations, implications for medical management, and the potential impact of results on biological relatives.

By signing this consent form, you acknowledge that:

You understand the purpose, benefits, and limitations of Hearing Impairment genetic testing.

You understand that testing may identify inherited genetic variants that have implications for your biological relatives.

You understand that results may identify unexpected or uncertain findings that could require additional medical evaluation or long-term clinical follow-up.

You understand that your results will be reviewed by your ordering healthcare provider and may be discussed with you by your physician, otolaryngologist (ENT specialist), audiologist, geneticist, or a qualified genetics professional.

You voluntarily consent to undergo the requested genetic testing.

Methodology Limitations

Next-generation sequencing (NGS) has inherent technical limitations. Certain types of genetic variants may not be detected by this methodology, including some structural rearrangements, repeat expansions (unless specifically included in the assay), low-level mosaic variants, or variants located in genomic regions with limited sequencing coverage.

Testing is limited to the genes and variant types validated for the ordered assay.

All test results are treated as confidential and are released only to the patient, the ordering healthcare provider, and other individuals authorized by the patient or required by law. No tests other than those authorized will be performed. Patient specimens will be retained for up to **60 days** following completion of testing and then destroyed according to laboratory policy unless otherwise required.

I have read this consent form, or it has been read to me. I have received satisfactory answers to my questions and understand that I may ask additional questions at any time. I voluntarily consent to the requested genetic test.

Consent / Insurance Release

By signing this form, I authorize **Revive Genetics** to submit medical information related to this testing to my designated insurance carrier for reimbursement when necessary. I authorize payment of benefits directly to **Revive Genetics**. I understand that I am responsible for any charges not covered by my insurance, including but not limited to non-covered, non-authorized, or denied services. I authorize a copy of this authorization to be used in place of the original.

Patient or Parent/Guardian Signature: _____ **Date:** _____

Print Name: _____

Optional Consent for Research and Future Contact

To improve genetic testing and advance the understanding of inherited hearing disorders, I voluntarily authorize **Revive Genetics** to use any remaining specimen, de-identified genetic data, and/or clinical information for research, educational, quality improvement, or laboratory development purposes. My identity will not be disclosed in any publication or research activity.

I also authorize **Revive Genetics** to contact me in the future regarding newly discovered disease-associated genes, updated variant classifications, research opportunities, or clinically relevant follow-up related to my testing.

Participation is voluntary and declining this consent will not affect the performance of my ordered test or my medical care.

Patient or Parent/Guardian Signature: _____ **Date:** _____

Print Name: _____

Medical Professional Signature: _____ **Date:** _____

Print Name: _____

Hearing Impairment Panel

Requisition & Consent Form

Website www.revivegenetics.us
Email info@revivegenetics.us

Address
9000 SW 152nd St Suite 107, Palmetto Bay, FL
33157

Lab Director
Uzma Farooq, MD

Phone Number
786-250-3419