

### **Genetics and Genomics Diagnostic Laboratory**

For courier service and/or inquiries, please contact 513-636-4474 • Fax: 513-636-4373 www.cincinnatichildrens.org/moleculargenetics • Email: LabGeneticCounselors@cchmc.org

#### Mailing Address:

3333 Burnet Avenue, Room R1042 Cincinnati, OH 45229

## **HEARING LOSS TESTING REQUISITION**

All Information Must Be Completed Before Sample Can Be Processed

| PATIENT INFORMATION   | ETHNIC/RACIAL BACKGROUND (Choose All)   |
|---|---|
| Patient Name:,,,,,  | <ul> <li>□ European American (White)</li> <li>□ African-American (Black)</li> <li>□ Native American or Alaskan</li> <li>□ Pacific Islander</li> <li>□ Ashkenazi Jewish ancestry</li> <li>□ Latino-Hispanic</li> </ul> |
| Home Phone:   | (specify country/region of origin)  ☐ Other (specify country/region of origin)  |
| BILLING INFORMATION (Choole REFERRING INSTITUTION  Institution:   | COMMERCIAL INSURANCE* Insurance can only be billed if requested at the time of service.  Policy Holder Name:  |
| * PLEASE NOTE:  • We will not bill Medicaid, Medicaid HMO, or Medicare except for the followi  • If you have questions, please call 1-866-450-4198 for complete details.  | ng: CCHMC Patients, CCHMC Providers, or Designated Regional Counties.   |
| SAMPLE/SPECIMEN INFORMATION   | REFERRING PHYSICIAN   |
| SPECIMEN TYPE: Amniotic fluid Blood Cord blood CVS  2 Cytobrushes (GJB2, GJB6, or mtDNA Panel only)  6 Cytobrushes (Tier 1, EYA1, and SLC26A4 only)  Note: Cytobrush samples are not acceptable for any panel except Tier 1 and mtDNA Panel  Other: | Physician Name (print):   |
| Specimen Date:// Time:  Specimen Amount:  Each test requires 3 mL of whole blood in EDTA tube. Please call before sending alternate tissue samples, and for free cytobrush or saliva collection kits.   | Genetic Counselor/Lab Contact Name:   |
| *Phlebotomist must initial tube of specimen to confirm sample identity  | Referring Physician Signature (REQUIRED)  |

 $\square$  Patient signed completed ABN

Medical Necessity Regulations: At the government's request, the Molecular Genetics Laboratories would like to remind all physicians that when ordering tests that will be paid under federal health care programs, including Medicare and Medicaid programs, that these programs will pay only for those tests the relevant program deems to be (1) included as covered services, (2) reasonable, (3) medically necessary for the treatment and diagnosis of the patient, and (4) not for screening purposes.



☐ Treacher Collins Syndrome and Mandibulofacial Dysostosis Gene Panel (sequencing of 10 genes including *DHODH*, *EDNRA*, *EFTUD2*, *POLR1A*,

POLR1B, POLR1C, POLR1D, SF3B4, TCOF1, TXNL4A)

☐ Reflex to Whole Exome Sequencing\*\*

| Patient Name: Date of Birth: |  |
|------------------------------|--|

#### **TEST(S) REQUESTED**

### AUDIOGRAM MUST BE ATTACHED. ATTACH CT/MRI IF AVAILABLE.

Please complete Required Clinical Information Sheet (page 4) prior to requisition submission.

Please indicate if Audiogram or CT/MRI is unavailable:  $\ \square$  Audiogram unavailable  $\ \square$  CT/MRI unavailable

| Hearing Loss Panels  ☐ Hearing Loss Panel Tier I* (GJB2 sequencing, GJB6 deletion analysis and 8 mitochondrial mutations) ☐ Reflex to deletion/duplication of GJB2  ☐ OtoSeq Hearing Loss Panel** (sequencing of 23 genes including ADGRV1, CDH23, CLRN1, EYA1, FOXI1, GJB2, GJB6, KCNJ10, MYO6, MYO7A, OTOF, PCDH15, POU3F4, SIX1, SIX5, SLC26A4, TMC1, TMIE, TMPRSS3, USH1C, USH1G, USH2A, WHRN) ☐ Reflex to deletion/duplication of entire panel' ☐ Reflex to deletion/duplication of single gene(s)' (specify): ☐ Hearing Loss Panel Tier I* with reflex to OtoSeq Hearing Loss Panel, if indicated ☐ Branchiootorenal Spectrum Disorder (BOR/BOS) Panel (sequencing of EYA1, SIX1, SIX5) | Single Gene Tests  □ CDH23 (USH1D and DFNB12) □ Reflex to deletion/duplication of CDH23 (USH1D and DFNB12) □ EYA1 (branchiootorenal spectrum disorder type 1) □ Reflex to deletion/duplication of EYA1 □ GJB2 (connexin 26) □ Reflex to deletion/duplication of GJB2 □ GJB6 (connexin 30) deletion analysis □ MYO7A (USH1B, DFNB2, DFNA11) □ Reflex to deletion/duplication of MYO7A (USH1B, DFNB2, DFNA11) □ OTOF (AUNB1, DFNB9) □ Reflex to deletion/duplication of OTOF (AUNB1, DFNB9) □ SLC26A4 (Pendred syndrome, DFNB4) |
|---|---|
| Reflex to deletion/duplication of entire panel  | ☐ Reflex to deletion/duplication of <i>SLC26A4</i> (Pendred syndrome, DFNB4)  |
| ☐ Reflex to deletion/duplication of single gene(s) (specify):   | ☐ Targeted (family specific) mutation analysis for gene  If testing was <b>not</b> performed at CCHMC, please include proband's report  |
| □ Branchiootorenal Spectrum Disorder (BOR/BOS) Panel with reflex to OtoSeq reanalysis, if indicated □ Hearing Loss mtDNA Panel (mtDNA 961, 1555, 1494, 3243, 3271, 7445, 7511, 8344) □ Pendred Syndrome Panel (FOXI1, KCNJ10, SLC26A4) □ Reflex to deletion/duplication of entire panel □ Reflex to deletion/duplication of single gene(s) (specify):   | and at least 100ng of proband's DNA to use as a positive control.  Proband's name  Proband's DOB  Proband's mutation  Please call 513-636-4474 to discuss any family-specific mutation analysis with genetic counselor prior to shipment.   |
| □ Pendred Syndrome Panel with reflex to OtoSeq reanalysis, if indicated □ Usher Syndrome Panel (sequencing of ADGRV1, CDH23, CLRN1, MYO7A, PCDH15, USH1C, USH1G, USH2A, WHRN) □ Reflex to deletion/duplication of entire panel' □ Reflex to deletion/duplication of single gene(s)' (specify):  | *Either Hearing Loss Panel Tier 1* or OtoSeq Hearing Loss Panel** is indicated for patients with sensorineural hearing loss of unknown etiology who have had no previous genetic testing. OtoSeq Hearing Loss Panel may also be used as follow-up testing in patients with normal <i>GJB2</i> or Hearing Loss Panel Tier 1 test results.  Please see our website, <a href="www.cincinnatichildrens.org/hearing-loss">www.cincinnatichildrens.org/hearing-loss</a> , for   |
| ☐ Usher Syndrome Panel with reflex to OtoSeq® reanalysis, if indicated  | complete information.  'Deletion/Duplication analysis of <i>WHRN</i> is not available at this time.   |
| Additional Gene Panels  |   |
| □ Stickler Syndrome Gene Panel (sequencing of 13 genes including <i>BMP4</i> ,  COL11A1, COL11A2, COL2A1, COL9A1, COL9A2, COL9A3, GZF1, LOXL3, LRP2,  PLOD3, SOX9, VCAN)  □ Reflex to Whole Exome Sequencing <sup>††</sup>  | **Whole exome sequencing (WES) orders require a signed WES Consent Form and completion of the WES Test Requisition. Also, inclusion of biological parental samples is strongly encouraged to assist with the analysis of WES and to increase test yield. Please visit our website at <a href="https://www.cincinnatichildrens.org/exome">www.cincinnatichildrens.org/exome</a> to obtain the required documents. WES testing will NOT be started until all forms are completed and received by the                            |
| CI TRACTIEL COURS SYNOTOME AND MANDINHOLACIAL DYSOSIOSIS (SENE PANEL  |   |



| Patient Name: | Date of Birth: |
|---------------|----------------|
|               |                |

## TEST(S) REQUESTED, CONTINUED

| CUSTOM GENE SEQUENCING  | DELETION AND DUPLICATION ASSAY  |
|---|---|
| Gene(s) to be sequenced (specify):  | Gene(s) to be analyzed (specify):   |
| Serie(3) to be sequenced (specify).   |   |
| Only genes with clear published functional relationship to rare diseases are accepted.    | Please see list of available genes at: www.cincinnatichildrens.org/deldup             |
| Suspected syndrome/ condition:  | Suspected syndrome/ condition:  |
| Please choose one of the following:   | Please choose one of the following:   |
| ☐ Full gene(s) sequencing   | $\square$ Deletion and duplication analysis of gene(s) specified above                |
| $\hfill\square$ Full gene(s) sequencing with reflex to deletion and duplication analysis, | $\square$ Deletion and duplication analysis of gene(s) specified above with reflex to |
| if indicated (please see list of genes available for del/dup at                           | sequencing, if indicated  |
| www.cincinnatichildrens.org/deldup)   | ☐ Analysis of gene(s) specified above from previously analyzed deletion               |
| ☐ Familial mutation analysis  | and duplication   |
| Proband's name:   | ☐ Familial deletion analysis  |
| Proband's DOB:  | Proband's name:   |
| Proband's mutation:   | Proband's DOB:  |
| Patient's relation to proband:  | Proband's mutation:   |
| If testing was <u>not</u> performed at CCHMC, please include proband's report             | Patient's relation to proband:  |
| and at least 100ng of proband's DNA to use as a positive control.                         | If testing was <b>not</b> performed at CCHMC, please include proband's report         |
|   | and at least 100ng of proband's DNA to use as a positive control.                     |



| Patient Name: | Date of Birth: |
|---------------|----------------|
|               |                |

# **HEARING LOSS TESTING PROGRAM**

### **REQUIRED CLINICAL INFORMATION**

| Test indication:                                       | Syndromic Associations:  |
|--|--|
| ☐ Diagnosis in symptomatic patient                     | BOR/BOS:   |
| ☐ Family study (please attach proband's report)        | □ None   |
| ☐ Prenatal testing (by previous arrangement only)      | ☐ Ear tags/pits  |
| ☐ Carrier testing                                      | ☐ Ear abnormalities  |
| Audiologic History: Audiogram (MUST BE ATTACHED)       | ☐ Branchial clefts/cysts   |
| Congenital Hearing Loss:                               | ☐ Renal abnormalities  |
| Yes  | Pendred Syndrome:  |
| □ No   | □ None   |
| f <b>NOT</b> congenital, age at onset of hearing loss: | ☐ Abnormal perchlorate test (>15%)                                   |
|  | ☐ Goiter   |
| Type of Hearing Loss:                                  | ☐ Enlarged vestibular aqueduct (EVA)                                 |
| ☐ Sensorineural  | $\square$ Cochlear hypoplasia (Mondini malformation/dysplasia)       |
| ☐ Conductive   | Usher Syndrome:  |
| ☐ Mixed  | □ None   |
| ☐ Auditory neuropathy                                  | ☐ Retinitis pigmentosa   |
| Progression:   | If yes, age at diagnosis:  |
| ☐ Stable   | Other syndromic features or medical problems:                        |
| ☐ Progressive  |  |
| ☐ Fluctuating  |  |
| □ Unknown  |  |
| /estibular Problems:                                   | Aminoglycoside exposure:   |
| □None  | ☐ Yes  |
| □Unknown   | □ No   |
| □ Delayed walking                                      |  |
| ☐ Dizziness/vertigo                                    | Previous Genetic Testing:  |
| ☐ Balance abnormalities                                | ☐ Yes  |
| Radiologic Evaluation: (PLEASE ATTACH)                 | □No  |
| CT scan/MRI of temporal bones?                         | If Yes; specify gene and results including variants:                 |
| ☐ Yes  |  |
| □ No   | Family History:  |
| ☐ Ordered  | Relative(s) with hearing loss?                                       |
|  | □Yes   |
| f Yes, Dilated vestibular aqueducts/EVA?               | □No  |
| ⊒ Yes  | If yes, please specify relationship to patient:                      |
| No   | Parental consanguinity?  |
| Mondini malformation/inner ear dysplasia?              | ☐ Yes  |
| ⊒ Yes  | □No  |
| □No  |  |
|  | All information must be completed before testing will be undertaken. |