

## PEDIATRIC/ADULT GENETIC TEST REQUISITION

All Information Must Be Completed Before Sample Can Be Processed

### PATIENT INFORMATION

Patient Name: \_\_\_\_\_  
Last First MI  
Address: \_\_\_\_\_  
Home Phone: \_\_\_\_\_  
MR# \_\_\_\_\_ Date of Birth \_\_\_\_/\_\_\_\_/\_\_\_\_  
Gender:  Male  Female

### SPECIMEN INFORMATION

**SPECIMEN TYPE:**  Peripheral blood  Skin biopsy (site): \_\_\_\_\_  
 Bone Marrow  Saliva  Cytobrushes  Other: \_\_\_\_\_  
**\*See Page 3 for a list of acceptable specimen types for each test**  
Specimen Date: \_\_\_\_/\_\_\_\_/\_\_\_\_ Time: \_\_\_\_\_  
Specimen Amount: \_\_\_\_\_  
DRAWN BY: \_\_\_\_\_

### INDICATIONS/DIAGNOSIS/ICD-10 CODE

- |  |   |
|--|---|
| <input type="checkbox"/> ADD/ADHD                                | <input type="checkbox"/> Failure to thrive              |
| <input type="checkbox"/> Acute myelogenous leukemia (AML)        | <input type="checkbox"/> Hydrocephalus                  |
| <input type="checkbox"/> Amenorrhea: 1' or 2'                    | <input type="checkbox"/> Hyper/Hypopigmentation         |
| <input type="checkbox"/> Aplastic Anemia                         | <input type="checkbox"/> Hypotonia                      |
| <input type="checkbox"/> Autism Spectrum Disorder                | <input type="checkbox"/> Immune deficiency              |
| <input type="checkbox"/> Broad thumbs and/or halluces            | <input type="checkbox"/> Intellectual disability        |
| <input type="checkbox"/> Congenital heart anomaly                | <input type="checkbox"/> Language disorder              |
| <input type="checkbox"/> Developmental Delay                     | <input type="checkbox"/> Limb malformation              |
| <input type="checkbox"/> Dysmorphic features                     | <input type="checkbox"/> MRI, abnormal                  |
| <input type="checkbox"/> Encephalopathy                          | <input type="checkbox"/> Macrocephaly                   |
| <input type="checkbox"/> Eye anomaly                             | <input type="checkbox"/> Microcephaly                   |
| <input type="checkbox"/> Erythematous "butterfly" lesion on face | <input type="checkbox"/> Myelodysplastic syndrome (MDS) |

- PDD-NOS  
 Pancytopenia  
 Seizures, convulsions  
 Short stature  
 Other: \_\_\_\_\_

#### Newborn Indications:

- Abnormal NIPT/prenatal screen  
 Suspected trisomy 21  
 Suspected Turner's syndrome  
 Ambiguous genitalia  
 Other: \_\_\_\_\_

#### Family History

- Family history of genetic condition: \_\_\_\_\_  
 Consanguinity (describe relationship): \_\_\_\_\_  
 Known Chromosome Abnormality: \_\_\_\_\_  
 Other: \_\_\_\_\_

### BILLING INFORMATION (Choose ONE payment method)

#### REFERRING INSTITUTION

Institution: \_\_\_\_\_  
Address: \_\_\_\_\_  
City/State/Zip: \_\_\_\_\_  
Accounts Payable Contact Name: \_\_\_\_\_  
Phone: \_\_\_\_\_  
Fax: \_\_\_\_\_  
Email: \_\_\_\_\_

#### COMMERCIAL INSURANCE\*

Insurance can only be billed if requested at the time of service.

Policy Holder Name: \_\_\_\_\_  
Gender: \_\_\_\_\_ Date of Birth \_\_\_\_/\_\_\_\_/\_\_\_\_  
Authorization Number: \_\_\_\_\_  
Insurance ID Number: \_\_\_\_\_  
Insurance Name: \_\_\_\_\_  
Insurance Address: \_\_\_\_\_  
City/State/Zip: \_\_\_\_\_  
Insurance Phone Number: \_\_\_\_\_

#### \* PLEASE NOTE:

- We will not bill Medicaid, Medicaid HMO, or Medicare except for the following: CCHMC Patients, CCHMC Providers, or Designated Regional Counties.
- If you have questions, please call 1-866-450-4198 for complete details.

### PROVIDER INFORMATION

Provider Name (print): \_\_\_\_\_  
Address: \_\_\_\_\_  
Phone: ( \_\_\_\_\_ ) \_\_\_\_\_ Fax: ( \_\_\_\_\_ ) \_\_\_\_\_  
Email: \_\_\_\_\_  
Genetic Counselor/Lab Contact Name: \_\_\_\_\_  
Phone: ( \_\_\_\_\_ ) \_\_\_\_\_ Fax: ( \_\_\_\_\_ ) \_\_\_\_\_  
Email: \_\_\_\_\_  
Date: \_\_\_\_/\_\_\_\_/\_\_\_\_

#### Referring Physician Signature (REQUIRED)

**Contact information for results/questions (if different than ordering provider):**  
Name and Title: \_\_\_\_\_  
Phone: ( \_\_\_\_\_ ) \_\_\_\_\_ Fax: ( \_\_\_\_\_ ) \_\_\_\_\_  
Email: \_\_\_\_\_

### ETHNIC/RACIAL BACKGROUND (Choose All)

- |  |  |
|--|--|
| <input type="checkbox"/> European American (White)                                   | <input type="checkbox"/> African-American (Black)  |
| <input type="checkbox"/> Native American or Alaskan                                  | <input type="checkbox"/> Asian-American            |
| <input type="checkbox"/> Pacific Islander  | <input type="checkbox"/> Ashkenazi Jewish ancestry |
| <input type="checkbox"/> Latino-Hispanic _____<br>(specify country/region of origin) |  |
| <input type="checkbox"/> Other _____<br>(specify country/region of origin)           |  |

#### 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.

**TEST(S) REQUESTED**

**Cytogenetic Testing**

**Chromosome Analysis**

- Routine chromosome analysis\*
- High resolution chromosome analysis\*
  - Chromosome mosaicism study\*
  - Reflex to SNP Microarray if chromosome results are normal\*

\*For chromosome analysis: reflex STAT prelim results on infants <1 month. Reflex to mosaicism study when sex chromosome/mosaic aneuploidy abnormality suspected by laboratory based on indications provided.

\*Additional charge for reflex testing. If SNP Microarray is denied by insurance, Chromosome Analysis will be performed as the first test in the algorithm.

**SNP Microarray**

- SNP Microarray - Constitutional

**FISH (Fluorescent In Situ Hybridization)**

- 22q11.2 del (VCFS) (metaphase FISH)
- SRY (Xp11.1q11.1/Yp11.2) (metaphase FISH)
- X/Y centromeres (Xp11.1q11.1/Yp11.1q11.1) (interphase FISH)
- Other FISH (please call lab): \_\_\_\_\_

**Other Testing**

- Special study: \_\_\_\_\_
- Cell Culture, storage & freezing
- Other: \_\_\_\_\_

**Neurodevelopmental Reflex Genetic Test\*\***

Tests will be run sequentially based on your selection below:

- Patient is macrocephalic: SNP Microarray → Fragile X → PTEN
- Male patient with normal or small head circumference:
  - SNP Microarray → Fragile X
- Female patient with normal or small head circumference:
  - SNP Microarray → Fragile X → MECP2

\*\*See page 3 for additional information

\*\*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 [www.cincinnatichildrens.org/exome](http://www.cincinnatichildrens.org/exome) to obtain the required documents. WES testing will NOT be started until all forms are completed and received by the lab.

**Molecular Genetic Testing**

- Fragile X DNA testing
- MECP2* sequence analysis (Rett syndrome)
- MECP2* deletion/duplication analysis by MLPA
- Prader-Willi/Angelman - by methylation-sensitive MLPA
- PTEN* Autism Spectrum Disorder sequencing
- Rubinstein-Taybi and Related Syndromes Gene Panel (*CREBBP, EP300, HNRNPH1, HNRNPH2, SIN3A, SIN3B, SRCAP* with *CREBBP* and *EP300* deletion/duplication analysis by MLPA)
  - REFLEX to Whole Exome Sequencing\*\*** (See additional details below)
- Spinal Muscular Atrophy - *SMN1/SMN2* Copy Number Analysis by MLPA
- Other: \_\_\_\_\_

**Fanconi Anemia Testing**

- Fanconi Anemia (FA) Chromosome Breakage Study
- Fanconi Anemia Gene Sequencing Panel (*BRCA1, BRCA2, BRIP1, ERCC4, FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, MAD2L2, PALB2, RAD51, RAD51C, RFWWD3, SLX4, UBE2T, XRCC2*)

If **both** FA Breakage Study and FA Gene Seq Panel are ordered, testing will be run sequentially (breakage study then molecular sequencing if breakage study is positive) unless concurrent testing is selected here:

- Concurrent FA testing requested**

**Single Gene Sequencing**

- FANCA* full gene sequencing     *FANCG* full gene sequencing
- FANCC* full gene sequencing

**Chromosome Breakage Disorders Testing**

- Bloom Syndrome - Sister Chromatid Exchange (SCE) analysis
- Chromosome Breakage Disorders Gene Sequencing Panel (*ATM, BLM, BRCA1, BRCA2, BRIP1, ERCC4, FANCA, FANCB, FANCC, FANCD2, FANCE, FANCF, FANCG, FANCI, FANCL, LIG4, MAD2L2, MYSM1, NBN, NHEJ1, NSMCE3, PALB2, RAD51, RAD51C, RFWWD3, SLX4, UBE2T, XRCC2*)

**CUSTOM GENE SEQUENCING**

Gene(s) to be analyzed (specify): \_\_\_\_\_

*Only genes with clear published functional relationship to rare diseases are accepted.*

Suspected syndrome/ condition: \_\_\_\_\_

Please choose one of the following:

- Full gene(s) sequencing
  - Full gene(s) sequencing with reflex to deletion and duplication analysis, if indicated (please see list of genes available for del/dup at [www.cincinnatichildrens.org/deldup](http://www.cincinnatichildrens.org/deldup))
  - Familial mutation analysis
    - Proband's name: \_\_\_\_\_
    - Proband's DOB: \_\_\_\_\_
    - Proband's mutation: \_\_\_\_\_
    - Patient's relation to proband: \_\_\_\_\_
- If testing was **not** performed at CCHMC, please include proband's report and at least 100ng of proband's DNA to use as a positive control.

**DELETION AND DUPLICATION ASSAY**

Gene(s) to be analyzed (specify): \_\_\_\_\_

Please see list of available genes at: [www.cincinnatichildrens.org/deldup](http://www.cincinnatichildrens.org/deldup)

Suspected syndrome/ condition: \_\_\_\_\_

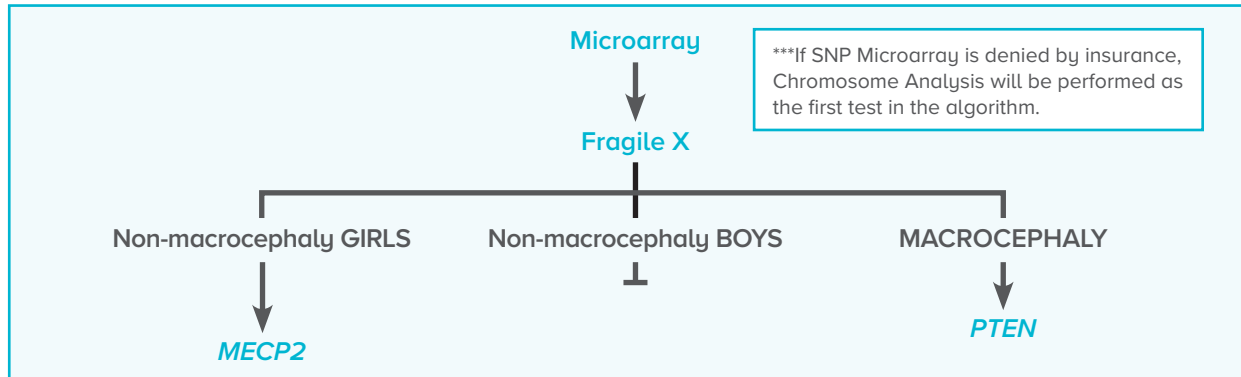
Please choose one of the following:

- Deletion and duplication analysis of gene(s) specified above
  - Deletion and duplication analysis of gene(s) specified above with reflex to sequencing, if indicated
  - Analysis of gene(s) specified above from previously analyzed deletion and duplication
  - Familial deletion analysis
    - Proband's name: \_\_\_\_\_
    - Proband's DOB: \_\_\_\_\_
    - Proband's mutation: \_\_\_\_\_
    - Patient's relation to proband: \_\_\_\_\_
- If testing was **not** performed at CCHMC, please include proband's report and at least 100ng of proband's DNA to use as a positive control.

## ADDITIONAL INFORMATION

### NEURODEVELOPMENTAL REFLEX GENETIC TESTING ALGORITHM

Tests will be performed sequentially based on the path that your patient follows in the Neurodevelopmental Reflex Test algorithm. Charges will apply to the tests completed in the patient's defined Neurodevelopmental Reflex Test algorithm. Testing will only proceed to the next step if the previous test result is uninformative.



### SPECIMEN REQUIREMENTS

**Routine & High Resolution Chromosome Analysis:**

3–5 mL blood (NaHep)

**SNP Microarray:**

3 mL blood (NaHep) **and** 3 mL blood (EDTA)

**FISH Tests:** 1–3 mL blood (NaHep)

**Neurodevelopmental Reflex Genetic Testing:**

3 mL blood (NaHep) **and** 3–5 mL blood (EDTA)

**Fragile X DNA Testing:**

3 mL blood (EDTA)

**MECP2 del/dup by MLPA, Prader-Willi/Angelman - by MLPA, Spinal Muscular Atrophy - SMN1/SMN2 Copy Number Analysis & Deletion/Duplication Assay:**

3 mL blood (EDTA)

**Fanconi Anemia Chromosome Breakage Study:**

5–10 mL blood (NaHep), 5–10 mL bone marrow (NaHep), or Skin biopsy (3-4 mm tissue in sterile transport media)

**FANCA, FANCC, FANCG, MECP2, PTEN & Custom Gene Sequencing:**

3 mL blood (EDTA), saliva collection kit\*, or 6 cytobrushes

**Bloom syndrome – Sister Chromatid Exchange(SCE) analysis:**

3–5 mL blood (NaHep)

**Chromosome Breakage Disorders Gene Seq, Fanconi Anemia Gene Seq & Rubinstein-Taybi and Related Syndromes Gene Panels:**

3 mL blood (EDTA) or saliva collection kit\*

*\*Call the office at 513-636-4474 to obtain saliva collection kits*