



Submitter/billing information
DO NOT LEAVE BLANK

(PLEASE PRINT USING CAPITALS- FIELDS IN RED ARE REQUIRED)

(1) Patient Last Name		First Name		Middle Name	
(2) Name Change- Former Last Name					
(3) Patient Address					
(4) City		State		Zip	
				County of Residence	
(5) Date of Birth		(6) Age		(7) Sex <input type="checkbox"/> Male <input type="checkbox"/> Female	
(8) Ethnicity <input type="checkbox"/> Hispanic/Latino <input type="checkbox"/> Non-Hispanic/Latin		(9) <input type="checkbox"/> Amer Indian <input type="checkbox"/> Asian <input type="checkbox"/> Other _____		<input type="checkbox"/> Black/African Amer <input type="checkbox"/> Pacific Islander <input type="checkbox"/> White	
(10) Chart #/ Patient ID Number		(11) Submitter Specimen ID Number		(15) NPI # _____	

(13) ADDITIONAL REPORT COPIES NEEDED?
Please check this box AND
Enter the clinician's name and address on the back of this form

(16) Attached copies of front and back of insurance card(s)?

(18) MEDICAID# _____
 PRIVATE INSURANCE# _____ Bill to Submitter MEDICARE# _____

(20) Please write the letter corresponding to the appropriate ICD-10 Code to the left of the test name below (where applicable)

(A) ICD-10 Code _____ (B) ICD-10 Code _____ (C) ICD-10 Code _____ (D) ICD-10 Code _____ (E) ICD-10 Code _____

(21) Date of collection _____ (22) Time of collection _____

Specimen Source Amniotic Fluid Blood Chorionic Villus Sample Paraffin Section (tissue type _____)
 Cord Blood Buccal Saliva

Products of Conception (tissue type _____) Tissue Biopsy (tissue type _____)

Gestational Age: _____
Ultrasound Age: _____
Estimated Date of Delivery: ___ / ___ / ___

Reason for Referral (please provide in addition to ICD-10 code above):

CHROMOSOME ANALYSIS	MOLECULAR ANALYSIS	FISH ANALYSIS
<input type="checkbox"/> 801 Chromosome Analysis, Blood <input type="checkbox"/> 803 Chromosome Analysis, Blood, Abridged Examination for Familial Chromosome Rearrangements <input type="checkbox"/> 850 Chromosome Analysis, Amniotic Fluid <input type="checkbox"/> 852 Chromosome Analysis, Amniotic Fluid, Abridged (must also order 890) <input type="checkbox"/> 855 Chromosome Analysis, Chorionic Villus Sample <input type="checkbox"/> 857 Chromosome Analysis, CVS, Abridged (must also order 890) <input type="checkbox"/> 831 Chromosome Analysis, Products of Conception/ Tissue Biopsy <input type="checkbox"/> 860 Tissue culture and shipment for additional testing	<input type="checkbox"/> 828 Molecular Analysis, Fragile-X, Genetic Diagnosis <input type="checkbox"/> 889 Methylation-Specific PCR, SNRPN gene, 15q11.2 <input type="checkbox"/> 890 Chromosomal Microarray Analysis (CMA), SNP-based <input type="checkbox"/> Tiered (Hold array pending chromosome analysis) <input type="checkbox"/> 890FAM Targeted Microarray Analysis (Family Studies) <input type="checkbox"/> 895M47 PlainSEEK Panel <input type="checkbox"/> 895M64 Exome Sequencing- Proband Analysis Only <input type="checkbox"/> 895M64a Exome Sequencing- Proband with Parental TVAR <input type="checkbox"/> 895M65 Exome Sequencing- Duo Analysis <input type="checkbox"/> 895M66 Exome Sequencing- Trio Analysis <input type="checkbox"/> 895M68 Sudden Unexplained Death in the Young (SUDY) Exome <input type="checkbox"/> 895M50 Targeted Variant Analysis via Sanger DNA Sequencing (variant nomenclature: _____)	<input type="checkbox"/> 873 Prenatal Aneuploidy Panel, Amniotic Fluid <input type="checkbox"/> 875 Stillbirth Aneuploidy Panel, Paraffin Embedded <input type="checkbox"/> 871F28 DiGeorge /Velo-cardio-facial / Schpritzen /Conotruncal anomaly Syndrome, Deletion 22q11.2, TUPLE1 <input type="checkbox"/> 871F34 SRY (Sex determining Region of Y), Yp11.3 <input type="checkbox"/> 870F52 X and Y sex chromosomes

A sample processing fee is charged for blood samples that have only FISH analysis

WSLH contributes submitted clinical information and test results for molecular cytogenetic tests to a HIPAA-compliant, de-identified public database a part of the National Institutes of Health's effort to improve diagnostic testing and our understanding of the relationships between genetic changes and clinical symptoms. For information about the ClinVar database, visit their website at <http://www.ncbi.nlm.nih.gov/clinvar/>. Confidentiality of each sample is maintained. Patients may request to withdraw consent for the storage of their sample and/or use of the data by: 1) checking the box below, 2) calling the laboratory at (608) 262-0402 and asking to speak with a genetic counselor, or by 3) visiting our website at www.slh.wisc.edu/cytogenetics.

Refusal for inclusion in these efforts may be indicated by checking this box. (If the box is not checked, the data will be anonymized and used.)