

University of Wisconsin Collaborative Genomics Core 465 Henry Mall Madison, WI 53706-1578

Genetic Diagnosis (03/2019)	CGC# 131
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Madison, WI 53706-1578 Collaborative Genomics Core UNIVERSITY OF WISCONSIN-MADISON www.slh.wisc.edu/cytogenetics						
(PLEASE PRINT	USING CAPITALS- FIEL	DS IN RED ARE REQUIRED)				
1) Patient Last Name	First Name	Middle Name				
2) Name Change- Former Last Name	e					
3) Patient Address						
4) City State Zip		County of Residence (13)		(13) Additional Report Copies Needed?		
5) Date of Birth	(6) Age	(7) Sex		Please check this box		
8) Ethnicity		Black/African Amer □White Pacific Islander		(14) Ordering Provider		
10) Chart #/ Patient ID Number	(11) Submitter Specimen ID Nun	iber	(15) NPI #	(15) NPI #		
16) Attached copies of front and back of in	nsurance card(s)?					
18)						
□ PRIVATE INSURANCE# □ Bill to						
•		ode to the left of the test name below (where ap	•	(D) 10D 10 0 1		
(A) ICD-10 Code (B) 21) Date of collection	3) ICD-10 Code (C	(D) ICD-10 Code (D) ICD-10 C	ode	(E) ICD-10 Code	·	
Specimen Source	tic Fluid □ Blood □ Ch	orionic Villus Sample	ssue type)	Gestational Age:	
□ Products of Conception (tissue type) □ Tissue Biopsy (tissue type))		Ultrasound Age:		
Reason for Referral (please provide in a					Estimated Date of Delivery://	
CHROMOSOME ANAI	LYSIS	MOLECULAR ANALYSIS				
 □ 801 Chromosome Analysis, Blood □ 803 Chromosome Analysis, Blood, Abridged Examination for Familial Chromosome Rearrangements □ 850 Chromosome Analysis, Amniotic Fluid □ 852 Chromosome Analysis, Amniotic Fluid, Abridged (must also order 890) □ 855 Chromosome Analysis, Chorionic Villus Sample □ 857 Chromosome Analysis, CVS, Abridged (must also order 890) □ 831 Chromosome Analysis, Products of Conception/ Tissue Biopsy □ 860 Tissue culture and shipment for additional testing 		 □ 828Molecular Analysis, Fragile-X, Gen Diagnosis □ 889Methylation-Specific PCR, SNRPN 15q11.2 □ 890Chromosomal Microarray Analysis (CMA), SNP-based □ Tiered (Hold array pending chromosom analysis) □ Concurrent (with chromosome analysis (Family Studies) □ 895FAM Targeted Microarray Analysis (Family Studies) □ 895M64 Exome Sequencing- Proband Analysis Only □ 895M65 Exome Sequencing- Duo Anal 895M66 Exome Sequencing- Trio Anal 895M41 Cardiomyopathy/Arrhythmia □ 895M50 Targeted Variant Analysis via DNA Sequencing (variant nomenclature: 		[HHPCR] □ 842C92 Factor II Genotyping [PTPCR] □ 842C90 Factor V Genotyping [FVPCR] FISH ANALYSIS		
				Fluid 875 Stillbirth Aneuploidy Panel, Paraffin Embedded 871 F28 DiGeorge /Velo-cardio-facial / Schpritzen /Conotruncal anomaly Syndrome, Deletion 22q11.2, TUPLE1 NGS 871 F34 SRY (Sex determining Region of Y), Yp11.3 870 F52 X and Y sex chromosomes		

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.

 \square 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.)