

2016 PERIPHERAL BLOOD REFERRAL FORM FOR CHROMOSOME ANALYSIS 2016

Medical Genetics – Wake Forest University School of Medicine – Winston Salem NC

www.wfubmc.edu/medical genetics

Phone: 336-716-4321 Fax: 336-716-2554

Collection Date: _____ **Time:** _____ am/pm **WFU LAB #:** _____

Name: _____ / _____ / _____ / _____
(Please print) Last First Middle Maiden
Address: _____ / _____ / _____ / _____ **Daytime Phone:** (____) _____
Mailing Address City State Zip
Birth Date: _____ **SS#:** _____ **Sex:** male / female
Hospital: _____ **Hospital/Unit #:** _____ **Patient's Mother's first name:** _____
Type of Specimen: Blood PUBS Blood (POC) Cancer

Collection Technique: 5 – 10 mls blood collected in a green stoppered sodium heparin tube. **Keep at room temperature**

Physician/Provider Order	Statement of Financial Responsibility
Physician: Last, First / Phone / Beeper	I authorize any holder of medical or other information about me to release to my healthcare provider, third party processor, the Centers for Medicare and Medicaid Services or its intermediaries or carriers any information needed for this health care encounter or related claim. I permit a copy of this authorization to be used in place of the original, and request payment of authorized insurance benefits be made on my behalf to the WFU Physicians. I understand I am responsible for payment of these charges. I am also responsible for payment if my insurance carrier decides this is a non-covered service or requires prior authorization, which I did not obtain. Patient Signature: _____ Date: _____
1. _____	
X Physician Signature Required	
2. _____	
3. _____	

Billing Information

Bill: Forsyth Hospital/Novant Moses Cone Hospt clinical lab
 Women's Hospt clinical lab Wesley Long Hospital Other : _____
 Medicare # _____ Medicaid #: _____ Carolina Access #: _____
 Insurance: _____ Employer: _____ Policy #: _____

(Enclose copy of both sides of insurance card)

SIGNS/SYMPTOMS/INDICATION (ICD-10 CODES) FOR CHROMOSOME STUDY

Indicate all that apply. Codes do not represent entire listing of ICD-10 codes available. Please consult current ICD-10 code book for complete listing.

<input type="checkbox"/> Other chromosome conditions (Q99.9)	<input type="checkbox"/> Anomalies face/neck (Q18.8)	<input type="checkbox"/> Lack of nml physiol dev (R62.50)
<input type="checkbox"/> Other sex chromosome anomaly (Q97.8)	<input type="checkbox"/> Eye anomaly (Q15.9) NOS	<input type="checkbox"/> Brain anomalies (Q04.8) NOS
<input type="checkbox"/> Ambiguous genitalia (Q56.4)	<input type="checkbox"/> Cleft lip/palate (Q37.9) NOS	<input type="checkbox"/> Failure to thrive (R62.51)
<input type="checkbox"/> Multiple congenital anomalies (Q89.7)	<input type="checkbox"/> Hypotonia, cong. (P94.2)	<input type="checkbox"/> Mixed develop disorder (F82) NOS
<input type="checkbox"/> Unspec anom congen anomaly (Q89.9)	<input type="checkbox"/> Obesity (E66.9)	<input type="checkbox"/> Other spec cond. perinatal (P96.89)
<input type="checkbox"/> Other Specified anomalies (Q87.89) congenital	<input type="checkbox"/> Delay in sexual development (E30.0)	<input type="checkbox"/> Fam hx other cong anomal (Z82.79)
<input type="checkbox"/> Developmental delay (F81.9) NOS	<input type="checkbox"/> Abnormal ears (Q17.8)	<input type="checkbox"/> Family hx, mental retard (Z81.0)
<input type="checkbox"/> Delayed learning (F81.81) NOS	<input type="checkbox"/> Short Stature (R62.52)	<input type="checkbox"/> Fam hx other conditions (Z84.89)
<input type="checkbox"/> MR Moderate (F71) <input type="checkbox"/> Severe (F72) <input type="checkbox"/> Profound (F73)		<input type="checkbox"/> Congen heart anomaly (Q24.9)
<input type="checkbox"/> Infertility -> <input type="checkbox"/> female (N97.8) NOS / <input type="checkbox"/> male (N46.9) NOS		<input type="checkbox"/> Language Delay (F80.1)
<input type="checkbox"/> Multiple Miscarriages (habitual abortion) {G ____ P ____ A ____} [SAB ____ TAB ____]		
<input type="checkbox"/> Additional Clinical Information/ICD-10 codes:		

Test Requested Note: When ordering tests for which Medicare reimbursement will be sought, it is recommended that the Provider consult any Local Medical Review Policies (LMRP) or National Coverage Decisions (NCD) that may be applicable to the test(s) being ordered. Based on guidance issued in either of these policies it may be necessary to obtain an Advanced Beneficiary Notice (ABN) from the Medicare Patient. For Medicaid and other carriers a signed **Statement of Financial Responsibility** from the patient may be necessary. (See Statement of Financial Responsibility at top of form.)

SUSPECTED DIAGNOSIS OR CHROMOSOME ABNORMALITY (not for billing purposes)

<input type="checkbox"/> Trisomy 21 (Down Syndrome)	<input type="checkbox"/> Habitual Aborter	<input type="checkbox"/> Smith Magenis Synd. (17p-)
<input type="checkbox"/> Trisomy 13	<input type="checkbox"/> Prenatal confirmation	<input type="checkbox"/> Di-George/ VCF Synd. (22q-)
<input type="checkbox"/> Trisomy 18	<input type="checkbox"/> Prader Willi Synd. (15q-)	<input type="checkbox"/> Miller-Dieker Synd. (17p-)
<input type="checkbox"/> Turner Syndrome (45,X)	<input type="checkbox"/> Angelman Synd. (15q-)	<input type="checkbox"/> Klinefelter Syn. (47,XXY)
<input type="checkbox"/> Mental Retardation (for Fragile X Syndrome – purple top also required)		
<input type="checkbox"/> Other genetic/chromosome abnormality: _____		

TEST REQUESTED	FISH Specific Probes
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<input type="checkbox"/> Routine Chromosome / karyotype (7 - 10 days) <small>(88230, 88262, 88280, 88285)</small>	<input type="checkbox"/> Angelman 15q11 <input type="checkbox"/> +13 <input type="checkbox"/> +18 <input type="checkbox"/> +21
<input type="checkbox"/> STAT Chromosomes (48 hrs. - additional charges) <small>(88230 88262,88280, 88285, 88261)</small>	<input type="checkbox"/> Prader-Willi 15q12 <input type="checkbox"/> Sex - X&Y
<input type="checkbox"/> Chromosome / karyotype +Stat FISH (13/18/21/X/Y) <small>(88230, 88262, 88280, 88285,88271x5, 88274x5)</small>	<input type="checkbox"/> DiGeorge/VCF 22q11.2 <input type="checkbox"/> SRY Yp11.3
<input type="checkbox"/> Routine Chromosome / karyotype +FISH select → →	<input type="checkbox"/> Miller-Dieker 17p13 <input type="checkbox"/> 4p-
<input type="checkbox"/> FISH Only - chromosome previously done select → → →	<input type="checkbox"/> STS Xp22.3 <input type="checkbox"/> 5p-
<input type="checkbox"/> Culture Only – Microarray testing	<input type="checkbox"/> Kallmann Xp22.3 <input type="checkbox"/> _____
	<input type="checkbox"/> Smith-Magenis 17p11 <input type="checkbox"/> Subtelomere
	<input type="checkbox"/> Williams 7q11 <input type="checkbox"/> SHOX Xp22.3

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CYTOGENETIC LAB USE ONLY

Name: _____ **Lab #:** _____

Date Received: _____/_____/_____ **Time Received:** _____

Additional Specimen Evaluation: _____

Additional Samples Received: DNA: _____ FRAX FISH other _____

LABORATORY REPORT SUMMARY

Date culture initiated: _____/_____/_____ **Tech:** _____

Sample type: PB FISH FRAX PUBS BM stat

Culture: 48h. 72h. col. 72h.EtBr DBM 24h 96h.

Media: RPMI 1640 less folic acid RPMI 1640

REPORT OF RESULTS / SPECIMEN SUMMARY

Final Preliminary Read Back **Date** _____ **Tech** _____

To: _____

KARYOTYPE: 46,XY 46,XX

INTERPRETATION: normal male normal female

Abnormal: _____

Additional Studies / Results: NOR C-band R-band

FISH: normal male normal female

+13 +18 +21 -X +X +Y

- Williams 7q22
- DiGeorge/VCF 22q11
- Prader-Willi 15q12
- Angelman 15q12
- Smith-Magines 17p11
- Miller-Dieker 17p13
- STS Xp22.3
- chromosome paint # _____

To: _____ **By:** _____ **Date** _____