

Surname:			First name(s):			Date of birth:		Sex: <b>M/F</b>	
Address:			GP Address:			Consultant:			
						Hospital:		Department:	
						Hospital number:		Genetics number: <b>G</b>	
Postcode:			GP Postcode:						
LMP/Gestational age: <i>(for ongoing pregnancies only)</i>		Urgent: <b>YES / NO</b>	NHS / PP	NHS number:		Sample type:		Date & time of collection:	
<b>CHECK BLOOD TUBE TYPE and PLEASE USE CAPITAL LETTERS</b> <b>Incorrect tubes may be discarded and illegible or incomplete request forms will delay processing</b>					Family history / clinical information / reason for test				
<b>CHROMOSOME ANALYSIS</b> <u>4ml blood in LITHIUM HEPARIN</u> (1ml newborns / babies)			<b>DNA / GENE ANALYSIS</b> <u>4-8ml blood in EDTA / K2E / K3E</u> (1-2ml newborns / babies)						
<input type="checkbox"/> Routine chromosomes (karyotype)  <input type="checkbox"/> FISH for .....  <input type="checkbox"/> Other: .....			<input type="checkbox"/> array-CGH * <input type="checkbox"/> store DNA only  State tests required:  <i>Please note that DNA will be stored unless requested otherwise</i>						
<b>* ARRAY-CGH REFFERALS MUST HAVE THE ADDITIONAL "REQUEST FOR ARRAY-CGH ANALYSIS" SECTION OF THIS FORM COMPLETED</b>									
<b>LAB</b>		Date received:	Cyto sample number:	Lithium Heparin:	Sample condition:	Referring doctor: :		Bleep / extension / phone number:	
<b>USE</b>		Initials:	DNA sample number:	EDTA:	Sample condition:				

During the consultation we have discussed the following issues and you have agreed to the uses indicated below

(Circle as applicable)

- |  |          |
|--|----------|
| A) I agree to analysis of the sample for _____   | YES / NO |
| B) I am happy for further diagnostic testing on the stored sample if new tests become available, without being contacted | YES / NO |
| C) I agree that information and results can be shared to help other family members                                       | YES / NO |
| D) I agree to the sample being used anonymously for research   | YES / NO |
| E) I am aware that the tests may reveal unexpected information, including information about a child's biological parents | YES / NO |

SIGNATURE: \_\_\_\_\_ PRINT NAME: \_\_\_\_\_ DATE: \_\_\_\_\_

I have fully explained the nature of the requested test(s) to the patient / parent / legal guardian\*

SIGNATURE: \_\_\_\_\_ PRINT NAME: \_\_\_\_\_ DATE: \_\_\_\_\_

\* PLEASE DELETE AS APPROPRIATE

Samples and completed referral forms should be packaged appropriately and according to UN3373 guidelines where necessary.  
All samples should be sent by first class post, courier, hospital transport or taxi to:

**SW Thames Regional Genetics Laboratory, Specimen Reception, Jenner Wing, Lower Ground Floor,  
St. George's University of London, Cranmer Terrace, London, SW17 0RE**

For further information or advice, please telephone 020 8725 5332  
or email [swtrel@stgeorges.nhs.uk](mailto:swtrel@stgeorges.nhs.uk).

More information on our services can be found at [www.southwestthamesgenetics.nhs.uk](http://www.southwestthamesgenetics.nhs.uk)



This section of the form must be completed fully for array CGH analysis requests in addition to completion of the "Request for Chromosome & DNA Analysis" section. Failure to do so may result in delay or failure to process the sample.

Last name:	First name:	DOB	Sex M / F
Referring clinician:	Referring centre:	GEN NO.	DNA No.
Suspected syndrome: Please specify:			
Relevant family history:			

CLINICAL FEATURES	<i>Please circle / complete where applicable</i>
Developmental delay difficulties/Autism	Mild / Moderate / Severe
Learning difficulties	Mild / Moderate / Severe
Autism / behavioural problems	Mild / Moderate / Severe
Cleft Lip	Y / N
Cleft Palate	Y / N
Heart defects	Y / N If "Y", please specify:
Limb Anomaly	Upper / Lower / Hand / Foot Please give details:
Microcephaly	Y / N
Macrocephaly	Y / N
MRI brain anomaly	Y / N If "Y", please specify:
Overgrowth	Y / N
Growth below the third centile	Y / N
Dysmorphic	Y / N If "Y", please specify:
Other anomaly	Y / N If "Y", please specify:
Any previous genetic tests:	Y / N If "Y", please specify: