

**DEPARTMENT OF CYTOGENETICS & GENOMICS – REFERRAL LETTER**

Department Code: 29

Lab no. \_\_\_\_\_

<b>PATIENT DETAILS</b>			
Name:	<input type="text"/>	ID no.:	<input type="text"/>
Address:	<input type="text"/>	Tel no. home:	<input type="text"/>
	<input type="text"/>	Tel no. mobile:	<input type="text"/>
CING no.:	<input type="text"/>	Clinic File no.:	<input type="text"/>
		Hospital File no.:	<input type="text"/>
Date of Birth:	<input type="text"/>	Nationality:	<input type="text"/>
	Gender:	<input type="checkbox"/> Male <input type="checkbox"/> Female <input type="checkbox"/> Fetus	

Partner's name:	<input type="text"/>	Date of Birth:	<input type="text"/>	ID no.:	<input type="text"/>
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<b>PATIENT STATUS</b>			
Case type:	<input type="checkbox"/> Outpatient	<input type="checkbox"/> Inpatient	
GESY: <input type="checkbox"/>	Government-Non GESY: <input type="checkbox"/>	Hospital Card no: <input type="text"/>	Private-Non GESY: <input type="checkbox"/>

<b>REFERRING PHYSICIAN/SCIENTIST – STATUS</b>			
CING: <input type="checkbox"/>	Government (OKYπY): <input type="checkbox"/>	Private-GESY: <input type="checkbox"/>	GESY no: <input type="text"/>
			Private-Non GESY: <input type="checkbox"/>
Name:	<input type="text"/>		
Address:	<input type="text"/>	Tel no.:	<input type="text"/>
	<input type="text"/>	Fax no.:	<input type="text"/>
Hospital/Clinic:	<input type="text"/>	Email:	<input type="text"/>

*The referring physician undertakes and confirms understanding and compliance in respect of the mutual obligations as these are determined under the GDPR.*

<b>CLINICAL INFORMATION</b>			
Reason for Referral:	<input type="text"/>		
	<input type="text"/>		
	<input type="text"/>		
Sample type:	<input type="text"/>	Collection date: <input type="text"/>	Time: <input type="text"/>
		Biopsy date: <input type="text"/>	
LMP: <input type="text"/>	Gestational age in weeks: <input type="text"/>	Family no.: <input type="text"/>	

Delivered by:	<input type="text"/>	Received by:	<input type="text"/>	Time:	<input type="text"/>	Date: <input type="text"/>
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*The Cyprus Institute of Neurology and Genetics (CING) complies with the General Data Protection Regulation EE 2016/679. For further information on how we protect personal data and on how data subjects can exercise their rights please refer to our privacy policy at <https://www.cing.ac.cy/> or contact the CING Data Protection Officer at 22358600 or through the e-mail: [dpo@cing.ac.cy](mailto:dpo@cing.ac.cy).*

**TYPE OF SERVICE**

<b>CLINICAL CYTOGENETICS – CHROMOSOMAL ANALYSIS of</b>
<input type="radio"/> 1*. Chorionic Villus Sample (CVS) (Transport Medium) <input type="radio"/> 2*. Amniotic Fluid (AF) (20ml) <input type="radio"/> 3*. Fetal Blood (FB) (2ml in Sodium or Lithium Heparin) <input type="radio"/> 4*. Peripheral Blood (PB) (5ml in Sodium Heparin) <input type="radio"/> 5*. Skin Biopsy (SB) (Transport Medium/Normal Sterile Saline) <input type="radio"/> 6*. Examination of Products of Conception (POC) (Transport Medium/Normal Sterile Saline) <input type="radio"/> 7. Tissue Culture only – (TC) To facilitate other tests <input type="radio"/> 8*. Peripheral Blood for couples (PB) (5ml in Sodium Heparin)

<b>Patient Name:</b> _____	<b>Lab no.:</b> _____
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**MOLECULAR CYTOGENETICS**

- 20\*. Identification/Confirmation/Characterisation of chromosomal abnormality(ies) by Fluorescence In Situ Hybridization (FISH) (F) (5ml PB or 2ml FB in Sodium Heparin, AF, CVS)
- 21\*. Diagnosis of the following diseases/syndromes by FISH (F) (5ml PB or 2ml FB in Sodium Heparin, AF, CVS)
 

<input type="checkbox"/> Prader-Willi/Angelman (D15S10)	<input type="checkbox"/> Williams	<input type="checkbox"/> Miller-Dieker	<input type="checkbox"/> Cri du chat
<input type="checkbox"/> Prader-Willi/Angelman (SNPRN)	<input type="checkbox"/> X-Linked Ichthyosis	<input type="checkbox"/> 22q11.2 deletion	<input type="checkbox"/> Kallmann
<input type="checkbox"/> Smith-Magenis	<input type="checkbox"/> Retinoblastoma	<input type="checkbox"/> Wolf-Hirschhorn	<input type="checkbox"/> Other
- 22. Multiprobe detection centromeric/telomeric by FISH (MP) (5ml PB or 2ml FB in Sodium Heparin, AF, CVS)
- 25. Sperm FISH for chromosome aneuploidies (Semen)
- 26. Sperm DNA Fragmentation (SF) (Semen)
- 27. Sperm Oxidative Stress (OS) (Semen) (Private only)

**DNA ANALYSIS**

- 60\*. Prenatal Diagnosis of Fragile X Syndrome (FR) (AF, CVS, 2ml FB in EDTA)
- 61\*. Postnatal Diagnosis of Fragile X Syndrome (FR) (5ml PB in EDTA)
- 62. Investigation of abnormalities/syndromes with MLPA/qRT-PCR (NF) (5ml PB in EDTA)
 

<input type="checkbox"/> MLPA	<input type="checkbox"/> qRT-PCR	<input type="checkbox"/> UPD	<input type="checkbox"/> X-inactivation
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- 63. DNA Isolation (DNA) (5ml PB or 2ml FB in EDTA, AF, CVS)
- 64. Detection of Y chromosomal material (Y) (5ml PB in EDTA)
- 65\*. Screening for Y (AZF) chromosomal microdeletion (PWA) (5ml PB in EDTA)
- 66. Achondroplasia Mutations G1138A and G1138C analysis (AC) (5ml PB or 2ml FB in EDTA, AF, CVS)
- 67. Central Diabetes Insipidus (CDI) (5ml PB in EDTA)
- 68\*. Molecular Diagnosis of Prader Willi/Angelman Syndrome (PWA) (5ml PB in EDTA)
- 69\*. Rapid prenatal diagnosis of 13, 18, 21, X, Y aneuploidies with QF-PCR (QF) (AF, CVS, FB)
- 70\*. Detection of genomic imbalances with high-resolution microarray-CGH (MK) (5ml PB or 2ml FB in EDTA, AF, CVS)
- 71\*. Pre-implantation Genetic Testing with microarray-CGH or NGS (PGT) (Biopsied embryo in 2-3µl PBS)

**CELL LINES**

- 80. Establish Lymphoblastoid Cell line (CL) (10ml PB in Sodium Heparin)
- 81. Establish Fibroblast Cell Line (CL) (Transport Medium)
- 82. Thawing-Freezing-Expansion up to 2T-25 Flasks (CL)
- 83. Maintain Two Vials/Sample in N2/year (CL)

**NEXT GENERATION SEQUENCING – (NGS) – Postnatal referrals: 5ml P. Blood in EDTA from patient, and parents as needed.**

**Please tick the box if diagnostic NGS tests are being requested and fill in the following fields:**  Current NGS referral is for diagnostic purposes only  
 Please state if any other diagnostic tests have been performed for this patient YES / NO  
 If YES state which tests: \_\_\_\_\_  
 If NO state why patient is referred for NGS as a first-tier test: \_\_\_\_\_

- 110\*. Targeted Panel up to 50 genes (TNGS)
  - Rett/Angelman-like panel
- 111\*. Clinical Exome Sequencing (CES) (Single patient analysis) (CNGS)
- 112\*. Clinical Exome Sequencing (Trio analysis) (CNGS)
- 113\*. *In Silico* Panel from Clinical Exome Sequencing (INGS) (Single patient analysis)
 

<input type="checkbox"/> Brain Malformation panel	<input type="checkbox"/> Syndromic Autism panel
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- 114\*. *In Silico* Panel from Clinical Exome Sequencing (INGS) (Trio analysis)
 

<input type="checkbox"/> Brain Malformation panel	<input type="checkbox"/> Syndromic Autism panel
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- 115\*. Whole Exome Sequencing (WES) (ENGS) (Single patient analysis)
- 116\*. Whole Exome Sequencing (ENGS) (Trio analysis)
- 117\*. Open up the CES data after in silico panel (RNGS)
- 118\*. Open up the WES data after in silico panel (RNGS)
- 119\*. Sanger DNA sequencing for confirmation of variant detected by NGS/patient (SEQ)
- 99. Research Area (R): \_\_\_\_\_

**Consent**

I authorize the Department of Cytogenetics and Genomics to test my (or my child's/my fetus') sample for the designated genetic condition. My signature below constitutes my acknowledgment that the benefits, risks and limitations of this test have been explained to my satisfaction by a qualified health professional. I understand that all test results are treated with confidentiality and that results will only be provided to the referring clinician.

I authorize the department to store my/my child's/fetus' DNA sample, for future reference or use upon authorization.

Patient's/Guardian's Signature: \_\_\_\_\_ Date: \_\_\_/\_\_\_/\_\_\_

Clinician's Signature: \_\_\_\_\_ Date: \_\_\_/\_\_\_/\_\_\_