



Referral Letter

Department of Molecular Genetics, Function & Therapy

Head: Leonidas Phylactou PhD

P.O.BOX 23462, 1683 NICOSIA, CYPRUS

TEL.: +357-22392659, FAX: +357-22392615

Email: mgftlab@cing.ac.cy

Department Code: 27

DNA/MS/ST # _____ Dept. code _____

Patient details CING No.: _____ (if applicable)

Name: _____ Surname: _____

D.O.B.: ____/____/____ I.D. No.: _____

Nationality: _____ Gender: ☐ Male ☐ Female

Patient Status:

☐ GESY ☐ Private-Non GESY

☐ Government-Non GESY Hospital Card No.: _____

Case type: ☐ Outpatient ☐ Inpatient

Address: _____

City: _____ Code: _____ Country: _____

Phone: _____ e-mail: _____

Informed Consent

I authorize the Department of Molecular Diagnosis Function and Therapy to use my (or my child's/my foetus) sample (whole blood, serum or CSF) for test validation or research purposes after the personal details are removed making the procedure anonymous.

I have the right to refuse the above without having any consequence to the analysis. Samples will be stored for future reference or use.

- I can withdraw my consent at any time by contacting the laboratory at +35722392659.

Patient/Guardian Signature: _____ Date: ____/____/____

Sample Receipt (For Laboratory Internal Use)

Received by: _____ Signature: _____

Receipt Date: ____/____/____ Sample Type & Amount: _____

Referring clinician's / scientist's details

Name: _____ Surname: _____

Hospital / Clinic: _____

Address: _____

City: _____ Code: _____ Country: _____

Phone: _____ Fax: _____

e-mail: _____

Reason for Referral: _____

Clinician Signature: _____ Date: ____/____/____

Referring clinician status:

☐ CING ☐ Government (OKYπY) ☐ Private -Non GESY

☐ Private-GESY GESY No. _____

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

Please tick the box if diagnostic NGS tests are been 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: _____

Sample details (Please tick ☒ accordingly)

Date and Time of Sample Collection: _____

Sample: ☐ Blood (3-4ml) ☐ CSF (0.5ml) ☐ Urine ☐ Muscle

☐ CVS Other (please specify): _____

☐ First Investigation ☐ Repetition

For Genetic testing 3-4ml whole blood in EDTA is required.

For Multiple Sclerosis 0.5ml Serum and 0.5ml CSF are required.

Test Required (Please tick ☒ accordingly)

CFTR GENE, CYSTIC FIBROSIS (CF)

☐ CFTR analysis by NGS (1.1) *ISO 15189 (GESY)

Sweat Test: ☐ Sweat test (8.7) (GESY)

HEREDITARY RECURRENT FEVERS (HRFS)

☐ Hereditary recurrent fevers (HRFs) *in silico* panel from WES (24.3) *ISO 15189 (GESY)

MEFV Gene, Familial Mediterranean Fever *ISO 15189

☐ MEFV gene mutation analysis (sequencing Exons 2, 3, 5 & 10) (7.1) (GESY)

☐ MEFV gene analysis for known mutation (sequencing) (7.2) (GESY)

Mevalonate kinase Deficiency - MKD

☐ MVK gene full mutation analysis (sequencing) (18.1) (Non-GESY)

☐ MVK gene analysis for known mutation (sequencing) (18.2) (Non-GESY)

Tumour Necrosis factor associated periodic syndrome - TRAPS

☐ TNFRSF1A gene full mutation analysis (sequencing) (23.1) (Non-GESY)

☐ TNFRSF1A gene analysis for known mutation (sequencing) (23.2) (Non-GESY)

Cryopyrin-associated periodic syndrome - CAPS

☐ NLRP3 gene full mutation analysis (sequencing) (24.1) (Non-GESY)

☐ NLRP3 gene analysis for known mutation (sequencing) (24.2) (Non-GESY)

INHERITED DEAFNESS

☐ Hearing Loss *in silico* panel from WES (15.6) *ISO 15189 (GESY)

Connexin 26 & 30, Inherited Deafness

 *ISO 15189

☐ Connexin 26 full mutation analysis (sequencing) (13.1) (GESY)

☐ Connexin 26 analysis for known mutation (sequencing) (13.2) (GESY)

☐ Connexin 30, Del(GJB6-D13S1830) (15.1) (GESY)

HAEMOCHROMATOSIS

HFE Gene, Haemochromatosis *ISO 15189

HFE gene, Haemochromatosis analysis of individual mutations (10.7) (GESY)

☐ p.H63D ☐ p.C282Y

MULTIPLE SCLEROSIS

☐ Oligoclonal Bands detection for MS patients (14) *ISO 15189 (GESY)

EXOME SEQUENCING

 *ISO 15189

☐ Whole Exome Sequencing (WES) (43) (GESY) / ☐ WES (Trio) (44) (GESY)

☐ Sanger sequencing based confirmation of a variant identified by NGS (46) (GESY)

DISORDERS OF SEXUAL DIFFERENTIATION

☐ Disorders of Sexual Development *in silico* panel from WES (45) *ISO 15189 (GESY)

☐ CYP21A2 Gene, Congenital Adrenal Hyperplasia (CAH) *ISO 15189

☐ CYP21A2 gene full mutation analysis (sequencing & MLPA) (16.1) (GESY)

☐ CYP21A2 gene analysis for known mutation (sequencing) (16.2) (GESY)

SRD5A Gene, 5α Reductase Deficiency, Disorder of Sexual Differentiation *ISO 15189

☐ SRD5A2 gene, full mutation analysis (sequencing) (21.1) (Non-GESY)

☐ SRD5A2 gene analysis for known mutation (sequencing) (21.2) (Non-GESY)

PROP1 Gene, Growth Hormone Deficiency

☐ PROP1 full mutation analysis (sequencing) (22.1) (Non-GESY)

☐ PROP1 DNA extraction/storage (22.6) (Non-GESY)

PREMATURE AND DELAYED PUBERTY

☐ Premature and Delayed Puberty *in silico* panel from WES (38.4) *ISO 15189 (GESY)

Central Precocious Puberty (CPP)

☐ MKRN3 gene full mutation analysis (38.1) (Non-GESY)

☐ MKRN3 gene known mutation analysis (38.2) (Non-GESY)

GLUCOSE AND INSULIN HOMEOSTASIS (MODY AND OBESITY)

☐ Glucose and Insulin Homeostasis *in silico* panel from WES (20.7) *ISO 15189 (GESY)

GCK and HNF1α Genes, Maturity Onset Diabetes of the Young (MODY)

☐ GCK gene, MODY 2 full mutation analysis (sequencing) (19.1) (Non-GESY)

☐ GCK gene, MODY 2 analysis for known mutation (sequencing) (19.2) (Non-GESY)

☐ HNF1α gene, MODY 3 full mutation analysis (sequencing) (20.1) (Non-GESY)

☐ HNF1α gene, MODY 3 analysis for known mutation (sequencing) (20.2) (Non-GESY)

MC4R Gene, Obesity

☐ MC4R gene full sequencing (17.1) (GESY)

☐ Obesity, analysis for known mutation (sequencing) (17.2) (GESY)

THYROID FUNCTION

☐ Thyroid Function *in silico* panel from WES (39.1) *ISO 15189 (GESY)

RET-proto-oncogene, Multiple Endocrine Neoplasia type II *ISO 15189

☐ RET mutation analysis (sequencing Exons 10, 11 13 14, 15 & 16) (11.1) (GESY)

☐ RET analysis for known mutation (sequencing) (11.2) (GESY)

OTHER

☐ DNA extraction/storage (96)