

NHLS LAB NUMBER BARCODE

Prac	ctice number: 5200296 (Lab	contact nu	umber: 021 40	4 4449 /	021 650	1630)									
INF	IERITED METABOLIC D	ISEASE	GENETIC T	EST RE	QUEST	T FORN									
L	HOSPITAL / CLINIC		c ICD10 CODES			CLINICAL DIAGNOSIS									
0	WARD						L		•						
С	COPY REPORT TO						N								
	PATIENT ID NO													ID/Pas	sport
Р	HOSPITAL NUMBER														
Α	SURNAME														
T	FIRST NAME												SEX	M	F
E N	DATE OF BIRTH	D	D	/	M	M	/	,	Υ	Υ	Υ	Υ	AGE		
т	PATIENT ADDRESS														
	PATIENT TEL NO	H:					W:				(C:			
	TATIENT TEE NO														
Р	MEDICAL AID				PLA	N									
R	MEDICAL AID NO			DEP	DEP CODE										
v	MEMBER NAME								D NO						
T	MEMBER ADDRESS														
E	MEMBER TEL NO	H:					W:				(C:			
	CLINICIAN NAME								HPCSA / SANC NO						
Н	CONTACT NO						EMAIL ADDRESS								
w	CONSULTANT I/C							☐ I have taken informed consent from patien						atient	
	CONTACT NO/EMAIL				(see back of form)										
	EDTA blood	Urine (Early morning) Skin Biopsy site:							Diagnostic (All biochem/clin data provided))
S	Muscle						T .		Prenatal (Mutation details provided)						
P E	Other	Describe:						TYPE			testing (Fam history provided)				
С	Date collected	Time									(Fam history provided)				
	Collected by:								Other						
_	IMD LABOR	ATOR'	Y GENET	TIC TE	ESTS	(excl	uding	mito	ochor	ndria	l dise	ase –	see ov	erleaf	
١Ļ	Alpha-1 Antitrypsin Deficiency (<i>Serpina1</i> : S and Z alleles) AMP Deaminase deficiency (<i>AMPD1</i> Full gene)			es)	☐ Glutaric aciduria Type 1 (GCDH) ☐ Common Indigenous African mutation (p.A293T) ☐ Full sequencing (UOA results required)										
۱⊨															
-	-	tinidase deficiency (<i>BTD</i>) Common European mutation (p.Cys33PhefsTer36)					Glutathione synthetase deficiency (GSS)								
	☐ Full gene sequer						Glycogen storage disease								
	Centronuclear myopathy, AR (RYR1)						☐ Type 1A (<i>G6PC</i>) full sequencing								
	☐ Common Indigenous African mutations (2 mutations)				ons)	☐ Type 5, McArdles (<i>PYGM</i>) p.R50X (European)									
	Common SA Caucasian mutations (4 mutations)					Hyperbilirubinemia (<i>UGT1A1</i>)									
_	☐ Mixed ancestry panel (5 mutations)					Gilbert syndrome common variant only									
╽┝	Congenital adrenal hyperplasia (17-OHP required)					☐ Criggler-Najjar / Gilbert full sequencing ☐ Isovaleric Acidemia (<i>IVD</i>) Afrikaner mutation (p.G123R)									
-	☐ Cystinosis (C7NS) ☐ Common Indigenous African mutation (c.971-12G>A)					Lesch Nyhan Syndrome (<i>HPRI</i>) (send 2 EDTA tubes on ice)									
	☐ Full gene sequencing (AA/UOA results required)					Mevalonic aciduria (MVK) (UOA results required)									
	Disorders of sexual disorientation (DSD) (Discuss with lab)					Primary hyperoxaluria (<i>AGXI</i>)									
	☐ Androgen receptor insensitivity (<i>AR</i>)						Common Indigenous African mutation (p.A112D)								
_	☐ Steroid 5-alpha reductase deficiency (<i>SRD5A2</i>)						Full sequencing								
	Galactosaemia (GALI)				Urea cycle disorders (AA/UOA results required)										
	☐ Common Indigenous African mutation (p.S135L) ☐ Full sequencing (Enzyme results required)				☐ N-Acetylglutamate synthase (<i>NAGS</i>) ☐ Ornithine transcarbamylase (<i>OTC</i>) gene										
-	·	EHZYTTIE	resuits reqt	iiied)			П.					-	_	ie	
	Full sequencing (Other:	Enzyme	results requ	uired)			□×					ylase (0 ophy (A	_	ne	

IMD LABORATORY - MITOCHONDRIAL GENETICS							
Mitochondrial DNA (mtDNA) genetics: mtDNA full sequencing (muscle preferred, blood accepted) Clinical scenario: MELAS; MERRF; LHON Leigh syndrome; Other: mtDNA large deletion screen for: CPEO (muscle ONLY) Kearn Sayre syndrome (muscle ONLY in >20 year old) Pearsons disease (blood ONLY) Multiple deletion screen (muscle ONLY) mtDNA point mutation screens (single gene/mutation only): MELAS m.3243A>G only (blood and urine/muscle) MERRF MT-TK only (blood and urine/muscle) MIDD m.3243A>G (blood and urine/muscle) NARP m.8993T>C/G (blood/urine) LHON screen (3 common mutations) (blood) mtDNA depletion screen (arrange with lab): Other (specify):	Nuclear DNA (nDNA) genetics: Leigh syndrome SURF1 PDH deficiency (PDHA1) Full Leigh disease gene panel (discuss with lab) CPT2 deficiency ETHE1 deficiency (UOA results required) mtDNA maintenance disorders: MPV17 neurohepatopathy Common Indigenous African mutation (p.Q36X) Sequencing (mRNA, arrange) POLG sequencing (costly, discuss with lab) Thymidine kinase 2 (TK2) Twinkle (c10ORf2) sequencing (arrange) MNGIE (TYMP) sequencing (arrange) mtDNA depletion gene panel (discuss with lab)						
Clinical Details (Mitochondrial disease): If yes, then which of the following? Alpers' syndrome CPEO CPEO Leigh syndrome Limm Melas If no, then which of the following clinical features are present? Anaemia Diabetes Cardiomyopathy Dysphagia Central apnoea Dystonia Constipation Encephalopathy Deafness Endocrinopathy Failure to thrive Failure to thrive Fatigue Muscle biopsy results (if available): Enzymology: Clinical investigations: Blood Lactate	, ,	NARP/MILS Pearson's syndrome Pure myopathy SANDO SNHL Nystagmus Renal disease Retinopathy Optic atrophy Seizures Stroke/stroke-like episodes					
Metabolic / Biochem findings (i.e. UOA / AA results): Other Relevant clinical findings and family history:							
Ethnic origin of patient's motherFather_		For lab use only Received by: Labelled by: Date: Time:					

CONSENT FOR DNA ANALYSIS AND INCLUSION IN RESEARCH STUDIES PLEASE MAKE A COPY OF THIS PAGE FOR THE PATIENT

Research Lab contact number: 021 650 1630

1.	. I,, request DNA testing on myself \Box / my ch	ild ☐ / my unborn child ☐ for disease-					
	causing mutations in the gene/genes responsible for:						
2.	I understand that the genetic material (DNA) for analysis is to be obtained from: blood ☐, sk tissue ☐ (specify:) provided today.	kin □, muscle □, or any other body					
3.	I request that a portion of the sample be stored indefinitely for (Tick where applicable):						
	 ☐ Possible re-analysis. ☐ Further diagnostic testing / analysis when more information about my disorder becomes a ☐ Analysis for the benefit of members of my immediate family. OR ☐ I request that no portion of the sample be stored for later use. 	available.					
4.	If clinically relevant, I authorise that the results may be made known to family members \square .						
5.	I give consent $\ \ \ \ \ \ \ \ \ \ \ \ \ $						
	Information about consent for research: Research into rare metabolic disorders is param conditions that affect patients in South Africa and find new and effective ways to diagnose at some instances taking part in such studies may provide diagnostic answers to individuals / in that this is not always the case. Agreeing to make your biological samples and health inform have no direct benefits to you, but may go a long way in ensuring better diagnostic and treat What are the risks to me / my child / my family? The main concerns people have with get their genetic or clinical information may become known to third parties, such as insurance of discrimination. Rest assured that every attempt is made to reduce the risks of such confider samples and information are stored securely with controlled access which is strictly limited to staff from the IMD laboratory who are bound by confidentiality agreements. If at any stage yet such research you may withdraw your consent with no questions asked by contacting the IM.	and treat these conditions. Although in advividual families, it is important to note ation available for such research may ment outcomes for future patients. In the cresearch is that, although unlikely, ompanies, possibly resulting in stigma or atiality breaches occurring. All patient to the authorised research and diagnostic outchange your mind about taking part in					
6.	I have been informed that (Delete where not applicable): (a) The analysis procedure is specific to the genetic condition and cannot determine the complete genetic makeup of an individual. (b) The IMD laboratory is under obligation to respect medical confidentiality and results are only reported to the specified doctor(s). (c) Genetic analysis may not be informative for some families or family members. (d) Even under the best conditions, current technology of this type is not perfect and could lead to incorrect results. (e) Where biological material or data is used for research purposes, there may be no direct benefit to me.						
7.	understand that I may withdraw my consent for any aspect of the above at any time without this affecting my future medical care.						
8.	ALL OF THE ABOVE HAS BEEN EXPLAINED TO ME IN A LANGUAGE THAT I UNDERSTAND AND MY QUESTIONS ANSWERED BY:						
	Doctor / Consultant Signature:	Date					
	Patient (or Parent in case of a minor) Signature:	Date					
	Witness Signature:	Date					

RECOMMENDED MITOCHONDRIAL DISEASE GENETICS TESTING STRATEGIES Adapted from Meldau et al. (2016) SAMJ; 106(3):234-236

