NAME, First name:	
(CAPITAL LETTERS, please)	

Laboratory	only
DM-	



DIAGNOSTIQUE / Service de Médecine Génétique

Centre d'accueil des prélèvements (CAP) Bâtiment des Laboratoires (BATLab), local 8D-0-850.1 4 rue Gabrielle-Perret-Gentil, 1211 Genève 14

Genomic and Molecular Diagnostics Laboratory Accredited since 2003, formerly STS 0382



DIAGMOL

http://www.hug-ge.ch/feuilles-de-demande

Head of Genetic Medicine Division: Prof. Marc ABRAMOWICZ Head of Laboratory, medical genetics FAMH Dr. sc. J.-L. BLOUIN, jean-louis.blouin@hcuge.ch Biologists laboratory managers, medical genetics FAMH

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Mr. Mrs. (IN UPPER CASE, please)
Name:
Maiden Name:
First name:
Date of birth: / /
Legal representative (for minors) : □father □mother
Name/first name :
Street/N°:
Town, ZIPCODE:
Hospitalisation Unit: Physician :
N° EdS:
Insurance : Insured N :

Lab direct or results: Phone/FAX: +41 (0) 22 37 21 826 / 21 860 Sample Entrance Center (CAP) : Phone +41 (0) 22 37 21 800	
PHYSICIAN	
PHYSICIAN (NAME/First name - Street/N°- Town, ZIPCODE - Phone/FAX. IN UPPER	CASES, PLEASE)
COPY TO OTHER PHYSICIAN (NAME/First name - Street/N°- Town, ZIPCODE - P.	Phone/FAX. IN UPPER CASES, PLEASE)
« The laboratory is granted permission by the Physician/Patient to transmit copies of the ☐ Opposition of the patient to the registration of this request results in t	
If the patient belongs to a family already known to the laboratory, please	indicate index case NAME:
CLINICAL INFORMATIONS given by the physician:	
oznino iz nin oram triono givon by the physiciani	
given by the physician	

Ethnic origins Father Mother **Currently pregnant** Date of last menses Number of weeks of amenorrhea SAMPLE(S)

Most of our tests work from 4 ml of blood in EDTA (children <2 ans : 1 ml: ok) or from **purified DNA** (some exceptions apply for some tests) Please contact us for any other type of sample.



On every and single tube. Mandatory!

NAME First name Date of birth

Sampling date : Time (optional) : Sample Number:

TA)

☐ DNA from external source

☐ Saliva (sampled only in tube Oragene-DNA)

□ Other type of sample Please indicate type:

☐ DNA already in bank at our laboratory Reference number:

Prenatal □ Amniotic fluid □ Chorionic villi □Fetal tissue

□Other

O sang/EDTA O ADN déjà en banque O Tissus fixés O Liq. Amnio.

Quantité, Remarques

NAME, First name : (CAPITAL LETTERS, please)	Laboratory only
	DM-
(if known)	
TEST LIST : SEE OVER (PAGES 3 ET 4)	
PHYSICIAN'S SIGNATURE AND INFORMED CONSENT	
PHYSICIAN (NAME/First name – Street/N°- Town, ZIPCODE - Phone/FAX): * By signing here, the physician confirms having informed the patient/the legal according to the current legal requirements (LAGH in Switzerland, http://www. (including on the cost of tests that are not covered by health insurances), that the legal representative had enough time to ask questions and take his/her decand having received the patient's/the legal representative's informed consent.	sgmq.ch) the patient/
The patient has given his informed consent for the checked a sample specified at the first page.	nalyses that are listed at the pages 3-4, to be done on the
The following questions marked by a star * must be checked (MA	NDATORY!).
Decision of the patient regarding the storage and use of his/h *mandatory	er remaining biological sample(s) and raw analytical data :
 He/she agrees that the remaining biological material and raw informed consent will be necessary for any further additional In case of a negative answer the remaining biological sample will be destroyed 	
He/she agrees that his/her biological sample and raw analytical	al data are used anonymously for quality testing. * □YES □NO
Decision of the patient regarding the transmission of results	OUGHPUT SEQUENCING OF WHOLE EXOME (SEE PAGE 4) not directly related to the testing requested (so called "incidental
findings") *mandatory	
He/she wishes to be informed about genetic results belonging to t	ne following categories :
 Carrier of a disorder for which preventive and/or therapeutic m 	easures are available *:
Person incapable of discernment: □YES □NO	
The following questions do NOT apply for persons incapable	
 Carrier of a disorder for which no preventive / therapeutic mea 	sures are yet available*: □YES □NO
Healthy carrier of a recessive disorder which could concern the	e following generation or other family members*: □YES □NO
Other decisions :	
OPTIONAL	
The use of his/her sample and data for research purposes.	
Should he/she agree in principle to participate in research studies	and the disease this below. Obsailed this be the seese be/she are all
be contacted at a later stage with details concerning the research participation in any actual research projects.	project(s). A positive answer below is not yet consent for the

NAME, First name:		Laboratory only
(CAPITAL LETTERS, please)		DM-

REQUESTED ANALYSIS / ANALYSES

NB: IF REQUEST IS FOR A HIGH THROUGHPUT SEQUENCING (GENOME CLINIC), PLEASE GO DIRECTLY TO PAGE 4.

* Test not included in the Swiss federal list of laboratory tests (OFSP, BAG, FOPH). The out-of-list tests are not automatically reimbursed by Swiss health insurances.

na Test not accredited; @ Please contact us in advance. The rates applied are those of the list of analyzes of the OFSP / BAG. The laboratory reserves the right to select the most appropriate technique (traditional or high throughput sequencing, cf. page 4) based on efficiency and cost effectiveness. cl Depending on the choice of technique, the analysis may or may not be accredited.

General tests	Cystic Fibrosis (CF, CFTR)	
☐ Alpha-tryptasemia (TPSAB1 and TPSB2)	Please indicate the ethnic origins of the patient at page 1 (for	Mitochondriopathies (ADNmt)
☐ Amyloidosis (familial, <i>TTR</i>)	residual risk calculation) Screening frequent mutations (with / without	Leber optic neuropathy (LHON)
☐ AS, Angelman syndrome ^{na}	IVS8 5T, <i>CFTR</i> -related disorders)	☐ Cytopathy MELAS, MERRF, NARP ☐ Deletions/Dup (muscle biopsy only) *
☐ APECED (AIRE)	☐ Full CFTR analysis (sequencing+ del/dup by	Chromosome Mit sequencing
☐ Beckwith-Wiedemann (BWS) na	MLPA) ^{na} Hyperophogonia fatal howel	Monogenic Diabetes (MODY, NDM) *
☐ BPES (FOXL2) ×	Hyperechogenic fetal bowel (frequent mutations CFTR in parents)	Whole Exome Sequencing and Targeted Gene
☐ CMM2 Cutan. Malign. Melanoma (CDKN2A) *na	Hyperechogenic fetal bowel	Panel Analysis: see next page (page 4) Or
☐ EGFR mutations (T790M and others) on ctDNA	(frequent mutations CFTR +del/dup by MLPA in	☐ HNF4A gene (MODY 1) d ☐ GCK gene (MODY 2) d
(only in <u>Streck BCT</u> or <u>PAXgene DNA</u> tubes) ≭	parents)	☐ HNF1A gene (MODY 3) cl
☐ FG (Keller syndrome, <i>MED12</i>) **na	Deafness ☐ DFNB1, congenital (locus DFNB1) ★	PDX1 gene (MODY 4) d
☐ HBLRG, Gilbert syndrome (<i>UGT1A1</i>) *	☐ Mitochondrial mutations **na	☐ HNF1B gene (MODY 5) d ☐ INS gene (MODY 10) d
☐ HDGC, Her. Diff. Gastric Cancer (CDH1) **na	Endocrine Neoplasias, Pheochromocytoma,	☐ INS gene (MODY 10) ^d ☐ KCNJ11 gene ^d ☐
☐ HED, Hypohidrotic Ectodermal Dysplasia (EDA) *	Paraganglioma (MEN, PCC, PGL)	Neurological and Neuromuscular
☐ HFE-HH, Hered. Hemochromatosis (<i>HFE</i>)	☐ MEN1, Multi, Endoc, Neopl, type I (MEN1)	SMA, Spinal Musc. Amyotrophy (SMN1)
☐ HSCR, Hirschsprung (<i>RET</i>) x na		CADASIL (NOTCH3) *
XLI, Ichtyosis, X-linked type (STS)	Full sequencing (+MLPA) or	☐ C9orf72 (ALS, FTD) * ☐ DOPA-responsive dystonia (<i>GCH1</i>) * ^{na}
☐ PFIC3, Intrahepatic Cholestasis (ABCB4) x ^{cl}	☐ SDHB gene ☐ RET gene	☐ Dravet syndrome (SCN1A) *na
SMAX1, Kennedy (SBMA, AR) na	☐ SDHC gene cl ☐ SDHD gene	☐ EPM1, Unverricht-Lundborg (CSTB) ★
☐ KNO1, Knobloch, (COL18A1) ^{na}	Individual prices available upon request Von Hippel Lindau (VHL)	Southern+sequencing
☐ Lactose intolerance (LCT) ×	Familial Pneumothorax (Birt-Hogg-Dubé, BHD) **	FSHD1, Facio-Scapulo-Humeral Dystrophy,
☐ LWD, Leri-Weill (SHOX)	☐ Frequent mutation, FLCN, exon 11	type1 ★ ^{na} (only from fresh EDTA blood) ☐ GLUT1 (SLC2A1) ★ ^{na}
☐ LFS, Li Fraumeni, (TP53)	FLCN full gene analysis	☐ SPAST, Hered. Spastic Paraparesis (SPG4)
☐ Marfan (FBN1)	Familial Adenomatous Polyposis (FAP)	☐ STARTLE (Hyperekplexia, GLRA1) * na
☐ NF1, Neurofibromatosis type I (<i>NF1</i>) na	☐ Full Screening APC + MUTYH or	DMD, BMD, Muscular Dystroph Duchenne/Becker
Non-invasive prenatal diagnostic of monogenic	☐ APC gene ☐ MAP (MUTYH, ex. 7,13) ^{na}	(DMD), deletions
diseases (contact us in advance) *	FGFR3 (syndromes linked to)	DM1, Myotonic Dystrophy of Steinert (<i>DMPK</i>)OPMD, Oculopharyngeal Muscular Dystrophy
☐ PJS, Peutz-Jeghers (STK11) na	Achondroplasia	(PABPN1) **na
☐ PTEN Hamartoma Tumor syndrome (PHTS, Cowden,	Craniosynostosis or Muenke	☐ TOR1A, Torsion Dystonia (DYT1) ** na
Hamartomas, BRRS, Proteus, PTEN)	☐ Hypochondroplasia☐ Thanatophoric dysplasia, types I, II	Pancreatitis
PWS, Prader-Willi na	SADDAN	☐ CFTR+ IVS8 5T (frequent mutations) ☐ SPINK frequent mutation
☐ Rendu-Osler-Weber (ROW) * ^{cl} @	Fibrinopathies *	 □ SPINK frequent mutation □ PRSS1 frequent mutations *
☐ RETT syndrome (<i>MECP2</i>)	Afibrinogenemia (FGA, FGB, FGG)	Primary Ciliary Dyskinesia (PCD) *
RSS, Russell-Silver syndrome (11p15) na	Dysfibrinogenemia (FGA, FGG)	Whole Exome Sequencing and Targeted Gene
☐ Sickle cell anemia (Drepanocytosis, HBB)	☐ Hypofibrinogenemia (<i>FGA</i> , <i>FGG</i>) Fragile X (FRAXA, <i>FMR1</i>) na	Panel Analysis: see next page (page 4)
■ UPD, Uniparental Disomy, Chr*	Diagnostic	Waardenburg (WS) ■ Types I and III (PAX3 gene)
∇WF, all types, ★na @	☐ Carrier testing	Type II (MITF gene) na
WAGR, Wilms tumor (WT1) ★na	Premature ovarian failure (POI)	_ ',F' (g)
Alpha-1-antitrypsin deficiency (A1AT)	Genetic sex Genetic sex determination	Miscelleanous (cf. additional informations)
Genotyping PI*S/Z	☐ SRY search in a Turner	DNA Extraction + Banking
☐ Full sequencing of SERPINA1 Ashkenazi mutations (rare disease carrier)	☐ SRY sequencing	☐ RNA Extraction + Banking na (only in PAXgene blood RNA tubes)
☐ Full screening * or	Hemophilias	☐ Circulating cell-free DNA Extraction
☐ CFTR ☐ Fragile-X	☐ HA, inversions F8 (IVS22, IVS1) ★	(only in <u>Streck BCT</u> or <u>PAXgene blood DNA</u> tubes)
☐ Tay-Sachs+ FD+Fanconi+Canavan ★	☐ HA, <i>F8</i> , complete analysis ^d ☐ HB, <i>F9</i> , complete analysis	☐ Out-of-list (OFAS) gene na * per exon
☐ von Gierke+Bloom+Niemann-Pick+ML-IV Individual prices available upon request Individual prices available	Huntington disease (HD, <i>HTT</i>)	☐ Specific/ Known familial mutation *
Ataxias	Diagnostic	mRNA analysis **** @ (blood only in PAXgene blood RNA tubes)
☐ Full screening	Presymptomatic (2 tubes please)	Exclusion of maternal contamination in fetal
□ SCA1 na □ SCA2 na □ SCA3 na	Hereditary Periodic Fevers (HRF) *	sample (amnio-, choriocentesis) *
SCA6 na SCA7 na SCA17 na	☐ Full Sequencing (8 genes) ☐ Full Screening Frequent Mutations (4 genes)	☐ Transfer of DNA to an external laboratory
☐ Friedreich ☐ DRPLA ^{na} ☐ FXTAS ^{na} Cardiac Arrhythmias (Channelopathies, CCP) *	FMF, MEFV gene	(please provide specifics below) and complete
☐ SCN5A gene (Brugada) cl	FMF, MEFV gene (complete sequencing)	the ECA forms for internal requests. Transfer of DATA of NGS to an external
☐ KCNQ1 gene (QT-long) cl	CAPS, NLRP3 gene	laboratory (please provide specifics below),
KCNH2 gene (QT-long) d	☐ HIDS, <i>MVK</i> gene ☐ TRAPS, <i>TNFRSF1A</i> gene	complete the DATA exchange form and
☐ KCNE1, KCNE2, KCNJ2 genes d Whole Exome Sequencing and Targeted Gene	Whole Exome Sequencing and Targeted Gene	complete the consent level 3.
Panel Analysis: see next page (page 4)	Panel Analysis: see next page (page 4)	
Cardiomyopathies (HCM, DCM, NC, CMR,) *	HNPP (tomaculous neuropathy)	
Whole Exome Sequencing and Targeted Gene	Deletion CMT1A	
Panel Analysis: see next page (page 4)	☐ PMP22 gene sequencing Lynch syndrome (HNPCC)	
Carrier Screening Full screening *	Full screening by NGS see next page (page 4) or	
Charcot-Marie-Tooth (CMT)	☐ MLH1+PMS2 genes	
Duplication CMT1A	☐ MSH2+MSH6 genes	
PMP22 gene sequencing (CMT1A)	☐ MSI (on tumor biopsy)	
☐ MPZ gene sequencing (CMT1B)☐ GJB1 gene sequencing (CMTX)	☐ BRAF1 V600E (on tumor biopsy) na Male infertility	
Chromosomal Microdeletions *	CFTR+5T (frequent mutations)	
22q11, MLPA	☐ Y chromosome microdeletions (DAZ)	
☐ Screening, recurrent microdeletions, MLPA		

-	_	
NAME, First name:		
(CAPITAL LETTERS, please)		DM-
•	_	

REQUESTED ANALYSIS / ANALYSES

HIGH THROUGHPUT SEQUENCING OF TARGETED OR WHOLE EXOME AND BIOINFORMATIC ANALYSIS (GENOME CLINIC) NB: IF THE REQUEST CONCERNS A CLASSICAL ANALYSIS, WITHOUT HIGH THROUGHPUT SEQUENCING, SEE PAGE 3 * Test not included in the Swiss federal list of laboratory tests (OFSP, BAG, FOPH). The out-of-list tests are not automatically reimbursed by Swiss health insurances.

"a Test not accredited; @ Please contact us in advance (availability, TAT,...). The rates applied are those of the list of analyzes of the OFSP / BAG. The laboratory reserves the right to select the most appropriate technique (high throughput sequencing or traditional, cf. page 3) based on efficiency and cost effectiveness. DNA extraction and banking HIGH THROUGHPUT SEQUENCING FOLLOWED BY BIOINFORMATIC ANALYSIS OF 1 TO 10 GENES @ Prescription by a physician with a federal postgraduate FMH diploma in medical genetics or related to the examined pathology, according to the Swiss federal list of laboratory tests (Chapter 2.Genetics na 2.2.2.Moleculare genetics analyses). ☐ Cardiac Channelopathies (Arrhythmias, CCP) * ☐ Lynch syndrome (HNPCC) ☐ Cardiomyopathies ***** ■ Neurofibromatosis type I ☐ Growth disorder syndromes ☐ Diabetes, monogenic (MODY, NDM) * (Beckwith-Wiedemann, Silver-Russel, Sotos, etc) ☐ Primary Ciliary Dyskinesia (PCD) * ☐ FGF receptor-associated dysplasias Other disease (please specify below the genes to analyze and the ☐ Duchenne and Becker dystrophinopathies and muscular position in the Swiss list of laboratory tests) dystrophies (protein disorders associate with dystrophin) ☐ Hereditary periodic fevers * ☐ Other orphan diseases ***** (please specify below the genes to analyze) ☐ Wilson's disease An "Orphan disease" reimbursement request must be filled by a physician with a federal postgraduate FMH diploma ☐ Hereditary neoplasia HIGH THROUGHPUT SEQUENCING FOLLOWED BY A BIOINFORMATIC ANALYSIS OF MORE THAN 10 GENES @ Prescription only by a physician with a federal postgraduate FMH diploma in medical genetics, according to the Swiss federal list of laboratory tests. Mitochondriopathies Diseases related to urogenital system, fertility / sterility ☐ 11-100 genes ☐ 11-100 genes ☐ > 100 genes ☐ > 100 genes Cardiac Channelopathies (Arrhythmias, CCP) * Hereditary neoplasia 11-100 genes ☐ 11-100 genes Cardiomyopathies *na ☐ > 100 genes ☐ 11-100 genes Sensorimotor neuropathies Primary Ciliary Dyskinesia (PCD) * (CMT, HNPP, amyloid polyneuropathy) ☐ 11-100 genes ☐ 11-100 genes Diabetes, monogenic (MODY, NDM) * Kallman syndrome ☐ 11-100 genes ☐ 11-100 gènes **Ehlers-Danlos** Marfan syndrome and other thoracic aorta syndromes ☐ 11-100 genes ☐ 11-100 genes Mendelian syndrome with growth disorder Epilepsy * ■ 11-100 genes ☐ 11-100 genes ☐ > 100 genes ☐ > 100 genes Diseases related to coagulation, blood and immune system disorders Developmental disorders * ☐ 11-100 genes ☐ 11-100 genes ☐ > 100 genes ☐ > 100 genes Neuromuscular et neurodegenerative diseases Other diseases (please specify below the genes to analyze and the ☐ 11-100 genes position in the Swiss list of laboratory tests) ☐ > 100 genes ☐ 11-100 genes Diseases related to skin, connective tissue or bones ☐ > 100 genes ■ 11-100 genes ☐ > 100 genes Other orphan diseases * (please specify below the genes to analyze) Metabolic and endocrine diseases ursement request must be filled by a physician w An "Orphan disease" reimbursemen a federal postgraduate FMH diploma ■ 11-100 genes ☐ > 100 genes 11-100 genes Ophtalmologic diseases ☐ > 100 genes ☐ 11-100 genes ☐ > 100 genes **ADDITIONAL ANALYSES @** Additional bioinformatic analysis ☐ 1-10 genes ☐ 11-100 genes ☐ more than 100 genes Comments: INFORMATIONS ABOUT BIOINFORMATIC ANALYSES Gene panels: http://www.hug-ge.ch/medecine-genetique/liste-panels-genes Gene panel to analyze (please contact us in advance): (Please specify the investigated pathology, the number of genes and the requested gene panel (if available) or else provide your gene list in an Excel file) **SEARCH FOR VARIANTS IN THE PARENTS** In case of request of search of variants in the parents, please send us for each of them the sample and a request of DNA extraction and banking. ☐ Consanguineous parents Precisions/comments :__ Sample available : Tyes No Will be sent Father: Last name :__ ____ First name :__ __ Date of birth :___ _ Sample available : ☐ Yes ☐ No ☐ Will be sent _ First name :_ _ Date of birth :_ Mother: Last name:

Complementary information /comments: