

General Information

Fetus of		Sender / Clinic			
Surname of mother:		Surname:			
First name of mother:		First name:			
Birthday of mother:	_				
Sex of Fetus: ☐ male ☐ f	female 🛘 unknown	Institution:			
Has MCC testing been performed?	☐ Yes ☐ No	Street:			
Material		Postcode/City:			
☐ Amniotic fluid ☐ Chorionic villi	Starting material has been cultivated	Country:			
□ Abortion material		Phone:			
Extracted DNA μg (min. 1-2 μg from DNA-Nr.:		Email:			
		VAT:			
Forte we all ID		If applicable, please inclu	ude a VAT number or a copy of your bu	usiness registration certificate) .
		Invoice	to sender / clinicto patient / other (KVA-N	No.:)
Pregnancy week and estimated due of		Surname:	F ((
Samples can be sent by mail in a cardbox or air cut to direct sunlight. Dried blood spot cards can be o	shion envelope. Samples should not be exposed	First name:			
Declaration of consent		Street:			
By signing this form, I declare that I have rece the genetic background related to the disease limitations of molecular genetic testing. I unde	in question, as well as the possibilities and	Postcode/City: Country:			
consent for genetic analyses.					
I have been informed, and agree, that my period analysis will be recorded, evaluated or stored tabases, and that further, in accordance with the request, or parts thereof, may be transmitted.	I in an pseudonymized form in scientific dadata protection and medical confidentiality,	Email: If you do not check	these boxes, your answer	will be recorded as "	'No".
I consent to the re-evaluation of my test resucant alterations become apparent, my physical	ilts within the data storage period. If signifi-	I consent to the storage and/or quality control (fo	of my genetic material for addition max. 10 years).		□ No
have been informed, and agree to the electro sion of all data collected by CeGaT GmbH.	nic storage, processing, use, and transmis-	I consent to the storage 10 years (as required by	of my test results beyond the tim		□ No
For more detailed information on data priva www.cegat.com/privacy-policy.	acy as well as your rights please refer to		nymous storage and use of surpluults for scientific research and in	scientific	□ No
Please Note			ondary findings I would	☐ Yes 〔	□ No
All genes, including the complete mtDNA are sequenced when exome diagnostics is performed. The diagnostic evaluation is limited to variants in genes relevant to the provided		like to be informed:			
phenotypic information. Correct family relation analysis using data from several family memb This declaration of consent can be comple	pers (e.g., trio exome analysis).	of the requested gene these variants is limite	sometimes be identified, which tic analysis (so-called seconda and to pathogenic alterations (AC	ary findings). The report CMG classes 4 and 5)	ting of within
have had sufficient time to consider givin	, , ,	family (according to t	nich a treatment or course of the decurrent guidelines of the	American College of M	/ledica
I, the referring physician, confirm that I am a above-mentioned patient. For predictive testii I have fulfilled the requirements, to request the	ng, I confirm that I am authorized, and that	www.cegat.com/acmg-g	s; details on genes and associa <u>enes/</u>). There is no claim of a of of secondary findings cannot be	comprehensive analysis	of this
the consent of all legal guardians. If the patient did not sign this order form: patient received genetic counseling and agr			netic Diagnostic Act (GenDG) we we asse indicate here the contact email		
consent has been obtained in writing.		Email:			
		Physican's stamp	o / Barcode	(DAkkS	
				Deutsche Akkreditieru D-ML-13206	
				⊕ CAP	
Patient / Legal Guardian (Block letters)	Physican (Block letters)			ACCREDITED COLLEGE of AMERICAN PATHOLOG	_
•	,			CLIA CERTIFIED ID: 99D2130	
X Patient / Legal Guardian	XPhysican			CeGaT is accredited by DAkkS according to DIN EN ISO 15189:20 the College of America	14,
(Date, Signature)	(Date, Signature)			Pathologists (CAP) and	



Indication

Please attach copies of medical reports (including ultrasound or MRI reports, if available). The variant interpretation is based on clinical information available at the time of analysis.			
Indication / Suspected diagnosis	3:		
Ultrasound medical report avail	able? ☐ Yes (please attach copy) ☐ No		
Clinical symptoms:			
Preliminary genetic diagnostics	for fetus or parents?		
	s: ☐ Yes (please attach copy) ☐ No		
Array-CGH:	☐ Yes (please attach copy) ☐ No		
Other			
Pedigree	Consanguinity: ☐ Yes ☐ No	Ethnic origin:	
			○ □ not affected
			● ■ affected
			• known carrier
			Ø deceased
			unrelated parents
			consanguine parents
			unborn child
			↓ abortion, stillborn child
			person of unknown sex
			identical twins (monozygous)
			fraternal twins (dizygous)
Are there other family members w	n pregnant in the past, were there any anom ho currently have or have had a disease or	alies during pregnancy' disorder relevant for the	?
If yes, please list the affected fami			
(not required)	Relationship to the fetus (e.g., mother)	Age of onset	Diagnosis / Symptoms



Inquiry

Inquiry – Exome			
☐ Single Exome: Exome diagnostics of the fetus including medical report (EXM01)			
☐ Maternal cell contamination (MCC) testing (please provide material from the mother of the fetus, EDTA blood recommended)			
☐ Trio Exome: Comparative exome diagnostics between fetus and parents incl. medical report (EXM02)			
☐ Genes to be considered in the context of exome diagnostics:			
The analysis of the fetus and both non-affected parents (Trio Exome) a	llows a more efficient evaluation of the variants identified in the fetus and leads		
to an increased chance of positive identification of the disease causing	variants.		
Additional analysis (additional fees may apply)	For further information and advice please do not he-		
☐ Please perform array-CGH diagnostics	sitate to contact our Diagnostic Support team.		
☐ prior or	www.cegat.com/diagnostic-support		
☐ parallel	diagnostic-support@cegat.com		
to exome diagnostics.	Phone +49 7071 565 44-55		



Additional Information

Please use this space to provide any additional relevant information.			



Declaration of consent Parent 1

Personal data (Family membe	r)		
Surname:	First name:		
Date of birth:	Sample ID:		
Relationship to the patient			
☐ Father ☐ Mother	☐ Other; please state:		
Does the family member suffer t	from an illness or disorder with (suspected)	genetic cause?	
☐ No ☐ Yes, sy	mptoms are:		
of Medical Genetics and Genomic A negative "ACMG genes" report may not be performed for disease scope of the primary medical indi	cs. The analysis is restricted to the sequence data, cannot be used to rule out (genetic) disease risk. es which have an onset in adulthood. Therefore, so	nes for secondary analysis, according to the current guidel re-sequencing of regions with poor sequence coverage ware Additional fees may apply. According to German legislation me genes will not be analyzed for minors, unless the pherenes!	rill not typically be performed. on, predictive tests for minors
□ Pharmacogenetics (PGX) (3 ABCG2, CACNA1S, CYP2B6, C SLCO1B1, TPMT, UGT1A1, VKC	CYP2C19, CYP2C9, CYP2D6, CYP3A4, CYP3A5	i, CYP4F2, DPYD, G6PD, HLA-A, HLA-B, IFNL3, MT-RI	NR1, NUDT15, POR, RYR1,
I would like to receive an addit	ional report analyzing known variants in 22 ge	nes that are involved in the metabolism of pharmace	utical products.
Declaration of consent		If you do not check these boxes, your answer will be	
the genetic background related to the di	re received comprehensive information regarding sease in question, as well as the possibilities and I understand that I have the right to withdraw my	I consent to the storage of my genetic material for additi- and/or quality control (for max. 10 years). I consent to the storage of my test results beyond the tin	☐ Yes ☐ No
I have been informed, and agree, that analysis will be recorded, evaluated or tabases, and that further, in accordance	my personal data and the data obtained in the stored in an pseudonymized form in scientific date with data protection and medical confidentiality, ansmitted to a specialized cooperating laboratory.	10 years (as required by German law). I consent to the pseudonymous storage and use of surplumaterial and/or test results for scientific research and in s literature.	cientific
I consent to the re-evaluation of my tes cant alterations become apparent, my p	at results within the data storage period. If signifi- physican will be informed by email.	With regard to secondary findings I would like:	☐ Yes ☐ No
I have been informed, and agree to the estion of all data collected by CeGaT Gm	electronic storage, processing, use, and transmisbH.	☐ to be informed ☐ to NOT be informed Genetic variation may sometimes be identified, which do	es not fit within the scope of the
For more detailed information on data www.cegat.com/privacy-policy.	a privacy as well as your rights please refer to	requested genetic analysis (so-called secondary finding ants is limited to pathogenic alterations (ACMG classes for which a treatment or course of action exists for you	gs). The reporting of these vari- 4 and 5) within selected genes,
diagnostic evaluation is limited to variants in	sequenced when exome diagnostics is performed. The genes relevant to the provided phenotypic information. omparative exome analysis using data from several family	current guidelines of the American College of Medical Ge genes and associated diseases can be found at www.ce no claim of a comprehensive analysis of this gene set. Ar cannot be used to indicate a reduced disease risk.	netics and Genomics; details on gat.com/acmg-genes/). There is
This declaration of consent can be c I have had sufficient time to conside	ompletely or partially withdrawn at any time. r giving my consent.	Targeted analysis of the ACMG genes according to obe requested as "additional analyses".	current recommendations can
above-mentioned patient. For predictive	am authorized to request genetic testing for the e testing, I confirm that I am authorized, and that uest this testing. For minors, I declare that I have		
	form: I, the referring physician, confirm that the d agrees with the genetic testing. The patient's		
		Physican's stamp / Barcode	DAKKS Deutsche Akkreditierungsstelle D-ML-13206-01-00
Patient (Block letters)	Physican (Block letters)		ACCREDITED COLLEGE of AMERICAN PATHOLOGISTS CLIA CERTIFIED ID: 99D2130225
Y	Y		CeGaT is accredited by
Patient (Date, Signature)	Physican (Date, Signature)		DAkkS according to DIN EN ISO 15189:2014, the College of American Pathologists (CAP) and CLIA.



Declaration of consent Parent 2

Personal data (Family membe	r)		
Surname:	First name:		
Date of birth:	Sample ID:		
Relationship to the patient			
☐ Father ☐ Mother	☐ Other; please state:		
Does the family member suffer t	rom an illness or disorder with (suspected)		
,	mptoms are:	•	
	,		
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I would like to receive an addit	ional report analyzing known variants in 22 ge	nes that are involved in the metabolism of pharmaceu	utical products.
Declaration of consent		If you do not check these boxes, your answer will be	
the genetic background related to the di	re received comprehensive information regarding sease in question, as well as the possibilities and I understand that I have the right to withdraw my	I consent to the storage of my genetic material for additional and/or quality control (for max. 10 years). I consent to the storage of my test results beyond the time.	☐ Yes ☐ No
	my personal data and the data obtained in the	10 years (as required by German law).	☐ Yes ☐ No
tabases, and that further, in accordance	stored in an pseudonymized form in scientific da- e with data protection and medical confidentiality, insmitted to a specialized cooperating laboratory.	I consent to the pseudonymous storage and use of surplu material and/or test results for scientific research and in so literature.	s genetic cientific
I consent to the re-evaluation of my tes cant alterations become apparent, my p	t results within the data storage period. If signifi- physican will be informed by email.	With regard to secondary findings I would like:	□ Yes □ No
I have been informed, and agree to the ession of all data collected by CeGaT Gm	electronic storage, processing, use, and transmisbH.	☐ to be informed ☐ to NOT be informed Genetic variation may sometimes be identified, which do	es not fit within the scope of the
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diagnostic evaluation is limited to variants in	sequenced when exome diagnostics is performed. The genes relevant to the provided phenotypic information. omparative exome analysis using data from several family	current guidelines of the American College of Medical Ge genes and associated diseases can be found at www.ce ; no claim of a comprehensive analysis of this gene set. Ar cannot be used to indicate a reduced disease risk.	<u>gat.com/acmg-genes/)</u> . There is
This declaration of consent can be c I have had sufficient time to conside	ompletely or partially withdrawn at any time. r giving my consent.	Targeted analysis of the ACMG genes according to c be requested as "additional analyses".	urrent recommendations can
above-mentioned patient. For predictive	am authorized to request genetic testing for the e testing, I confirm that I am authorized, and that uest this testing. For minors, I declare that I have		
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