

General Information

Patient	
Surname:	_____
First name:	_____
Date of birth:	_____
Sex:	<input type="checkbox"/> male <input type="checkbox"/> female
External ID:	_____

Declaration of consent

By signing this form, I declare that I have received comprehensive information about the genetic background related to the disease in question, as well as the possibilities and limitations of molecular genetic testing. I understand that I have the right to withdraw my consent to genetic analyses.

I have been informed, and agree, that my personal data and the data obtained in the analysis will be recorded, evaluated or stored in an pseudonymised form in scientific databases, and further, in accordance with data protection and medical confidentiality, that the request, or parts thereof, may be transmitted to a specialized cooperating laboratory.

I consent to the re-evaluation of my test results within the data storage period. If significant alterations become apparent, my doctor will be informed by e-mail.

I have been informed, and agree, that all data collected by CeGaT GmbH is electronically stored, processed, used and transmitted.

For more detailed information on data privacy as well as your rights please refer to www.cegat.de/en/privacy-policy

Please Note

Our panels are regularly updated to reflect current scientific research. It should therefore be recognized that there is the possibility that the list of genes on the order form may have changed slightly (genes added or removed) by the time the sample is analyzed in the laboratory. By signing this form, the physician accepts that the list of genes actually analyzed may be slightly different from what is currently listed. When NGS is utilized more than the requested genes are sequenced for each sample.

This consent includes the permission to request tumor sample materials and reports from external sources.

This declaration of consent can be completely or partially withdrawn at any time. I have had sufficient time to consider giving my consent.

I, the referring physician, confirm that I am qualified to request genetic testing for the above-mentioned patient. For minors, I declare that I have the consent of all legal guardians.

If the patient did not sign this order form: I, the referring physician, confirm that the patient received genetic counseling and agrees with the genetic testing. The patient's consent has been obtained in writing.

Sender / Clinic	
Surname:	_____
First name:	_____
Institution:	_____
Street:	_____
Postcode/City:	_____
Country:	_____
Phone:	_____
Email:	_____
VAT:	_____
If applicable, please include a VAT number or a copy of your business registration certificate.	
Invoice	<input type="checkbox"/> to sender / clinic <input type="checkbox"/> to patient / other (KVA-No.: _____)
Surname:	_____
First name:	_____
Street:	_____
Postcode/City:	_____
Country:	_____
Email:	_____

If you do not check these boxes, your answer will be recorded as "No".

I consent to the storage of my genetic material for additional tests and/or quality control (for max. 10 years). Yes No

I consent to the storage of my test results beyond the timespan of 10 years (as required by German law). Yes No

I consent to the pseudonymous storage and use of surplus genetic material and/or test results for scientific research and in scientific literature. Yes No

With regard to secondary findings I would like to be informed: Yes No

Genetic variation may sometimes be identified, which does not fit within the scope of the requested genetic analysis (so-called secondary findings). The reporting of these variants is limited to pathogenic alterations (ACMG classes 4 and 5) within selected genes, for which a treatment or course of action exists for you or your family (according to the current guidelines of the American College of Medical Genetics and Genomics; details on genes and associated diseases can be found at <https://www.cegat.com/acmg-genes/>). There is no claim of a comprehensive analysis of this gene set. An absence of secondary findings cannot be used to indicate a reduced disease risk.

As part of this analysis we also examine germline changes present in leukocyte DNA. Even there is no known family history, it is possible that a clinically relevant germline variant is detected. This may be of relevance for the therapy, but possibly also for tumor follow-up, prevention or for at-risk family members. Therefore, we generally report clinically relevant germline variants (variants with therapeutic relevance or pathogenic/likely pathogenic variants only) in selected genes, unless explicitly contradicted. The results should be discussed as part of a genetic counseling.

According to German Genetic Diagnostic Act (GenDG) we will issue the medical report to the counselling physician. Please indicate here the contact email of the counselling physician:

Email: _____

_____	_____
Patient / Legal Guardian (Block letters)	Doctor (Surname, First name)
X _____	X _____
Patient / Legal Guardian (Date, Signature)	Doctor (Date, Signature)

Doctor's stamp / Barcode



CeGaT is accredited by DAkkS according to DIN EN ISO 15189:2014, the College of American Pathologists (CAP) and CLIA.

Indication

For targeted and effective processing, please complete the medical history form with as much detail as possible and include a copy of all existing reports.

Indication / Suspected diagnosis / Course of disease / Pedigree

 index patient

not affected

affected

known carrier

deceased

 unrelated parents

 consanguine parents

 unborn child

 abortion, stillborn child

 person of unknown sex

 identical twins (monozygous)

 fraternal twins (dizygous)

Already initiated / carried out somatic genetic analyses

Clinical report(s) added

Laboratory report(s) of Pathology / Cytology / Cytogenetics / Flow Cytometry added

Transplants (bone marrow, tissue, stem cells) No Yes, (please specify) _____

Sample material: Liquid biopsy (cfDNA)

Liquid Biopsy samples are specimens that can only be withdrawn using special collection tubes that stabilize the cell-free DNA. If you are planning a diagnostic examination based on cfDNA, please use such collection tubes. We gladly provide such special collection tubes. Please contact us in time at tumor@cegat.de to order the tubes.

3x 10ml cfDNA Tubes

Inquiry

All relevant variants in a named exon are analysed. Exon numbers refer to coding exons (CDS) of the respective gene. The diagnostic is not limited to the listed example hotspot mutations. Exons not named and all variants within are not part of the analysis.

Gene	NM_Nr.	Enriched region (incl. example hotspot (HS)-variants)	Gene	NM_Nr.	Enriched region (incl. example hotspot (HS)-variants)
AKT1	NM_005163	Exon 2 (HS E17)	IDH2	NM_002168	Exon 4 (HS R140, R172)
ALK	NM_004304	Exons 21-25 (incl. HS F1174)	JAK2	NM_004972	Exon 12 (HS V617)
ARAF	NM_001654	Exon 6 (HS S214)	KIT	NM_000222	Exons 9, 11, 13, 14, 17, 18 (incl. HS W557_K558del, D816)
BRAF	NM_004333	Exons 11 and 15 (incl. HS V600)	KRAS	NM_004985	Exons 1-3 (incl. HS G12, Q61)
CTNNB1	NM_001904	Exon 2 (incl. HS S37, S45)	MAP2K1	NM_002755	Exon 3 (HS P124)
EGFR	NM_005228	Exons 18-21 (incl. HS E746_A750del, T790, L858)	MET	NM_001127500	Exon 18 (incl. HS Y1248, Y1253)
ERBB2	NM_004448	Exon 8, 19-21 (incl. HS V842)	MYCN	NM_005378	Exon 1 (HS P44)
ERBB3	NM_001982	Exons 3, 6-9, 23 (incl. HS V104, E928)	NRAS	NM_002524	Exons 1-3 (incl. HS G12, Q61)
ERBB4	NM_005235	Exon 12 (incl. HS E452)	PDGFRA	NM_006206	Exons 4, 9, 11, 13, 17 (incl. HS D842)
ESR1	NM_000125	Exons 4-8 (incl. HS K303, Y537, D538)	PIK3CA	NM_006218	Exons 4, 7, 9, 13, 20 (incl. HS E542, E545, H1047)
FGFR2	NM_000141	Exons 6, 8, 11 (incl. HS S252, N549)	PTEN	NM_000314	Exons 5-7 (incl. R130, R233)
FGFR3	NM_000142	Exon 12 (HS V555)	RAC1	NM_018890	Exon 2 (HS P29)
GNA11	NM_002067	Exon 5 (HS Q209)	RAF1	NM_002880	Exon 6 (incl. HS S257, S259)
GNAQ	NM_002072	Exon 5 (HS Q209)	RET	NM_020975	Exon 10, 11, 13-16 (incl. HS C634)
GNAS	NM_000516	Exon 8 (HS 201) and Exon 9 (HS Q227)	STAT5B	NM_012448	Exon 15 (HS N642)
H3-3A	NM_002107	Exon 1 (HS K27 and G34)	TERT	NM_198253	Promotor HS c.-124 (C228), c.-146 (C250)
H3-3B	NM_005324	Exon 1 (HS K37)	TP53	NM_000546	Entire coding region
HRAS	NM_005343	Exons 1-3 (incl. HS G12, Q61)			
IDH1	NM_005896	Exon 2 (HS R132)			

Remarks:

For further information and advice please do not hesitate to contact our Diagnostic Support team.
www.cegat.de/en/diagnostic-support · diagnostic-support@cegat.de
 Phone +49 7071 56544-55