



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:

Ultrasound medical report available? ☐ Yes (please attach copy) ☐ No

Clinical symptoms:

Preliminary genetic diagnostics for fetus or parents?

Chromosome / Karyotype analysis: ☐ Yes (please attach copy) ☐ No

Array-CGH: ☐ Yes (please attach copy) ☐ No

Other

Pedigree

Consanguinity: ☐ Yes ☐ No

Ethnic origin:

☐ 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)

Family medical history

If the mother of the fetus has been pregnant in the past, were there any anomalies during pregnancy? ☐ Yes ☐ No

Are there other family members who currently have or have had a disease or disorder relevant for the clinical indication of the fetus? ☐ Yes ☐ No

If yes, please list the affected family members:

Name (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) allows 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)**

- ☐ Please perform array-CGH diagnostics
- ☐ prior **or**
- ☐ parallel
- to exome diagnostics.

**For further information and advice please do not hesitate to contact our Diagnostic Support team.**

**www.cegat.com/diagnostic-support**  
**diagnostic-support@cegat.com**  
**Phone +49 7071 565 44-55**

Please use this space to provide any additional relevant information.

# Order Form Prenatal ExomeXtra® - Conspicuous Ultrasound



## Declaration of consent Parent 1

### Personal data (Family member)

Surname: \_\_\_\_\_ First name: \_\_\_\_\_

Date of birth: \_\_\_\_\_ Sample ID: \_\_\_\_\_

### Relationship to the patient

☐ Father ☐ Mother ☐ Other; please state: \_\_\_\_\_

Does the family member suffer from an illness or disorder with (suspected) genetic cause?

☐ No ☐ Yes, symptoms are: \_\_\_\_\_

### ☐ ACMG genes diagnostics

I would like to be informed of relevant alterations within the list of recommended genes for secondary analysis, according to the current guidelines of the American College of Medical Genetics and Genomics. The analysis is restricted to the sequence data, re-sequencing of regions with poor sequence coverage will not typically be performed. A negative "ACMG genes" report cannot be used to rule out (genetic) disease risk. Additional fees may apply. According to German legislation, predictive tests for minors may not be performed for diseases which have an onset in adulthood. Therefore, some genes will not be analyzed for minors, unless the phenotypic spectrum is within the scope of the primary medical indication of the patient.

Details on genes and associated diseases can be found at [www.cegat.com/acmg-genes/](http://www.cegat.com/acmg-genes/).

### ☐ Pharmacogenetics (PGX) (22 genes)

ABCG2, CACNA1S, CYP2B6, CYP2C19, CYP2C9, CYP2D6, CYP3A4, CYP3A5, CYP4F2, DPYD, G6PD, HLA-A, HLA-B, IFNL3, MT-RNR1, NUDT15, POR, RYR1, SLC01B1, TPMT, UGT1A1, VKORC1

I would like to receive an additional report analyzing known variants in 22 genes that are involved in the metabolism of pharmaceutical products.

### Declaration of consent

By signing this form, I declare that I have received comprehensive information regarding 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 for 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 pseudonymized form in scientific databases, and that further, in accordance with data protection and medical confidentiality, 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 physician will be informed by email.

I have been informed, and agree to the electronic storage, processing, use, and transmission of all data collected by CeGaT GmbH.

For more detailed information on data privacy as well as your rights please refer to [www.cegat.com/privacy-policy](http://www.cegat.com/privacy-policy).

### Please Note

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 phenotypic information. Correct family relationships are assumed for comparative exome analysis using data from several family members (e.g., trio exome analysis).

**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 authorized to request genetic testing for the above-mentioned patient. For predictive testing, I confirm that I am authorized, and that I have fulfilled the requirements, to request this testing. 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.

**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 ☐ to NOT be informed

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 [www.cegat.com/acmg-genes/](http://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.

**Targeted analysis of the ACMG genes according to current recommendations can be requested as "additional analyses".**

Physician's stamp / Barcode



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

Patient  
(Block letters)

Physician  
(Block letters)

X  
Patient  
(Date, Signature)

X  
Physician  
(Date, Signature)

# Order Form Prenatal ExomeXtra® - Conspicuous Ultrasound



## Declaration of consent Parent 2

### Personal data (Family member)

Surname: \_\_\_\_\_ First name: \_\_\_\_\_

Date of birth: \_\_\_\_\_ Sample ID: \_\_\_\_\_

### Relationship to the patient

☐ Father ☐ Mother ☐ Other; please state: \_\_\_\_\_

Does the family member suffer from an illness or disorder with (suspected) genetic cause?

☐ No ☐ Yes, symptoms are: \_\_\_\_\_

### ☐ ACMG genes diagnostics

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(Date, Signature)

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