labdia labordiagnostik
St. Anna Forschungsinstitut

REFERRAL FORM
PAEDIATRIC SOLID TUMOURS
DEPARTMENT FOR TUMOUR BIOLOGY

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Area for stamps

Referring provider (please use **stamp,** alternatively blockletters)

Name/Clinic: .....
Department: .....
City/Country: .....
Phone: .....

Invoice address:
☐ Referring provider ☐ Patient/Other (declaration of cost coverage)

Referring clinician

Name: .....
Phone: .....
E-Mail: .....

Required patient information (please use **label**, alternatively blockletters)

Last name: .....
First name: .....
Gender: ☐ F ☐ M Insurance number/Date of birth: ..... / dd mm yyyy
Study: ..... Patient Study ID: .....
EUPID: .....

Reason for referral / Indication

☐ Suspected diagnosis
☐ Initial diagnosis
☐ During treatment
☐ Relapse

☐ Neuroblastoma ☐ Nephroblastoma ☐ Germ cell tumor
☐ Ewing sarcoma ☐ Rhabdomyosarcoma ☐ Osteosarcoma
☐ .....

Remarks: .....

Sample

Date sample taken: dd mm yyyy
Time sample taken: hh : mm

Remarks: .....

☐ Tissue

☐ **Primary tumour** ☐ **Metastasis**

☐ Native
☐ Frozen
☐ Touch imprints
☐ Paraffin material (FFPE)

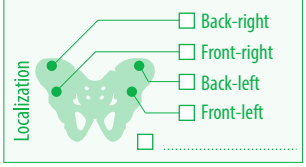
Histology No.

Tumour cell content %

Localization / Remarks (e.g. when several tumour pieces are available):

☐ Bone marrow

☐ EDTA
☐ Cytospin slides
☐ PAXgene *(to be requested)*



☐ .....

☐ Peripheral blood

☐ EDTA
☐ PAXgene *(to be requested)*
☐ .....

☐ Apheresis product*

☐ EDTA
☐ .....

**Please also provide peripheral blood before and after apheresis*

☐ Body fluid

☐ Ascites
☐ CSF
☐ Pleural effusion

☐ Other samples

☐ .....
.....

The primary sample collection manual can be found on https://www.labdia.at.

Moleculargenetic and immunocytologic testing

☐ Neuroblastoma

☐ *MYCN* amplification (FISH)
☐ *ALK* amplification / rearrangement (Break apart FISH)
☐ *ALK* mutations (ddPCR, sequencing)
☐ Segmental und numerical chromosomal aberrations (SCAs & NCAs) (SNP-Array)
☐ .....

☐ Bone marrow infiltration / Minimal residual disease (MRD) (GD2/CD56 staining)

☐ Rhabdomyosarcoma

☐ *FOXO1 (FKHR)* rearrangement / fusion with:
  ☐ *PAX3* (FISH, qPCR)
  ☐ *PAX7* (FISH, qPCR)
☐ Rare *FOXO1* oder *PAX3* rearrangements (Break apart FISH)
☐ Genome-wide profile (CN/LOH) (SNP-Array)
☐ .....

☐ Bone marrow infiltration / Minimal residual disease (MRD) (GD2/CD56 staining)

☐ Ewing Sarcoma

☐ *EWSR1* rearrangement / fusion with:
  ☐ *FLI1* (FISH, qPCR)
  ☐ *ERG* (FISH, qPCR)
☐ Rare *EWSR1* rearrangements (Break apart FISH)
☐ .....

☐ Bone marrow infiltration / Minimal residual disease (MRD) (qPCR)
information on the fusion incl. exons:
.....

☐ Nephroblastoma (Wilms-Tumor)

☐ 1q gain (FISH)
☐ Genome-wide profile (CN/LOH) (SNP-Array)
☐ .....

☐ Other tests

☐ .....
☐ .....
☐ .....

Asservation

☐ asservation for now, feedback on desired testing follows
☐ only for asservation in biobank / for a study:
Study: .....