

SEND TO

Department of Pathology UMC Utrecht
Attn Molecular Pathology
Roomnumber H04.312
Heidelberglaan 100
3584 CX Utrecht
The Netherlands

Patient information

Name + initials:

Adress:

Date of birth:

Gender:

Citizen Service Number (BSN):



UMC Utrecht

Patient administration

Tel +31 (0)88 7557615
Fax +31 (0)88 7569588
E-mail administratie-pathologie@umcutrecht.nl

Molecular Pathology

Tel +31 (0)88 7574252
E-mail pathology-moleculardiagnosics@umcutrecht.nl

PATHOLOGY UMC UTRECHT:

Requesting laboratory and/or physician

Name : Date :
Department (code) : External PA- and/or reference number :
Institution/country :

Material

H&E stained slide
FFPE block :
Fresh frozen* :

*Tick the box in case of fresh frozen material for PMC:

*Tick the box in case of fresh frozen material for PMC **and** blood for PMC (FFPE will follow):

DNA :
Other :
Slides** :

**In case of slides: for non-FISH applications: 10 unstained sections 4µm (coated slides)
for FISH applications: 6 unstained sections 4µm (coated slides)

REMARKS:

NGS

- BRCA 1/2 (somatic)
 CLL (chronic lymphatic leukemia) TP53
 Colon BRAF/RAS
 Desmoid tumor CTNNB1
 TERT Glioma panel
 GIST KIT/PDGFRA/BRAF
 Glioma e.g. IDH1/IDH2/1p19q codeletion
 Hematological diseases e.g. SF3B1/SRSF2/U2AF1/TET2/SXL1/EZH2
 LPL (lymphoplasmacytic lymphoma) MYD88
 Melanoma BRAF/HRAS/NRAS/KIT
 Melanocytic lesions APC/BRAF/CTNNB1/GNA11/GNAQ/HRAS/IDH1/KIT/NRAS/TERT
 MPN (myeloproliferative neoplasms) CALR/MPL/JAK2
 Kidney panel (PMC) TP53/WT1/WTX/FBXW7/SMARCB1/SMARCA4
 Thyroid carcinoma BRAF/RAS
 TP53/Tumorclonality TP53
 Other Indicate genes of interest at remarks

Lung:

- Adenocarcinoma*/non-small cell lung cancer (NSCLC)** KRAS/EGFR/BRAF/HER2
 Squamous cell carcinoma*** KRAS/FGFR1 (mut + amp)
 cMET exon 14 skipping
 EGFR TKI resistance EGFR/HER2 (mut)/cMET (amp)
 ALK/ROS1 inhibitors ALK (mut)

Predictive IHC

- PD-L1 (lung) PD-L1 (bladder)
 PD-L1 (breast) Pan-NTRK

RT-PCR

- KIAA-BRAF duplicatie 7q34
 YWHAE-FAM22A/B t(10;17)
 EWSR1 FL1/ERG/WT1
 FGFR3-TACC3 duplicatie 4p16
- Other molecular analyses**
- Archer FusionPlex Lung
 BRAF Idylla**** V600E/D + V600K/R/M
 HPV m Allinity (RvA: M 208) High risk****
 HPV genotyping assay High risk + low risk****
 B-cel clonality
 T-cel clonality
 Tissue-identification
 Neuroblastoma SNP: 1p/17q/ALK, FISH: N-MYC, NGS: ALK
 PNET V medulloblastoma IHC: CTNNB1 + p53, FISH: cMYC/N-MYC/CEN6, NGS
- ddPCR**
- BRAF p.(V600E)
 EGFR exon 19 deletions
 EGFR p.(L858R)
 EGFR resistance p.(T790M)
 MYD88 p.(L265P)

Arrays

- SNP array e.g. Wilms tumor, clonality, etc.
 Methylation profiling Classification of CNS tumors

Please turn over for FISH, MSI, MLPA,
DNA-isolation and Chimerism

Fluorescence In Situ Hybridisation

Soft tissue

- | | |
|---------------------------------------|---------------------------|
| <input type="checkbox"/> CHOP | Break-apart |
| <input type="checkbox"/> EWSR1 | Break-apart + fusion FLI1 |
| <input type="checkbox"/> FKHR (FOXO1) | Break-apart |
| <input type="checkbox"/> FUS | Break-apart |
| <input type="checkbox"/> MDM2 | Break-apart |
| <input type="checkbox"/> SYT | Break-apart |

Lymphoma

- | | |
|--|--------------------------|
| <input type="checkbox"/> BCL-2 | Break-apart |
| <input type="checkbox"/> BCL6 | Break-apart |
| <input type="checkbox"/> cMYC | Break-apart + fusion IgH |
| <input type="checkbox"/> CCND1 (Cyclin-D1) | Break-apart |
| <input type="checkbox"/> MALT1 | Break-apart |

Chromosomes

- Centromere X/Y/18
 Centromere 13/18/21

Other

- | | |
|--|--------------------------------------|
| <input type="checkbox"/> ALK | Break-apart |
| <input type="checkbox"/> BCOR | Break-apart |
| <input type="checkbox"/> cMET | Amplification |
| <input type="checkbox"/> COL1A1-PDGFB | Fusion |
| <input type="checkbox"/> ERBB2 (Her2neu) | Amplification |
| <input type="checkbox"/> ETV6 | Break-apart |
| <input type="checkbox"/> HMGA2 | Break-apart |
| <input type="checkbox"/> MAML2 | Break-apart |
| <input type="checkbox"/> MUM1 (IRF4) | Break-apart |
| <input type="checkbox"/> MYB | Break-apart |
| <input type="checkbox"/> N-MYC | Amplification |
| <input type="checkbox"/> NRG1 | Break-apart |
| <input type="checkbox"/> NTRK 1/2/3 | Break-apart |
| <input type="checkbox"/> PLAG1 | Break-apart |
| <input type="checkbox"/> PLAG1/CTNNB1 | Fusion |
| <input type="checkbox"/> RET | Break-apart |
| <input type="checkbox"/> ROS1 | Break-apart |
| <input type="checkbox"/> TFE3 | Break-apart |
| <input type="checkbox"/> USP6 | Break-apart |
| <input type="checkbox"/> Research ISH | Indicate gene of interest at remarks |

MSI for Lynch syndrome (tumor and normal)
(IHC MMR proteins + Idylla)

- MSI Lynch (via Idylla)
 MLH1 hypermethylation and BRAF
 Mutation analysis (only V600E) for MSI

MSI for therapeutic purposes
(IHC MMR proteins)

- Only tumor tissue

DNA-isolation

- DNA-isolation tumor tissue
 DNA-isolation normal tissue
 DNA-isolation other, e.g. blood

MLPA

- MLPA 1p19q codeletion
 MLPA FGFR
 MLPA Her2neu
 MLPA MDM2/CDK4
 MLPA Trisomy 13/18/21/X/Y
 MLPA Wilms tumors

- MS-MLPA BRCA1 hypermethylation
 MS-MLPA MGMT promoter methylation

Chimerism

- Whole blood
 T-/non-T

* For lung applications: also send a distinctive staining, e.g. TTF1, together with your application

** If no mutations are found in KRAS and/or EGFR for adenocarcinoma/NSCLC, additional translocation analysis will be performed (according to agreement)

*** If no mutations are found in KRAS and/or FGFR1 for squamous cell carcinoma an additional ALK analysis will be performed

**** For all HPV applications:

Severe dysplasia/CIS/invasive: apply for HPV m Allinity

Low/moderate dysplasia/verruccous: apply for HPV genotyping assay

***** For BRAF Idylla please send a block or a 10µm tissue slide in a tube with a representative H&E staining for the appropriate tumor percentage (at least 20% is necessary)