

AOK	LKK	BKK	IKK	VdAK	AEV	Knappschaft	Privat	Re. an K-Haus
Name, Vorname des Versicherten <span style="float: right;">geb. am</span>								
Kassen-Nr.			Versicherten-Nr.			Status		
Vertragsarzt Nr.			VK gültig bis			Datum		

# Antrag auf molekularpathologische Begutachtung

(mit Einverständnis des Patienten)

**Zentrum für Pathologie Allgäu (ZfPA)**  
**Medizinisches Versorgungszentrum am Klinikum Kempten**  
**PD Dr. med. Konrad Aumann und KollegInnen**  
 Postfach 2122, 87411 Kempten      Tel. 0831 530-2180  
 Robert-Weixler-Str. 48, 87439 Kempten      Fax 0831 530-2170  
 E-Mail: pathologie-info@klinikverbund-allgaeu.de  
 www.patho-kempten.de  
**Zertifiziert nach DIN EN ISO 9001:2015**

Dieser Anforderungsbogen ersetzt nicht den Überweisungsschein bei Kassenpatienten.

Ambulant       Privat       Stationär

## Klinische Angaben:

## Diagnose (ICD 10):

### Prädiktive und prognostische Analysen (organspezifisch)

- |  |  |   |  |
|--|--|---|--|
| <ul style="list-style-type: none"> <li><input type="radio"/> <b>Lungenkarzinom</b></li> <li><input type="radio"/> ALK-Translokation [t(2q23)]</li> <li><input type="radio"/> RET-Translokation [t(10q11)]</li> <li><input type="radio"/> ROS-Translokation [t(6q22)]</li> <li><input type="radio"/> MET-Amplifikation</li> <li><input type="radio"/> BRAF-Mutationen (Exon 15)</li> <li><input type="radio"/> EGFR-Mutationen (Exon 18, 19, 20, 21)</li> <li><input type="radio"/> EGFR-T790M-Mutationen (Exon 20)</li> <li><input type="radio"/> HER2-Mutationen (Exon 20)</li> <li><input type="radio"/> KRAS-Mutationen (Exon 2, 3, 4)</li> <li><input type="radio"/> PIK3CA-Mutationen (Exon 9, 20)</li> <li><input type="radio"/> MET Ex 14 Skipping Variante</li> <li><input type="radio"/> Keap1-Mutationen (Exon 2-6)</li> <li><input type="radio"/> STK11-Mutationen (Exone 1-9)</li> <li><input type="radio"/> PD-L1 (IHC)</li> <li><input type="radio"/> <b>Kolorektales Karzinom</b></li> <li><input type="radio"/> KRAS-Mutationen (Exon 2, 3, 4)</li> <li><input type="radio"/> NRAS-Mutationen (Exon 2, 3, 4)</li> <li><input type="radio"/> BRAF-Mutationen (Exon 15)</li> <li><input type="radio"/> PIK3CA-Mutationen (Exon 9, 20)</li> <li><input type="radio"/> HER2-Analytik (17q12 [ERBB2])</li> <li><input type="radio"/> PD-L1 (IHC)</li> </ul> | <ul style="list-style-type: none"> <li><input type="radio"/> <b>HNPCC-Diagnostik</b></li> <li><input type="radio"/> MMR-Proteine (IHC: MLH1, MSH2, MSH6, PMS2)</li> <li><input type="radio"/> Mikrosatelliten-Instabilität (MSI Panel)</li> <li><input type="radio"/> BRAF-Mutationen (Exon 15)</li> <li><input type="radio"/> MLH1-Promotor Methylierung*</li> <li><input type="radio"/> <b>Gastrointestinaler Stromatumor / GIST</b></li> <li><input type="radio"/> KIT-Mutationen (Exon 9, 11, 13, 17)</li> <li><input type="radio"/> PDGFRA-Mutationen (Exon 12, 18)</li> <li><input type="radio"/> <b>Malignes Melanom</b></li> <li><input type="radio"/> BRAF-Mutationen (Exon 15)</li> <li><input type="radio"/> KIT-Mutationen (Exon 9, 11, 13, 17)</li> <li><input type="radio"/> NRAS-Mutationen (Exon 2, 3, 4)</li> <li><input type="radio"/> <b>Mammakarzinom</b></li> <li><input type="radio"/> HER2-Analytik 17q12 [ERBB2]</li> <li><input type="radio"/> PIK3CA-Mutationen (Exon 9, 20)</li> <li><input type="radio"/> BRCA1/2</li> <li><input type="radio"/> PD-L1 (IHC)</li> <li><input type="radio"/> <b>Magen-/Ösophaguskarzinom</b></li> <li><input type="radio"/> HER2-Analytik 17q12 [ERBB2]</li> <li><input type="radio"/> EBV (EBER-ISH)</li> <li><input type="radio"/> PD-L1 (IHC)</li> </ul> | <ul style="list-style-type: none"> <li><input type="radio"/> <b>Ovarialkarzinom</b></li> <li><input type="radio"/> BRCA1/2</li> <li><input type="radio"/> HRD-Analyse*</li> <li><input type="radio"/> <b>Prostatakarzinom</b></li> <li><input type="radio"/> BRCA1/2</li> <li><input type="radio"/> ATM-Mutation</li> <li><input type="radio"/> HRD-Analyse*</li> <li><input type="radio"/> <b>Pankreaskarzinom</b></li> <li><input type="radio"/> BRCA1/2</li> <li><input type="radio"/> <b>Kopf-Hals Tumoren</b></li> <li><input type="radio"/> 8p11 [FGFR1] FGFR-Analytik</li> <li><input type="radio"/> EGFR-Mutationen (Exon 18, 19, 20, 21)</li> <li><input type="radio"/> RAS-Mutationen (Exon 2, 3, 4)</li> <li><input type="radio"/> PIK3CA-Mutationen (Exon 9, 20)</li> <li><input type="radio"/> EBV (EBER-ISH)</li> <li><input type="radio"/> PD-L1 (IHC)</li> <li><input type="radio"/> <b>Schilddrüsenkarzinom</b></li> <li><input type="radio"/> t (10q11) [RET]</li> <li><input type="radio"/> RET-Mutation</li> <li><input type="radio"/> BRAF-Mutationen (Exon 15)</li> </ul> | <ul style="list-style-type: none"> <li><input type="radio"/> <b>Urothelkarzinom</b></li> <li><input type="radio"/> FGFR1/2/3-Fusionen</li> <li><input type="radio"/> FGFR1/2/3-Mutationen</li> <li><input type="radio"/> HER2-Analyse</li> <li><input type="radio"/> PD-L1 (IHC)</li> <li><input type="radio"/> <b>Cholangiokarzinom</b></li> <li><input type="radio"/> FGFR1/2/3-Fusionen</li> <li><input type="radio"/> IDH1/2-Mutationen</li> <li><input type="radio"/> BRAF-Mutationen</li> <li><input type="radio"/> KRAS-Mutationen</li> <li><input type="radio"/> BRCA1/2-Mutationen</li> <li><input type="radio"/> PD-L1 (IHC)</li> <li><input type="radio"/> <b>Endometriumkarzinom</b></li> <li><input type="radio"/> MSI-Diagnostik</li> <li><input type="radio"/> P53 (IHC)</li> <li><input type="radio"/> POLE-Mutationsanalyse*</li> </ul> |
|--|--|---|--|

*\*in Kooperation mit externen Laboren*

### Somatische Mutations-/Fusionsanalytik (NGS Panel-Diagnostik)

- Illumina Cancer Hotspot Panel v2**  
 ABL1, AKT1, ALK, APC, ATM, BRAF, CDH1, CDKN2A, CSF1R, CTNNB1, EGFR, ERBB2, ERBB4, EZH2, FBXW7, FGFR1, FGFR2, FGFR3, FLT3, GNA11, GNAQ, GNAS, HNF1A, HRAS, IDH1, IDH2, JAK2, JAK3, KDR, KIT, KRAS, MET, MLH1, MPL, NOTCH1, NPM1, NRAS, PDGFRA, PIK3CA, PTEN, PTPN11, RB1, RET, SMAD4, SMARCB1, SMO, SRC, STK11, TP53, VHL
- QIaseq Targeted DNA Panel (nNGM)**  
 ALK, BRAF, CTNNB1, EGFR, ERBB2, IDH1, IDH2, FGFR1, FGFR2, FGFR3, FGFR4, KRAS, MAP2K1, MET, NRAS, PIK3CA, PTEN, TP53, ROS1, NTRK1, NTRK2, NTRK3, RET, HRAS, STK11, KEAP1
- RNA-Fusionsanalyse (QIaseq Targeted RNAscan Panel (nNGM))**  
 ALK, MET, BAG4, MPRIIP, BRAF, MRPS14, CCDC6, NRG1, CD74, NTRK1, CIAO1, NTRK2, COPA, NTRK3, CUX1, RAD51, EGFR, RET, EML4, ROS1, ETV6, SDC4, EZR, SLC34A2, FGFR1, STRN, FGFR2, TACC3, FGFR3, TFG, GOPC, TPM3, HIP1, TPR, KIF5B, TRIM33, KLC1, UBE3C, LRIG

### In-situ-Hybridisierungen zur Evaluation von Genrearrangements / Sonstiges:

- Sonde (Beispiel)
- |  |  |  |
|--|--|--|
| <input type="radio"/> MDM2 [12q15] (Liposarkom)        | <input type="radio"/> HER2/CEP17 [17q11] (Mamma)           | <input type="radio"/> 19q13/19p13 (Gliome)                   |
| <input type="radio"/> DDIT3 bap [12q13] (Liposarkom)   | <input type="radio"/> SS18 bap [18q11] (synoviales Sarkom) | <input type="radio"/> 1p36/1q25 (Gliome)                     |
| <input type="radio"/> FUS bap [16p11] (Liposarkom)     | <input type="radio"/> MET-Amplifikation (Lungenkarzinom)   | <input type="radio"/> EBV EBER-ISH (Magenkarzinom, Lymphome) |
| <input type="radio"/> EWSR1 bap [22q12] (Ewing Sarkom) | <input type="radio"/> CCND1-bap [11q13] (Mantelzellymphom) |  |

Sonstiges: \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_  
 \_\_\_\_\_

Stempel Arzt / Klinik
Unterschrift