

Pat.-Name, Pat.-Vorname <hr/> Adresse (bei Privatversicherten) <hr/> <div style="text-align: right;"><input type="checkbox"/> w <input type="checkbox"/> m</div> Geburtsdatum: <hr/> Ihre Eingangsnummer: <hr/>	Versicherungsstatus <input type="checkbox"/> KV (Bitte Ü-Schein!) <input type="checkbox"/> Privat (Bitte Adresse angeben!) <input type="checkbox"/> Privat stationär <input type="checkbox"/> stationär (Bitte Kostenträgerangabe!) <hr/>	Absender/Praxisstempel <hr/>
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Organ <hr/> Diagnose <hr/>	Pathologie Göttingen Tumorzellgehalt in %: (Markiertes Areal) B-Zellgehalt in %: T-Zellgehalt in %: <hr/>	Pathologie Göttingen HZ: <hr/> Eingang <hr/>	Barcode Pathologie Göttingen <hr/>
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EILT

Lunge:

<input type="checkbox"/> Basispanel (AdenoCa.): BRAF*, EGFR*; ALK, ROS1	<input type="checkbox"/> NGS LUNG Panel*	<input type="checkbox"/> Archer CTL (RNA)	<input type="checkbox"/> PEC-Panel: BRAF*; MET/Cen7
<input type="checkbox"/> EGFR (Ex. 18, 19, 20, 21)*	<input type="checkbox"/> KRAS (Ex. 2)*	<input type="checkbox"/> BRAF (Ex. 15)*	<input type="checkbox"/> MET (Ex. 14 skip.)*
<input type="checkbox"/> ALK/EML4 Inversion	<input type="checkbox"/> AXL/19p13 Amp.	<input type="checkbox"/> ERBB2/Cen17 Amp.	<input type="checkbox"/> FGFR1/Cen8 Amp
<input type="checkbox"/> KRAS/Cen12 Amp.	<input type="checkbox"/> MET/Cen7 Amp.	<input type="checkbox"/> NRG1 Trans.	<input type="checkbox"/> KIF5B Trans.
<input type="checkbox"/> RET Trans.	<input type="checkbox"/> ROS1 Trans.	<input type="checkbox"/> NTRK1 Trans.	<input type="checkbox"/> NTRK2 Trans.
		<input type="checkbox"/> NTRK3 Trans.	

Darm:

<input type="checkbox"/> Darmpanel: KRAS*, NRAS*	<input type="checkbox"/> KRAS (Ex. 2, 3, 4)*	<input type="checkbox"/> NRAS (Ex. 2, 3, 4)*	<input type="checkbox"/> BRAF (Ex. 15)*	<input type="checkbox"/> Mikrosatelliteninstabilität (MSI)*	<input type="checkbox"/> MLH1-Methylierung
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Sarkome/GIST:

<input type="checkbox"/> KIT (Ex. 9, 11, 13, 17) (GIST)*	<input type="checkbox"/> PDGFRα (Ex. 12, 14, 18) (GIST)*
<input type="checkbox"/> Resistenz Panel GIST: KIT (Ex. 13, 14, 15, 17)*, PDGFRα (Ex. 18)*	<input type="checkbox"/> CTNNB1 (Ex. 3) (β-Catenin)*
<input type="checkbox"/> DDIT3 (CHOP) Trans. (myxoides LipoS.)	<input type="checkbox"/> ETV6 Trans. (infantiles FibroS.)
<input type="checkbox"/> EWSR1 Trans. (Ewing-S.)	<input type="checkbox"/> FOXO1 Trans. (RhabdomyoS.)
<input type="checkbox"/> FUS Trans. (myxoides LipoS., low grade FibromyoidS.)	<input type="checkbox"/> JAZF1 Trans. (endometriales StromaS.)
<input type="checkbox"/> MDM2/Cen12 Amp. (LipoS.)	<input type="checkbox"/> MYC/Cen8 Amp. (AngioS.)
<input type="checkbox"/> NR4A3 Trans. (extraskellettales myxoides ChondroS.)	<input type="checkbox"/> RMSI (PAX3/FOXO1) Fusion (RhabdomyoS.)
<input type="checkbox"/> RMSII (PAX7/FOXO1) Fusion (RhabdomyoS.)	<input type="checkbox"/> PDGFB Trans. (Dermatofibrosarcoma protuberans)
<input type="checkbox"/> SS18 (SYT) Trans. (SynovialS.)	<input type="checkbox"/> TFE3 Trans. (alveoläres S.)
<input type="checkbox"/> USP6 Trans. (noduläre Faszitis/ AKZ)	<input type="checkbox"/> WT1 Trans. (DSRCT)
<input type="checkbox"/> WWTR1 Trans. (epitheloides Hämangioendothelium)	<input type="checkbox"/> YWHAE Trans. (endometriales StromaS.)

Lymphom:

<input type="checkbox"/> JH (B-Zell-Klonalität)*	<input type="checkbox"/> γTCR (T-Zell-Klonalität)*	<input type="checkbox"/> MYD88 (Ex. 5)*	<input type="checkbox"/> EBER In-situ-Hybridisierung (EBV)
<input type="checkbox"/> BCL-2 Trans. (follikuläres L.)	<input type="checkbox"/> BCL-6 Trans. (follikuläres L.)	<input type="checkbox"/> CCND1 Trans. (Mantelzell-L.)	
<input type="checkbox"/> IGH Trans. (B-Zell-L.)	<input type="checkbox"/> MALT1 Trans. (Marginalzonen-L.)	<input type="checkbox"/> MYC Trans. (Burkitt-L., diffus großzell. B-Zell-L.)	

Hämatologische Erkrankungen:

<input type="checkbox"/> JAK2 (V617F; MPS)*	<input type="checkbox"/> BCR/ABL1 Fusion (CML)	<input type="checkbox"/> 4q12 Rearrangement (FIP1L1/PDGFRα) (HES/ CEL)
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Mamma-/Magenkarzinom:

<input type="checkbox"/> ERBB2/Cen17 Amp.	<input type="checkbox"/> FGFR2/Cen10 Amp. (Magen)
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Gliome:

<input type="checkbox"/> 1p36/1q25 Del. und 19q13/19p13 Del.	<input type="checkbox"/> MGMT-Methylierung*	<input type="checkbox"/> IDH1 (Ex. 4)*	<input type="checkbox"/> IDH2 (Ex. 4)*
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Melanom:

<input type="checkbox"/> Melanompanel: BRAF*, NRAS*, KIT*	<input type="checkbox"/> BRAF (Ex. 15)*	<input type="checkbox"/> NRAS (Ex. 2, 3, 4)*	<input type="checkbox"/> KIT (Ex. 9, 11, 13, 17)*	<input type="checkbox"/> GNAQ (Ex. 5)*	<input type="checkbox"/> GNA11 (Ex. 5)*
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weitere solide Tumoren:

<input type="checkbox"/> NGS BRCA1/2 (OvarialCa.)*	<input type="checkbox"/> CTNNB1 (Ex. 3) (β-Catenin)*	<input type="checkbox"/> FGFR3 (Ex. 10) (UrothelCa.)*	<input type="checkbox"/> PIK3CA (Ex. 9, 20)*
<input type="checkbox"/> CDKN2A (p16)/Cen9 Del. (Mesotheliom)	<input type="checkbox"/> FGFR1 Trans.	<input type="checkbox"/> FGFR3/4p11 Amp.	<input type="checkbox"/> FGFR3 Trans. (Harnblase)
<input type="checkbox"/> ERG-TMPRSS2 Inversion (ProstataCa.)	<input type="checkbox"/> VHL/Cen3 Del. (Niere)	<input type="checkbox"/> VHL/1p12/Cen7/Cen17 (Niere)	<input type="checkbox"/> MLH1-Methyl. (Endometr.)
<input type="checkbox"/> RB1/13q12 Del. (Retinoblastom)	<input type="checkbox"/> CCND1-BA/2q11/Cen6 (Niere)	<input type="checkbox"/> TFE3 Trans. (NierenzellCa., alveoläres WeichteilS.)	
<input type="checkbox"/> UroVysion CEP 3, 7, 17; LSI 9p21 (Harnblase)	<input type="checkbox"/> NTRK1 Trans.	<input type="checkbox"/> NTRK2 Trans.	<input type="checkbox"/> NTRK3 Trans.

Erregernachweis:

<input type="checkbox"/> HPV-Typisierung	<input type="checkbox"/> Mykobakterien (TBC, atyp. Myko.)	<input type="checkbox"/> EBER In-situ-Hybridisierung (EBV)
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