

## DEPARTMENT OF PATHOLOGY ANATOMIC MOLECULAR DIAGNOSTICS LABORATORY

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PATIENT INFORMATIONINPATIENTOUTPATIENT	PHYSICIAN/FACILITY/CLIENT INFORMATION
<u>.</u>	
Last name First name Middle	Contact Person/Physician ordering test UPIN
DOB Sex SSN	-
	Phone Fax Department
Street Address City State Zip	Address City State Zip
	Address City State Zip
I D. 'I IDM I	CDECIMEN INFORMATION C. II. (* D. )
Insurance Provider ID Number	SPECIMEN INFORMATION: Collection Date:/
	DIAGNOSIS:
Guarantor	
TESTS OFFERED (paraffin o	embedded tissue/cytologic smear)
BREAST CANCER	SOFT TISSUE TUMOR (SARCOMA)
1	Ewing Tumor Family
☐ HER2/neu (ERBB2) amplification by FISH	Ewing Tumor Tumity  EWS gene rearrangement by FISH
LUNG CANCER	Synovial Sarcoma
☐ EGFR mutations (including T790M) by PCR	SYT (SS18) gene rearrangement by FISH
	Alveolar Rhabdomyosarcoma
ALK gene rearrangement by FISH	FOXO1 (FKHR) gene rearrangement by FISH
ROS1 gene rearrangement by FISH	Myxoid/Round Cell Liposarcoma
☐ BRAF V600E mutation by PCR	DDIT3 (CHOP) gene rearrangement by FISH
KRAS mutation by PCR/DNA sequencing	DDITS (CHOF) gene rearrangement by Fish
GOV ON GANGER	LYMPHOMA
COLON CANCER	Mantle cell lymphoma
KRAS mutation by Real-Time PCR/DNA sequencing	CCND1-IGH t(11;14) gene fusion by FISH
NRAS mutation by PCR/DNA sequencing	Diffuse large B-cell lymphoma panel
☐ BRAF V600E mutation by PCR	☐ MYC gene rearrangement by FISH
☐ EGFR amplification by FISH	BCL2-IGH gene rearrangement by FISH
MMR deficiency by IHC (MSI equivalent)	BCL6 gene rearrangement by FISH
	Belo gene rearrangement by 11511
BRAIN TUMOR (GLIOMA)	LYNCH SYNDROME SCREEN (COLON, ENDOMETRIUM)
☐ 1p/19q deletion by FISH	MMR deficiency by IHC
BRAF V600E mutation by PCR	☐ BRAF V600E mutation by PCR *
☐ EGFR amplification by FISH	MLH-1 promotor methylation *
LOT R amplification by 1 1511	* BRAF mutation and/or MLH1 methylation may be needed for those
MELANOMA	with abnormal MMR expression.
☐ BRAF V600 mutation by PCR	OTHER
□ NRAS mutation by PCR/DNA sequencing	OTHER
- INAS mulation by I CIV DIVA Sequencing	