Molecular Diagnostics Labs, Laboratory of Pathology, National Cancer Institute, National Institutes of Health

Requisition for Molecular Diagnostic Tests

Center for Cancer Research Laboratory of Pathology	

Gender: Rirth				of Request:		
Octidet Dirti	n Date:	CRIS	Order # (if any)	# (if any):		
Pathology Case Number:	Medical Record Number:					
LP Resident/Fellow: Dr	Re	questing Physi	cian: Dr	Telephone #:		
Specimen Information						
Formalin-fixed, Pa	raffin-embedded	Tissue (see #5	on back page for	unstained slide order	ing code):	
Slides:	NIH recut	Outside Unsta	nined Slides			
				Tumor Content:	%	
		•	•	Tissue Source:		
_						
Other (specify): _		 Tissue	Source:			
Clinical Information						
Diagnosis / Preliminary Diag	nosis:					
Brief Clinical History:						
Molecular History (MD#, DN	IA#, molecular res	ult):				
Tests Requested (If mic	roarrays is high prior	rity for classification	n, please check the chec	k box. Otherwise, NGS will	be run first)	
Hematopathology	Sarcoma		COMPA	SS	High Prior	
B Cell Clonality: IGH@	EWSR1/	Desmoplastic Round Cell: EWSR1/WT1, t(11;22)		DNA Methylation Microarrays Classifier TruSight Oncology 500 (TSO500)		
IGK@ T Cell Clonality:	Ewings:	ERG, t(21;22)		3, 111 (1	,	
TRG@		FLI1 Type1,t(11;	22)			
_	EWSR1/	FLI1 Type2, t(11	.22)	lutation Test		
Tumor-associated Virus:		osarcoma: 0XO1, t(2;13)	_	V600E		
EBV (EBNA2)		0XO1, t(2,13) 0XO1, t(1;13)		R T790M		
HHV8	Synovial:			MYD88 L265P		
HTLV-I (Pol)		<1, t(X;18) <2, t(X;18)		TERT Promoter (C228T & C250T)		
	3117337	(2, 1(), 10)	DNA Ext	raction for DLM Micro	biology Lab	
MD #:	Date	Accession:				
		,		0 111		
DNA #: RNA #:	Concentration:	(ng.	(ul) Vol (ul)	Quality:		
Test Name	Concentration	(119	Test Result	Quality.		
			<u>10001100011</u>			
		Signature:		Date:		
Version 7.0 revised 10/06	(10000 5					

Instructions for Specimen Submission of Molecular Diagnostic Tests

- 1. Please type or clearly print all information and check appropriate boxes applied to your specimens.
- 2. Paraffin-embedded tissue should be formalin fixed. DNA extracted from tissues fixed in B-5 or other heavy metal containing fixatives does not amplify well.
- 3. For B and T-cell clonality assays and single mutation tests, the minimal amount of tissue required is 1 cm².

$$1 \text{ cm}^2 = \boxed{}$$
 or

- 4. For all NGS assays. please submit 1 H&E slide and 10 unstained slides for small biopsies and cytology specimens, or 5 unstained slides for larger excisional biopsies and resection specimens.
- 5. Unstained slide ordering codes in SoftPath:

Hematopathology, BRAF V600E, MYD88
TERT Promoter, Methylation Classifier, TSO500

- UNST Sarcoma,

- UNMOL

Pulling archived unstained slides - YMOL

- 6. Please include the Outside Case #, SoftPath Block ID, and Tissue Source on your request. This information is required for QM/QA in the lab and during the molecular sign-out.
- 7. Always submit the completed requisition form with the specimen. For cases from the Clinical Center, you must place a CRIS order as well.
- 8. Please provide a molecular diagnostic history with LP MD# if a prior molecular test was performed in house. To compare B & T cell clonality between the current case and a previous case that was signed out before 6/1/2009 (gel detection method), please submit unstained slides from the previous case for reanalysis with our current CE detection method.

Contacts:

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