MLH1 Promotor Hypermethylation REQUEST FORM

Sarah Cannon Molecular Diagnostics Ground Floor, Shropshire House 1 Capper Street, London WC1E 6JA



Part of HCA Healthcare UK

TEL: +44 (0)203 794 1920		EMAIL: info@sarahcannon-md.co.uk			Web: <u>v</u>	Web: www.sarahcannon-md.co.uk			
FOR LABORATORY USE ONLY									
SCMD No:	Receive			Receiv (Date/Tir					
INDIVIDUAL AUTHORISING REQUEST (e.g. Clinician / Pathologist)									
Name:		Address:							
Phone:									
DESTINATION FOR ANALYSIS REPORT (ESSENTIAL – Results may be delayed if not completed)									
Name:		Address:							
Phone:		Note: If as above please tick here							
Required Method(s) for Report Delivery (please tick all that apply): Post Fax Email									
Results Fax number(s):		Results	e-mail(s):						
INVOICING DETAILS (ESSENTIAL – Results may be delayed if not completed in full)									
Contact name:		Full Organisation Name and Postal Address:							
Phone:									
Email:		Note: An authorisation code is mandatory if providing private medical insurance details							
PATIENT DETAILS (At least 3 unique identifiers are mandatory)									
Surname:		Forename:			DOB (DD/MM/YYYY): Gender: M F				
Hospital Name: Surgical C		ase ID: including block number(s) Hospital I			Number:	Requester	Ref: (if ap	pplicable)	
SAMPLE / PATHOLOGY DETAILS (Please provide as much information as possible) NOTE: This assay requires both tumour and normal tissue for comparative purposes. Single blocks/slides may be accepted if suitable for macro-dissection (i.e. yield sufficient normal and tumour components with >20% tumour nuclei in the latter).									
Material Supplied: (see we	_				mour nucle	ei in tumour	compone	ent	
FFPE Single Block	☐ FFPE	E Paired Blocks	< 20% (reject) 21-50%			51-75%			
Unstained Slides		0 T				>75%			
		Tumour Sub-Type: (e.g. Adenocarcinoma)		Tissue Sam	Tissue Sample(s) Supplied: (e.g. colon biopsy,				
Address for return of FF				e analysis re	eport.				
Microsatellite / Mismatch Repair / BRAF Mutation Status (if known)									
☐ MSI-High		MMR Defective (by IHC)		BRAF	codon 600	wild-ty	ре	