

SOMATIC VARIANT ANALYSIS REQUEST FORM (NGS based tests for targeted therapy stratification*)

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*Please note: Whilst much more comprehensive, these assays will have longer turnaround times than our more limited non-NGS based tests. Please consider appropriate clinical needs when choosing.



SARAH CANNON

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FOR LABORATORY USE ONLY

SCMD No:	Received by:	Prepared by:	Received: (Date/Time)
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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 <input type="checkbox"/>	

Required Method(s) for Report Delivery (please tick all that apply): Post Fax Email

Results Fax number(s):	Results e-mail(s):
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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/YY):	Sex: M <input type="checkbox"/> F <input type="checkbox"/>
Hospital Name:	Surgical Case ID:	Hospital Number:	Requester Ref (if applicable):

SAMPLE / PATHOLOGY DETAILS (Please provide as much information as possible)

Material Supplied: (see website for tissue requirements)	Estimated % of tumour nuclei (required for curls/DNA)		
FFPE Block <input type="checkbox"/>	Unstained Slides <input type="checkbox"/>	5-20% <input type="checkbox"/>	51-75% <input type="checkbox"/>
Sections 'curls' <input type="checkbox"/>	Other: <input type="checkbox"/>	21-50% <input type="checkbox"/>	>75% <input type="checkbox"/>
Primary Tumour Type: e.g. Lung	Tumour Subtype: e.g. Adenocarcinoma	Tissue Supplied: e.g. Liver	

Address for return of FFPE block or other unused material:

Note: If left blank, material will be returned to the same address as specified for the analysis report.

ANALYSIS REQUIRED (See website for full details of each assay)

MGP-4 DNA (Small variant/CNG panel) <input type="checkbox"/> Analysis of 'Hotspot' SNVs & small indels in: AKT1, ALK, BRAF, CCNE1, CDK4/6, CDKN2A, CTNNB1, CXCR4, EGFR, ERBB2, ESR1, FBXW7, FGFR1/2/3, GNA11, GNAQ, GNAS, HRAS, IDH1, IDH2, KIT, KRAS, MAP2K1, MET, MTOR, MYD88, NRAS, NTRK1/3, PDGFRA, PIK3CA, POLD1, POLE, PTEN, RET, ROS1, SMAD4, STK11, TP53. Includes analysis of CNGs (copy number gains >2 fold in samples with >20% tumour) in: EGFR, ERBB2, KRAS, KIT, MET, PIK3CA.	MGP-4 RNA (Fusion panel) <input type="checkbox"/> Analysis of inter-/intra-genic rearrangements (fusions) in: ALK, BRAF, EGFR, MET, NTRK1/3, RET, and ROS1. Please note that other gene fusions are included but are currently designated RUO.	Reflex test DNA > RNA <input type="checkbox"/> Reflex to NGS-based gene fusion panel (can reduce cost but will increase turnaround time). Reflex to rapid PCR based Lung fusion panel: ALK, ROS1, RET & MET exon 14 skipping (can reduce cost without excessive delay, but limited scope and restricted to NSCLC only).
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