

Deliver actionable information for personalized treatment selection with APEX

A 50-gene, targeted Next-Generation Sequencing (NGS) panel for the detection of sensitizing and resistance mutations for cancers in:

- ✓ Lung
- ✓ Breast
- Colorectal
- Gastrointestinal Stromal





Gene List (50 genes)

AKT1	♦	CHEK2	•		FGFR4	•	MAP2K1	•		PDGFRA	•	
AKT2	♦	CTNNB1	•		FLT3	•	MAP2K2	♦		PIK3CA	♦ ■	
AKT3	♦	EGFR	♦ I		GNA11	•	MET	♦ ■ •)	PTEN	♦ ■	
ALK	$\blacklozenge \blacksquare \bullet$	ERBB2	♦ I		GNAQ	•	MTOR	•		RAF1	•	
AR	$\blacklozenge \blacksquare \bullet$	ERBB3	♦ I		GNAS	•	NRAS	•		RET	•	
ARAF	•	ERBB4	•		HRAS	•	NRG1	•)	ROS1	•	
BRAF	•	ESR1	•	•	IDH1	•	NTRK1	•)	RSP02		
CD274		FGFR1	♦ I		IDH2	♦	NTRK2	• •)	RSP03		 SNVs/InDe
CDK4	♦	FGFR2	♦ I		KIT	•	NTRK3	•)	SMO	•	Amplificati
CDKN2A	◆ ■	FGFR3	•		KRAS	♦ ■	NUTM1	•)	TP53	•	Fusions

Patient profiles suitable for APEX



(美) Non-Small Cell Lung Cancer

Newly diagnosed, treatment-naïve patients with advanced NSCLC for 1L treatment selection. Relapsed patients who progressed on 1L/2L EGFR, ALK or ROS1 TKI.



Colorectal Cancer

Newly diagnosed, treatment-naïve patients with metastatic CRC, in which testing for KRAS/NRAS/BRAF status is intended.



Breast Cancer

HR+/HER2- BC patients who progressed on CDK4/6 inhibitor and AI, in which testing for resistance mutations is intended.



Gastrointestinal Stromal

Newly diagnosed, treatment-naïve patients with advanced or metastatic GIST in which testing for KIT/ PDGFRA mutation is intended.

Note: Not suitable for patients who have failed multiple treatments and exhausted most treatment options. A larger panel may be warranted in these patients.

Sample Requirement



Provide 10 unstained sections (5-10 μm thick) on uncoated/ uncharged slides*



Provide 1 matching H&E stained slide with tumour region marked out



Tumour cellularity: ≥ 30% Tumour region: ≥ 25 mm³



Place all the slides in the slide holders provided

^{*} Additional 5 sections are required if tumour region <25 mm²



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