

# Therapy Selection FOR SOLID TUMORS



Northstar Select™ detects mutations across 84 genes in cell-free tumor DNA in blood specimens from patients with advanced-stage solid tumors.

## INDICATIONS FOR USE

Northstar Select™ is a liquid biopsy that is appropriate for stage III/IV patients with solid tumor malignancies. It can provide insight into which therapies may be most appropriate based on the genomic profile of the patient’s tumor. The assay can provide therapeutic options for first-line therapy, immunotherapy, and/or therapies that may be beneficial upon progression and/or resistance.

## SUMMARY

This ultra-sensitive next generation sequencing (NGS) assay focuses on actionable mutations, including resistance mutations, associated with FDA-approved targeted and immunotherapies. The assay interrogates 82 genes for single nucleotide variants (SNVs) / indels, 19 genes for copy number amplification, 5 genes for copy number loss and 9 genes for fusions. The panel also covers genes relevant to early phase drug development efforts. The assay has been CLIA validated and requires at least one 8.5mL Streck tube of peripheral blood.

SNVs / Indels 82 genes						CNAs: Amplifications 19 genes			Fusions 9 genes
AKT1	CCNE1	EZH2	JAK2	NOTCH1	RB1	AR	ERBB2	MYC	ALK
AKT2	CD274 (PD-L1)	FANCA	JAK3	NPM1	RET	BRAF	ESR1	PDGFRA	BRAF
ALK	CDH1	FBXW7	KIT	NRAS	RHOA	CCNE1	FGFR1	PIK3CA	FGFR2
APC	CDK12	FGFR1	KRAS	NTRK1	RIT1	CD274 (PD-L1)	FGFR2	RAF1	FGFR3
AR	CDK4	FGFR2	MAP2K1 (MEK1)	PALB2	ROS1	CDK4	KIT	RET	NTRK1
ARAF	CDK6	FGFR3	MAP2K2 (MEK2)	PDGFRA	SF3B1	CDK6	KRAS		NTRK2
ARID1A	CDKN2A	FGFR4	MET	PIK3CA	SMAD4	EGFR	MET		NTRK3
ATM	CDKN2B	GATA3	MLH1	PMS2	SMO				RET
BRAF	CHEK2	GNA11	MPL	PTEN	STK11				ROS1
BRCA1	CTNNB1	GNAQ	MSH2	PTPN11	TERT				
BRCA2	DDR2	GNAS	MSH6	RAD51C	TP53				
BRIP1	EGFR	HRAS	MTOR	RAD51D	TSC1				
CCND1	ERBB2 (HER2)	IDH1	MYC	RAF1	VHL				
CCND2	ESR1	IDH2	NF1						

  

CNAs: Losses 5 genes		Biomarker
ATM	CDKN2A	MSI
BRCA1	PTEN	
BRCA2		

## PERFORMANCE SPECIFICATIONS

Utilizing BillionToOne’s proprietary cfDNA enrichment and bioinformatic processes, Northstar Select™ can detect SNVs / indels and fusions down to two mutated molecules in a single tube of blood. This translates to variant detection at a 0.02 - 0.04% allele fraction even with limited DNA input quantities (i.e. 15 - 30ng of cfDNA). Clinical sequencing is performed to a depth ~40,000 - 60,000x coverage.

Alteration Type	Reportable Range	Allele Fraction / Copy Number	Analytical Sensitivity	Analytical Specificity
SNVs / Indels	≥0.01%	>0.31% 0.17 - 0.30% 0.13 - 0.16% 0.06 - 0.12%	100% 98.9% 95.5% 78.4%	>99.9%
CNAs: Amplifications	≥2.1 copies*	2.125 - 2.160 copies**	95%	
CNAs: Losses	≤1.9 copies	≥1.77 copies	>95%	
Fusions	≥0.02%	>1%	>99%	
MSIs	n/a	0.07%** >0.4%	95% 100%	

\* 2.5% tumor fraction at 6 copies of amplification. \*\* Impacted by biological variability.

Internal data on file, Jun 2023.