

# ClearSeq AML

ClearSeq –NGS Cancer Research Panels  
Next Generation Comprehensive Cancer Profiling



## Benefits

### Expert-identified and Content

- Designed in collaboration with Dr. Robert Ohgami and Dr. Daniel Arber at the Stanford Department of Pathology, Stanford University
- Targets **20 genes** relevant to AML

### Premium Performance You Can Trust

- Provides comprehensive coverage of genetic regions
- From sample to report in less than 48 hours

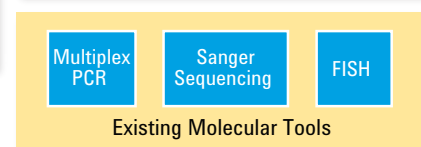
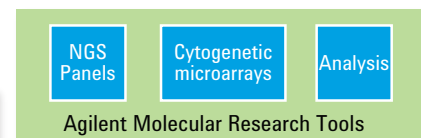
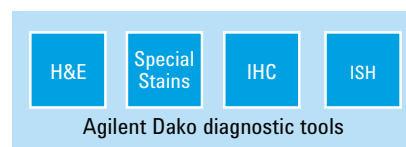
### Easy to implement and to integrate into existing workflows

- From sample to mutation report in 3 simple steps using SureCall
- Obtain all reagents for sample preparation, QC and automation tools from one trusted partner

## Agilent's NGS pathology solutions

Focused cancer gene panels enable you to analyze more mutations with higher accuracy and sensitivity compared to today. Specific panels of genes are being widely adopted to elucidate genomic alterations in cancer and as alternatives to first-generation sequencing techniques as well as replacing time-consuming multi-platform validation iterations.

### Anatomical-to-Molecular Solutions



An end-to-end anatomical-to-molecular genomics portfolio that includes solutions from IHC for cancer diagnosis to NGS for cancer research.

ClearSeq AML developed together with clinical research thought leaders is a proven and reliable solution. NGS can be set up quickly and easily in a lab with comprehensive Agilent solutions for sample preparation, QC, and analysis. Our NGS cancer research solutions compliment existing workflows to provide comprehensive insights.

## Acute myeloid leukemia (AML)

Acute myeloid leukemia (AML) is the most common myeloid neoplasm affecting adults and the role of chromosomal structural variations in its molecular pathogenesis have been well-established. In recent years, next generation sequencing has led to a revolution in the study of hematological malignancies and shown that indels and mutations play an essential part in the pathogenesis of AML. Genetic information coupled with standard morphological findings, provides a deeper characterization and classification of AML.

The ClearSeq AML, designed in collaboration with Dr. Robert Ohgami and Dr. Daniel Arber at the Stanford Department of Pathology, targets 20 genes found to be commonly mutated in AML. These genes are also implicated in other hematological malignancies such as MDS and MPN.



### Easier, faster, more accurate and sensitive performance

- ClearSeq AML is superior in sensitivity and accuracy
- Efficient workflow enables you to identify all mutations with efficiency, accuracy and confidence
- ClearSeq AML can also be readily customized with adding or deleting genes using SureDesign

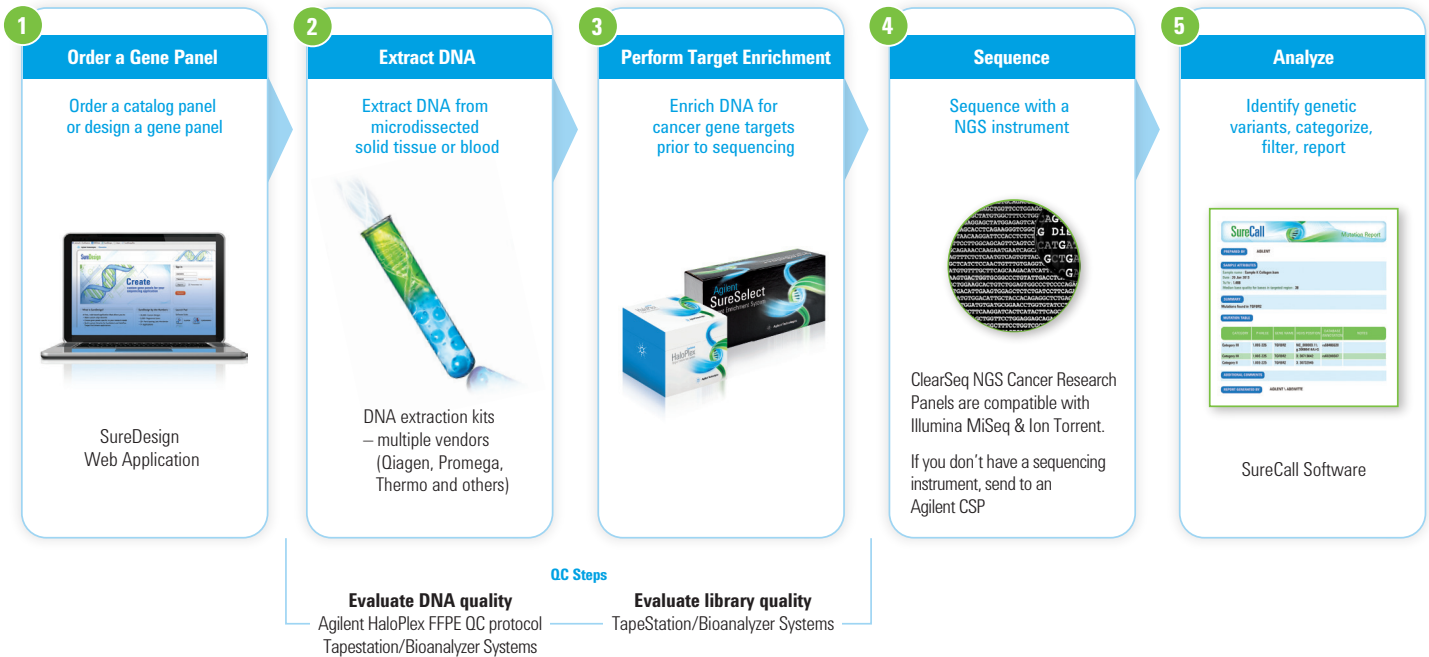
ClearSeq AML will be followed by the release of a series of cancer-type specific and all-cancer comprehensive panels. They will provide coverage of mutations and enable researchers to progress from sample to report in less than 48 hours.

#### Gene List (targeted exons)

<b>ASXL1</b>	12	<b>MPL</b>	10
<b>CSF3R</b>	14, 17	<b>NPM1</b>	11
<b>CBL</b>	8, 9	<b>NRAS</b>	2, 3
<b>CEBPA</b>	1	<b>RUNX1</b>	3, 4, 8
<b>DNMT3A</b>	4, 8, 13, 15, 16, 18 19, 20, 22, 23	<b>SETBP1</b>	3
		<b>SF3B1</b>	13–15, 17
<b>EZH2</b>	8, 17, 18	<b>SRSF2</b>	1
<b>FLT3</b>	14, 20	<b>TET2</b>	3, 9, 10, 11
<b>IDH1</b>	4	<b>TP53</b>	5–8
<b>IDH2</b>	4	<b>U2AF1</b>	2, 6
<b>JAK2</b>	12, 14		

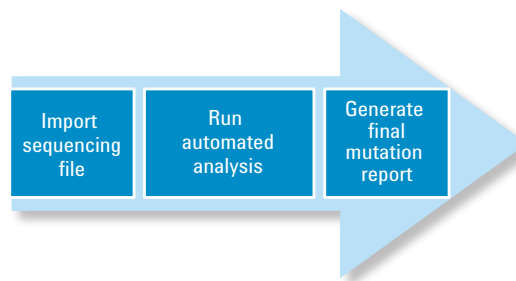
## ClearSeq - NGS Gene Research Panels

5-step process compliments existing pathology tools



## SureCall: Sample-to-Analysis Solution

The ClearSeq AML is ideally suited for analysis using SureCall, a NGS data analysis tool developed to address the critical bioinformatics needs of clinical researchers. Using pre-loaded cancer analysis workflows, go from sample to a mutation report containing all or known variants of interest within minutes, significantly reducing time-to-results. Together with robust sample QC and throughput scalability by automation, ClearSeq AML accelerates the confident detection of disease-associated variants.



Accelerate time-to-results with a simplified 3-step analysis workflow using SureCall. Obtain a report of categorized variants within minutes.

**SureCall Mutation Report**

PREPARED BY: AGTONA02

**SAMPLE ATTRIBUTES**  
Sample name: Sample\_42\_KnownVariant  
Date: 08 Jul 2014

Chr	Start	Stop	Allele	Type	HGVS(Genomic)
CHR11	43615872	43615873	A	SNP	NC_000011.10.g.43615872G>A
CHR11	115258747	115258748	T	SNP	NC_000011.10.g.115258747C>T
CHR9	5073770	5073771	T	SNP	NC_000009.11.g.5073770G>T
CHR17	74732785	74732786	T	SNP	17:74732785>T
CHR18	42531813	42531814	A	SNP	NC_000018.9.g.42531813G>A
CHR20	31023395	31023396	A	SNP	NC_000020.10.g.31023395G>A
CHR21	36164432	36164433	C	SNP	NC_000021.8.g.36164432T>C

**DESCRIPTION/DETAILS: Found Variants**

**NC\_000018.9.g.42531813G>A Category 1**

**Pathology Lab Comment:** Introduction of Stop Codon

**Mutation Impact:** This Variant has effect on ASXL1. Transcript NM\_015338 of ASXL1 are affected. The Variant is HIGH impact and STOP\_GAINED for NM\_015338. This results in codon change of tgg/tga and Amino acid change of W969\*

Known variants of interest



## Agilent 2100 Bioanalyzer, 2200 TapeStation: Sample and Library QC

[www.agilent.com/genomics/bioanalyzer](http://www.agilent.com/genomics/bioanalyzer)

[www.agilent.com/genomics/tapestation](http://www.agilent.com/genomics/tapestation)

## SureDesign – Create Custom Panels in 3-Steps

[www.agilent.com/genomics/suredesign](http://www.agilent.com/genomics/suredesign)

### Product Numbers

Part Number	Description
G9913A	ClearSeq AML Illumina Sequencer 16 reactions
G9913B	ClearSeq AML Illumina Sequencer 16 reactions
G9914A	ClearSeq AML Ion Torrent Sequencer 96 reactions
G9914B	ClearSeq AML Ion Torrent Sequencer 96 reactions

Request more information at [www.Agilent.com/genomics](http://www.Agilent.com/genomics) or call your Agilent service representative for a demo.



To learn more visit: [www.agilent.com/genomics/clearseq-panels](http://www.agilent.com/genomics/clearseq-panels)

Find an Agilent customer center in your country:

[www.genomics.agilent.com/contactUs.jsp](http://www.genomics.agilent.com/contactUs.jsp)

U.S. and Canada: 1-800-227-9770 | [cag\\_sales-na@agilent.com](mailto:cag_sales-na@agilent.com)

© Agilent Technologies, Inc. 2014

Printed in USA, September 5, 2014

5991-5156EN

For Research Use Only. Not for use in diagnostic procedures.

