

# Arima SV Detection Assay

An end-to-end solution for the detection and discovery of structural variants and rearrangements.

## Unlocking the Potential of Clinical and Translational Research Samples

Human clinical research samples, whether cultured, isolated, frozen, or formalin-fixed paraffin-embedded (FFPE), play an indispensable role in biomedical and translational research. In particular FFPE samples often present challenges due to DNA damage; however, the Arima SV Detection Assay is designed to overcome these barriers.

Leveraging the new Arima Bioinformatics Pipeline, researchers can go from data generation to .

### Advantages of the Arima SV Detection Assay



**Maximum Discovery:** Use Arima's proven HiC+ technology to generate high-quality data from clinically relevant samples



**Depth Without the Depth:** superior sensitivity compared to WGS with proximity-linked reads

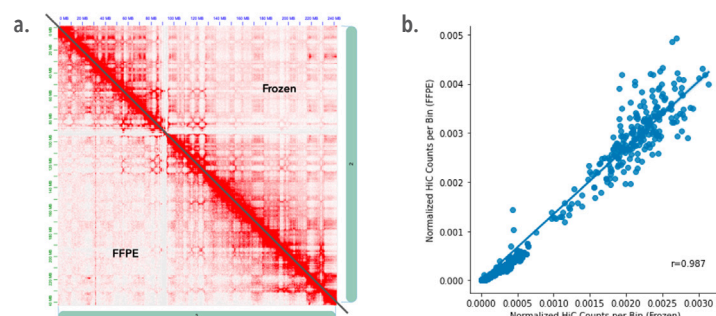


**Lower Investment:** This assay uses well-established NGS technologies; so no need for specialized optical genome mapping or long-read equipment



### High-Quality Data for Comprehensive Insights

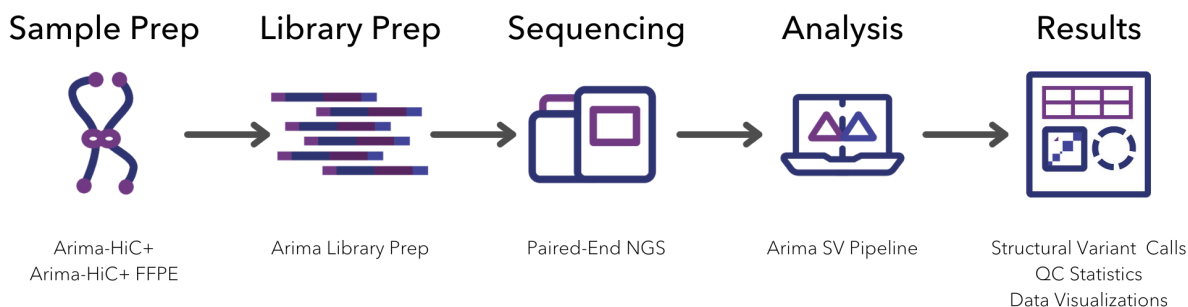
Generate dependable results from FFPE tissue samples to detect both coding and non-coding structural variants in this hard-to-use sample type.



**FFPE Sample Results are Highly Correlated to Frozen Tissues.** Matched frozen and FFPE tissue of K562 cell cultures were processed with the Arima-HiC+ and Arima-HiC+ FFPE kits, respectively. Libraries were generated following Arima protocols and sequenced using Illumina next generation sequencing (57M and 27M reads were generated, respectively). Representative data for Chromosome 2 at 500 Mb are shown in a. Hi-C contact heat map (500Mb resolution) and b. correlation plot of the normalized Hi-C counts per bin for frozen versus FFPE samples showing a high correlation ( $r=0.987$ ).

# Arima SV Detection Assay Workflow

The Arima SV Detection Assay includes reagents for Arima-HiC+ sample preparation and library preparation, in addition to access to the Arima SV pipeline hosted on the Arima Bioinformatics Platform. This assay is compatible with human FFPE and fresh/frozen tissues samples, cell lines and cultures, and cells isolated from liquid biopsies.



## Product List

Product	Description	Size	SKU
Arima SV Detection Assay	Includes: - Arima-HiC+ FFPE kit(s) - Arima Library Prep module - Arima SV Pipeline Analysis.	8 sample 16 sample	A203080 A203081
Arima SV Pipeline Analysis	Analysis of Arima-HiC+ NGS data on the Arima Bioinformatics Platform.	1 sample 8 sample 16 sample	A202082 A202080 A202081

Arima offers end-to-end service with all our products for any sample type.

## Specifications

Category	Specification
Sample Input	5 mm <sup>3</sup> of tissue (~5 x 5µm FFPE tissue sections) 50 mg of fresh frozen tissue 2 mL whole blood ≥1 million cells (standard input method) ≤1 million cells (low input method)
Sequencing Requirements	2 x 150 paired-end reads from Illumina® or Element® A minimum of 100M reads for SV detection is recommended*
Analysis Pipeline	FASTQ Read 1 and Read 2 data 50 -500 Million sequencing reads

\*Performance results vary based on sample quality and purity.

Contact an Arima Genomics scientist for a quote or project consultation. 