

# Product Insert VARIANTPlex™ CFTR v2 Panel

## **VARIANT**Plex CFTR v2

### **Description**

The VARIANT*Plex* CFTR v2 panel is a balanced pool of gene-specific primer (GSP) oligonucleotides that is optimized for use with VARIANT*Plex* reagents and molecular barcode (MBC) adapters to produce targeted NGS libraries. This product insert should be used in conjunction with VARIANT*Plex* standard protocol (RA-DOC-057).

VARIANT*Plex* CFTR v2 contains **208** GSPs with coverage of all exons of the Cystic Fibrosis Conductance Regulator (CFTR) gene, including 5' and 3' untranslated regions (UTRs) and select intronic variants.

Description	Part number	Storage	
VARIANTPlex CFTR v2 GSP1, 8 reactions	SA24000081	-20°C ± 10°C	
VARIANTPlex CFTR v2 GSP2, 8 reactions	SA24000082	-20 C ± 10 C	

## **Recommended PCR cycling**

	Step	Temperature (°C)	Time	Cycles
First PCR reaction	1	95	3 min	1
	2	95	30 sec	
	3	60	10 sec	 14
	4	65	5 min (100% ramp rate)	
	5	72	3 min	1
	6	4	Hold	1

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RA-DOC-466 / REV01

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#### Recommended PCR cycling (cont.)

	Step	Temperature (°C)	Time	Cycles
Second PCR reaction	1	95	3 min	1
	2	95	30 sec	
	3	60	10 sec	22 <sup>†</sup>
	4	65	5 min (100% ramp rate)	
	5	72	3 min	1
	6	4	Hold	1

<sup>†</sup>The number of PCR2 cycles may be decreased if you regularly experience library yields greater than 200 nM.

### Recommended reads and multiplexing

VARIANT*Plex* CFTR v2 libraries should be sequenced to a minimum of **500,000 reads**. Starting read depth recommendations for standard profiling may be adjusted based on user needs.

#### **Archer™ Analysis settings**

Sequencing data should be processed using Archer Analysis (v7.2 Version, or greater). The VARIANT*Plex* CFTR v2 panel requires selection of the *SNV/Indel and Structural Variation* pipeline, found under the *DNA* Input Type (see the Archer Analysis User Guide for more details on setting up your analysis). Selection of the DNA Target Coverage pipeline is optional.

Processing of VARIANT*Plex* CFTR v2 libraries requires a one-time upload of the Panel GTF. When performing DNA Target Coverage analysis, users must also select a Region of Interest BED file. Users may optionally add a Targeted Mutations VCF file for targeted SNV/Indel detection. Files can be obtained by contacting <a href="mailto:archer-tech@idtdna.com">archer-tech@idtdna.com</a>

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#### Limitations of use

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Safety data sheets pertaining to this product are available upon request.

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#### **Revision History**

Document Number	Date	Description of change
RA-DOC-466/REV01	December 2023	Initial release.

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