

## VARIANT*Plex* Solid Tumor Focus v2

### Description

The VARIANT*Plex* Solid Tumor Focus 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* HS/HGC protocol for Illumina® (RA-DOC-056).

VARIANT*Plex* Solid Tumor Focus v2 contains **575** GSPs targeting **20** genes commonly mutated in solid tumors as well as microsatellite instability (**MSI**).

Description	Part number	Storage
VARIANT <i>Plex</i> Solid Tumor Focus v2 GSP1, 8 reactions	SA20121081	
VARIANT <i>Plex</i> Solid Tumor Focus v2 GSP2, 8 reactions	SA20121082	-20°C ± 10°C
PreSeq™ DNA QC Assay Standard, 32 µL	SA0597	
PreSeq™ DNA QC Assay 10X Primer Mix, 120 µL	SA0598	

### Required reagent volumes

Protocol reference	Protocol step	Reagent	Volume per reaction (µL)
A	Ligation Step 2 Elution	5mM NaOH	36
B	First PCR	VARIANT <i>Plex</i> Solid Tumor Focus v2 GSP1	4
C	First PCR	10mM Tris-HCl pH 8.0	38
D	First PCR	Purified PCR1 eluate	36
E	Second PCR	VARIANT <i>Plex</i> Solid Tumor Focus v2 GSP2	4

## Recommended PCR cycling

	Step	Temperature (°C)	Time	Cycles
First PCR reaction	1	95	3 min	1
	2	95	30 sec	
	3	60	10 sec	15
	4	65	10 min (100% ramp rate)	
	5	72	3 min	1
	6	4	Hold	1
Second PCR reaction	1	95	3 min	1
	2	95	30 sec	
	3	60	10 sec	20 <sup>†</sup>
	4	65	10 min (100% ramp rate)	
	5	72	3 min	1
	6	4	Hold	1

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

## Recommended reads and multiplexing

VARIANT*Plex* Solid Tumor Focus v2 libraries should be sequenced to a minimum of **1.5M 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 (v7.0, or greater). The VARIANT*Plex* Solid Tumor Focus v2 panel requires selection of the **SNV/Indel, Structural Variation, Copy Number Variation, and MSI** pipelines, 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* Solid Tumor Focus 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 [archer-tech@idtdna.com](mailto:archer-tech@idtdna.com)

## Assay targets

Gene	Accession	Exon
<i>AKT1</i>	NM_005163	2,3,6,11
<i>BRAF</i>	NM_004333	11,15
<i>EGFR</i>	NM_005228	3,7,12,15,18,19,20,21,22
<i>EGFR</i>	NM_201282	16
<i>EGFR</i>	NM_201283	10
<i>ERBB2</i>	NM_004448	8,10,17,19,20,21,22,24
<i>FOXL2</i>	NM_023067	1(p.C134)
<i>GNA11</i>	NM_002067	5
<i>GNAQ</i>	NM_002072	4,5
<i>GNAS</i>	NM_000516	6,7,8,9
<i>HRAS</i>	NM_005343	2,3
<i>IDH1</i>	NM_005896	3,4
<i>IDH2</i>	NM_002168	4
<i>KIT</i>	NM_000222	2,8,9,10,11,12,13,14,15,17,18
<i>KRAS</i>	NM_004985	2,3,4,5
<i>MET</i>	NM_000245	2,11,14,15,16,19,20,21
<i>NRAS</i>	NM_002524	2,3,4,5
<i>PDGFRA</i>	NM_006206	7,10,11,12,14,15,16,18,23
<i>PIK3CA</i>	NM_006218	2,3,5,7,8,9,10,14,19,21

Gene	Accession	Exon
<i>RET</i>	NM_020630	10,11,13,14,15,16
<i>TERT</i>	NM_198253	3,6,10
<i>TERT</i>	NM_198253	Promoter (chr5:1295148-1295374)
<i>TP53</i>	NM_000546	1,2,3,4,5,6,7,8,9,10,11
<i>TP53</i>	NM_001276696	10

### Genes targeted for CNV

<i>AKT1</i>	<i>EGFR</i>	<i>KIT</i>	<i>MET</i>	<i>PDGFRA</i>	<i>RET</i>
<i>BRAF</i>	<i>ERBB2</i>	<i>KRAS</i>	<i>NRAS</i>	<i>PIK3CA</i>	<i>TERT</i>

Please contact [archer-tech@idtdna.com](mailto:archer-tech@idtdna.com) to inquire about enabling additional genes for CNV detection.

### SNPs and sites targeted for sample tracking

rs560681	rs430046	rs987640	rs10776839	rs12393891
rs740598	rs8078417	rs6444724	rs6530357	chrX:4429309
rs1498553	rs9951171	rs6811238	rs5971553	chrX:11314433
rs10773760	rs576261	rs13182883	rs5953060	chrY:6738552
rs1058083	rs1109037	rs214955	rs6524626	chrY:19490214
rs4530059	rs1523537	rs321198	rs5940270	
rs1821380	rs221956	rs4606077	rs722847	

SNPs may be used in combination to uniquely tag and track samples over time. Contact [archer-tech@idtdna.com](mailto:archer-tech@idtdna.com) for further details.

### 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-032/REV01	June 2023	Initial release.
RA-DOC-032/REV02	November 2023	Updated First and Second PCR cycling conditions to include separate anneal and extended steps.  Added MSI pipeline information to the “Archer Analysis settings” section.  Updated branding.