# **Product Insert**LIQUID*Plex*<sup>TM</sup> Universal Solid Tumor Panel

## LIQUIDPlex Universal Solid Tumor

#### **Description**

The LIQUID *Plex* Universal Solid Tumor panel is a balanced pool of gene-specific primer (GSP) oligonucleotides that is optimized for use with LIQUID *Plex* reagents and molecular barcode (MBC) adapters to produce targeted NGS libraries. This product insert should be used in conjunction with LIQUID *Plex* protocol for Illumina® (RA-DOC-052).

Description	Part number	Storage
LIQUIDPlex Universal Solid Tumor GSP1, 8 reactions	cSA5061081	− −20°C ± 10°C
LIQUID <i>Plex</i> Universal Solid Tumor GSP2, 8 reactions	cSA5061082	— −20 C ± 10 C

### Required reagent volumes

Protocol reference	Protocol step	Reagent	Volume per reaction (µL)
Α	Ligation Step 2 Elution	5mM NaOH	36
В	First PCR	LIQUIDPlex Universal Solid Tumor GSP1	4
С	First PCR	10mM Tris-HCl pH 8.0	38
D	First PCR	Purified PCR1 eluate	36
Е	Second PCR	LIQUID <i>Plex</i> Universal Solid Tumor GSP2	4

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#### **Recommended PCR cycling**

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

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

#### Recommended reads and multiplexing

LIQUID Plex Universal Solid Tumor libraries should be sequenced to a minimum of **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 LIQUID *Plex* Universal Solid Tumor panel requires selection of the *SNV/Indel, Structural Variation*, *and Copy Number Variation* pipelines, found under the *cfDNA* Input Type (see the Archer Analysis User Guide for more details on setting up your analysis).

Processing of LIQUID*Plex* Universal Solid Tumor libraries requires a one-time upload of the Panel GTF. 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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#### **Assay targets**

Gene	Accession	Exon*
AKT1	NM_005163	Select hotspots
ALK	NM_004304	Select hotspots
AR	NM_000044	Select hotspots
BRAF	NM_004333	Select hotspots
CTNNB1	NM_001904	Select hotspots
EGFR	NM_005228	Select hotspots
ERBB2	NM_004448	Select hotspots
ERBB3	NM_001982	Select hotspots
ESR1	NM_000125	Select hotspots
FGFR1	NM_015850	Select hotspots
FGFR2	NM_000141	Select hotspots
FGFR3	NM_000142	Select hotspots
HRAS	NM_005343	Select hotspots
IDH1	NM_005896	Select hotspots
IDH2	NM_002168	Select hotspots
KIT	NM_000222	11
KRAS	NM_004985	Select hotspots
MAP2K1	NM_002755	Select hotspots

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Gene	Accession	Exon*
MET	NM_000245	Select hotspots
NRAS	NM_002524	Select hotspots
NTRK1	NM_002529	Select hotspots
NTRK2	NM_006180	Select hotspots
NTRK3	NM_002530	Select hotspots
PDGFRA	NM_006206	Select hotspots
PIK3CA	NM_006218	Select hotspots
RET	NM_020630	Select hotspots
ROS1	NM_002944	Select hotspots
TP53	NM_001276698	6
TP53	NM_000546	1,2,3,4,5,6,7,8,9,10,11

<sup>\*</sup>Contact archer-tech@idtdna.com for a target BED file of the targeted genomic regions

### **Genes targeted for CNV**

AR	CDK6	ERBB2	FGFR1	FGFR3
BRAF	EGFR	ERBB3	FGFR2	MET

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#### Limitations of use

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#### **Revision History**

Document Number	Date	Description of change
RA-DOC-039/REV01	June 2023	Initial release.
RA-DOC-039/REV02	November 2023	Updated branding.

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