

VARIANT*Plex* MPN Focus

Description

The VARIANT*Plex* MPN Focus 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).

| Description | Part number | Storage |
|--|-------------|--------------|
| VARIANT <i>Plex</i> MPN Focus GSP1, 8 reactions | dSA09814081 | -20°C ± 10°C |
| VARIANT <i>Plex</i> MPN Focus GSP2, 8 reactions | dSA09814082 | |
| PreSeq [™] DNA QC Assay Standard, 32 µL | SA0597 | |
| PreSeq [™] DNA QC Assay Standard, 32 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> MPN Focus 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> MPN Focus 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 | 5 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† |
| | 4 | 65 | 5 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* MPN Focus libraries should be sequenced to a minimum of **800,000** 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 (v7.0, or greater). The VARIANT*Plex* MPN Focus panel requires selection of the **SNV/Indel and DNA Structural Variation** 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* MPN Focus 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

Assay targets

| Gene | Accession | Exon |
|---------------|--------------|---|
| <i>ASXL1</i> | NM_015338 | 11,12,13 |
| <i>CALR</i> | NM_004343 | 8,9 |
| <i>CSF3R</i> | NM_000760 | 10,14,15,16 |
| <i>CSF3R</i> | NM_156039 | 17 |
| <i>CSF3R</i> | NM_172313 | 18 |
| <i>DNMT3A</i> | NM_022552 | 2,3,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20,21,22,23 |
| <i>DNMT3A</i> | NM_153759 | 1,2 |
| <i>DNMT3A</i> | NM_175630 | 4 |
| <i>EZH2</i> | NM_004456 | 2,3,4,5,6,7,8,9,10,11,12,13,14,15,16,17,18,19,20 |
| <i>IDH1</i> | NM_005896 | 3,4 |
| <i>IDH2</i> | NM_002168 | 4,6 |
| <i>JAK2</i> | NM_004972 | 12,13,14,15,16 |
| <i>KIT</i> | NM_000222 | 2,8,9,10,11,12,13,14,15,17,18 |
| <i>MPL</i> | NM_005373 | 10,12 |
| <i>SETBP1</i> | NM_015559 | 4 (p.799-p.950) |
| <i>TET2</i> | NM_001127208 | 4,5,6,7,8,9,10,11 |

| Gene | Accession | Exon |
|------|-----------|------|
| TET2 | NM_017628 | 3 |

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 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-038/REV01 | June 2023 | Initial release. |
| RA-DOC-038/REV02 | November 2023 | Updated First and Second PCR cycling conditions to include separate anneal and extended steps. Updated branding. |