

Specifications

The CleanPlex™ TP53 Panel contains 29 pairs of PCR primers targeting the full TP53 gene exon. The panel kit contains primers, multiplex PCR reagent, digestion reagent and other components necessary to construct amplicon libraries for Next-Generation Sequencing on Illumina Sequencers.

100% coverage of TP53 gene with superior uniformity

The panel covers 100% of the coding regions of the TP53 gene. The observed uniformity of this panel (at 0.2x mean coverage) is over 98%.

Simplify your workflow

The entire library preparation workflow can be completed in 2.5 hours with only 30-minute hands-on time from sample DNA to sequencing-ready libraries. No need for ligation, end repair, DNA fragmentation, overnight hybridization, or microfluidic devices.

Take on difficult samples with limited DNA input

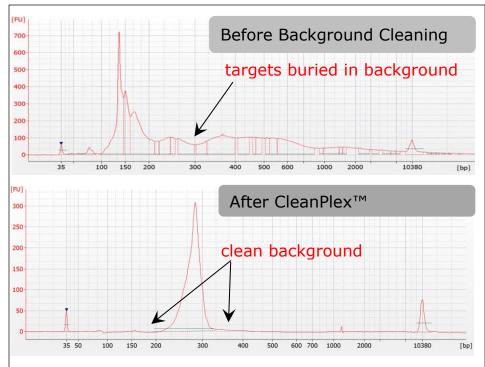
With an average amplicon size of 133 bp, this panel is compatible with degraded samples such as formalin-fixed, paraffin-embedded (FFPE) tissue DNA. Obtain high quality sequencing data for germline genotype calling with only 200 pg of input DNA.

Ordering information		
Product Name	Sku	
CleanPlex™ TP53 Panel (8 rxns)	916008	
CleanPlex™ TP53 Panel (96 rxns)	916009	
CleanPlex™ TP53 Panel (384 rxns)	916010	

Sequencing Illumina Sequencers Platform (MiniSeq, MiSeq, NextSeq, Hiseq) Enrichment Multiplex PCR Method # of Primer Pools 2 pools # of Primer Pairs 29 pairs # of Target Genes 1 gene Target Region 2080 bp Size Average 133 bp Amplicon Size (from 107-160 bp) Species Human For germline genotype calling: minimum 200 pg; Recommended Input DNA For somatic mutation calling with Amount an LOD of 1%: minimum 20 ng (10 ng / pool) Genomic DNA, FFPE DNA, and DNA from Tissue, Cell Culture, and Fine Sample Type Needle Aspirate (FNA) MiSeq 2×150 read length: Sample ~384 samples Multiplexing (at ~2000x mean NextSeg mid output 2×150 read coverage) length: ~2800 samples

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Most target enrichment kits do not provide effective background cleaning, resulting in sequencing of non-specific PCR products post amplification, which translates into the generation of excess reads.

CleanPlex™ By using technology, background noise is greatly reduced and only the targets of interest are This proprietary multiplex sequenced. DNA PCR technology eliminates fragmentation, hybridization and ligation resulting in higher target coverage, on-target rates and lower assay failure.

Important advantages to NGS lab operations and data quality

Uniformity	
Specificity	
Time	
Minimum Sample input	
Workflow	

Competitor X	Paragon Genomics CleanPlex™ Solution
85 - 95%	>98%
85 - 97% on-target bases	>97% on-target bases
6 hours	2.5 hours
20 – 40 ng	0.1 ng
5 steps	3 steps

