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QIAseq Targeted DNA Panels are intended for molecular biology applications. These products are not intended for the diagnosis, prevention, or treatment of a disease.

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QIAseq Targeted Methyl Panels offer a single-day solution capable of targeting thousands of CpG sites in the genome. As with targeted DNA panels, QIAseq solutions are based on single-primer extension (SPE) technology that provides a sensitive and specific solution for detecting regions all across the genome.

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According to the QIAseq Targeted DNA panel Handbook and protocol, and depending on the input DNA for the library preparation, the ideal value for Average reads per group should be 2 to 4, with 4 being the best value for the highest DNA input (i.e., 40ng).

[Quality Control for the Identify QIAseq DNA Variants workflow](#)

QIAseq Pan-cancer Multimodal Panel for comprehensive genomic profiling of solid tumors and heme malignancies . Unlock insights into multiple cancers with just 10 ng total nucleic acid input. Go to product page Experience our innovative, single-day sample to sequencing workflow for simultaneous enrichment and profiling of 605 relevant DNA variant and RNA fusion biomarkers found across multiple ...

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disclaimers, see the respective QIAGEN kit handbook or user manual. QIAGEN kit handbooks and user manuals are available at ... Comprehensive Cancer panel 33 QIAseq Targeted DNA Panels delivered consistently high sequencing specificity and coverage uniformity with different DNA inputs. 94,0 94,7 90,5 99,5 99,6 99,9 97,7 99,2 97,8 99,4 97,4 99,8 0 20 40 60 80 100 120 101Z-1 ng 101Z-10 ng 101Z-20 ...

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Massively Parallel Sequencing (MPS) allows to sequence the total mitochondrial DNA (mtDNA) through specific panels. Human Mitochondria Panel is one of the QIAseq Targeted DNA panels developed by QIAGEN for MPS, which have an optimized chemistry based on the integration of unique molecular indices (UMIs).

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QIAseq 16S/ITS Panel Handbook 04/2019 9 Data analysis Sequencing is performed on an Illumina MiSeq NGS system using a v2 kit with 251 x 2 paired end run or a v3 kit with 276 x 2 paired end run. It is highly recommended to perform sequencing data analysis with the CLC Genomics Workbench with Microbial Genomics Module and QIAseq 16S Panel Analysis.

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QIAseq Targeted NGS for Cancer Research, 10.10.2016 • Offers specificity that beats capture-based approaches Features Benefits • Lets you use sequencing capacity on regions targeted by the panel, with minimal off-target sequencing • Lets you achieve more uniform enrichment for more sequencing efficiency It delivers unmatched specificity and uniformity (compared to capture-based methods)

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