

Truseq Enrichment Guide

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NextSeq 500.

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Prepare 5% extra reagent mix if you are preparing multiple samples. 2 Remove the PCR plate from magnetic stand 3 Add 30 μ l of the elution pre mix to each well of the PCR plate using a single channel or multichannel pipette. 4 Gently pipette the entire volume of each well up and down 10 – 20 times to mix thoroughly. Ensure that the beads are fully resuspended. 5 Seal the PCR plate with a Microseal ' B ' adhesive seal.

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The TruSeq Exome Kit supports 12-plex pre-enrichment library pooling, enabling researchers to maximize sequencing throughput and variant identification by sequencing up to 12 libraries per flow cell lane. This enables sequencing of more exomes per run, so researchers can maximize their budgets.

TruSeq DNA Exome (formerly TruSeq Exome Kit)

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Procedure. 1 [Plate]ArrangeIndex1(i7)adaptersincolumns1 – 12oftheTruSeqIndexPlate Fixture. 2 [Plate]ArrangeIndex2(i5)adaptersinrowsA – HoftheTruSeqIndexPlateFixture. 3 [Plate]PlacetheplateontheTruSeqIndexPlateFixture. 4 Add5 µ l ofeachIndex1 (i7)adapterasfollows.

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Illumina RNA Prep with Enrichment is designed to provide exceptional capture efficiency and coverage uniformity, minimizing required sequencing depth for accurate detection without bias. Only 2.5 µL probes are needed per enrichment reaction.

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TruSeq RNA Exome, previously known as the TruSeq RNA Access Library Prep Kit, converts total RNA into template molecules of known strand origin, followed by sequence-specific capture of coding RNA. This provides a low-cost solution for analyzing human RNA isolated from formalin-fixed, paraffin-embedded (FFPE) tissues and other low-quality samples. Affordability and focus—Isolating human transcriptome coding regions maximizes discovery power at a fraction of the sequencing depth.

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