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The Change In An Anatomic Pathology Report Is Usually In The Diagnosis Or Narrative, But Occasionally May Involve A Change In A Number Or Some Other Quality. Analytical Validation - The Process Used To Confir 23th, 2024

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RevisionHistory Document Date DescriptionofChange Document#15045738v04 November 2019 Corrected 3th, 2024

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MiSeq Software Release Notes Part Number: 15059913 Rev A Release Date: September 15, 2014 Page 3 Of 5 Perform A Template Line Wash. This Warning Does Not Apply To Other Workflows. 20th, 2024

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TruSeq Stranded mRNA Illumina Stranded mRNA Prep TruSeq Stranded mRNA Illumina Stranded mRNA Prep % RRNA (28S/18S) 5.1 1.8 5.3 1.5 % Strandedness 99.6 99.4 99.6 99.4 Median CV Of Coverage 0.49 0.46 0.50 0.47 % Duplicates 5.4 3.7 8.5 3.3 % Aligned 97.1 97.8 96.8 97.8 Results Are From RNA Se 23th, 2024

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Sciclone NGS: Illumina TruSeq™ RNA Introduction Preparation Of Samples For mRNA Sequencing On The Illumina, Inc. Platform Requires A Series Of Manipulations To Isolate And Fragment mRNA, Convert The RNA To Double-stranded CDNA, And Efficiently Ligate Appropriate Indexed Adapters To Pro 23th, 2024

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(previously Available With TruSeq™ Stranded Total RNA And TruSeq Stranded mRNA). Methods Sample Preparation Several Microbial Isolates And Mock Community Standards Were Obtained From ATCC For Evaluation (Table 1). Total RNA Was Extracted Using The RNeasy PowerMicrobiome Kit (QIAGEN, Cata 15th, 2024

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