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| <h2>Lyo-Ready™ 1-Step RT-qPCR Mix, 2x</h2> <p>For research or further manufacturing use only</p> | Catalog No: | MDX062 |
| | Lot No: | EM078-B112760 |
| | Storage Conditions: | -20°C |
| | Component Lot No: | LY1S-222411A |
| | Expiry date: | December 2024 |

Quality Control Parameters

Ready-to-use, glycerol-free RT-qPCR mastermix formulated with a specialized blend of excipients for lyophilization into beads or cakes

| Analysis | Specification | Result |
|---------------------|--|--------|
| Functional | <p>Quantitative RT-qPCR analysis amplifying 3 genes from a dilution series of mouse RNA under standard conditions.</p> <p><u>Pass Criteria:</u></p> <p>Amplification profiles must be consistent for the test and reference sample within ± 0.5 Cq variance.</p> | Passed |
| DNA contamination | <p>Quantitative PCR analysis with no template. Presence of <i>E. coli</i> and mouse genomic DNA checked. Test sample must amplify in concordance with control sample.</p> <p><u>Pass Criteria:</u></p> <p>Amplification traces must overlay with the negative control.</p> | Passed |
| DNase contamination | <p>DNase contamination is measured as DNA substrate degradation against a DNase I dilution series by agarose gel electrophoresis.</p> <p>Limit of detection: 6.25×10^{-4} KU DNase I.</p> <p><u>Pass Criteria:</u></p> <p>No detectable degradation.</p> | Passed |

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| | | |
|---------------------|---|--------|
| RNase contamination | Quantitative PCR analysis with high and low RNase standards. Limit of detection: 9.7×10^{-3} ng/ μ L RNase <u>Pass Criteria:</u> Test sample must show less RNase activity than the limit of detection. | Passed |
|---------------------|---|--------|

QA / QC Representative:



Alberta Newton

Date: 25th November 2022

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Virus MMLV-RT

For research or further manufacturing use only

| | |
|---------------------|---------------|
| Catalog No: | MDX062 |
| Lot No: | EM078-B112760 |
| Storage Conditions: | -20°C |
| Component Lot No: | VRT1-122311A |
| Expiry date: | December 2024 |

Quality Control Parameters

MMLV-RT (Moloney Murine Leukemia Virus Reverse Transcriptase) is a reverse transcriptase that can be used for cDNA synthesis and subsequent PCR or qPCR in a one-step or two-step assay.

| Analysis | Specification | Result |
|---------------------|---|--------|
| Functional | Activity is measured as reverse transcriptase units by primer extension analysis against a reference enzyme. <u>Pass Criteria:</u> Activity must be equal to or greater than 200 U/μL | Passed |
| Specific Activity | The Specific activity must be $\geq 300,000$ U/mg | Passed |
| Purity | Densitometric analysis of SDS-Page. Purity must be higher than 95% | Passed |
| DNA contamination | Quantitative PCR analysis with no template. Presence of <i>E. coli</i> and mouse genomic DNA checked. Test sample must amplify in line with control sample. | Passed |
| DNase contamination | Incubation of a 1Kb double stranded DNA fragment. Incubation for 4 hours at 37°C with dilution series of DNase I. Analysed by agarose gel electrophoresis. Test sample must show less degradation than the limit of detection 2.5×10^{-3} U DNase I. | Passed |
| RNase contamination | Quantitative PCR analysis with high and low RNase standards. Test sample must show less RNase activity than the limit of detection 9.7×10^{-3} ng/μL RNase. | Passed |

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Enzyme Dilution Buffer

For research or further manufacturing use only

| | |
|---------------------|---------------|
| Catalog No: | MDX062 |
| Lot No: | EM078-B112760 |
| Storage Conditions: | -20°C |
| Lot number: | TDB-222111A |
| Expiry date: | December 2024 |

Quality Control Parameters

Enzyme Dilution Buffer is a glycerol-free 10x buffer used for the dilution of Taq antibody or reverse transcriptase.

| Analysis | Specification | Result |
|------------------------|--|--------|
| Functional | A 3Kb fragment was amplified with a dilution series of Enzyme Dilution Buffer, using standard conditions and 30 cycles. A 3Kb fragment was amplified with a dilution series of <i>Taq polymerase</i> , using standard conditions and 30 cycles. Single distinct bands were observed with agarose gel electrophoresis (ethidium stained). | Passed |
| Glycerol determination | Glycerol content is < 0.2% determined by spectrophotometric analysis and comparison to a standard curve. | Passed |
| DNA contamination | Quantitative PCR analysis with no template. Presence of <i>E. coli</i> and mouse genomic DNA checked. Test sample must amplify in line with a reference sample. | Passed |
| DNase contamination | Incubation of a 1Kb double stranded DNA fragment. Incubation for 4 hours at 37°C with dilution series of DNase I. Analysed by agarose gel electrophoresis. Test sample must show less degradation than the limit of detection 6.25×10^{-4} KU/ μ L. | Passed |
| RNase contamination | Quantitative PCR analysis with high and low RNase standards. Test sample must show less RNase activity than the limit of detection: 9.7×10^{-3} ng/ μ L RNase. | Passed |

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