

*be* INSPIRED *drive* DISCOVERY *stay* GENUINE

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## New England Biolabs Certificate of Analysis

| Product Name:          | NEBNext® Poly(A) mRNA Magnetic Isolation Module |
|------------------------|---|
| Catalog Number:        | E7490L  |
| Packaging Lot Number:  | 10112752  |
| Expiration Date:       | 04/2023   |
| Storage Temperature:   | 4°C   |
| Specification Version: | PS-E7490S/L v1.0                                |

| NEBNext® Poly(A) mRNA Magnetic Isolation Module Component List |                                      |            |                      |  |
|--|--------------------------------------|------------|----------------------|--|
| NEB Part Number  | Component Description                | Lot Number | Individual QC Result |  |
| E7499AAVIAL  | NEBNext® Oligo dT(25) Magnetic Beads | 10103528   | Pass                 |  |
| E7496AAVIAL  | NEBNext® Tris Buffer                 | 10103527   | Pass                 |  |
| E7495AAVIAL  | Nuclease-free Water                  | 10103526   | Pass                 |  |
| E7493AAVIAL  | NEBNext® Wash Buffer                 | 10103525   | Pass                 |  |
| E7492AAVIAL  | NEBNext® RNA Binding Buffer (2X)     | 10103524   | Pass                 |  |

| Assay Name/Specification  | Lot # 10112752 |
|---|----------------|
| * Individual Product Component Note<br>Standard Quality Control Tests are performed for each component included in NEBNext®   | Pass           |
| Poly(A) mRNA Magnetic Isolation Module and meet the designated specifications.  |                |
| <b>Functional Testing (Poly(A) Isolation)</b><br>The NEBNext® Poly(A) Isolation Module is functionally validated using commercially<br>available human RNA (e.g. UHRR). After treatment using the kit's minimum and maximum<br>input requirements, RNA yield is assessed by Bioanalyzer. Libraries made from<br>previous and current lots, using the minimum and maximum input amounts are sequenced<br>together on the same Illumina® flow cell and compared across various metrics<br>including library yield, individual transcript abundance correlations, 5'-3'<br>transcript coverage, and fraction of reads mapping to a reference. This method<br>produces libraries with less than 10% rRNA. | Pass           |

This product has been tested and shown to be in compliance with all specifications.

One or more products referenced in this document may be covered by a 3rd-party trademark. Please visit www.neb.com/trademarks for additional information.





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Christine Sumner Production Scientist 24 Jun 2021

Michae m. l

Michael Tonello Packaging Quality Control Inspector 24 Jun 2021

