

Nov 14, 2024

Quantitative bulk transcriptomics

DOI

<https://dx.doi.org/10.17504/protocols.io.3byl4w7rzvo5/v1>

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Protocol Citation: Livia Hecke Moraes 2024. Quantitative bulk transcriptomics. **protocols.io**
<https://dx.doi.org/10.17504/protocols.io.3byl4w7rzvo5/v1>

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Protocol status: Working

We use this protocol and it's working

Created: November 14, 2024

Last Modified: November 14, 2024

Protocol Integer ID: 112065

Keywords: ASAPCRN, quantitative bulk transcriptomics quantitative, quantitative bulk transcriptomics quantitative bulk transcriptomic, bulk transcriptomic, total striatum tissue rna, rna, sequencing

Funders Acknowledgements:

ASAP

Abstract

Quantitative bulk transcriptomics used in Morais et al. for total striatum tissue RNA sequencing.



RNA extraction

- 1 Homogenize the tissue in TRIzol.
- 2 200 μ l chloroform per 1 ml of TRIzol.
- 3 Shake vigorously 15 sec. (DO NOT VORTEX)
- 4 Incubate at RT for 3 min.
- 5 Spin 15 min at 4C, 12,000g to separate phases.
- 6 Carefully transfer 400 μ l of the clear, colorless supernatant to a clean sterile 1.5 ml tube. Avoid the middle, white layer. Dispose of the remaining liquid in the liquid waste.
- 7 To the 1.5 ml tube, add an equal volume (400 μ l) of 70% ethanol to the collected supernatant, mix well through pipetting.
- 8 Transfer 700 μ l to an RNeasy spin column placed in a 2mL collection tube. Centrifuge at room temperature for 1 minute at 10 000 rpm.
- 9 Discard the flow-through and transfer any remaining solution to the column and centrifuge again.
- 10 Transfer the flow through column to a new collection tube.
- 11 Wash with 350 μ l Buffer RW1.
- 12 Centrifuge at room temperature for 1 minute at 10 000 rpm. Discard the flow through with a pipette.



- 13 Separate tube, add 10 μ L DNase I Stock Solution to 70 μ L Buffer RDD. Mix by gently inverting the tube, and centrifuge briefly to collect residual liquid from the sides of the tube
- 14 Carefully add the 80 μ L of DNase I mix directly to the RNeasy spin column membrane, and place on the benchtop (20–30°C) for 15 min
- 15 Add 350 μ L Buffer RW1 to the RNeasy spin column.
- 16 Add 350 μ L Buffer RW1 to the RNeasy spin column.
- 17 Centrifuge for 1 minute at 10 000 rpm. Discard the flow through with a pipette.
- 18 Wash with 500 μ L Buffer RPE
- 19 Centrifuge at room temperature for 1 minute at 10 000 rpm. Discard the flow through with a pipette
- 20 Centrifuge at room temperature for 1 minute at 10 000 rpm. Discard the flow through with a pipette
- 21 Centrifuge at room temperature for 2 minutes at 13 000 rpm. This step removes any ethanol or other washes.
- 22 Transfer the column to a clean, sterile 1.5 ml Eppendorf tube.
- 23 Elute RNA with 35 μ L RNase free water. Incubate at room temperature for 5 minutes.
- 24 Centrifuge at room temperature for 1 minute at 13000 rpm.
- 25 Immediately store at -20 or -80 until further use.



Library prep using Lexogen QuantSeq 3' mRNA-Seq Library Prep Kit FWD HT

- 26 Mix 200 ng of total RNA in a volume of 5 µl, with 5 µl First Strand cDNA Synthesis Mix 1 (**FS1**) in a PCR plate or 8-well strip tubes. If necessary, adjust the total volume to 10 µl with RNase-free water. Mix well by pipetting.
- 27 Denature the RNA / **FS1** mix for 3 minutes at 85 °C in a thermocycler and then cool down to 42 °C.
- 28 Prepare a mastermix containing 9.5 µl First Strand cDNA Synthesis Mix 2 (**FS2**) and 0.5 µl Enzyme Mix 1 (**E1**) per reaction. Mix well, spin down, and pre-warm the mastermix for 2 - 3 minutes at 42 °C.
- 29 Quickly spin down the denatured RNA / **FS1** samples from step 2 at room temperature to make sure all liquid is collected at the bottom of the wells. Place the samples back onto the thermocycler at 42 °C.
- 30 Add 10 µl of the **FS2** / **E1** mastermix to each reaction, mix well, and seal the plate.
- 31 Quickly spin down at room temperature and incubate the reactions for 15 minutes at 42 °C.
- 32 Add 5 µl RNA Removal Solution (**RS**) directly to the first strand cDNA synthesis reaction. Mix well and reseal the plate.
- 33 Incubate 10 minutes at 95 °C, then cool down to 25 °C. Spin down the plate at room temperature and carefully remove the sealing foil.
- 34 Add 10 µl Second Strand Synthesis Mix 1 (**SS1**) to the reaction. Mix well by pipetting,
- 35 Incubate the plate for 1 minute at 98 °C in a thermocycler, and slowly cool down to 25 °C by setting the ramp speed to 10 % (0.5 °C/second). Incubate the reaction for 30 minutes at 25 °C.
- 36 Quickly spin down the plate at room temperature before removing the sealing foil.
- 37 Prepare a mastermix containing 4 µl Second Strand Synthesis Mix 2 (**SS2**) and 1 µl Enzyme Mix 2 (**E2**). Mix well.



- 38 Add 5 μ l of the SS2 / E2 mastermix per reaction. Mix well. REMARK: Use a pipette set to 30 μ l for efficient mixing.
- 39 Incubate the reaction for 15 minutes at 25 °C. Safe stopping point. Libraries can be stored at -20 °C at this point.
- 40 Add 16 μ l of properly resuspended Purification Beads (PB) to each reaction, mix well, and incubate for 5 minutes at room temperature.
- 41 Place the plate onto a magnetic plate, and let the beads collect for 2 - 5 minutes or until the supernatant is completely clear (depends on the strength of your magnet).
- 42 Remove and discard the clear supernatant without removing the PCR plate from the magnetic plate. Make sure that accumulated beads are not disturbed.
- 43 Add 40 μ l of Elution Buffer (EB), remove the plate from the magnet, and resuspend the beads properly in EB.
- 44 Incubate for 2 minutes at room temperature.
- 45 Add 56 μ l of Purification Solution (PS) to the beads / EB mix to reprecipitate the library.
- 46 Mix thoroughly, and incubate for 5 minutes at room temperature.
- 47 Place the plate onto a magnetic plate, and let the beads collect for 2 - 5 minutes or until the supernatant is completely clear.
- 48 Remove and discard the clear supernatant without removing the PCR plate from the magnetic plate. Make sure that accumulated beads are not disturbed.
- 49 Add 120 μ l of 80 % EtOH, and incubate the beads for 30 seconds. Leave the plate in contact with the magnet as beads should not be resuspended during this washing step.
- 50 Remove and discard the supernatant.



- 51 Repeat this washing step once for a total of two washes. Make sure to remove the supernatant completely as traces of ethanol can inhibit subsequent PCR reactions.
- 52 Leave the plate in contact with the magnet, and let the beads dry at room temperature for 5 - 10 minutes or until all ethanol has evaporated only. Do not let the beads dry too long (visible cracks appear), this will negatively influence the elution and the resulting library yield.
- 53 Add 20 μ l of Elution Buffer (EB) per well, remove the plate from the magnet and resuspend the beads properly in EB.
- 54 Incubate for 2 minutes at room temperature.
- 55 Place the plate onto a magnetic plate and let the beads collect for 2 - 5 minutes or until the supernatant is completely clear.
- 56 Transfer 17 μ l of the clear supernatant into a fresh PCR plate. Make sure not to transfer any beads. Safe stopping point.
- 57 Prepare a mastermix containing 7 μ l of PCR Mix (PCR) and 1 μ l Enzyme Mix 3 (E3) per reaction.
- 58 Add 8 μ l of this PCR / E3 mastermix to 17 μ l of the eluted library.
- 59 Add 5 μ l of the respective i7 index. Mix well by pipetting.
- 60 Seal the PCR plate and quickly spin down to make sure all liquid is collected at the bottom of the well.
- 61 Conduct 14 cycles of PCR with: Initial denaturation at 98 °C for 30 seconds, 14 cycles of 98 °C for 10 seconds, 65 °C for 20 seconds and 72 °C for 30 seconds, and a final extension at 72 °C for 1 minute, hold at 10 °C.
- 62 Safe stopping point. Libraries can be stored at -20 °C at this point.
- 63 Add 30 μ l of properly resuspended Purification Beads (PB) to each reaction, mix well, and incubate for 5 minutes at room temperature.



- 64 Place the plate onto a magnetic plate, and let the beads collect for 2 - 5 minutes or until the supernatant is completely clear.
- 65 Remove and discard the clear supernatant without removing the PCR plate from the magnetic plate. Make sure that accumulated beads are not disturbed.
- 66 Add 30 μ l of Elution Buffer (EB), remove the plate from the magnet, and resuspend the beads properly in EB. Incubate for 2 minutes at room temperature.
- 67 Add 30 μ l of Purification Solution (PS) to the beads / EB mix to reprecipitate the library.
- 68 Mix thoroughly, and incubate for 5 minutes at room temperature.
- 69 Place the plate onto a magnetic plate, and let the beads collect for 2 - 5 minutes or until the supernatant is completely clear.
- 70 Remove and discard the clear supernatant without removing the PCR plate from the magnetic plate. Make sure that accumulated beads are not disturbed.
- 71 Add 120 μ l of 80 % EtOH, and incubate the beads for 30 seconds. Leave the plate in contact with the magnet as beads should not be resuspended during this washing step. Remove and discard the supernatant.
- 72 Repeat this washing step once for a total of two washes. Make sure to remove the supernatant completely.
- 73 Leave the plate in contact with the magnet, and let the beads dry for 5 - 10 minutes or until all ethanol has evaporated.
- 74 Add 20 μ l of Elution Buffer (EB) per well, remove the plate from the magnet, and resuspend the beads properly in EB. Incubate for 2 minutes at room temperature.
- 75 Place the plate onto a magnetic plate and let the beads collect for 2 - 5 minutes or until the supernatant is completely clear.
- 76 Transfer 15 - 17 μ l of the supernatant into a fresh PCR plate. Do not transfer any beads.
- 77 Libraries are now finished and ready for quality control.



78 Libraries can be stored at -20 °C at this point.