A quick and cost-effective method for DNA-free total RNA isolation using magnetic silica beads

Aniruddha Das[1,2,#], Debojyoti Das[1,2,#], Arundhati Das[1,2], and Amaresh C. Panda[1,*]
Institute of Life Sciences, Nalco Square, Bhubaneswar, Odisha, India;

[1] Institute of Life Sciences, Nalco Square, Bhubaneswar, Odisha, India;

[2] School of Biotechnology, KIIT University, Bhubaneswar, India;

[#] Contributed equally to this work

*Correspondence: Amaresh C Panda; Tel: +91 674 2304 314; Fax: +91 674 2300 728
Email: amaresh.panda@ils.res.in

Keywords: TRIzol; Phenol-chloroform; RT-qPCR; RNA isolation; gene expression

ABSTRACT

Current RNA purification methods widely use silica-based columns that allow quick isolation of high quality and right quantities of RNA. However, the major limitations include high cost, the requirement of different kits for small RNA isolation, genomic DNA contamination, and not being flexible. Here, we used the in-house RNA isolation reagent for cell lysis, followed by precipitation of RNA using isopropanol resulted in a similar quantity and quality of RNA compared to the commercial TRIzol. The commercial RNA isolation kits with silica-based columns recommend genomic DNA digestion during or after RNA purification adding time and cost to RNA purification. Here, we developed an optimized in-house protocol for isolating high-quality RNA free of genomic DNA contamination using magnetic silica beads without the need for DNase digestion. Additionally, our method purifies total RNA along with the small RNA fraction, including miRNAs, which usually require a separate kit for extraction. Additionally, the RNA prepared with our method was equally suitable for mRNA and miRNA expression analysis using RT-qPCR. Together, the in-house method of RNA isolation using a magnetic bead exhibited comparable or better total RNA extraction compared to commercial kits at a fraction of the cost and across various cells and tissues.

INTRODUCTION

The quality and quantity of RNA is the first and most crucial step for ensuring downstream gene expression analysis accuracy. The integrity of the total RNA is checked before its use in the downstream applications, including quantitative RT-PCR, microarrays, and RNA-sequencing, which are expensive, time-consuming, and labor-intensive [1]. The acceptable total RNA for downstream applications must fulfill a few requirements, including the RNA preparation must be free from protein, salts, alcohol, phenol, and genomic DNA (gDNA) which may interfere with downstream enzymatic reactions and gene expression analysis [1-4]. Additionally, the total RNA must be free of nucleases and stored appropriately to maintain integrity.

Various RNA extraction protocols have been developed to isolate high-quality RNA from a variety of cells and tissues. The RNA extraction methods broadly fall into three categories: organic extraction followed RNA precipitation, purification through a silica-membrane column, and silica-coated magnetic particles. Although the classical acid guanidinium thiocyanate-phenol-chloroform (AGPC) method of RNA extraction followed by RNA precipitation is the most commonly used method for decades, the isolated RNA samples are often contaminated with proteins, gDNA, organic solvents, and salts [1, 3-6]. However, the popular commercial kits use guanidinium salt-based lysis buffers without toxic organic solvents followed by RNA purification with silica column and magnetic silica beads [7, 8]. These commercial kits using the silica column are effective, relatively simple, and yield the right quantity of intact RNA. However, the silica columns often lose the small RNA fractions and recommend using specific RNA isolation kits to analyze small RNAs, including miRNAs. The magnetic bead-based RNA purification method is relatively new, and the kits are mostly available for specific applications such as viral RNA isolation [9, 10]. Additionally, the magnetic bead-based kits are often supplied with the automatic high-throughput RNA isolation system, which is very expensive. Additionally, both the silica column and magnetic bead-based purification methods often result in gDNA contamination in the purified RNA [1, 2, 11]. Since all these RNA isolation methods do not remove DNA, DNase treatment is recommended to remove the contaminating gDNA before proceeding to downstream applications, which add significant time and cost to the RNA purification [11].

In this work, we demonstrated a method where cells are lysed with the in-house prepared AGPC based cell lysis followed by depletion of the gDNA using a silica spin column. After gDNA depletion, the cell lysate was mixed with alcohol and allowed to bind silica-coated magnetic beads, followed by washing and elution of DNA-free total RNA. This method also isolates the small RNAs, including miRNAs, for expression analysis. In sum, the proposed method is a quick, practical, and cost-effective approach for isolating DNA-free total RNA, including small RNAs.

MATERIALS AND METHODS

HeLa, C2C12, and βTC6 cell culture

Human HeLa cell and mouse C2C12 and βTC6 cells were cultured in DMEM supplemented with 10/15% FBS and antibiotics. The cells were maintained in a 5% CO_2 humidified atmosphere at 37°C.

The cells were washed with 1X PBS and trypsinized with 0.25% Trypsin-EDTA, followed by pelleting the desired number of cells for RNA isolation.

RNA isolation reagents and kits

The commercial RNA isolation kits were procured from different companies, including TRIzol (cat no. 15596026, Thermo Fisher Scientific), miRNeasy Mini Kit (cat no. 217004; Qiagen), PureLink miRNA isolation kit (cat no. K157001; Thermo Fisher), and HiPurA™ Total RNA Miniprep Purification Kit (cat no. MB602-250PR; HiMedia). The on-column DNase I digestion set (cat no. DNASE70) was procured from Sigma for genomic DNA (gDNA) digestion during RNA purification. The gDNA depletion columns are EconoSpin silica columns (cat no. 1920) procured from Epoch life sciences. The in-house RNA isolation reagent (RIR) comprises 38 % phenol, 0.4 M ammonium thiocyanate, 0.8 M guanidine thiocyanate, 0.1 M sodium acetate, and 5 % glycerol. RNA wash buffer comprised of 10 mM Tris-HCl pH-7.5 and 75% ethanol. All reagents for buffer preparation were procured from Sigma, SRL, or HiMedia.

RNA isolation

The first method is the classical AGPC based RNA isolation method described previously [6, 12]. Briefly, HeLa cells were lysed in 500 µl TRIzol (Thermo Scientific) or in-house RIR by pipetting followed by mixing with 100 µl of chloroform. The mixture was centrifuged at 12,000x g for 10 min at 4 °C, followed by a collection of 200 µl of the aqueous layer in a new tube. The aqueous layer was mixed with 200 µl of isopropanol and incubated for 10 min at room temperature followed by centrifugation at 12,000x g for 10 min at 4 °C. The supernatant was discarded, and the pellet was washed with the wash buffer (75% ethanol) and centrifuged 12,000x g for 5 min at RT. The pellet was air-dried for 3-5 min followed by dissolving it in 50 µl nuclease-free water.

In the in-house method, two to five million cells were lysed in 500 µl RIR by vigorous pipetting followed by mixing with 100 µl of chloroform and centrifuged at 12,000x g for 10 min at 4 °C. The 200 µl aqueous layer was transferred into a new tube. The flowthrough was passed through a silica column by centrifugation at 12,000 xg for 1 min at RT. The flowthrough was mixed with 300 µl ethanol and 10 µl of MagPrep silica beads. The mix was incubated for 5 min at RT, followed by putting the tube on a magnetic stand to hold the beads to the side of the tube, and the supernatant was discarded. For DNase treatment, the beads were digested for 15 min with DNase I following the manufacturer's protocol. The beads were washed twice with 500 µl wash buffer. The wash buffer was discarded by holding the beads on a magnetic stand followed by keeping the tube at RT for 2-3 min to allow the ethanol to evaporate entirely. The beads were then mixed with 60 µl nuclease-free water and incubated for 2-3 min before putting the tube on the magnetic stand. The dissolved RNA was collected in a tube for quality and quantity assessment.

Also, we used the HiPurA™ Total RNA Miniprep Purification Kit (cat no. MB602-250PR; HiMedia), miRNeasy Mini Kit (cat no. 217004; Qiagen), and PureLink miRNA isolation kit (cat no. K157001; Thermo Fisher) for preparing the total RNA and miRNA following the manufacturer's instructions. Since the PureLink miRNA isolation kit isolates the long RNAs in

column 1 and small RNAs (miRNA) in column-2, we mixed equal volume of both RNAs and considered that as total RNA for downstream analysis.

Assessment of RNA quantity and quality

The quantity and quality of RNA samples were analyzed using NanoDrop 2000 or Multiskan Sky (Thermo Scientific) by determining the A_{260}/A_{280} and A_{260}/A_{230} ratios [13]. Additionally, the RNA was assessed for gDNA contamination in the Qubit4 using DNA broad range reagent. The RNA integrity and gDNA contamination were evaluated by resolving the RNA on an SYBR-Gold stained 1.2 % denaturing formaldehyde-agarose gel and visualizing under a UV transilluminator. Furthermore, RNA samples were analyzed on a TapeStation 4200 (Agilent) using an RNA ScreenTape to determine the RNA integrity number (RIN).

Reverse transcription followed by quantitative PCR (RT-qPCR) and agarose gel analysis

For long RNA analysis, one μg of total RNA isolated using different commercial kits and the MagPrep silica beads method were used to synthesize cDNA using High Capacity cDNA Reverse Transcription Kit (Thermo Scientific)following the manufacturer's protocol. The cDNA was diluted to 500 μl with nuclease-free water and 2X PowerUP SYBR Green Master Mix (Applied Biosystems) was used for real-time quantitative PCR (RT-qPCR) analysis using gene-specific primers for *Gapdh* mRNA (AACTTTGGCATTGTGGAAGG & GGATGCAGGGATGATGTTCT), Actb mRNA (TGTTACCAACTGGGACGACA & GGGGTGTTGAAGGTCTCAAA), circular *Samd4* (*circSamd4*; GCCCCTCTACTCTTCCTCGT & CAAAGGCAGGTGAGTTAGCA), and circular *Rras2* (circRras2; AAGGACCGTGATGAGTTTCC & GGTCATCGATCACACATTGC), and circular *Tcf20* (circTcf20; ACCATTACCCATGTGCCATT & CTCCTGTGGGTAGCTCTGCT). For analyzing the small RNA fractions, total RNA was used for cDNA synthesis with the Mir-X miRNA First-Strand Synthesis Kit (Clontech) following the manufacturer's protocol. The levels of small RNAs were analyzed by RT-qPCR analysis using small RNA specific primers for *U6* snRNA (provided in the kit), *U1* snRNA (TACTTACCTGGCAGGGGAGATAC & GAACGCAGTCCCCCACTAC, miR-27b (TTCACAGTGGCTAAGTTCTGC), and miR-181a-5p (AACATTCAACGCTGTCGGTGAGT), and the universal reverse primer provided in the kit. RT-qPCR amplification was performed at 95 °C for 2 minutes and subsequently 40 cycles of 95 °C for 2 s and 60 °C for 20 s on a QuantStudio 3/6 Real-Time PCR System (Thermo Fisher Scientific). The levels of different RNAs isolated with different methods were analyzed using their Ct values in RT-qPCR. Regular RT-PCR amplification was performed using the DreamTaq PCR mastermix with a cycle set up of initial 95 °C for 2 minutes, followed by 40 cycles of 95 °C for 2 s and 60 °C for 20 s on a thermal cycler. The amplified PCR products were resolved in an SYBR-Gold stained 2.5 % agarose gel and visualized by UV transilluminator gel documentation system. The relative RNA levels were calculated using the comparative Ct method.

RESULTS AND DISCUSSION

RNA preparation with in-house RIR and magnetic beads require gDNA removal

First, we wanted to analyze the quantity and quality of RNA extracted with the classical organic extraction method using the in-house RIR [5, 14]. We used HeLa cells to isolate total RNA using in-house RIR and TRIzol with the standard RNA isolation protocol using isopropanol precipitation of the aqueous layer. As expected, the quality and quantity obtained with the in-house RIR were comparable with the commercial TRIzol (**Figure 1**).

Figure 1. Comparison of in-house RNA Isolation Reagent (RIR) and with commercial TRIzol. A. The relative amount of total RNA recovered using RIR and TRIzol from HeLa cells. **B.** The purity of total RNA prepared using RIR and TRIzol from mentioned cell lines measured by the ratios of A_{260}/A_{280} and A_{260}/A_{230}. **C.** Total RNA prepared with RIR and TRIzol from mentioned cells resolved on a denaturing 1.2% agarose gel stained with SYBR-Gold. Data in panel A and B represent the mean ± SEM from 6 independent experiments.

Here, we wanted to use the aqueous layer from RIR to prepare RNA using magnetic silica beads. As shown in **Figure 2A**, the aqueous layer was incubated with MagPrep silica magnetic beads to adsorb the RNA, followed by digestion with on-column DNase I. The DNase digested beads were washed, and the bound RNA was eluted from the beads. HeLa and C2C12 cells were used for RNA extraction using MagPrep silica beads. However, DNase I treatment lead to a decrease in the total RNA extracted from HeLa and C2C12 cells (**Figure 2B**).

Figure 2. DNase digestion during the RNA preparation with MagPrep silica beads decreases the RNA yield. A. Flow chart of crucial steps of in-house optimized RNA extraction method using magnetic silica beads. **B.** Amount of total RNA recovered with or without DNase digestion from HeLa and C2C12 cells using an in-house optimized MagPrep silica beads RNA extraction protocol, as shown in panel A. Data represent the mean ± SEM from 3 independent experiments.

Removal of genomic DNA with silica column eliminates the time-consuming and costly DNase digestion step

As shown in **Figure 3A**, we passed the aqueous layer from the RIR-chloroform step through a silica column to remove the genomic DNA, followed by RNA purification using the MagPrep silica magnetic beads. Unlike reduction in RNA yield after DNase treatment, gDNA depletion using the silica column did not affect the RNA yield (**Supplementary Figure S1A**). Also, gDNA depletion completely removed the gDNA contamination from total RNA isolated using MagPrep silica beads from HeLa and C2C12 cells (**Supplementary Figure S1B**). We found that cell lysis with RIR followed by gDNA removal using the silica column quickly and in-expensively removes gDNA without the need for the expensive and time-consuming DNase digestion. Isolation of DNA-free total RNA using the gDNA depletion column followed by RNA isolation with MagPrep silica beads could isolate ~22 µg of total RNA from 5 million HeLa cells (**Figure 3B**). The isolation of DNA-free total RNA from HeLa, C2C12, and βTC6 cells showed clean 28s, 18s, and 5s rRNAs on the gel, suggesting isolation of good quality RNA for downstream analysis (**Figure 3C**).

Furthermore, the purity of DNA-free total RNA prepared with our in-house method using MagPrep silica beads was analyzed by calculating the ratio between the absorbance at 260 vs. 280 nm (A_{260}/A_{280}) and 260 vs. 230 nm (A_{260}/A_{230}). As shown in **Figure 3D**, all the 260/280 ratios for RNA prepared using silica beads from HeLa, C2C12, and βTC6 cells were around 2.0, suggesting that the eluent was RNA without DNA or protein contaminants. Additionally, the 260/230 ratios were between 2-2.3, indicating high-quality RNA with no contaminants that can be used in downstream applications [13]. Although the spectroscopy method can determine the quality and quantity of RNA, it cannot determine the integrity of the RNA. Our TapeStation analysis of RNA integrity showed little to no sign of degradation with RIN values ~ 9 for RNAs isolated from HeLa and C2C12 cells (**Figure 3E, F**). Total RNAs with RIN value more than 6.4 are recommended for transcriptome analysis using RNA-sequencing [15]. Together, The DNA-free total RNA prepared with the MagPrep beads are suitable for downstream analysis, including reverse transcription, RT-PCR, and RNA-sequencing.

RNA prepared with MagPrep silica beads is equally good for gene expression analysis compared to commercial RNA isolation

To compare the functionality and suitability of the RNAs extracted from C2C12 cells using our MagPrep silica beads, we prepared good quality RNAs using the TRIzol, miRNeasy Mini Kit (Qiagen), PureLink miRNA isolation kit (Thermo Fisher), and HiPurA™ Total RNA Miniprep Purification Kit (HiMedia) commercial kits (**Figure 4A**). Since the PureLink miRNA isolation kit prepares long and small RNA separately, we mixed an equal volume of long RNA and small RNA to obtain the total RNA. Additionally, the gel image indicated that RNA purified with our magnetic bead-based method successfully isolated the small RNAs, which was hardly visible in some of the RNA prepared with the commercial kit (**Figure 4A**). The total RNAs were reverse transcribed, and specific mRNA and circRNAs were successfully amplified, suggesting that the RNA prepared

with the in-house MagPrep RNA isolation method is as good as the commercial kit for RT-PCR analysis (**Figure 4B**). Furthermore, real-time (RT)-qPCR analysis of *Gapdh* mRNAs, *circSamd4, circRras2,* and *circTcf20* showed similar expression levels in RNA prepared with our MagPrep method compared to the commercial kits (**Figure 4C**). Furthermore, RT-qPCR analysis showed that the RNA prepared with the MagPrep method isolates miRNAs with similar or better efficiency than some commercial RNA isolation kits (**Figure 4D**). Together, total RNA preparation with the in-house method using magnetic silica beads could be used for the expression analysis of all types of RNAs such as mRNAs, circRNAs, and miRNAs.

Figure 3. RNA preparation with magnetic silica beads requires gDNA depletion with DNase or silica column. A. Flow chart of crucial steps of in-house optimized DNA-free total RNA extraction method using MagPrep silica beads. **B.** DNA-free total RNA recovered using MagPrep silica beads from 2 and 5 million HeLa cells. **C.** Denaturing 1.2% agarose gel depicting the total RNA from HeLa, C2C12, and βTC6 cells prepared using MagPrep silica RNA beads with or without gDNA depletion. **D.** The ratio of A_{260}/A_{280} and A_{260}/A_{230} of total RNA recovered using MagPrep silica beads from HeLa, C2C12, and βTC6 cells. **E.** Tapestation analysis showing representative electropherogram of DNA-free total RNA extracted from HeLa, C2C12, and βTC6 cells using MagPrep silica beads. The RIN values represent mean ± SEM for 3-6 different RNAs. Data in panel B and D represent the mean ± SEM from at least 3 independent experiments.

Figure 4. RNA prepared with MagPrep kit is equally good or better than the commercial RNA isolation kits for gene expression analysis. A. Denaturing 1.2% agarose gel depicting the total RNA or miRNA prepared with different commercial kits using silica columns. **B.** The RT-PCR products of *Gapdh*, *Actb*, *circSamd4, circRras2,* and *circTcf20* were visualized in an SYBR Gold stained 2.5 % agarose gel. **C.** RT-qPCR analysis and Ct values of mRNAs and circular RNAs including, *Gapdh*, *Actb*, *circSamd4, circRras2,* and *circTcf20* in 10 ng of total RNA prepared with different commercial kits and in-house MagPrep method. **D.** RT-qPCR analysis and Ct values of small RNAs, including *U1* snRNA, *U6* snRNA, *miR-27b*, and *miR-218-5p* in 2.5 ng of total RNA prepared with different commercial kits and in-house MagPrep method. Data in panels C and D represent the means ± SEM from three independent experiments.

CONCLUSIONS

The isolation of intact and pure RNA is critical for all downstream applications, including cDNA synthesis, PCR, real-time PCR, northern blotting, microarray, and RNA-sequencing [1]. There are three commonly used methods of RNA purification, such as acid-phenol extraction, silica column, and magnetic silica bead-based isolation. First, the classical organic extraction of RNA using TRIzol is the cost-effective and straightforward gold standard method for RNA isolation [5].

However, it has a few disadvantages, including the use of hazardous reagents, time-consuming, laborious protocol, and often contaminated with salts and other biomolecules [2, 3]. Second, the silica-based RNA purification column is the most commonly used in the commercial kits allowing isolation of high-quality and right amounts of RNA without using the toxic-reagents [7]. Third, the RNA isolation with magnetic silica beads is available for specific applications, including viral RNA isolation, automatic high-throughput applications [9]. The magnetic silica bead-based high-throughput RNA isolation kits require the expensive instrument and consumables to automate the process.

Furthermore, the major limitation of the silica column or magnetic bead-based commercial RNA isolation kits is the incomplete lysis, contamination of RNA with gDNA, and high cost [2, 3]. Since the downstream gene expression analysis application strictly demands DNA-free RNA, some commercial kits include the on-column DNase digestion step, and some recommend DNase treatment after purification. However, DNase treatment adds significant time and cost to the experiment. DNase treatment often leads to a loss of RNA amount and might affect RNA integrity due to high-temperature exposure during heat inactivation of commercial DNases. Furthermore, the silica columns in commercial kits often lose the small RNA fractions, and specific kits are available to isolate the small RNA factions, including miRNAs [7].

To overcome all these issues with the commercial kits, we developed a quick, efficient, flexible, and economical RNA extraction method. The user has the flexibility to choose the amount of TRIzol or RIR or commercial RNA lysis buffer depending on the input sample type and amount. The aqueous phase is passed through the gDNA depletion silica column removing the gDNA without using time-consuming and expensive DNase treatment procedure (**Figure 2**). Then, the lysate was mixed with ethanol and MagPrep silica beads to bind the RNA in the sample. Since the MagPrep silica beads bind more than 50 µg RNA per 10 µl beads (**Supplementary Figure S1C**), the user has the flexibility to choose the number of beads depending on the expected amount of RNA in the sample. The proposed method requires only one wash buffer and magnetic collection and resuspension steps that reduces the hands-on time and simplify the protocol. The commercial MagPrep silica beads can be replaced with in-house prepared silica-coated magnetic beads [16].

Furthermore, RNA purity analysis indicated that the RNA prepared with this method is free of gDNA, salt, and protein contaminant (**Figure 3**). The analysis of RNA integrity on gel indicated the isolation of high-quality, intact RNAs for downstream applications. The RT, followed by PCR, shows that the RNA prepared with our proposed method is compatible with downstream applications like reverse transcription and PCR (**Figure 4**). Furthermore, RT-qPCR analysis suggested similar or better recovery and amplification of smaller size RNAs with our MagPrep RNA than some commercial column-based kits. Furthermore, RNA samples isolated through our method could efficiently isolate small RNA fractions for miRNA analysis, which often requires a separate kit. Our RNA isolation procedure can be completed in about 30 min, which is significantly less than the commercial kits to get DNA-free RNA, including miRNAs. In summary,

we believe that the MagPrep RNA isolation method is by far a quick, effective, and cost-effective in-house method for the preparation of DNA-free total RNA compared to the existing commercial kits.

Advantages of using the MagPrep-RNA isolation method are,
1. This method is a quick and inexpensive method for isolating high-quality total RNA, including miRNAs for downstream expression analysis.
2. The total RNA does not need the time-consuming and expensive DNase treatment for gDNA removal.
3. The method takes ~30 min for DNA-free total RNA isolation, which is significantly quicker than commercially available kits with DNase treatment.
4. The method is scalable, depending on the amount of input sample.
5. The total RNA can be eluted in a small volume ≥ 20 µl to have a concentrated RNA.
6. This method can also be adapted to isolate RNA from yeast, bacteria. animal tissues, and plant samples.

Acknowledgments
DD, AD, and AD were affiliated to KIIT University, Bhubaneswar, India. The authors thank our colleagues at the Institute of Life Sciences, Bhubaneswar, for helpful discussions and suggestions to improve the method.

Funding
This research was supported by the Wellcome Trust/DBT India Alliance Intermediate Fellowship (IA/I/18/2/504017) provided to Amaresh Panda. DD, AD, and AD were supported by the Junior Research Fellowship from University Grant Commission, India.

Conflicts of Interest

The authors declare no conflict of interest.

Reference:

1. Tan, S.C. and B.C. Yiap, *DNA, RNA, and protein extraction: the past and the present.* J Biomed Biotechnol, 2009. **2009**: p. 574398.
2. Bustin, S.A. and T. Nolan, *Pitfalls of quantitative real-time reverse-transcription polymerase chain reaction.* J Biomol Tech, 2004. **15**(3): p. 155-66.
3. Wilson, I.G., *Inhibition and facilitation of nucleic acid amplification.* Appl Environ Microbiol, 1997. **63**(10): p. 3741-51.
4. Schrader, C., et al., *PCR inhibitors - occurrence, properties and removal.* J Appl Microbiol, 2012. **113**(5): p. 1014-26.
5. Rio, D.C., et al., *Purification of RNA using TRIzol (TRI reagent).* Cold Spring Harb Protoc, 2010. **2010**(6): p. pdb prot5439.
6. Chomczynski, P. and N. Sacchi, *The single-step method of RNA isolation by acid guanidinium thiocyanate-phenol-chloroform extraction: twenty-something years on.* Nat Protoc, 2006. **1**(2): p. 581-5.
7. Rio, D.C., et al., *Guidelines for the use of RNA purification kits.* Cold Spring Harb Protoc, 2010. **2010**(7): p. pdb ip79.
8. Yaffe, H., et al., *LogSpin: a simple, economical and fast method for RNA isolation from infected or healthy plants and other eukaryotic tissues.* BMC Res Notes, 2012. **5**: p. 45.
9. Oberacker, P., et al., *Bio-On-Magnetic-Beads (BOMB): Open platform for high-throughput nucleic acid extraction and manipulation.* PLoS Biol, 2019. **17**(1): p. e3000107.
10. Berensmeier, S., *Magnetic particles for the separation and purification of nucleic acids.* Appl Microbiol Biotechnol, 2006. **73**(3): p. 495-504.
11. Rio, D.C., et al., *Removal of DNA from RNA.* Cold Spring Harb Protoc, 2010. **2010**(6): p. pdb prot5443.
12. Chomczynski, P., *A reagent for the single-step simultaneous isolation of RNA, DNA and proteins from cell and tissue samples.* Biotechniques, 1993. **15**(3): p. 532-4, 536-7.
13. Manchester, K.L., *Use of UV methods for measurement of protein and nucleic acid concentrations.* Biotechniques, 1996. **20**(6): p. 968-70.
14. Walker, S.E. and J. Lorsch, *RNA purification--precipitation methods.* Methods Enzymol, 2013. **530**: p. 337-43.
15. Gallego Romero, I., et al., *RNA-seq: impact of RNA degradation on transcript quantification.* BMC Biol, 2014. **12**: p. 42.
16. Oberacker, P., et al., *Simple Synthesis of Functionalized Paramagnetic Beads for Nucleic Acid Purification and Manipulation.* Bio-protocol, 2019. **9**(20): p. e3394.