

Design and Fabrication of a Nanobiosensor for the Detection of Cell-Free Circulating miRNAS-LncRNAS-mRNAS Triad Grid

Pooja Ratre,^{||} Nazim Nazeer,^{||} Arpit Bhargava, Suresh Thareja, Rajnarayan Tiwari, Vinay Singh Raghuwanshi, and Pradyumna Kumar Mishra*



showed that the sensor could detect the target up to 1 fg concentration. After appropriate validation, the developed nanobiosensor might prove beneficial to characterizing and detecting aberrant disease-specific cell-free circulating miRNAs-lncRNAs-mRNAs.

INTRODUCTION

The understanding of our genome and associated regulatory processes has increased as a result of the discoveries about the significance of noncoding RNAs (ncRNAs) in physiological and pathological processes. ncRNAs, particularly micro RNAs (miRNAs) and long noncoding RNAs (lncRNAs), are known to have a crucial role in post-transcriptional gene control and cellular functions. Increasing scientific research of ncRNAs has opened up new avenues for investigating the ongoing aberrant mechanisms of different chronic and age-associated diseases and identifying the molecular targets for diagnosis and therapeutics.¹ It is now evident that the aberrant ceRNA network plays a vital role in the onset and progression of several noncommunicable diseases and malignancies, including breast cancer (BC), the primary source of cancer-related complications in women around the world.^{2,3} As per recent estimates, one in every 20 women worldwide and one in eight women in countries with high incomes are affected by BC. Although the average mortality rate from BC has fallen due to the recent developments in therapeutic procedures, early detection and immediate treatment are two crucial factors for the cure of BC.⁴ As the metastatic disease causes more than 90% of BC-related deaths, numerous types of research have been performed to discover new and reliable biomarkers for BC, One of the recently developed biomarkers include ncRNAs such as lncRNAs, miRNAs, and mRNA, which circulates in the human body and has the potential to reflect the different steps of disease development and progression.^{5,6}

These ncRNAs are interrelated and communicate in an intricate way to regulate the expression of vital genes. For instance, lncRNAs have potential miRNA-binding motifs, which can prohibit miRNAs from adhering to their appropriate mRNA targets.⁷ On the contrary, miRNAs can adversely influence the production of lncRNAs by interacting with the corresponding regions. Both miRNAs and lncRNAs can jointly influence the expression of the common mRNAs.⁸ In addition, by interfacing through miRNA regulators or interacting directly with miRNAs, lncRNAs can modify the expression of miRNA target genes, which may regulate the overall gene expression pattern and modulate the ongoing vital cellular processes.⁹ Moreover, such interaction and crosstalk offer new insights into the control of gene expression and disease etiology of aggressive and metastatic BCs. Therefore, it is essential to investigate the lncRNA-, miRNA-, and mRNA-based triple prognostic biomarker assay for early-stage detection of BC.¹⁰

Due to their high sensitivity, simplicity, and cost-effectiveness benefits, optical sensing technologies for detecting different disease-specific biomarkers have gained significant attention. These biosensors utilize the properties of light to identify and

Received: August 4, 2023 Accepted: October 3, 2023 Published: October 18, 2023





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s. no.	primer used	sequences $(\vec{5'3'})$
1.	miR-34a – FmiR-34a -R	CAGCTGTGAGTGTTTCTTTGCTGATTGCTTCCTTAC
2.	ANRIL – FANRIL – R	CAACATCCACCACTGGATCTTAACAAGCTTCGTATCCCCAATGAGATACA
3.	PTEN – FPTEN – R	TGGGCCCTGTACCATCCCAAGTTGTGGCAACCACAGCCATCGT
Uracil-Modified Oligonucleotide Sequence		
34a – CCGGTCGACACTCACAAAGAAACCGTGAGA		
ANRIL – GTTGTAGGTGGTGACCTAGAATTGTATCTG		
PTEN – ACCCGGGACATGGTAGGGTTCAGGAAACAT		

Table 1. Primer and Probe Detail Sequences Used for the Profiling

analyze biological substances and biochemical reactions in a given sample.^{11,12} The detection signals rely on the specific interaction between the target analytes and a recognition component immobilized on the sensor's surface. The binding of the target molecule with the recognition element generates optical signals, which quantitively correlate with the number of analytes present in the sample. These optical sensors are equipped with different light-yielding materials, such as nanomaterials, antibodies, and probes. One of the widely accepted nanomaterials used for optical sensing is quantum dots (QDs).^{13,14}

Compared with other metallic NPs, QDs have enhanced biocompatibility, stable fluorescence characteristics, and limited toxicity, making them preferred moieties for a number of biological applications. In addition, they possess outstanding optoelectric excitation-dependent fluorescent emission, sizetunable optical properties, and photoinduced electron transfer and minimal or no spectral overlap. Due to the important role of RNA in various diseases, exploration of their molecular interaction and transport mechanisms in live cells is highly important in the field of biomedicine. Fewer studies using QDbased electrochemical biosensors as RNA detection tools have been reported in the past few years. For instance, a DNAmediated silver electrochemical biosensor based on GQDs was reported to have potential applications in detecting micro-RNAs.¹⁵ Similarly, an analytical electrochemical RNA sensor with a reaction catalyst assisted by horseradish peroxidase was also assessed for detection.¹⁶ However, apart from electrochemical biosensing applications of QDs, the present study investigates the fluorescence aspects of QDs in simultaneous detection of three different RNA classes, i.e., mRNA, lncRNAs, and miRNAs. For this, we designed a specific probe against targeted RNA and assessed a triple sensor using three differentsized and color-based QDs to detect lncRNAs, miRNAs, and mRNAs in biological samples.

MATERIAL AND METHODS

RNA Isolation and Expression Analysis. Cell-free circulating RNAs (lncRNAs and mRNAs) were isolated from the plasma of blood samples using a QIAamp circulating nucleic acid isolation kit (Qiagen, Hilden, Germany) in the described manner (IRB No.: IEC/ICMR-NIREH/ITR-GIA/BC/2022-23/12-01-23). A PCR-based method was used to carry out expression analysis. First, Takara BioInc. PrimeScript first strand cDNA Synthesis kit (Shiga, Japan) synthesized cDNA from the extracted RNAs. The 4 μ L of synthesized cDNA was added to a 10 μ L qPCR Master Mix (Luna universal qPCR master mix, New England Biolabs, Ipswich, Massachusetts, USA). After the addition of specific primers of targeted mRNA (PTEN) and lncRNAs (ANRIL) (Table 1), a PCR reaction was performed on an Insta Q96 Real-Time PCR instrument (Himedia Laboratories, Mumbai, MH, India) under the proper thermal cycling

conditions of the target gene. For the expression analysis' internal control, GAPDH was used. Ct values were acquired, and dCt values were calculated following scheduled PCR cycles.¹⁷

miRNA Isolation and Expression Analysis. A Nucleospin miRNA plasma kit manufactured by Mangeley Nagel (Duren, Germany) was used to isolate miRNAs. The isolated miRNAs were then processed for polyadenylation using the Poly-A tailing kit from Invitrogen-Thermofischer Scientific in Greenville, NC, USA. The polyadenylated miRNAs were then processed into cDNA synthesis using a cDNA Synthesis kit (Takara BioInc. PrimeScriptTM first strand cDNA Synthesis kit, Shiga, Japan), mainly containing oligo-dT primers and reverse transcriptase. The 10 μ L of qPCR Master Mix was then added 4 μ L to cDNA strands with the primers of target miRNAs (34a), and the reaction mixture was placed into a series of amplification cycles on a thermocycler an Insta Q96 Real-Time PCR (Himedia Laboratories, Mumbai, MH, India). U6 was used as an internal control. Ct values were recorded, and dCt values calculated.^{18,19}

Preparation of Triple Sensor. Preparation of Biotinylated Oligonucleotide Probes. Selected complementary uracilmodified oligonucleotide probes for miRNAs, lncRNAs, and mRNAs (34a, ANRIL, and PTEN) were bought from Integrated DNA Technologies (Coralville, Iowa, USA) and attached at the 3' position of biotin by using a Pierce 3'-end biotinylation kit purchased from Thermo Fisher Scientific (Waltham, MA, USA). Briefly, uracil-modified oligonucleotide probes (50 pmol) were incubated for 5 min at 85 °C. Following incubation, probes were swiftly cooled and combined with the appropriate amounts of 10X RNA ligation buffer, biotinylated cytidine (Bis) phosphate, RNase inhibitor, T4 RNA Ligase enzyme, and 30% PEG. The mixture was incubated overnight at 4 °C. After incubation, nuclease-free water (NFW) and chloroform/isoamyl alcohol (in a 24:1 ratio) were added, and high-speed centrifugation was performed to remove the RNA ligase. The resultant layer of water was combined with ice-cold 100% ethanol, glycogen, and 5 M NaCl and incubated to precipitate for 1 h at -20 °C. The obtained pellet was carefully collected and washed with 70% ethanol. The pellet that formed in the next step was resuspended in NFW (20 μ L), set to air-dry, and then kept at -20 °C.²⁰

Preparation of the Streptavidin-Conjugated QDs. Three different QDs, both organic and inorganic types (Qdots 525, QD 705, and GQDs), were procured from ThermoFisher Scientific, Waltham, MA, USA, and Merch KGaA, Darmstadt, Germany. The carboxyl group on the surface of these three QDs was first activated using carbodiimide chemistry comprising EDC (1-ethyl-3-[3-(dimethylamino)propyl] carbodiimide hydrochloride) and NHS (*N*-hydroxysuccinimide) from Thermo-Fisher Scientific (Waltham, MA, USA). First, an aliquot of QDs was prepared by mixing 10 μ L of each QDs in three different tubes containing 250 μ L of 1× PBS, followed by the addition of 25 μ L of EDC (1 mg/mL solution) and 25 μ L of NHS (1 mg/mL solution) for surface charge activation. The above mixture

was then incubated for 1 h at room temperature (RT). A 100 μ L aliquot of streptavidin (1 mg/mL) was added to the previously activated QDs and incubated for 20 min at RT. The activated carboxyl groups on the surface of QDs were then coupled with amine groups of streptavidin via the formation of an amide linkage. To improve the efficacy of the conjugation, a second aliquot of 25 μ L of EDC and NHS was added to the tube containing the QDs streptavidin conjugate. This was followed by 1 h of vortexing at RT. Centrifuging at 6000 rpm for 15 min separated the streptavidin coated QDs from the free QDs. The streptavidin coated QDs conjugates were pelleted and stored at 4 °C until further use.

Preparation of Final Nanoconjugates. After streptavidin was bound to the surface of QDs, biotinylated oligonucleotide probes were attached via the streptavidin—biotin coupling reaction to detect targeted miRNAs, lncRNAs, and mRNAs. For conjugation, the Streptavidin-attached QDs were added to 200 μ L of 1× PBS (pH 7.4), vortexed, and combined with the biotinylated probes in the following ratios: 3:1, 1:1, and 1:3. The reaction mixture was then vortexed for 1 h at RT and incubated overnight at 4 °C for further investigation. The prepared conjugates have target complementary strands on their surface for desired targets (34a, ANRIL, and PTEN) in the isolated sample (Figure 1).²¹

Targeted Detection of miRNA, LncRNA, and mRNA In Biological Samples. Applicability of Developed Triple Nanohybrid. Using fluorometry (Spark multimode microplate reader, TECAN, Seestrasse 103, Mannedorf, Switzerland), we detected three RNA markers in plasma. To achieve this, plasma samples were first filtered through 0.45 m filters to remove mesenchymal cells and cellular debris such as extracellular micro vesicles, lipid rafts, apoptotic/necrotic bodies, and exosomes and then with 0.22 m filters to obtain apparent plasma free of any substantial genetic debris. The 10 μ L of each nanoconjugate (QDs 705, QDs 525, and GQDs) were mixed with 90 μ L of filtered plasma in a microplate reader, followed by addition of different intercalating (propidium iodide and SYBR Safe) dye in each tube, the solution was incubated for 10 min and then analyzed in fluorometry under different channels. The method was repeated for the isolated RNA samples. These samples were examined and compared to a blank solution (NFW), which served as the negative control and did not include any capture or detector components. To assess the repeatability of the nanohybrid, experiments were carried out in triplicate in 15 samples. After the detection applicability assessment of the designed nanoconjugates, the results were validated by performing probe RT-PCR and agarose gel electrophoresis.

Microscopic Evaluation of Developed Triple Sensor. Fluorescence microscopy was used to evaluate the fluorescence properties of prepared nanoconjugates. The nanoconjugates mixed with amplicons of all different targeted RNAs and intercalating dye were added in the same volume as that in the applicability assessment. Different filters according to the excitation and emission of QDs used (FITC, DAPI, and TRITC) were selected. Two fluorescence pictures were uploaded to the ImageJ platform for the colocalization studies. The JaCop plugin was then used to analyze overlapping between the previous two images, which included Pearson's coefficient, overlap coefficient, intensity correlation analysis, Manders' coefficients M1 and M2, and Van Steensel's cross-correlation coefficient.

Selectivity Analysis of the Triple Sensors. The fluorometry method evaluated the prepared conjugates' selectivity toward



Figure 1. Image showing the fabrication of a triple nanohybrid system (A) showing the preparation of the streptavidin-conjugated with QDs; (B) biotinylation of oligonucleotide probes; (C) conjugation of streptavidin-conjugated QDs with biotinylated probes and sensing.

targeted RNAs. The 10 μ L prepared nanoconjugates were mixed in 1 μ L of PCR amplicons of the target RNAs (34a, ANRIL, PTEN) and 1 μ L of intercalating dye, and the fluorescence intensity was recorded using fluorometry (Spark multimode microplate reader, TECAN, Seestrasse 103, Mannedorf, Switzerland). The nanoconjugates were next tested against PCR amplicons of other nonspecific RNAs (7a, 29a, Gas-5, PVT, PIK3CA, and FOXO3).

Sensitivity Analysis of the Triple Sensors. To check the sensitivity and limit of detection (LOD) of developed nanoconjugates, different dilutions of targeted RNAs amplicons (miRNAs, lncRNAs, and mRNAs), ranging from 1 μ g to 1 fg, were prepared in NFW. The 10 μ L of prepared nanoconjugates were mixed in 1 μ L in 90 μ L NFW of all prepared dilutions, and the difference in the fluorescence intensity was recorded using fluorometry (Spark multimode microplate reader, TECAN, Seestrasse 103, Mannedorf, Switzerland).

Statistical Analysis. Each result's mean and standard error were calculated (n = 3). The GraphPad Prism Version 5.03 was

used for statistical analysis. One-way or two-way variance analysis was used to calculate the statistical differences.

RESULTS AND DISCUSSION

Biomedical sensing is one of the most promising areas that emerged in the field of nanosensors, as it offers novel methods for rapid and accurate diagnosis of different life-threatening diseases, including BCs. With continuous evolution in molecular biology, various disease-specific biomarkers have been identified. The clinical diagnosis of BCs is now focused on developing novel analytical methods capable of specific and parallel detection of biomarkers.²² The competitive endogenous RNAs (ceRNAs) are essential transcripts that interact with each other to create a regulatory network that is vital for regulating various ongoing biological processes. The interaction between lncRNAmiRNA-mRNA has been suggested to play a significant role in the onset of diseases, including BCs. Substantial investigations have shown that these RNAs' regulatory networks can regulate the oncogenesis process.⁴ Considering this fact, we have developed fluorescent-based triple sensors for the detection of miRNAs, lncRNAs, and mRNAs collectively. The existing methods for detection of these RNAs require additional labeling protocols, amplification techniques, and data normalization processes; however, the method discussed here is simple, specific, and has the ability to perform simultaneous detection of the three triad biomarkers (ccf-miRNA, lncRNAs, and mRNAs) without involving the additional complicated labeling protocols.

Initially, expression profiling of the target RNAs was performed using the qPCR method in 15 samples. The graph shown here represents an upregulated expression of 34a, ANRIL, and PTEN in most samples (Figure 2). Following the



PTEN targeted RNAs in biological samples through quantitative realtime PCR.

expression analysis, the triple sensor comprising three different QDs (QD-705, QD-525, and GQDs) was designed and assessed for their ability to detect biomarkers (34a, ANRIL, and PTEN) involved in the progression of BC. For the fabrication of the triple sensor, a set of probes highly specific toward target molecules was selected and biotinylated. This was followed by the surface activation of QDs and their subsequent labeling with streptavidin. The three different QDs used showed red (QDs 705), green (QDs 525), and blue (GQDs) color fluorescence with probe-bound mRNA (PTEN), lncRNAs (ANRIL), and miRNA (34a) respectively. The size-dependent spectrum characteristics of the QDs, along with the excellent quantum yield, strong molar extinction coefficient, and broad range of absorption spectrum, uniform fluorescence spectrum, significant Stokes shift, and strong photochemical stability are few of the





Figure 3. Graph showing the size distribution pattern of QDs.

The binding of the target RNA with the biotinylated probe was determined using intercalating dyes (Sybr safe and propidium iodide). Moreover, the presence of nanoparticle clusters attached to streptavidin and their ability to bind with the target were evaluated using fluorescence microscopy. To confirm their location on the same plain, a merged or colocalized image was taken, which shows in the integrated yellow and green color (Figure 4).

The quantitative change in the fluorescence intensity of functionalized QDs upon binding with the target was analyzed by fluorimetry, which showed that the intensity initially decreases when it binds with streptavidin (Figure 5). This may be due to the phenomenon of fluorescence quenching. In general, it has been shown that the interaction involving the surface atoms of QDs and any molecules or atoms on the surface causes the recombination of electron-hole pairs to be inhibited, which results in fluorescence quenching of QDs. The electronhole recombination process can be hampered, and changes to the surface or surroundings of QDs can reduce the fluorescence intensity. It is also known that photoinduced electron transfer from an excited QDs to a nearby compound with an energy intermediate to the valence and conduction band edge states causes the fluorescence quenching of QDs and results in a decrease in fluorescence intensity.²

The developed nanoconjugates were then analyzed for selective target detection and applicability in filtered plasma samples spiked with target RNA amplicons. The nanoconjugates have oligonucleotide probes on the surface, which are complementary to the targeted miRNA, lncRNA, and mRNA and ensure selective binding. The efficient assimilation of targeted RNAs led to the increased fluorescence intensity of RNA-attached nanoconjugates in contrast to the nanoconjugates attached with streptavidin and probe. Yet, a surge in the fluorescence intensity of intercalating dye was noted, which may be associated with the expression and amount of RNAs present in the plasma sample (Figure 6). The validation of nanoconjugates was performed after detection of target. The PCR results show the amplification of a selected probe against all three targets in different samples. For further confirmation the agarose gel electrophoresis was performed, and the desired bands appeared in the gel images (supplementary Figure 1).

After confirming the applicability of the developed sensor, we further looked for the sensitivity of all three nanoconjugates. The



Figure 4. Figure depicting the fluorescence microscopic images evaluation of QDs (A) QDs 705 cluster in red color, green SYBR safe dye confirming the presence of uracil modified probe of mRNA PTEN and merged image of both the images in yellow with colocalized scatter plot. (B) QDs 525 cluster in green color, red PI confirming presence of lncRNAs ANRIL probe, and merged image in yellow with colocalized scatter plot. (C) GQDs cluster in blue color, green SYBR safe dye confirming the presence of miRNA 34a and merged image in blue-green color with colocalized scatter plot.



Figure 5. Graph shows the fluorescent comparison analysis of Plain QDs and conjugated QDs in fluorometry.

nanoconjugates were added in different ranges of target RNAs ranging from 1 μ g to 1fg. The results observed via fluorometry, developed nanosensors that can analyze the target up to 1fg concentration and showed the limit of detection (LOD) value of 0.9882 fg (QDs 705), 0.314 fg (QDs 525), and 0.313 fg (GQDs) (Figure 7).

In order to perform specificity assessment, the nanoconjugates were added in the well containing targeted RNA amplicon and with nonspecific RNA amplicons. The fluorescence intensity was highest in the case of the selective RNA amplicon, and low or no fluorescence signal was observed in the nonselective amplicon. The increase in fluorescence intensity on hybridization with specific RNA was observed in all three conjugates due to the fluorescence quenching effect of QDs, and the same increase in fluorescence intensity was observed on addition of intercalating dyes due to their binding with major grooves on nucleotides (Figure 8).

CONCLUSIONS AND TRANSLATIONAL PERSPECTIVE

In today's world of advanced medicine and intelligent healthcare systems, nanotechnology has led to the development of a wide range of laboratory-based tests that have accelerated their broader translational purpose leading to the early detection of noncommunicable diseases such as BC.²⁵ In this scenario, making timely medical decisions based on fast, efficient, reliable, and reproducible diagnostic data has become critical in quality health management and patient care. The increasing use of nanobiosensors will help capture disease-associated marks and provide new insights into signaling events and pathways at the level of both the genome and epigenome. ncRNAs and their interactions have significantly affected BC development, metastasis, and treatment. Different miRNAs, lncRNAs, and mRNAs act alone or in combination on mRNA alone or as competing endogenous RNA (CeRNA) networks for regulation, as they compete for miRNAs at the post-transcriptional stage. Understanding the processes underlying these connections, which impact the expression of genes linked to BC, can help create intelligent analytical techniques for early disease identification. The expanding knowledge about the role of miRNA-lncRNAmRNA (as a triad grid) in the development and progression of BCs has made them excellent candidates as disease biomarkers. With the rapid growth of QDs in cancer imaging and diagnosis, there has been an increased interest in developing effective cancer diagnostic procedures using these nanomoieties. As we learn more about the significance of ncRNAs and their interactions in BC, we can better understand how to use QDs for early detection. We developed herein a unique triple sensor for the simultaneous assessment of miRNAs-lncRNAs-mRNAs

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Figure 7. Graph demonstrating the sensitivity of developed nanoconjugates upon hybridization with targeted RNAs in different concentrations (A) Relative fluorescence unit (RFU) of QDs noted in fluorometry (B) RFU of intercalating dyes.



Figure 8. Graph demonstrating the specific binding efficiency of developed nanoconjugates toward specific RNA sequences in comparison to other nonspecific RNAs. (A) QDs 705 specificity toward PTEN, (B) QDs 525 Specificity toward ANRIL, and (C) GQDs specificity toward 34a.

using three QDs (QD-705, QD-525, and GQDs). The developed sensor efficiently detected the target RNAs in isolated and plasma samples. We assume that after appropriate validation, the assay might prove beneficial to characterize and detect aberrant cell-free circulating miRNA-lncRNA-mRNA.

ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/acsomega.3c05718.

Expression profiling of targeted oligonucleotide probe, i.e., miRNA (34a), lncRNAs (ANRIL), and mRNA (PTEN) (PDF)

AUTHOR INFORMATION

Corresponding Author

Pradyumna Kumar Mishra – Division of Environmental Biotechnology, Genetics & Molecular Biology, ICMR-National Institute for Research in Environmental Health, Bhopal 462030, India; o orcid.org/0000-0002-0795-2819; Email: pkm 8bh@yahoo.co.uk

Authors

- **Pooja Ratre** Division of Environmental Biotechnology, Genetics & Molecular Biology, ICMR-National Institute for Research in Environmental Health, Bhopal 462030, India
- Nazim Nazeer Division of Environmental Biotechnology, Genetics & Molecular Biology, ICMR-National Institute for Research in Environmental Health, Bhopal 462030, India
- Arpit Bhargava Division of Environmental Biotechnology, Genetics & Molecular Biology, ICMR-National Institute for Research in Environmental Health, Bhopal 462030, India; Faculty of Science, Ram Krishna Dharmarth Foundation University, Bhopal 462030, India
- Suresh Thareja Department of Pharmaceutical Sciences and Natural Products, Central University of Punjab, Bathinda 151001, India
- Rajnarayan Tiwari Division of Environmental Biotechnology, Genetics & Molecular Biology, ICMR-National Institute for Research in Environmental Health, Bhopal 462030, India
- Vinay Singh Raghuwanshi Division of Environmental Biotechnology, Genetics & Molecular Biology, ICMR-National Institute for Research in Environmental Health, Bhopal 462030, India

Complete contact information is available at: https://pubs.acs.org/10.1021/acsomega.3c05718

Author Contributions

^{II}P.R. and N.N. contributed equally.

Notes

The authors declare no competing financial interest.

ACKNOWLEDGMENTS

The authors are thankful to the Indian Council of Medical Research (ICMR), Department of Health Research (DHR), Ministry of Health & Family Welfare (MoHFW), Government of India, New Delhi, for extramural project (DHR-GIA) funding support to the laboratory of Prof. (Dr.) Pradyumna Kumar Mishra.

ABBREVIATIONS:

LncRNAs-long noncoding ribonucleic acids miRNAs-micro ribonucleic acids GQDs-graphene quantum dots NHS-N-hydroxysuccinimide EDC-1-ethyl-3-(3-(dimethylamino)propyl) carbodiimide QDs-quantum dots ID-intercalating dye PI-propidium iodide NFW-nuclease free water LOD-limit of detection RFU-relative fluorescence unit

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NOTE ADDED AFTER ASAP PUBLICATION

This paper published ASAP on October 18, 2023 with an error in the title. The error was corrected and the paper reposted on October 20, 2023.