Check for updates

OPEN ACCESS

EDITED AND REVIEWED BY Jean Marie François, Institut Biotechnologique de Toulouse (INSA), France

*CORRESPONDENCE Jaimie Marie Stewart, Ims@seas.ucla.edu Kirill A. Afonin, Ims@kafonin@charlotte.edu

RECEIVED 23 December 2024 ACCEPTED 20 January 2025 PUBLISHED 04 February 2025

CITATION

Stewart JM, Yingling YG, Afonin KA, Hubé F and Kataoka N (2025) Editorial: Recent advancements in RNA technologies, diagnostics, and therapeutics. *Front. Bioeng. Biotechnol.* 13:1550225. doi: 10.3389/fbioe.2025.1550225

COPYRIGHT

© 2025 Stewart, Yingling, Afonin, Hubé and Kataoka. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

Editorial: Recent advancements in RNA technologies, diagnostics, and therapeutics

Jaimie Marie Stewart¹*, Yaroslava G. Yingling², Kirill A. Afonin³*, Florent Hubé⁴ and Naoyuki Kataoka⁵

¹Department of Bioengineering, University of California, Los Angeles, CA, United States, ²Department of Materials Science and Engineering, North Carolina State University, Raleigh, NC, United States, ³Nanoscale Science Program, Department of Chemistry, University of North Carolina Charlotte, Charlotte, NC, United States, ⁴Laboratoire Biologie du Développement, Institut de Biologie Paris-Seine, Transgenerational Epigenetics and Small RNA Biology, Sorbonne Université, CNRS, Paris, France, ⁵Laboratory of Cellular Biochemistry, Department of Animal Resource Sciences, Graduate School of Agriculture and Life Sciences, The University of Tokyo, Tokyo, Japan

KEYWORDS

RNA technologies, miRNA, circRNA, RNAi, theranostics, tissue engineering, RNA circuits, synthetic biology

Editorial on the Research Topic

Recent advancements in RNA technologies, diagnostics, and therapeutics

Introduction

RNA technology is an emerging field that exploits the unique structural and functional properties of RNA to build nanoscale structures and regulate complex biological systems (Stewart, 2024). RNA has been shown to assemble into structures with various shapes, sizes, and complexities, enabling applications in molecular sensing, drug delivery, immunomodulation, and cellular activity regulation (Chandler et al., 2021). This foundational work shows the significant potential of RNA molecules and their chemical analogs as a biomaterial for developing personalized diagnostic and therapeutic applications, as evidenced by numerous *in vitro* and *in vivo* studies and exemplified by several FDA approved formulations. However, critical challenges such as nuclease stability, targeted delivery of RNA therapies, regulation of their immune response, and lowering detection limits that must be further addressed to fully translate RNA nanotechnology into clinical applications.

This Research Topic highlights recent advancements and innovative work in RNA technologies for diagnostics and therapeutics of various classes of RNAs. This Research Topic features six review and research articles curated by international leaders in the fields of nucleic acid technologies, drug delivery, and computational studies. All manuscripts present a wide range of innovative technologies encompassing the design and optimization of gene therapies, production of RNAs, logic gating, tissue engineering and verifications of new therapeutic targets.

MicroRNA for regenerative medicine

Over 30 years ago, the first microRNA (miRNA) was identified in Caenorhabditis elegans (Lee et al., 1993; Wightman et al., 1993) and provided insights into how RNA can regulate gene expression. MicroRNAs (miRNAs) are small non-coding RNAs, ~22 nucleotides in length, that function as key regulators of gene expression. Through their regulatory activity, miRNAs can modulate several biological processes, including cellular differentiation, proliferation, and apoptosis (O'Brien et al., 2018). miRNAs are a promising strategy for advancing regenerative medicine, however, it is imperative to further elucidate miRNA regulatory mechanisms and pathways for clinical applications. Shahin et al. investigated the regulatory role and regenerative functions of microRNA (miR-155) in skin wound repair. The authors performed computational analysis of differentially expressed miRNAs in adipose-derived mesenchymal stem cells (AD-MSCs) and keratinocytes. hsa-miR-155 was identified and experimentally validated as having an enhanced immunomodulatory effect of AD-MSCs through regulating key wound healing proteins FGF2, FGF7, CCL2, and VCAM1. Castañón-Cortés et al. highlighted recent advancements in integrating miRNA into tissue-engineering scaffolds for optimized tissue repair and regeneration. The authors summarize the use of tissue-engineered scaffolds in combination with miRNAs applied to skin, musculoskeletal, nervous, and cardiovascular systems. Additionally, the study emphasizes the need for further experiments to understand fundamental miRNA-mediated mechanisms of action for specific tissue regeneration and minimal off-target effects. Lastly, the current challenges of cellular uptake and localized delivery are outlined.

Optimizing design approaches for RNA synthesis and gene expression

Engineering molecular machinery and RNA sequences are a practical approach for efficient RNA synthesis and gene regulation capabilities. Circular RNAs (circRNAs) are a highly stable class of non-coding RNAs with diverse biological roles, including acting as miRNA sponges, transcriptional regulation, protein recruitment, and enhancing protein activity (Kristensen et al., 2019). He et al. developed a one-pot process for circRNA synthesis by introducing specific mutations to T7 RNA polymerase (RNAP) to yield a thermostable variant that combines transcription and cyclization in a single reaction. The authors used consensus and folding free energy calculations for hotspot selections to construct a multisite mutant T7 RNAP. The engineered polymerase demonstrated stable activity at 45°C for over an hour, introducing new techniques for efficient circRNA production via a one-pot transcription and cyclization. RNA sequence design is another approach that can be used to tune gene expression. Codons are trinucleotide sequences that determine a specific amino acid. Due to the redundancy of the genetic code, synonymous codons can encode the same amino acid. However, these synonymous codons are not used equally, causing a codon bias (Novoa and de Pouplana, 2012). Paremskaia et al. presented a comprehensive analysis of current metrics for codon optimization for clinical use for gene therapy. The authors categorize methods to generate optimized sequences variants and protocols to experimentally verify mRNA stability and protein expression. The study concludes with persistent challenges of unintended effects on protein function and complexities in evaluating codon effectiveness.

Synthetic circuits

RNA molecules such as RNA aptamers, ribozymes, and riboswitches can organize to form networks that can perform complex functions including gene regulation, signal amplification, and logic operations for diagnostics and therapeutics (Pfeifer et al., 2023). Tian et al. engineered Boolean logic gates in the yeast Saccharomyces cerevisiae by reintroducing the naturally absent RNA interference (RNAi) pathway. The authors found that promoter leakage of pGAL1 inhibited logic behavior. Promoter leakage was reduced when the DNA fragment was placed between the two promoter regions, vastly improving the circuit reliability and performance. Armstrong and Isalan discussed using RNA-based circuits for the development of bacterial theranostics. The authors emphasized the use of mRNA and riboregulatory to construct circuits and program bacteria to sense and respond to physiochemical signals. The study closes with current issues of safety and proving a promising outlook for developing precise and versatile therapeutic systems.

Outlook

RNA technologies offer innovative approaches for therapeutic and diagnostic applications. However, broad application of RNA-based technologies in clinical settings will require focused research efforts in key areas: increasing target specificity and delivery efficiencies, improving stability and functional retention at ambient temperatures, ensuring precise patient-specific regulation of toxicities, and decreasing production and handling costs. Addressing these critical factors will be essential in realizing the full potential of RNA technologies for widespread clinical use.

Author contributions

JS: Writing-original draft, Writing-review and editing. YY: Writing-review and editing. KA: Writing-original draft, Writing-review and editing. FH: Writing-review and editing. NK: Writing-review and editing.

Funding

The author(s) declare that financial support was received for the research, authorship, and/or publication of this article. This work was supported by the Alfred P. Sloan Foundation through Award G-2023-21044 (JS) and the National Institute of General Medical

Sciences of the National Institutes of Health under Award Number R35GM139587 (KA). The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institutes of Health.

Conflict of interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

The author(s) declared that they were an editorial board member of Frontiers, at the time of submission. This had no impact on the peer review process and the final decision.

References

Chandler, M., Johnson, B., Khisamutdinov, E., Dobrovolskaia, M. A., Sztuba-Solinska, J., Salem, A. K., et al. (2021). The International Society of RNA Nanotechnology and Nanomedicine (ISRNN): the present and future of the burgeoning field. *ACS Nano* 15 (11), 16957–16973. doi:10.1021/acsnano.0c10240

Kristensen, L. S., Andersen, M. S., Stagsted, L. V., Ebbesen, K. K., Hansen, T. B., and Kjems, J. (2019). The biogenesis, biology and characterization of circular RNAs. *Nat. Rev. Genet.* 20, 675–691. doi:10.1038/s41576-019-0158-7

Lee, R. C., Feinbaum, R. L., and Ambros, V. (1993). The C. elegans heterochronic gene lin-4 encodes small RNAs with antisense complementarity to lin-14C. elegans heterochronic gene lin-4 encodes small RNAs with antisense complementarity to lin-14. *Cell* 75, 843–854. doi:10.1016/0092-8674(93)90529-y

Novoa, E. M., and de Pouplana, L. R. (2012). Speeding with control: codon usage, tRNAs, and ribosomes. 28, 574–581. doi:10.1016/j.tig.2012.07.006

Generative AI statement

The author(s) declare that no Generative AI was used in the creation of this manuscript.

Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

O'Brien, J., Hayder, H., Zayed, Y., and Peng, C. (2018). Overview of microRNA biogenesis, mechanisms of actions, and circulation. *Front. Endocrinol. (Lausanne).* 9, 402. doi:10.3389/fendo.2018.00402

Pfeifer, B. A., Beitelshees, M., Hill, A., Bassett, J., and Jones, C. H. (2023). Harnessing synthetic biology for advancing RNA therapeutics and vaccine design. *npj Syst. Biol. Appl.* 9, 60. doi:10.1038/s41540-023-00323-3

Stewart, J. M. (2024). RNA nanotechnology on the horizon: self-assembly, chemical modifications, and functional applications. *Curr. Opin. Chem. Biol.* 81, 102479. doi:10. 1016/j.cbpa.2024.102479

Wightman, B., Ha, I., and Ruvkun, G. (1993). Posttranscriptional regulation of the heterochronic gene lin-14 by lin-4 mediates temporal pattern formation in C. elegans, *C. elegans. Cell* 75, 855–862. doi:10.1016/0092-8674(93)90530-4