

Enhancing mRNA Analysis Using Ultra-wide Pore Size Exclusion Chromatography Columns

Haoran David*

Department of Bioprocess Engineering and Biotechnology, São Paulo State University, Araraquara, Brazil

Introduction

The analysis of Messenger Rna (mRNA) is critical for understanding gene expression and its regulation in biological systems. With the increasing use of mRNA-based therapies, such as the COVID-19 vaccines, the need for accurate and efficient mRNA analysis has become more pressing. Messenger RNA serves as the intermediary between DNA and protein synthesis, and its stability, processing, and translation play crucial roles in cellular function and disease progression. Advances in analytical techniques for mRNA are essential to ensure the quality, stability, and efficacy of mRNA-based therapeutics and to explore the underlying molecular mechanisms in cellular processes. One of the challenges in mRNA analysis is its structural complexity. mRNA molecules are composed of a sequence of nucleotides that encode proteins, with additional regulatory elements such as the 5' cap, 3' poly-A tail, and untranslated regions. These elements influence mRNA stability, translation efficiency, and cellular localization. Analyzing the size, purity, and integrity of mRNA, along with its modifications, is essential for understanding its functional properties and ensuring its proper delivery in therapeutic applications. Traditional methods, such as gel electrophoresis or reverse transcription-quantitative polymerase chain reaction have limitations when it comes to accurately measuring the size distribution or integrity of mRNA, particularly in complex mixtures or in the presence of degradation products. Consequently, there is a growing demand for improved analytical techniques that can provide more detailed and reliable insights into the characteristics of mRNA.

Description

Ultra-wide pore Size Exclusion Chromatography (SEC) columns represent one such advancement in the field of mRNA analysis. Size exclusion chromatography, also known as gel filtration chromatography, separates molecules based on their size. Traditional SEC columns typically utilize narrower pore sizes, which are ideal for separating smaller molecules such as proteins or low-molecular-weight compounds. However, mRNA molecules, especially those in therapeutic formulations, can be quite large and heterogeneous, requiring the use of ultra-wide pore size columns to accurately separate and analyze them. Ultra-wide pore SEC columns provide a larger pore size matrix, allowing for the separation of macromolecules like mRNA, oligonucleotides, and their complexes, based on their size and molecular weight [1]. This method has the potential to improve the accuracy and resolution of mRNA analysis, offering valuable insights into its structural integrity, size distribution, and other key characteristics. One of the key benefits of using ultra-wide pore SEC columns for mRNA analysis is their ability to resolve different mRNA species in complex mixtures. mRNA samples can often be heterogeneous, containing various isoforms, truncated versions,

or degraded fragments. The use of ultra-wide pore columns allows for better separation of these different species, enabling researchers to assess the full range of mRNA species present in a given sample. This is particularly important when analyzing mRNA-based therapeutics, where the presence of impurities or degradation products can significantly impact the efficacy of the final product. By ensuring that only intact, full-length mRNA is present in the therapeutic formulation, the purity and quality of the product can be guaranteed, minimizing potential risks to patient safety [2].

Another advantage of ultra-wide pore SEC columns is their ability to analyze mRNA in its native form, without the need for extensive sample preparation or denaturation. This is crucial for understanding the functional properties of mRNA and its interactions with other molecules in vivo. For example, mRNA's 5' cap and 3' poly-A tail play critical roles in its stability, translation efficiency, and cellular localization. By using ultra-wide pore SEC, researchers can analyze the mRNA in its native, unaltered state, providing more accurate information about how these elements contribute to mRNA function. This method also allows for the characterization of mRNA-lipid nanoparticle (LNP) formulations, which are commonly used for mRNA delivery in therapeutics. LNPs protect mRNA from degradation and facilitate its delivery into cells, but their size, charge, and composition must be carefully optimized for maximum therapeutic efficacy. Ultra-wide pore SEC can help evaluate the stability and integrity of mRNA-LNP complexes, ensuring that they are well-formed and capable of efficiently delivering the mRNA payload to target cells. The use of ultra-wide pore SEC columns also enables the analysis of mRNA stability under various conditions. mRNA molecules are inherently unstable and prone to degradation by exonucleases, which can shorten their poly-A tails or remove nucleotides from their coding regions. This instability presents a significant challenge for the development of mRNA-based therapeutics, where mRNA needs to be delivered intact to target cells to exert its therapeutic effects [3].

By monitoring the degradation of mRNA in real-time using ultra-wide pore SEC, researchers can identify conditions or formulations that enhance mRNA stability, thereby improving the quality of mRNA-based products. This can lead to the development of more effective mRNA vaccines and therapies, with greater shelf life and more reliable performance in vivo. Moreover, ultra-wide pore SEC can be combined with other analytical techniques, such as mass spectrometry, fluorescence detection, or multi-angle light scattering, to provide even more comprehensive insights into the properties of mRNA. For instance, mass spectrometry can be used to identify specific modifications or impurities within the mRNA, such as base modifications or chemical contaminants introduced during synthesis or formulation. Fluorescence detection can be employed to monitor the binding of mRNA to specific probes or interactions with other molecules, such as proteins or RNA-binding factors. Multi-angle light scattering can provide additional information about the size and shape of mRNA molecules in solution, offering insights into their conformation and structural integrity. By combining these methods with ultra-wide pore SEC, researchers can obtain a more holistic view of mRNA quality and functionality, which is essential for ensuring the success of mRNA-based therapeutics [4].

In addition to improving mRNA analysis, ultra-wide pore SEC can also be beneficial in the development of RNA-based therapeutics. mRNA vaccines have emerged as a revolutionary tool in the fight against infectious diseases, most notably the COVID-19 pandemic. However, the development of safe and effective mRNA vaccines requires careful optimization of the mRNA formulation, including the choice of lipid nanoparticles, stabilizers, and other excipients. Ultra-wide pore SEC can be used to evaluate the formulation of mRNA-LNP complexes, ensuring that they maintain their stability and integrity

*Address for Correspondence: Haoran David, Department of Bioprocess Engineering and Biotechnology, São Paulo State University, Araraquara, Brazil, E-mail: haorandavid@gmail.com

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during storage, transport, and administration. This optimization process is crucial for ensuring that the mRNA vaccines are both safe and effective when administered to patients. Furthermore, ultra-wide pore SEC has applications beyond mRNA vaccines, extending to other forms of RNA-based therapies, such as messenger RNA therapeutics for cancer, gene editing, and protein replacement. These therapies require the precise delivery of RNA molecules to target cells, and ultra-wide pore SEC can aid in the characterization of RNA delivery systems, ensuring that they function as intended. The ability to separate and analyze RNA molecules based on their size and integrity provides valuable information for optimizing RNA-based therapeutics, reducing the risk of off-target effects, and enhancing therapeutic outcomes [5].

Conclusion

Despite the many advantages of ultra-wide pore SEC for mRNA analysis, there are some challenges to consider. One of the limitations of this technique is its relatively low resolution compared to other high-throughput methods, such as next-generation sequencing or microarray analysis. While ultra-wide pore SEC provides valuable insights into mRNA size, integrity, and purity, it does not offer detailed information about the sequence or structure of the mRNA. Thus, it is often used in combination with other complementary techniques to obtain a more comprehensive understanding of mRNA characteristics. Additionally, ultra-wide pore SEC columns can be expensive and may require specialized equipment and expertise, which can limit their accessibility for some research labs or clinical settings. ultra-wide pore size exclusion chromatography represents a powerful tool for enhancing the analysis of mRNA, particularly in the context of mRNA-based therapeutics. By providing a more accurate and detailed assessment of mRNA size, integrity, and stability, this technique can contribute to the development of higher-quality mRNA-based products. Its ability to separate mRNA species, evaluate mRNA-LNP complexes, and assess mRNA stability under various conditions makes it an essential tool for optimizing RNA-based therapeutics and ensuring their success in clinical applications. As the field of RNA therapeutics continues to evolve, the integration of ultra-wide pore SEC with other advanced analytical techniques will play a critical role in advancing our understanding of mRNA and its potential as a therapeutic agent.

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Conflict of Interest

None.

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