

Current issues and potential solutions for the electrospinning of major polysaccharides and proteins: A review

Murtaza Haider Syed^a, *Md Maksudur Rahman Khan*^b, *Mior Ahmad Khushairi Mohd Zahari*^{a*}, *Mohammad Dalour Hossen Beg*^c, *Norhayati Abdullah*^{a*}

^a Faculty of Chemical and Process Engineering Technology, Universiti Malaysia Pahang Al-Sultan Abdullah, Gambang, Pahang, Malaysia

^b Petroleum and Chemical Engineering Programme Area, Faculty of Engineering, Universiti Teknologi Brunei, Gadong BE1410, Brunei

^c School of Engineering, University of Waikato, New Zealand

ABSTRACT

Biopolymers, especially polysaccharides and proteins, are the promising green replacement for petroleum based polymers. Due to their innate properties, they are effectively used in biomedical applications, especially tissue engineering, wound healing, and drug delivery. The fibrous morphology of biopolymers is essentially required for the effectiveness in these biomedical applications. Electrospinning (ES) is the most advanced and robust method to fabricate nanofibers (NFs) and provides a complete solution to the conventional methods issues. However, the major issues regarding fabricating polysaccharides and protein nanofibers using ES include poor electrospinnability, lack of desired fundamental properties for a specific application by a single biopolymer, and insolubility among common solvents. The current review provides the main strategies for effective electrospinning of the major biopolymers. The key strategies include blending major biopolymers with suitable biopolymers and optimizing the solvent system. A systematic literature review was done to provide the optimized solvent system of the major biopolymers along with their best possible biopolymeric blend for ES. The review also highlights the fundamental issues with the commercialization of ES based biomedical products and provides future directions to improve the fabrication of biopolymeric nanofibers.

KEYWORDS

Biopolymers; Electrospinning; Nanofibers; Biological macromolecules

ACKNOWLEDGEMENTS

The authors would like to thank the Ministry of Higher Education Malaysia for providing financial support under Fundamental Research Grant Scheme (FRGS) No. FRGS/1/2019/TK05/UMP/02/13 (University reference RDU1901176), Universiti Malaysia Pahang as well as for the additional financial support under PGRS grant PGRS220361 and Doctoral research scheme (DRS).