



Therapeutic Potential of Mesenchymal Stem Cell-derived Exosomes Isolated with the PS Affinity Method

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I. INTRODUCTION

Mesenchymal -derived Stem Cells (MSCs) have attracted much attention as a potential cellular therapeutic tool due to their regenerative and immunomodulatory capabilities. Their therapeutic effects are largely mediated by paracrine factors including Extracellular Vesicles (EVs) that are nanosized lipid bilayer particles. EVs derived from mesenchymal stem cells (MSC) carry lipids, proteins, and nucleic acids derived from their producing cells are emerging as a promising therapeutic strategy in regenerative medicine.

II. OBJECTIVE

To examine whether MSC-derived EVs isolated by the PS method have similar functions as those isolated by ultracentrifugation, we isolated EVs from the supernatant of bone marrow-derived MSCs and analyzed the effects of anti-inflammatory and antifibrosis.





IV. MATERIALS AND METHODS





Figure 1. (A) Principle of PS sandwich ELISA and Exosome concentrations determined by CD9, CD63 or CD81 antibodies. (B) Particle size distribution analysis using NanoSight. (C) Western blot analysis on the expression of the exosome markers CD9, CD63, and CD81.

2. BM-MSC-derived EVs decreases the expression of proinflammatory cytokines and increases the anti-inflammatory cytokines in LPS-treated human PBMC-monocyte.



Figure 2. The relative expression levels of the inflammatoryassociated genes were evaluated by qRT-PCR after stimulation by LPS and then subsequently after MSC-EVs treatment with LPS: TNF- α , IL-6, and TGF β .

VI. CONCLUSION AND DISCUSSION

We have developed the Exosome Isolation Kit using the PS affinity. Our study showed that the PS affinity method can isolate MSC-derived EVs with high recovery efficiency while maintain the high activity of them. We hope the PS affinity method would be a powerful tool for the development of future MSC-derived EVs therapies.



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CD9 25kDa

CD63 50kDa

CD81 25kDa

BM-MSC-derived EVs decreases the expression of fibrosis

Figure 3. The relative expression levels of the fibrocystic genes were

evaluated by gRT-PCR after stimulation by TGF^β and then

subsequently after MSC-EVs treatment with TGFB: Collagen III,

markers in TGF_β-treated lung fibroblastic cells, TIG3.

αSMA, and Collagen V.