Supplemental Figures

TLR3-/4-Priming Differentially Promote Ca²⁺ Signaling and Cytokine Expression and Ca²⁺-Dependently Augment Cytokine Release in hMSCs

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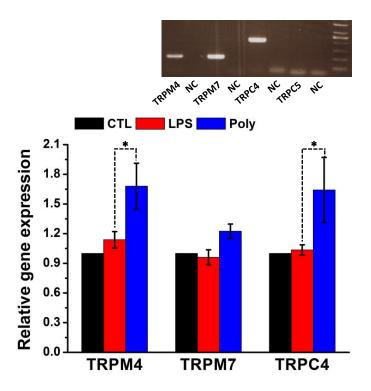


Figure S1. Representative RT-PCR blot illustrating mRNA expression of TRPM4, TRPM7 and TRPC4 in control hMSCs. NC denotes negative control with distilled water. Real time RT-PCR quantification showing relative mRNA expression levels of TRPM4, TRPM7 and TRPC4 in control (CTL; n = 5), LPS- (n = 5) and poly (I:C)-treated hMSCs (n = 5).

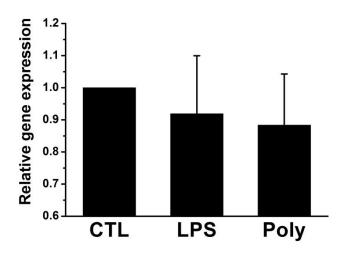


Figure S2. Real time RT-PCR quantification showing relative mRNA expression levels of MaxiK in control (CTL; n = 3), LPS- (n = 3) and poly (I:C)-treated hMSCs (n = 3).

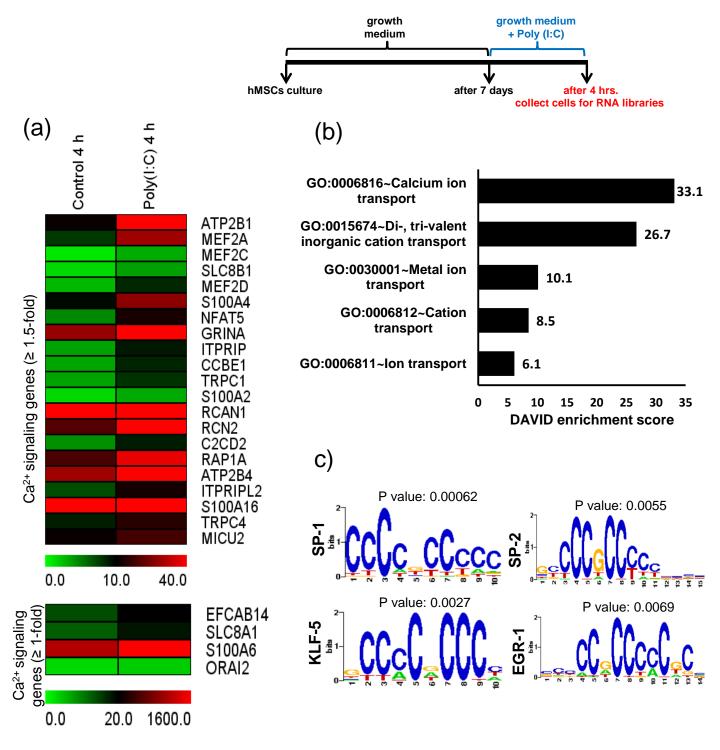


Figure S3. Transcriptomic analysis of the Poly (I:C)-stimulated Ca²⁺ signaling gene expression in hMSCs

(a) A heat map represents differential expression of Ca²⁺ signaling gene expression at 4 h after Poly (I:C) stimulation in hMSCs. (b) Gene ontology analysis of the functional annotations that were associated with Ca²⁺ signaling gene expression at 4 h after Poly (I:C) stimulation in hMSCs. (c) Patterns of transcription factor motif enrichments within the promoters of the Ca²⁺ signaling genes in Poly (I:C)-stimulated hMSCs.

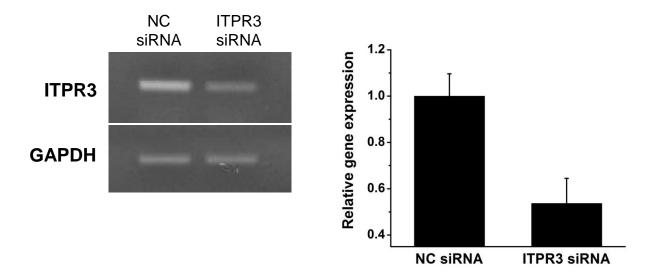


Figure S4. Representative RT-PCR blot and real time RT-PCR quantification showing relative mRNA expression levels of ITPR3 in scambled siRNA control (NC, n = 2) and ITPR3 siRNA treated hMSCs for 48 hours (n = 2).