

Supplementary Information

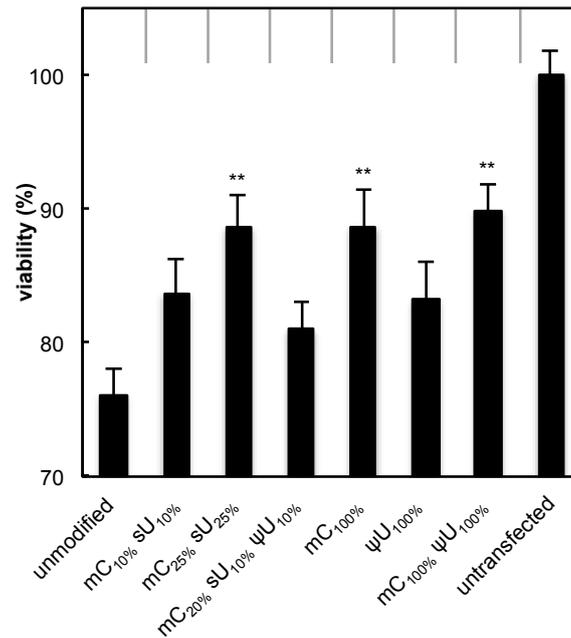


Figure S1. Cell viability after mRNA introduction to RAW264.7 cells. mRNA encoding GLuc was introduced to RAW264.7 cells using Lipofectamine™ LTX. After 4 h, cell viability was measured. Data are presented as the mean \pm standard error of the mean (S.E.M.) ($N = 6$). Statistical significance was assessed by one-way analysis of variance (ANOVA) followed by Dunnett's test. **, $p < 0.01$ versus unmodified mRNA.

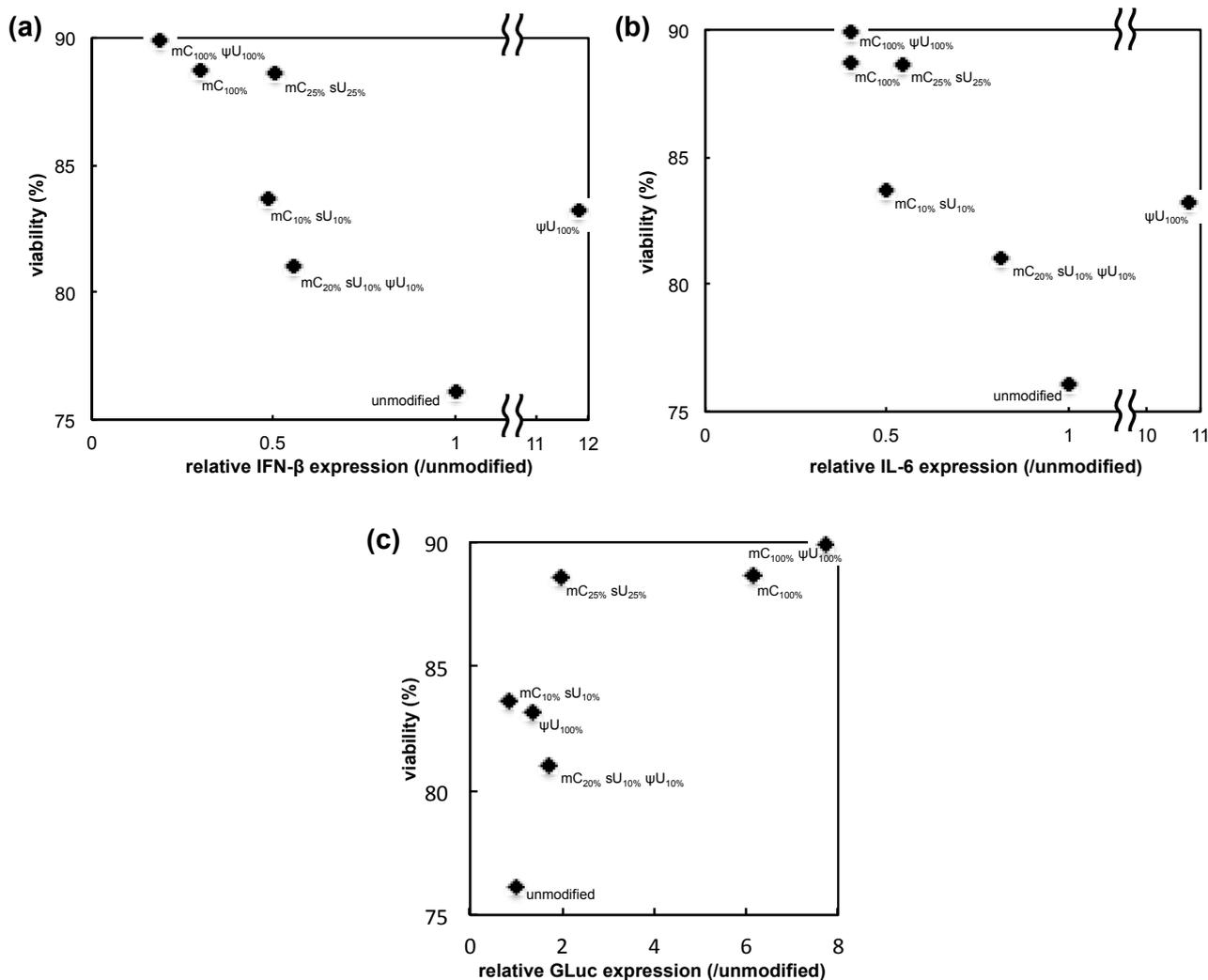


Figure S2. Correlation between cell viability, inflammatory responses, and GLuc expression. mRNA encoding GLuc was introduced to RAW264.7 cells using Lipofectamine™ LTX. After 4 h, cell viability (see Figure S1), expression of inflammatory molecules (see Figure 1), and GLuc expression (see Figure 2a) were measured. **(a,b)** Correlation between inflammatory responses and cell viability. **(c)** Correlation between GLuc expression and cell viability. The x-axis shows the expression of **(a)** interferon-β (IFN-β), **(b)** interleukin-6 (IL-6), and **(c)** GLuc. The y-axis showed the cell viability.

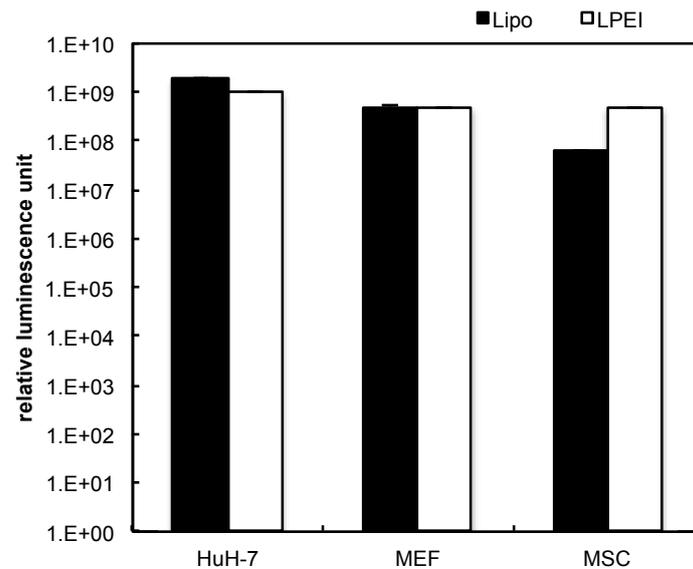


Figure S3. Comparison of GLuc expression efficiency by transfection of unmodified mRNA between two different transfection reagents. Unmodified mRNA was introduced to HuH-7 cells, MEFs and MSCs, using two types of transfection reagents, Lipofectamine™ LTX (Lipo, black bars) and linear polyethyleneimine (LPEI, white bars). The data are used in Figure 4 to calculate the “relative GLuc expression” by the modified mRNAs. Data are presented as the mean \pm standard error of the mean (S.E.M.) ($N = 6$).

© 2015 by the authors; licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution license (<http://creativecommons.org/licenses/by/4.0/>).