



Supplementary materials

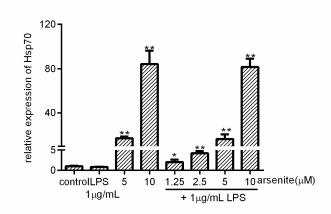


Figure S1. The expression of Hsp70 by 030D cells after arsenite and LPS stimulation. 030D cells were treated with various concentration of arsenite (1.25 μ M to 10 μ M) for 16 h as indicated. Then they were left unstimulated or stimulated with 1 μ g/mL LPS for 6 h. mRNA expression of Hsp70 was measured by qPCR. Data are shown as the mean \pm SD and are representative of three independent experiments. *p < 0.05, **p < 0.01, vs LPS alone group.

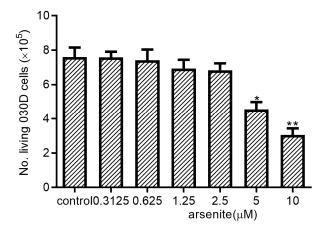


Figure S2. Effects of arsenite on the metabolic activity of 030D cells. 030D cells (2× 105 cells/well) were incubated with the indicated concentrations of arsenite for 24 h. Living 030D cells were counted. Untreated cells were used as control. Data are shown as the mean \pm SD and representative of three independent experiments. *p < 0.05, **p<0.01, vs control group.

Hsp70 (ENSCAFG00000000640) (3333 bp)

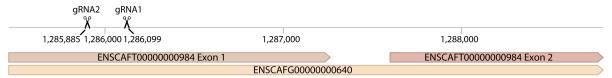


Figure S3.. Targeting sites of gRNA

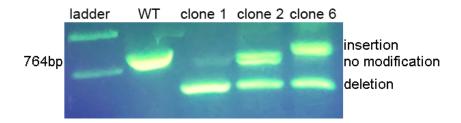


Figure S4. Validation of Hsp70 gene modification in 030D cell clones. Genomic DNA from these cells was isolated. PCR was performed using Hsp70 primers that cover the restriction sites and PCR products were visualized by gel electrophoresis on 1% agarose gel. Clone 1, clone 2 and clone 6 are gene-modified. WT = wild type.

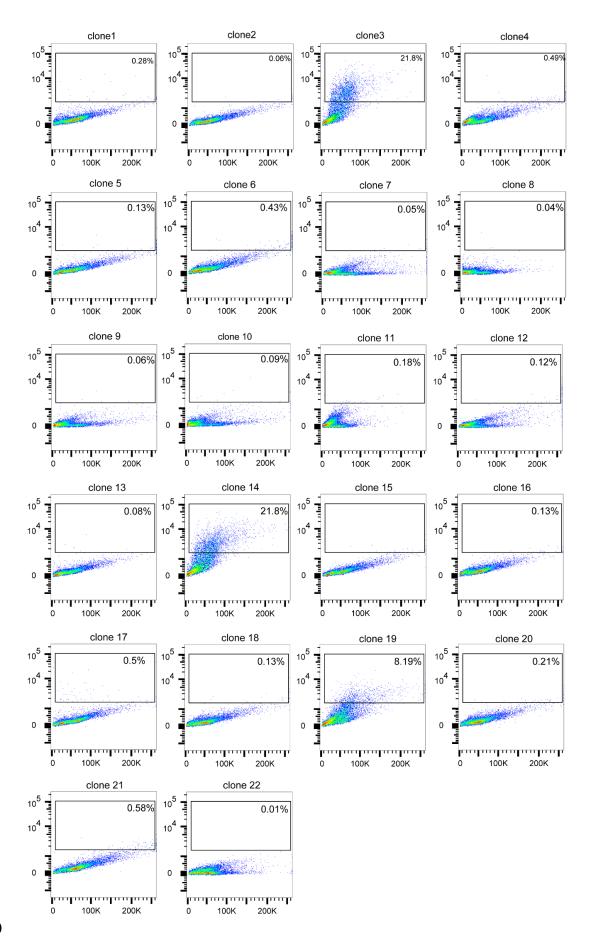


Figure S5. Flow cytometry analysis of gene-modified cell clones (1-22) to evaluate the expression of Hsp70 in 030D cells.

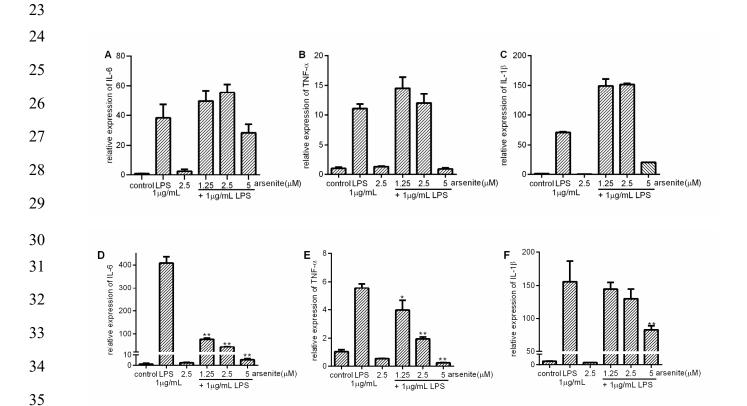


Figure S6. The effect of inducible Hsp70 on pro-inflammatory cytokine expression. A second Hsp70 knockout 030D clone, clone5 (**A**, **B**, **C**) and corresponding CRISPER/Cas9-treated wild type-like clones, clone 3 (**D**, **E**, **F**) were treated with different concentrations (1.25, 2.5 or 5 μ M) of arsenite or without for 16 h and then exposed to LPS. The cells were harvested after 6 h of LPS exposure. qPCR were performed to detect the expression of IL-6 (**A**), IL-1 β (**B**) and TNF- α (**C**) on clone 5 and clone 3 (**D**, **E**, and **F**, **respectively**). Data are shown as the mean \pm SD and are representative of three independent experiments. *P < 0.05, **P < 0.01, and ***P < 0.001, vs LPS alone group

44

45

46

47

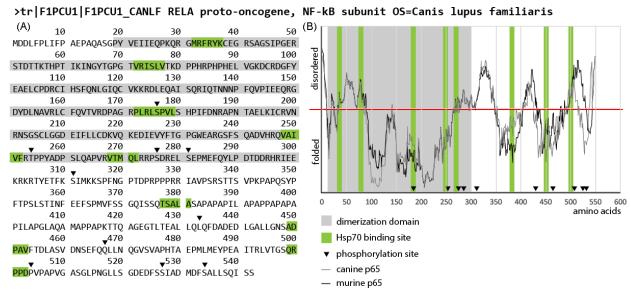


Figure S7. The prediction of the Hsp70 binding sites on p65 (RELA) protein. (**A**) shows the amino acid sequence of canine p65. (**B**) shows that the amino acid sequence of murine p65 (UniProt ID: Q04207) is strongly similar to canine p65 (UniProt ID: F1PCU1). Black triangle representative reported phosphorylation sites; green: predicted Hsp70 binding sites on p65.