

Supporting information

Table S1. Primers used for CHIP-qPCR.

	Forward	Reverse
iNOS	GGA CTTGGG ACCAGAAAGAGGTG	GCCATCCAGAGAGTTGTTTTTGC
MMP3	AAAATAGAGTAGCAGAGGCAGGTA	AGAGTGGTGGCAGTGATGTGAA
MMP13	CAACCATGGGGCTCAATCCT	CTTACGTGGCGACTTTTTCTTTTC

Table S2. Primers used for quantitative RT-PCR.

Gene	Forward	Reverse
<i>iNOS</i>	TGGCCATGGAACATCCCAA	AGGATGTTGTAGCGCTGGAC
<i>MMP3</i>	TGGGCCAGGGATTAATGGAG	CCGAGTCAGGTCTGTGAGTG
<i>MMP13</i>	AGGAGCATGGCGACTTCTAC	AGACCTAAGGAGTGGCCGAA
<i>SOX9</i>	GGAAGTCGGTGAAGAACGGG	CCTTGAAGATGGCGTTGGGG
<i>Col2a1</i>	TGACACTGGGACTGTCCTCT	TCTCCGAAGGGGATCTCAGG
<i>Aggrecan</i>	CAGTCGAAACAGCCACCTCC	TCTGTCTCCTTGCAGGTCCC
<i>18S rRNA</i>	GTAACCCGTTGAACCCATT	CCATCCAATCGGTAGTAGCG

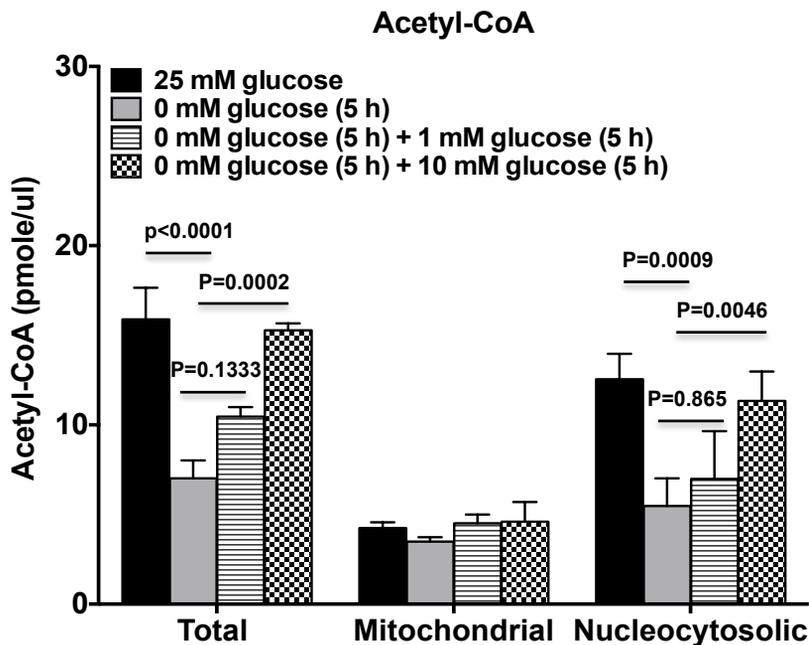


Figure S1. Effect of glucose concentration on acetyl-CoA generation in chondrocytes. Cultured primary human knee chondrocytes in DMEM media containing 25 mM glucose were subjected to glucose starvation in DMEM media without glucose for 5 hours, and then placed back in the DMEM media containing 1 or 10 mM glucose for 5 hours. The amounts of total and compartmental (mitochondrial and nucleocytosolic) acetyl-CoA were determined. Two-way ANOVA followed by Bonferroni's *post hoc* test was used for statistical data analysis. P values represent comparisons of the mean \pm SD.