

SUPPLEMENTAL MATERIAL

Table S1. Expression analysis of apoptosis-related genes. Expression of apoptosis-related genes as determined by DNA microarray are shown, which correspond to the top panel of Figure 6. See Excel file.

Table S2. Expression analysis of cell cycle-related genes. Expression of cell cycle-related genes as determined by DNA microarray are shown, which correspond to the second panel of Figure 6. See Excel file.

Table S3. Expression analysis of inflammatory response-related genes. Expression of inflammatory response-related genes as determined by DNA microarray are shown, which correspond to the third panel of Figure 6. See Excel file.

Table S4. Expression analysis of matrix metalloproteinase genes. Expression of genes for matrix metalloproteinases as determined by DNA microarray are shown, which correspond to the bottom panel of Figure 6. See Excel file.