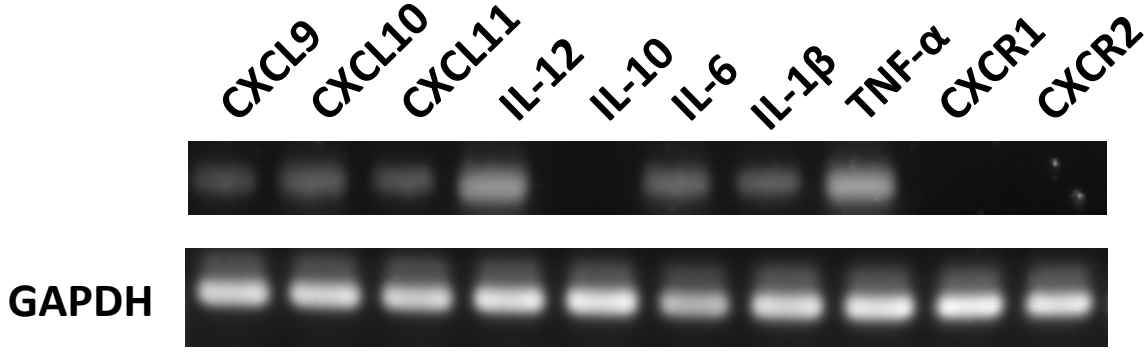

Additional file 3: mRNA level of M1 and M2 macrophage markers in RAW 264.7 cells.

RT-PCR analysis was performed for identification of RAW 264.7 cells phenotype. Briefly, RAW 264.7 cells were seeded in 6-well plate and incubated 24 h without stimulation. Cells were harvested using scraper and total RNA were extracted. GAPDH was used as the normalization control. The primer sequences are shown in methods and additional file 4.