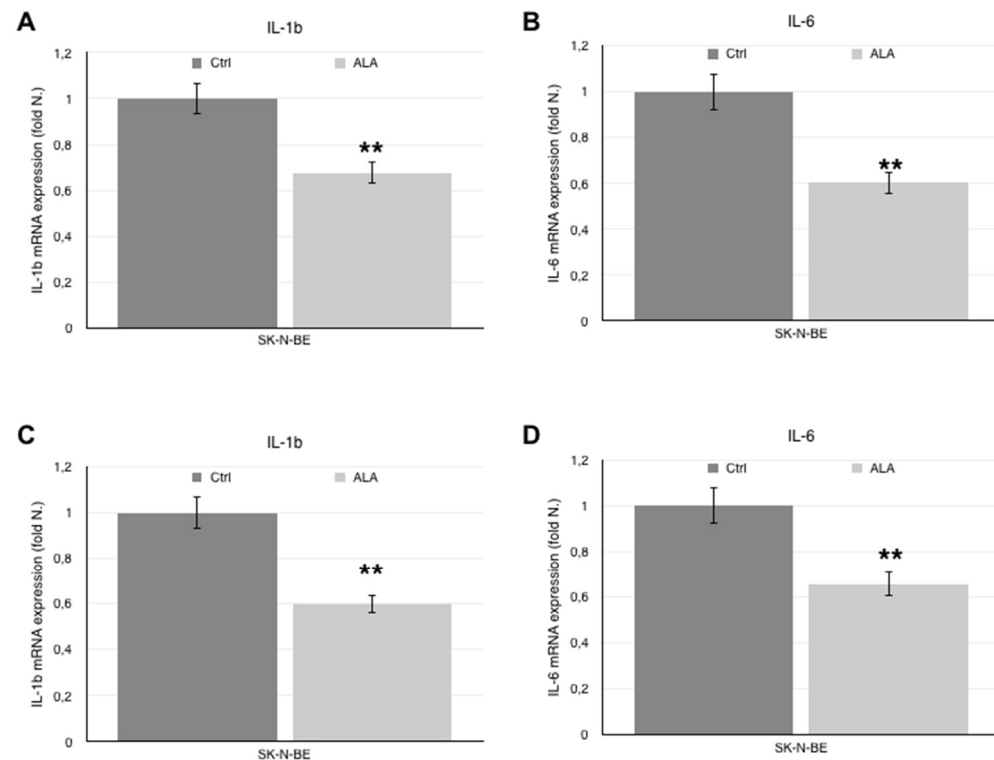




Supplementary Figure 1: mRNA expression levels, as determined by Real Time-PCR, in SK-N-BE cells treated with control (dark grey columns) and ALA supplemented (light grey columns) medium using GAPDH (A, B) and 18S (C, D) for normalization. Histograms indicate the mean value $\pm$ s.e.m. **: $p < 0.001$ vs. Ctrl.