



Figure S2. Expression of MAAs in MIC⁻ and MIC⁺ subsets. qRT-PCR analysis on 50 sorted cells (a) CD20^{-/+}, (b) CD24^{-/+}, (c) CD133^{-/+} and (d) ALDH^{low/high} cells, after pre-amplification with the AmpliGrid system; GOI – gene of interest; EF2 was used as a reference gene; each bar represents mean + SEM; * - p<0.05, *** - p<0.001 versus MIC⁻ cells.