Supplemental Figure 1. Phenotypic characterization of blood monocyte subsets

CCR2, CD206, CD163, CD86, TLR2, CD36 and MHC2 protein expressions at the surface of classical (CD14$^{\text{high}}$ CD16$^-$), intermediate (CD14$^{\text{high}}$ CD16$^{\text{low}}$) and non-classical (CD14$^{\text{low}}$ CD16$^{\text{high}}$) monocytes. Graphs represent geometric mean fluorescence quantification for the indicated protein. Results correspond to mean ± SEM. *$p < 0.05$, **$p < 0.01$ and ***$p < 0.001$ compared to classical monocyte subset in healthy subjects. $p$ values were determined using an analysis of variance followed by a Bonferroni’s multiple comparisons test. N.D = Non Detected.