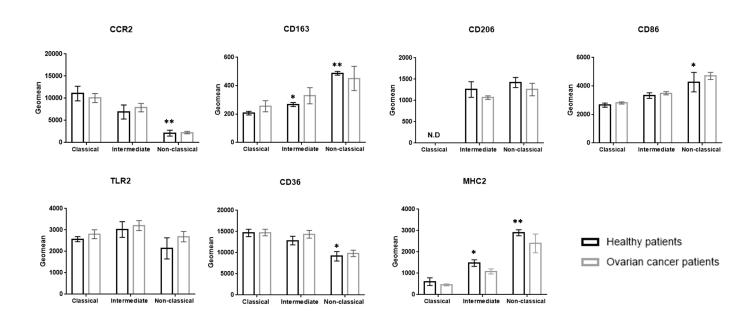
Supplementary material J Immunother Cancer



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^{high} CD16⁻), intermediate (CD14^{high} CD16^{high}) and non-classical (CD14^{low} CD16^{high}) monocytes. Graphs represent geomean fluorescence quantification for the indicated protein. Results correspond to mean \pm 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.