



SUPPLEMENTARY FIG. S2. Distribution of hBM subpopulations in CD18 versus CD31 plot. Backgating of lineage markers allowed associating specific hBM subpopulations to different regions of CD18 versus CD31 plot (**A**). FACS sorting of all hBM subpopulations double positive counterpart (sort gate displayed) confirmed expected morphologies (**B**). FACS, fluorescence-activated cell sorting.