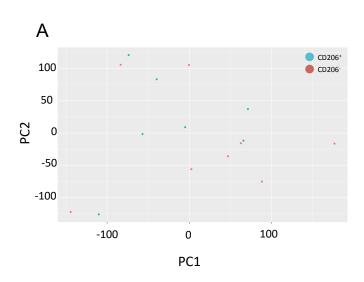
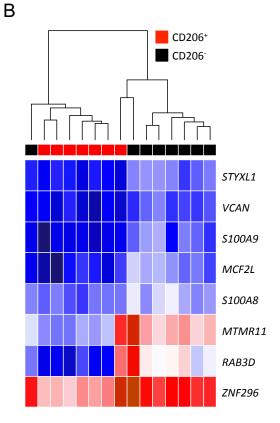


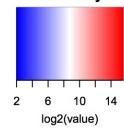
Supplementary Figure 1. Staining of CD45+ CD64+ CD14+ cells and CD45+ CD64+ MHC II+ cells to investigate the relationship between CD11c and CD206 staining.

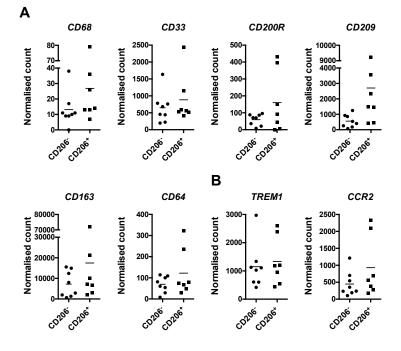




Color Key

Supplementary Figure 2. RNAseq analysis of FACS-purified CD206⁻ and CD206⁺ colonic human macrophage populations. (A) Principal component analysis (PCA) comparing CD206⁺ and CD206⁻ intestinal macrophage populations. (B) Eight genes with statistically significant (adjusted P-value <0.05) differential expression between CD206⁻ and CD206⁺ macrophage populations.

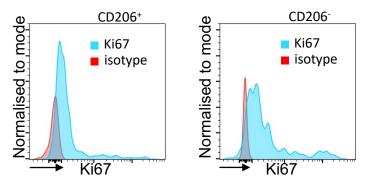




Supplementary Figure 3. RNAseq analysis of FACSpurified CD206⁻ and CD206⁺ colonic human macrophage populations. (A) Comparative expression of macrophage associated cell surface markers on CD206⁻ and CD206⁺ cells. (B) Comparative expression of monocyte-associated cell surface markers on CD206⁻ and CD206⁺ cells. Intestinal macrophages derived from healthy colon resections; n=8 and 7 for CD206⁻ and CD206⁺, respectively.

Gated live CD45⁺HLA-DR⁺CD64⁺

А



Supplementary Figure 4. Ki67 staining from CD206⁺ and CD206⁻ compared to their isotype control (red).