



Supplementary Figure S2. scRNA-seq profiling of ovarian ascites cells after *ex vivo* bexmarilimab treatment, related to Figure 1.

A, Average expression of main cluster marker genes in ovarian ascites cells analyzed by scRNA-seq ($n = 7$ patients). Dot size represents proportion of cells with marker gene expression. **B**, Expression levels of the indicated main cluster marker genes are shown on UMAP plots. **C**, Proportions of ovarian ascites cells belonging to each main cluster are displayed separately for each sample. **D**, Ovarian ascites cell UMAP plots colored by patient ID, *ex vivo* treatment and scRNA-seq experiment batch. BEX, bexmarilimab; IgG4, isotype control for bexmarilimab.