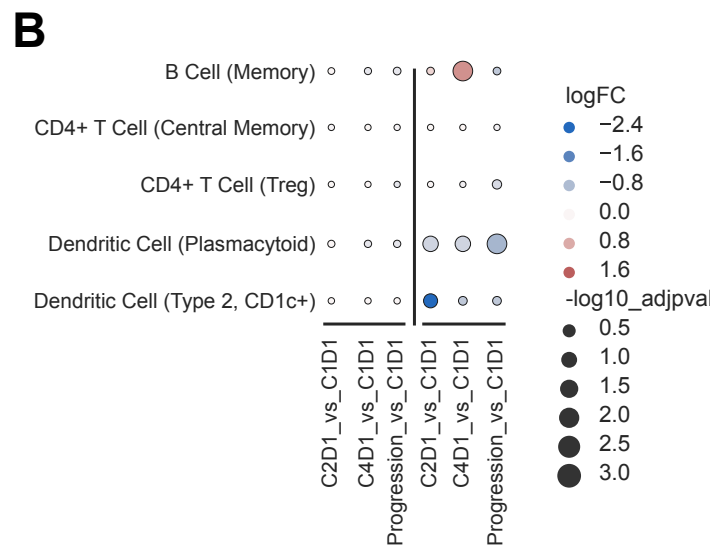
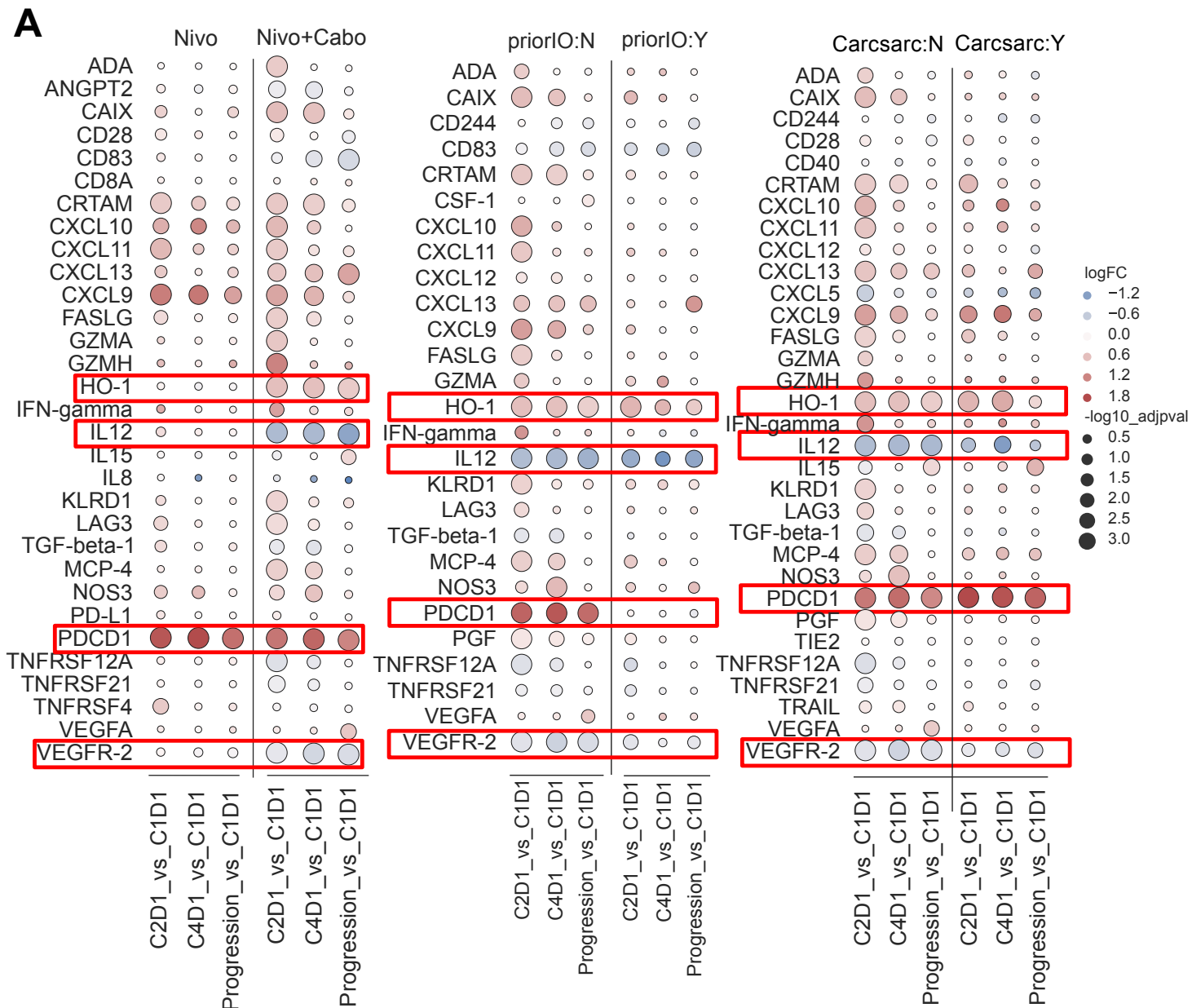




Supplementary Figure 4. Dotplot representation of fold changes (log2) and statistics (adjusted p values, -log10) for proteomic analytes (A) and cell populations (B) at C2D1, C4D1 and progression from baseline C1D1. Only statistically significant changes are shown. **A.** Dotplot for proteomic biomarkers comparing treatment arms, subjects with prior IO or carcinosarcoma treated with nivo+cabo. **B.** Cell frequency changes from baseline between treatment arms (left - nivo, right - nivo+cabo).