

Article

T-Cell Receptor Repertoire Characteristics Associated with Prognostic Significance in High-Grade Serous Ovarian Carcinoma

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Supplementary Figures:

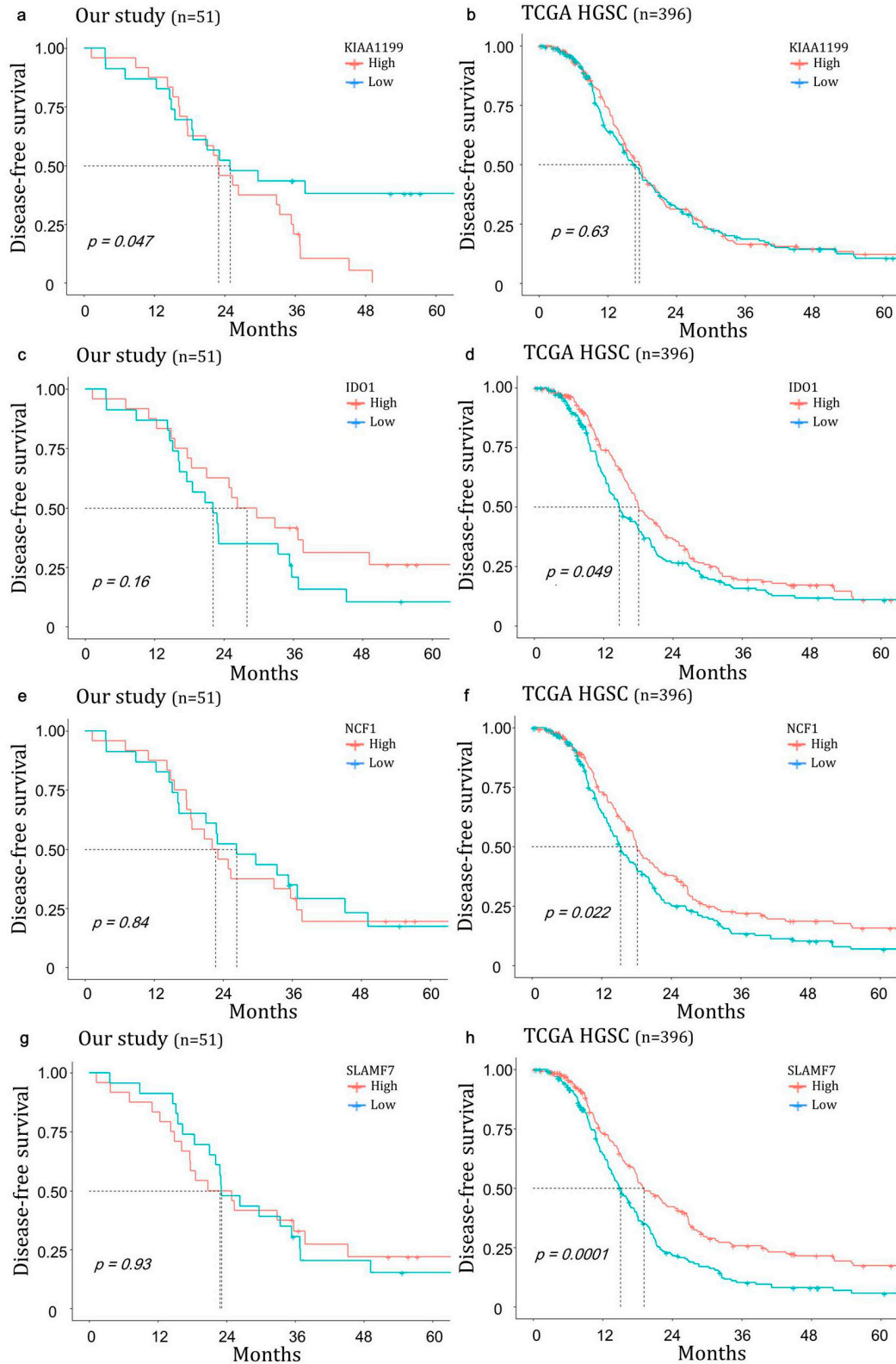


Figure S1. The survival analysis for 4 differentially expressed genes in our study and TCGA high-grade serous ovarian carcinoma (HGSC) data. For survival analysis, disease-free survival data were analyzed in both data. Survival results were shown for *KIAA1199* in our study (a) and TCGA HGSC (b), for *IDO1* in our study (c) and TCGA HGSC (d) for *NCF1* in our study (e) and TCGA HGSC (f) and for *SLAMF7* in our study (g) and TCGA HGSC data (f), respectively.

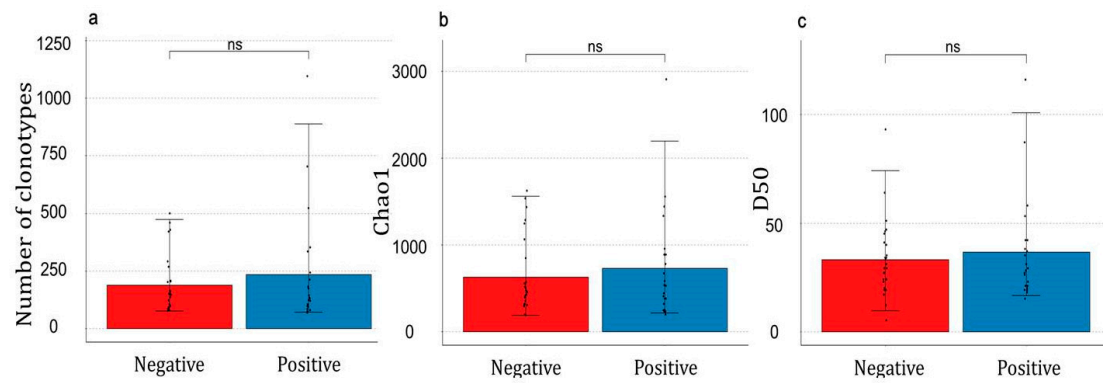


Figure S2. The expression pattern of the TCR repertoire in patient groups divided based on HRD mutation. **(a)** T-cell receptor counts of unique TCR clonotypes. **(b–c)** Richness and diversity of TCR estimated by Chao1, and D50 index in patient groups divided based on HRD mutation. **(a–c)** Non-significant p -values are indicated as ns.