

Figure S7

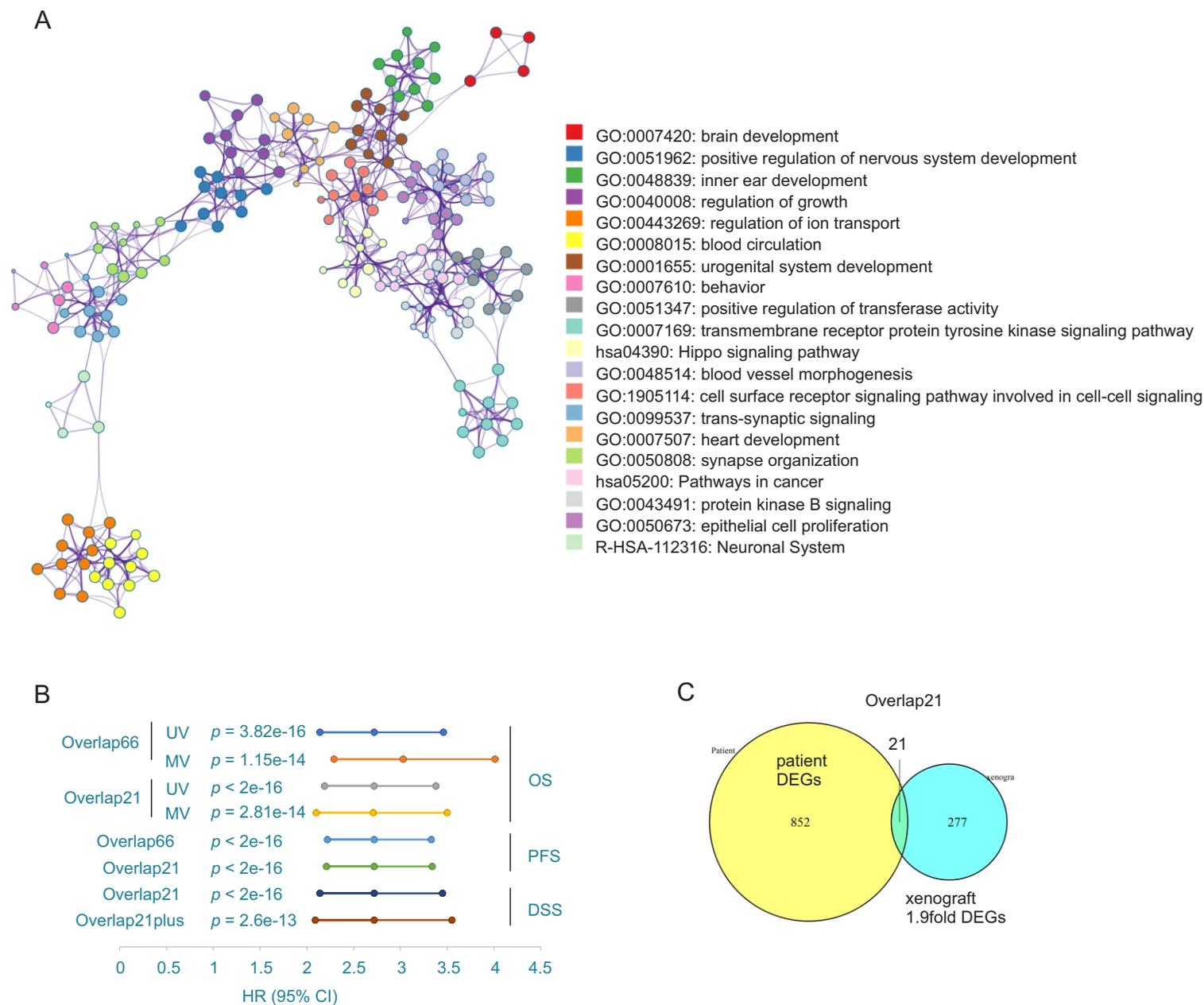


Figure S7. Prediction of OS, PFS, and DSS possibilities by Overlap66, Overlap21, and Overlap21plus. **(A)** Network connection of individual enriched terms within individual clusters (see Fig. 4a, Additional file 10: Table S4 for details). **(B)** The predictions of OS, PFS, and DSS possibility using Overlap66, Overlap21, and Overlap21plus risk scores. UV and MV: univariate and multivariate Cox analysis, HR: hazard ratio, and CI: confidence interval. **(C)** The number of overlapping genes between primary (patient) pRCC-derived DEGs and DEGs obtained from xenografts at fold change $\geq |1.9|$.