Description of Additional Supplementary Files

File Name: Supplementary Software 1

Description: Contains the R code to generate the graphs shown in Figure 6f and Supplementary Figures 6a and 6b. The code uses data downloaded from cBioPortal (https://www.cbioportal.org/) of patients with the indicated cancer type to determine the correlation between FGF pathway activation and MCL1 or BCL2 expression as well as patient survival stratified by FGF pathway activation and MCL1 or BCL2 expression.