Supplementary Figure S4
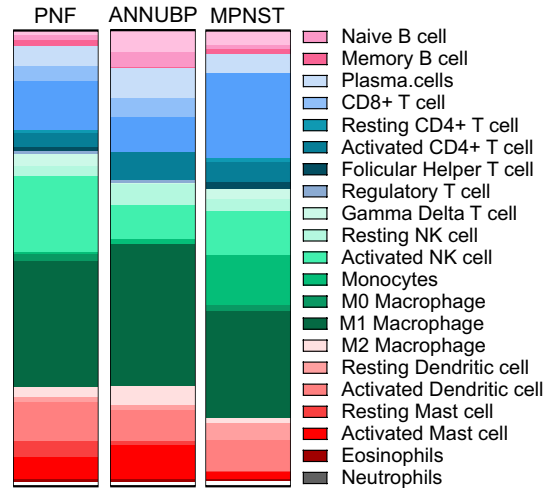

Supplementary Figure S4. Predicted immune cell composition of human PNFs, ANNUBPs, and MPNSTs based on CIBERSORT analyses. Patient-matched benign (PNF, ANNUBP) and malignant (MPNST) tumors had previously been examined by RNA sequencing (Kohlmeyer et al, *Clinical Cancer Research* 2020). Deconvolution of the gene expression data was performed using the CIBERSORT method to predict relative proportions of immune cells within the tumors.