



Supplemental Figure 3

Supplemental Figure 3. PNF, ANNUBP, and MPNST exhibit distinct global gene expression programs. Hierarchically clustered heatmap of the top 100 differentially expressed genes. Rows represent individual genes and their respective expression \log_2 normalized, z-score transformed expression values across each sample. Columns represent individual samples with tumor subtype annotated above. Rows and columns were clustered by correlation coefficient with average linkage.