

SPOT: Spatial Proteomics through On-site Tissue Proteins Labeling

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Supplementary Graphs

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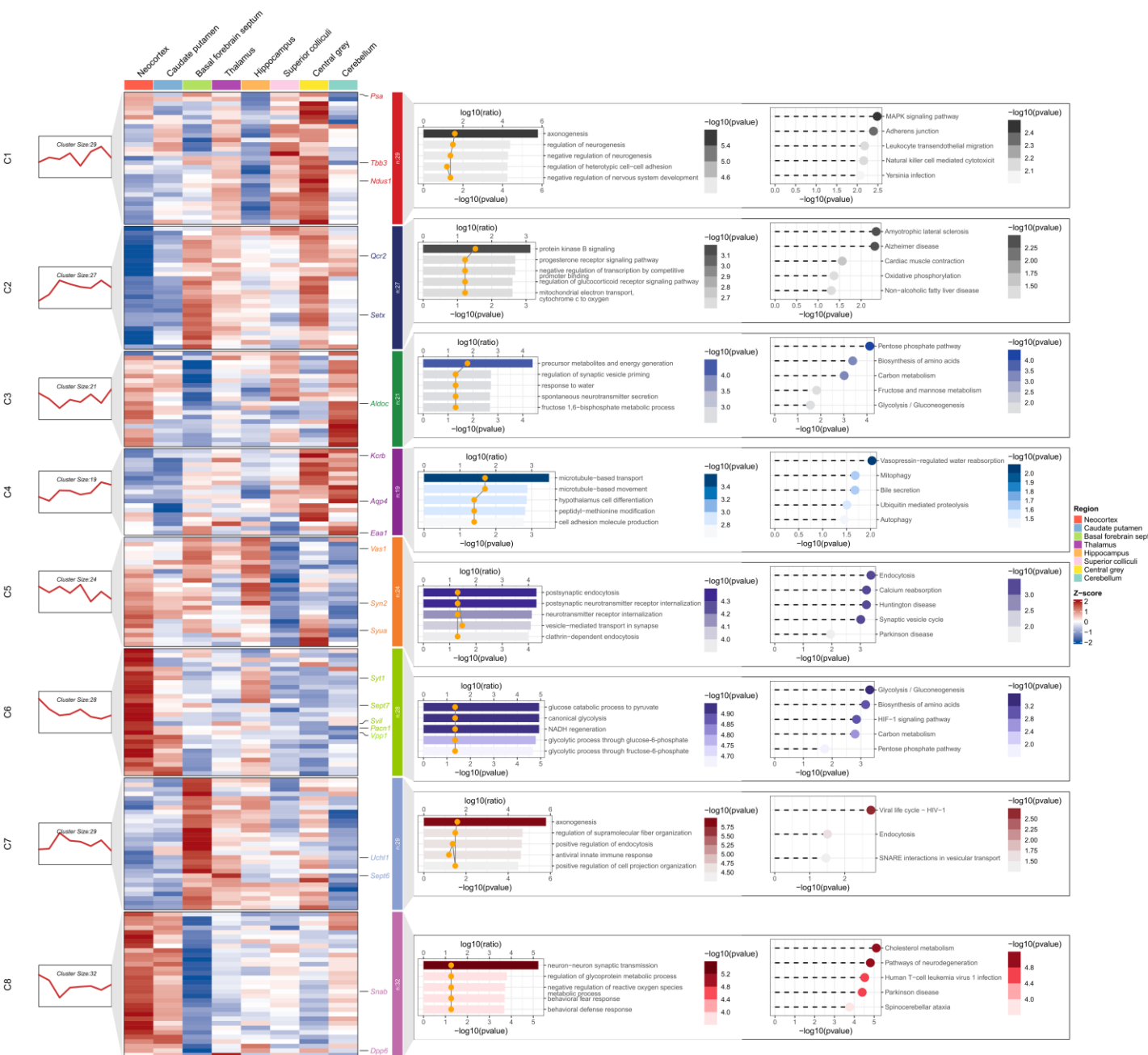
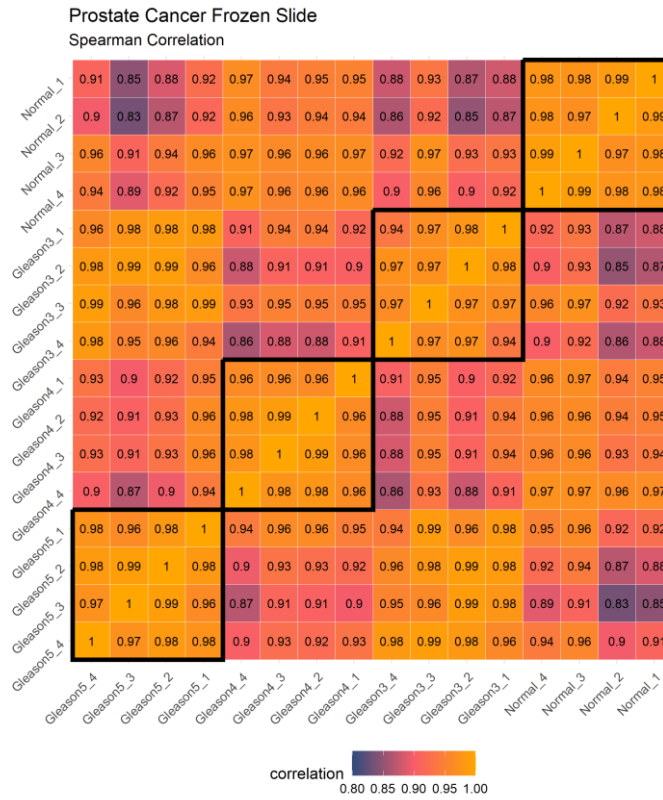


Figure S1. Detailed clustering results of the protein expressions visualized using heatmap-GO-KEGG combination graph. Proteins exclusively or abundantly expressed in mouse brain are marked for each cluster.

A



B

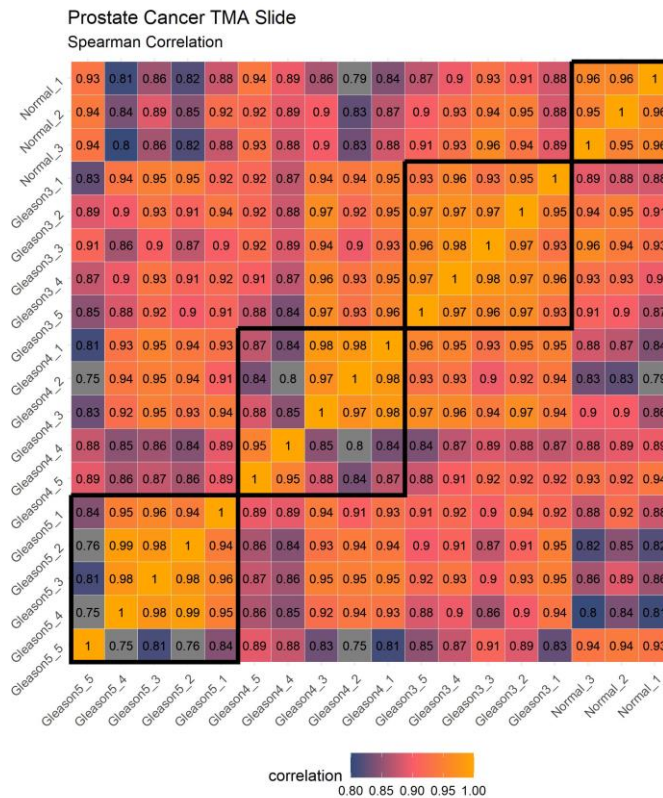


Figure S2. Spearman Correlation matrixes of the protein abundances (protein abundances were median-normalized) generated from the frozen prostate cancer tissue slides. A) Frozen slide. B) TMA slide.