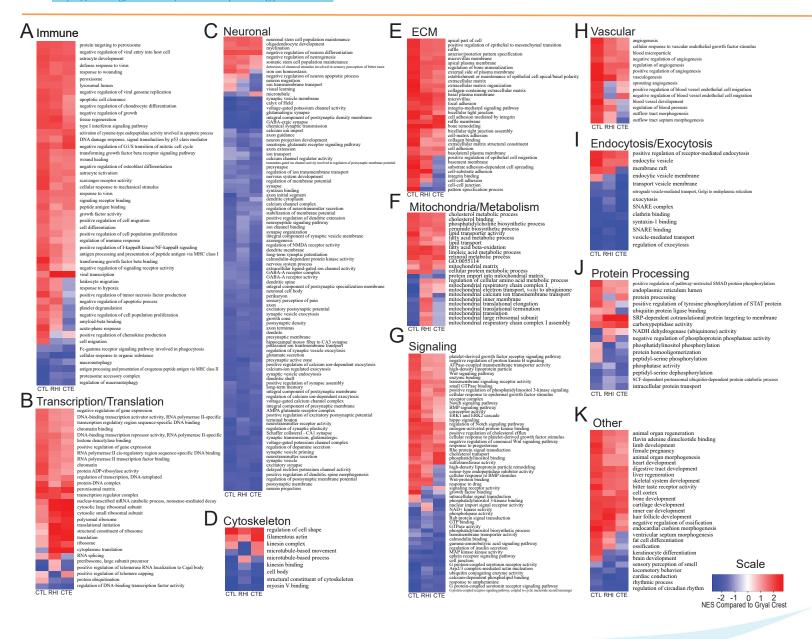
doi: https://doi.org/10.17879/freeneuropathology-2021-3453



## Figure 5. Heatmap of sulcal specific GSEA biological processes

Using the genes that were significantly altered in the sulcus compared to the crest, we used GSEA to identify biological processes. Each process was then assigned to a greater overall biological category: A) Immune, B) Transcriptional/translational, C) Neuronal, D) Cytoskeletal, E) Extracellular matrix, F) Mitochondria/metabolism, G) Signaling, H) Vascular, I) Endocytosis/exocytosis, J) Protein processing, K) Other.