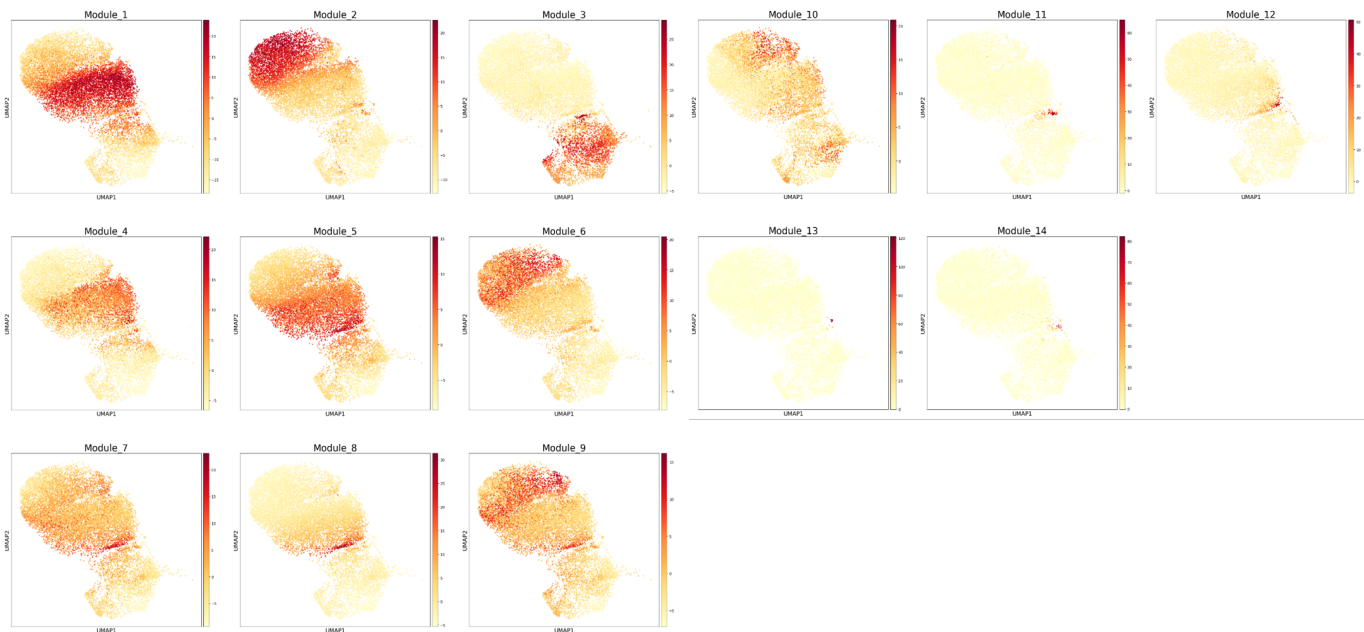
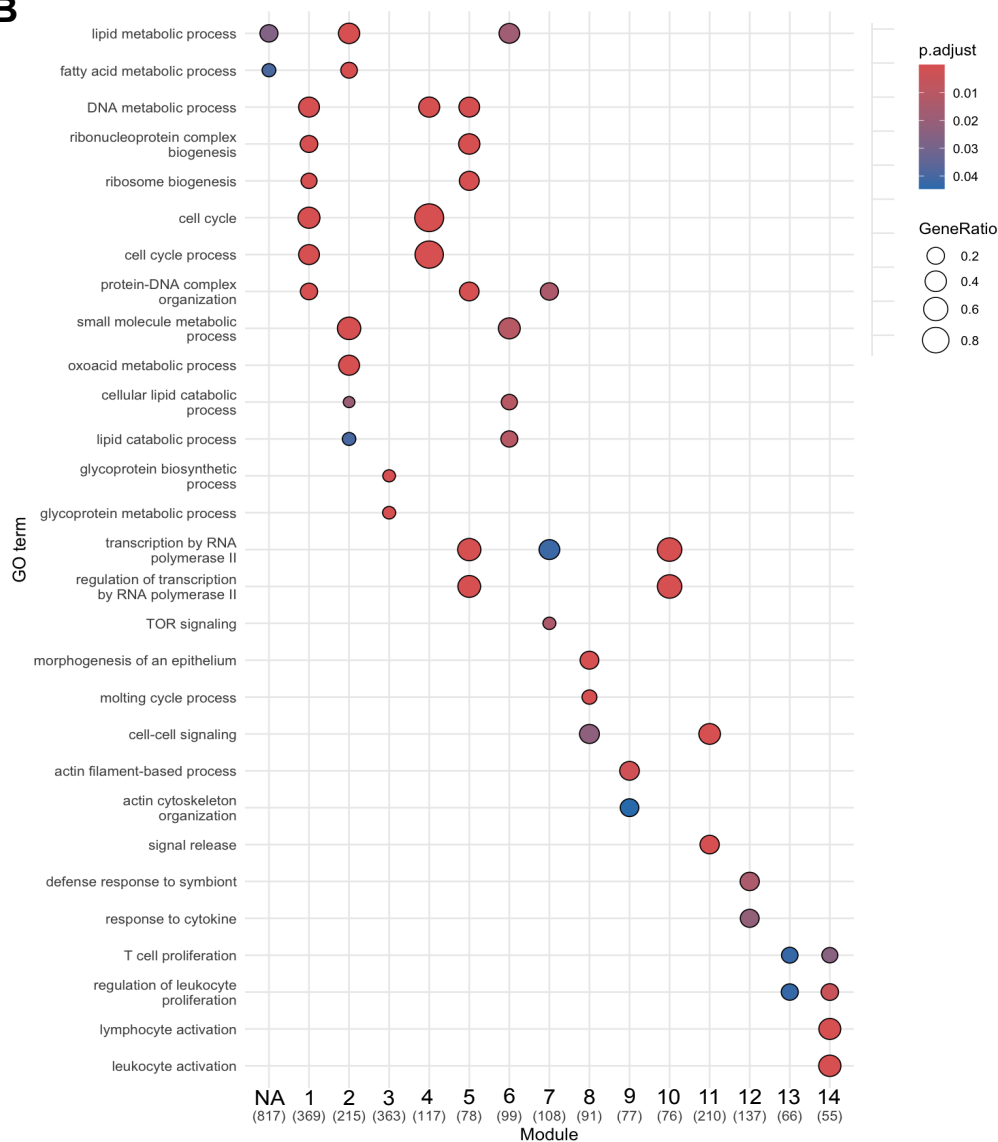


A



B



Supplementary Figure 3: Spatial and functional characterisation of Hotspot gene modules. A) Plots showing enrichment for each module detected through Hotspot in UMAP space. B) Dot plot representing the top enriched pathways in each Hotspot module (GSEA). Module number indicated on the x axis with number of genes in parentheses. NA = not classified in a specific module. Colour represents the adjusted p value, size of circles represents the ratio of genes upregulated compared to genes forming the pathway.