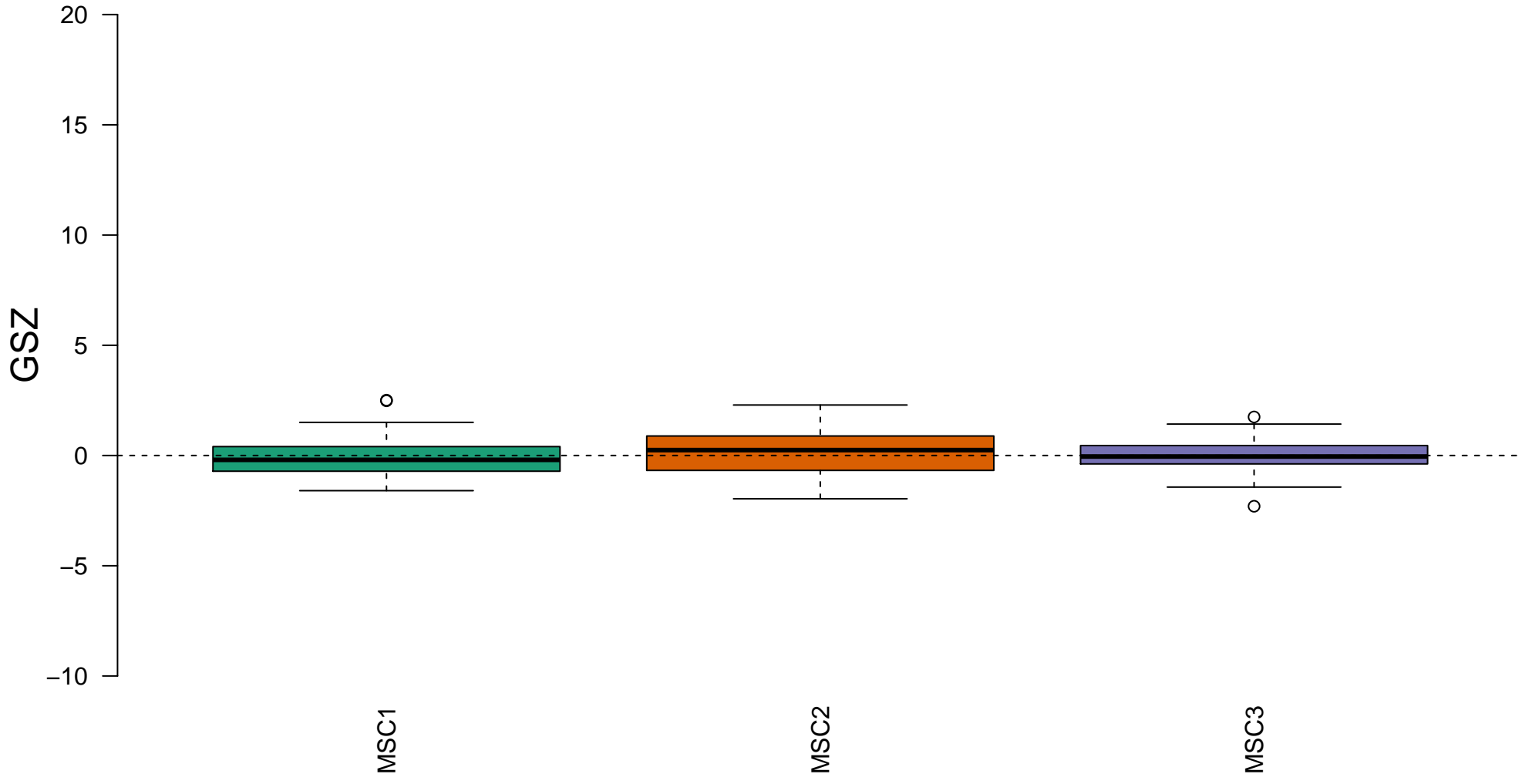
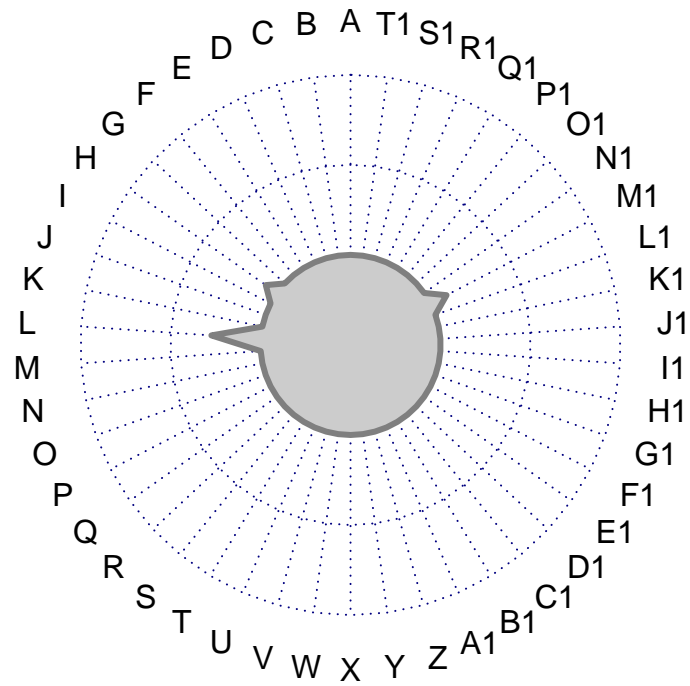




# BIOCARTA\_RANKL\_PATHWAY



Enrichment in overexpression spots



Enrichment in kMeans clusters

