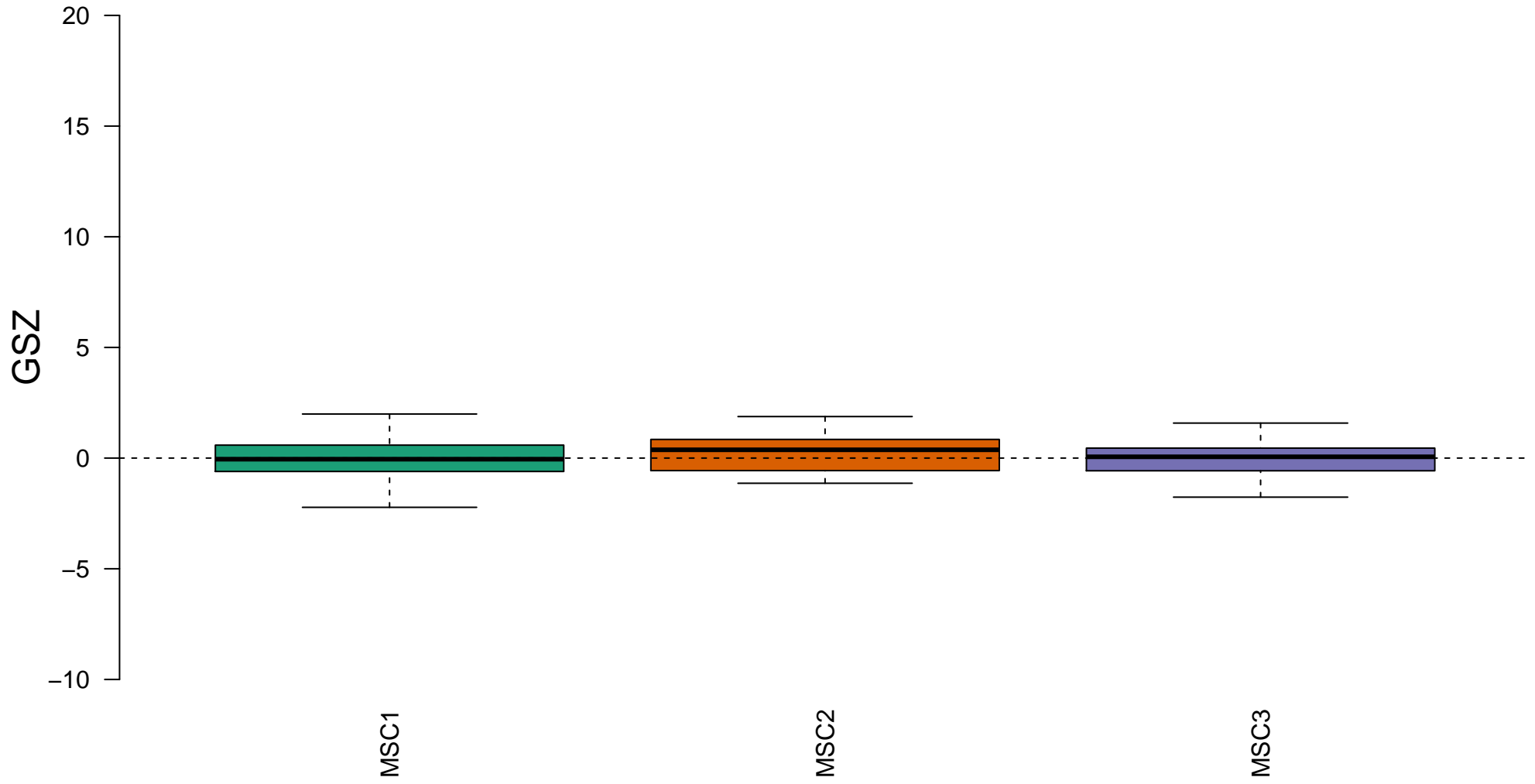
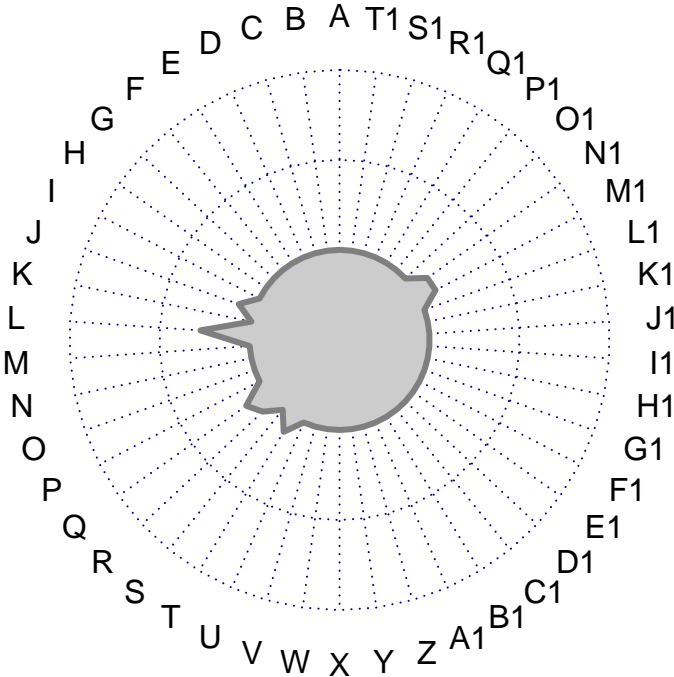




# BIOCARTA\_TALL1\_PATHWAY



Enrichment in overexpression spots



Enrichment in kMeans clusters

