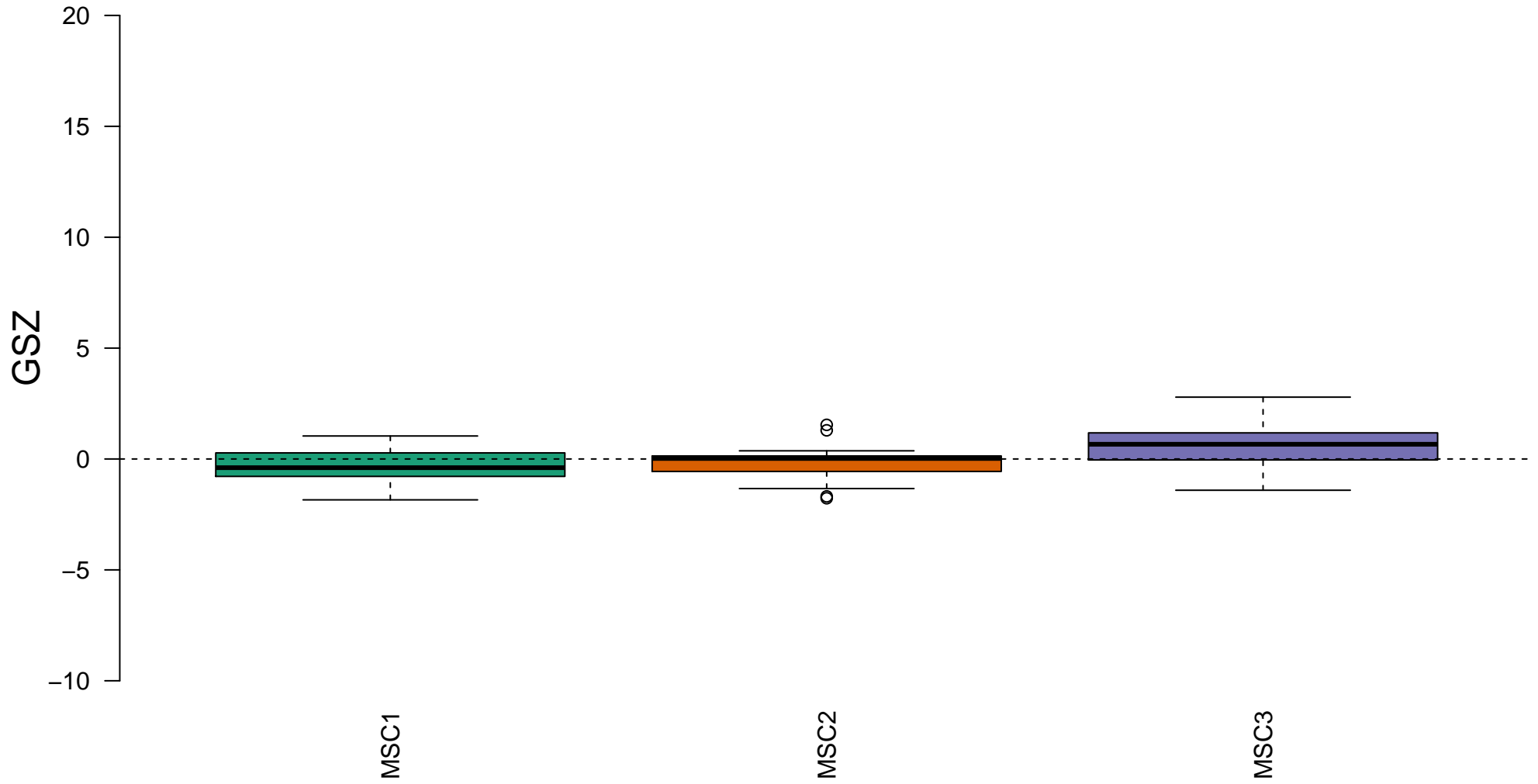
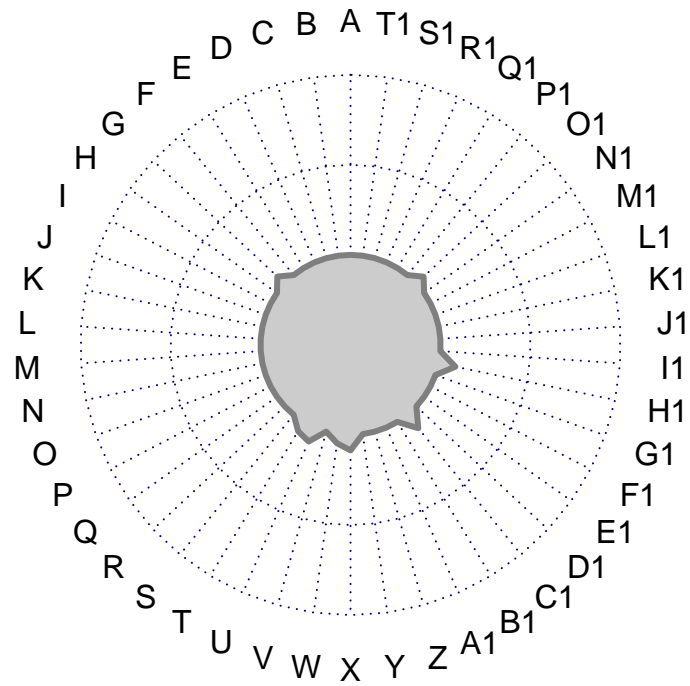




# BIOCARTA\_AGR\_PATHWAY



Enrichment in overexpression spots



Enrichment in kMeans clusters

