



Methylation UMAP plot

Close-up of UMAP for 209547730116_R08C01

Neighbors in UMAP for 209547730116_R08C01

A pie chart illustrating the distribution of GBM types. The chart is divided into two segments: a large red segment representing GBM_MES at 73.3%, and a smaller blue segment representing GBM_RTK_II at 26.7%. A legend to the right of the chart identifies the colors: red for GBM_MES (73 %) and blue for GBM_RTK_II (27 %).

GBM Type	Percentage
GBM_MES	73.3%
GBM_RTK_II	26.7%

Figure 1: A Manhattan plot showing the association of 10,000 SNPs with the expression of 100 genes. The y-axis represents the association p-value on a log scale from -1.2 to 1.2. The x-axis shows chromosomes from chr1 to chrY. Red dots indicate SNPs associated with down-regulation, and blue dots indicate SNPs associated with up-regulation. Genes are labeled with their names and corresponding SNPs. A legend indicates that red dots represent SNPs associated with down-regulation and blue dots represent SNPs associated with up-regulation.