

Differential Methylation Regions on Genome of Gastric Cancer

bioinformatic group@AMS

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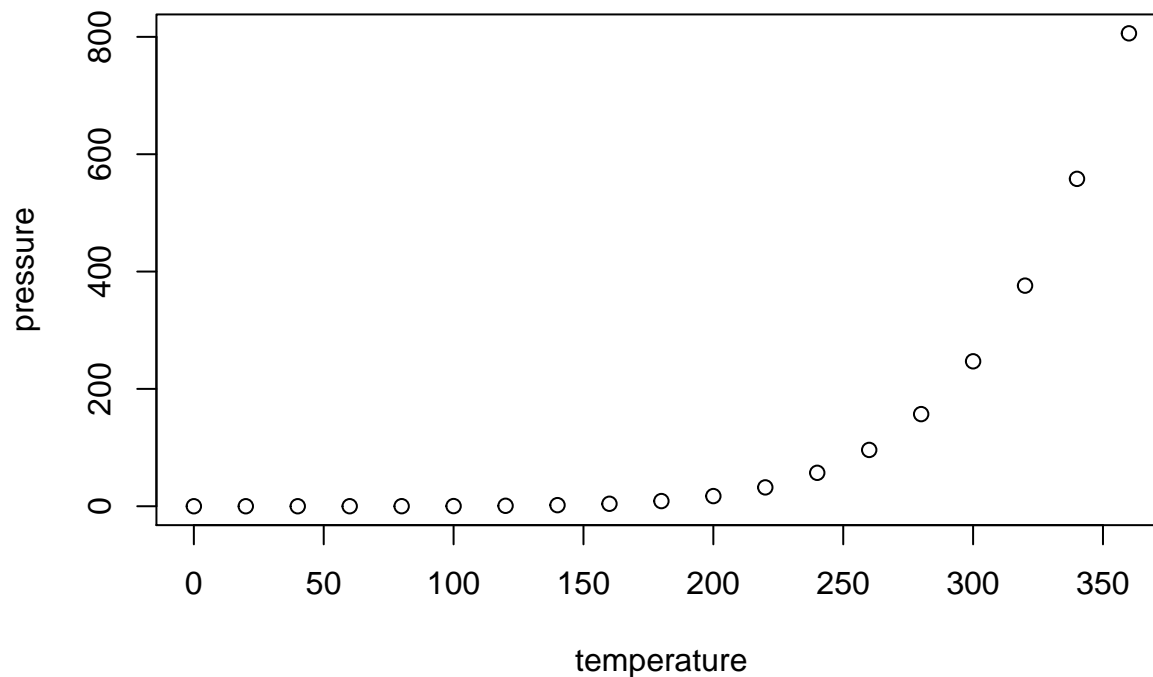
Datasets used in this study

Only two samples match gastric cancer in the methylation dataset of TCGA (The Cancer Genome Atlas), this requires us to look for more other datasets. After searching the GEO repository of NCBI , we found the following 4 studies focus on the methylation status of gastric cancer genome.

access	platform	normalNumber	cancerNumber	reference
GSE30601	GPL8490	94	203	1
GSE25869	GPL8490	32	32	2
GSE99553	GPL13534	84	0	3
GSE103186	GPL13534	61	130	4

Including Plots

You can also embed plots, for example:



Note that the `echo = FALSE` parameter was added to the code chunk to prevent printing of the R code that generated the plot.