Run by chosen

Bed12  
to  
Bed6

Load Input Gene

Load CGmap

Two groups difference

Metaplot

Two groups difference

Chromosome View

Differentially Methylated Gene

Fold Enrichment

Heatmap & Principle Component Analysis

Generate merge bed files

body bed

igr bed

Find

promoter bed

Extract transcript regions

Calculate Differentially Methylated Region

1. Generate Common Region File
2. Generate UnionSite File

CHG

Calculate Common Region

CHH

CG

Combine CGmap