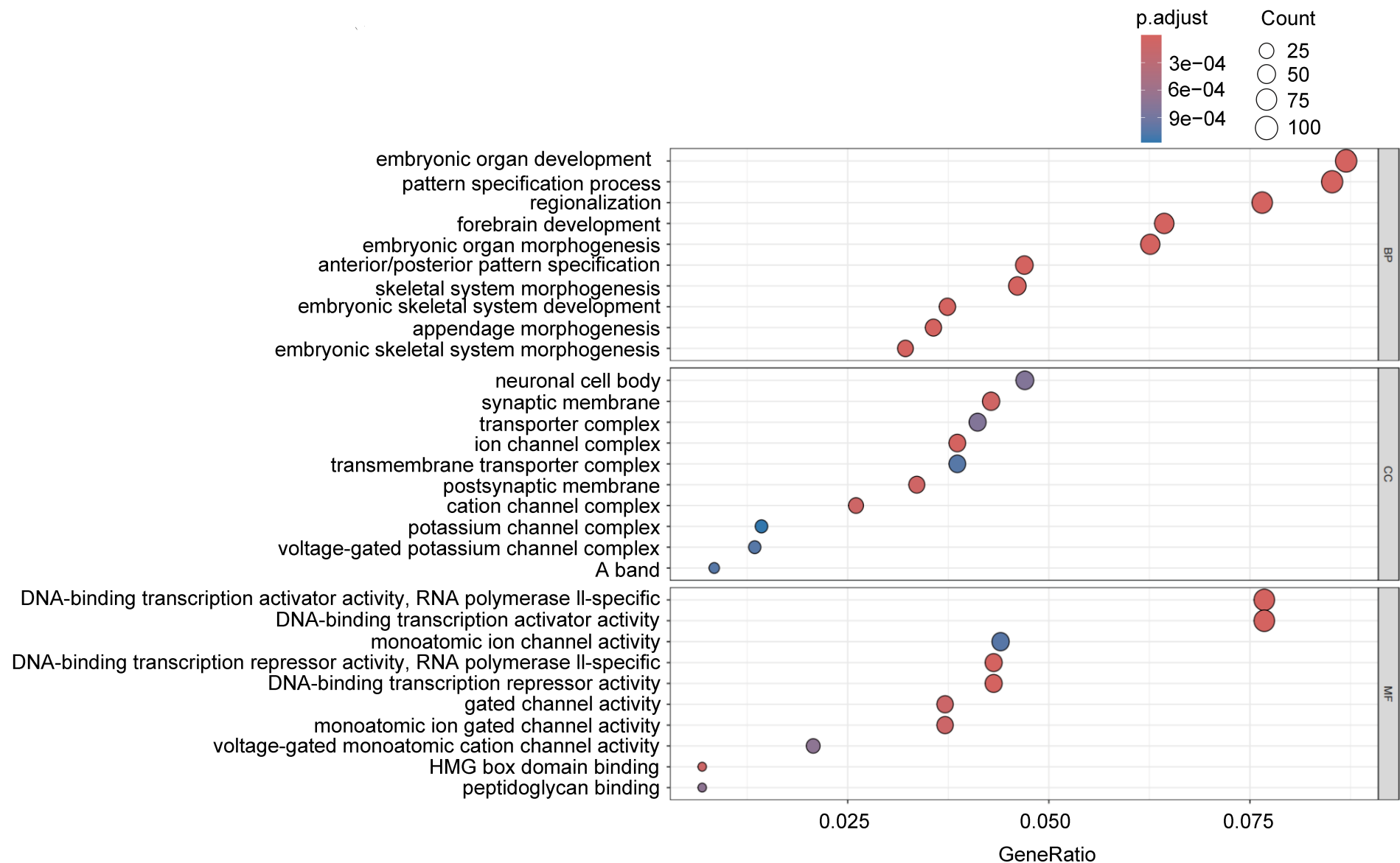




Supplementary Figure 2. Gene ontology analysis revealing the overrepresentation of genes with remarkable hypomethylation or hypermethylation among the DMPs.