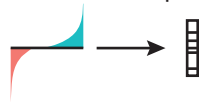
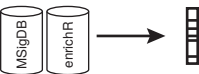


(i) Assemble gene list of interest

Differential expression



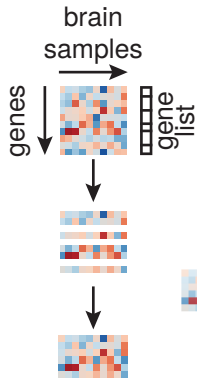
Gene list databases



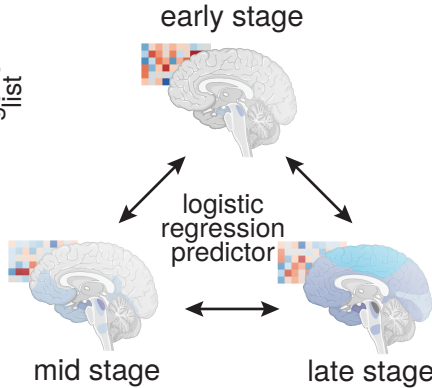
Functional gene list



(ii) Subset AMP-AD RNA-seq data



(iii) Training disease stage predictor for each gene list



(iv) Evaluate performance of gene list vs background

