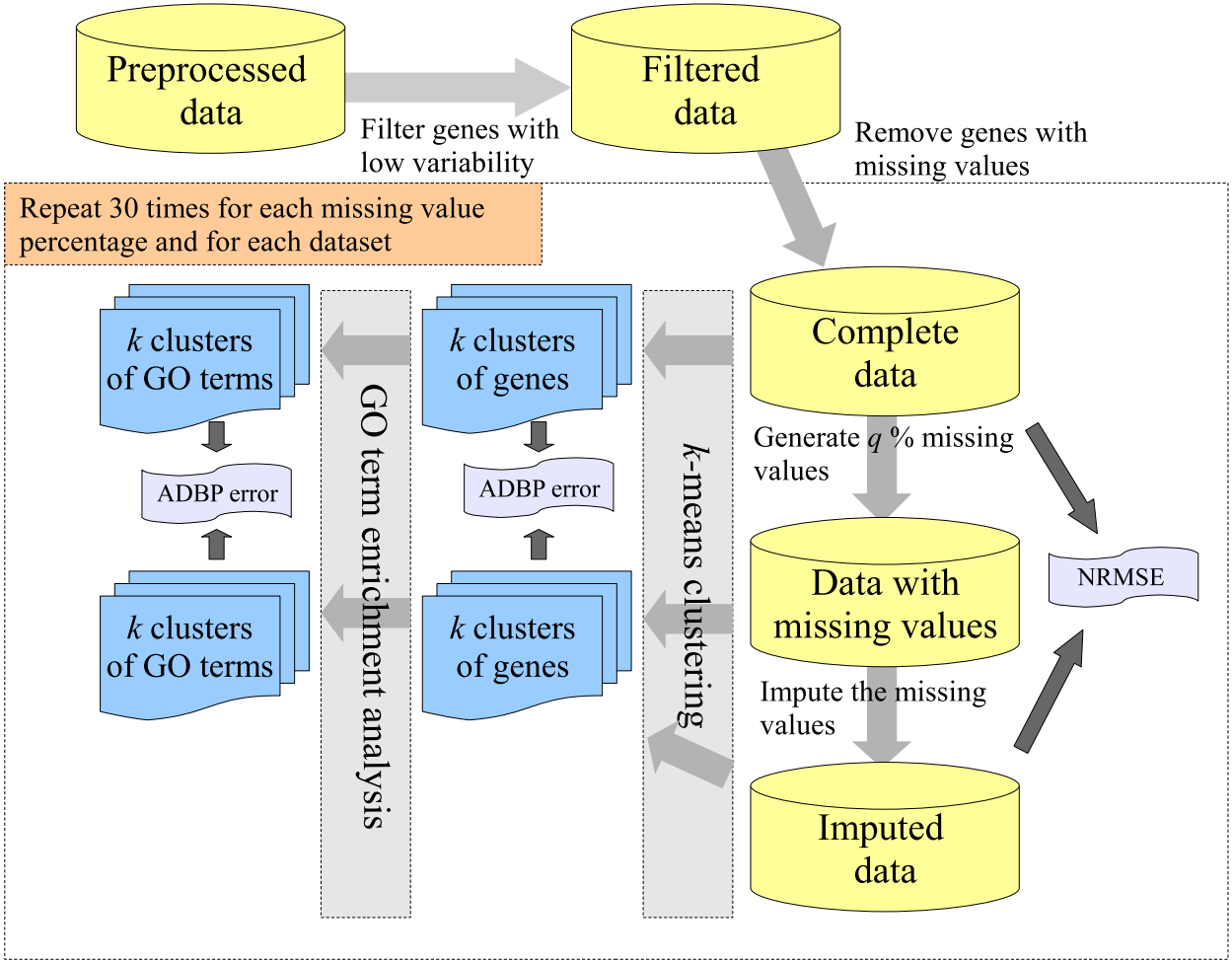
Preprocessed data
Filter genes with low variability
Filtered data
Remove genes with missing values
Repeat 30 times for each missing value percentage and for each dataset
Complete data
Generate q % missing values
Data with missing values
Impute the missing values
Imputed data
NRMSE
k-means clustering
k clusters of genes
ADBP error
k clusters of genes
GO term enrichment analysis
k clusters of GO terms
ADBP error
k clusters of GO terms