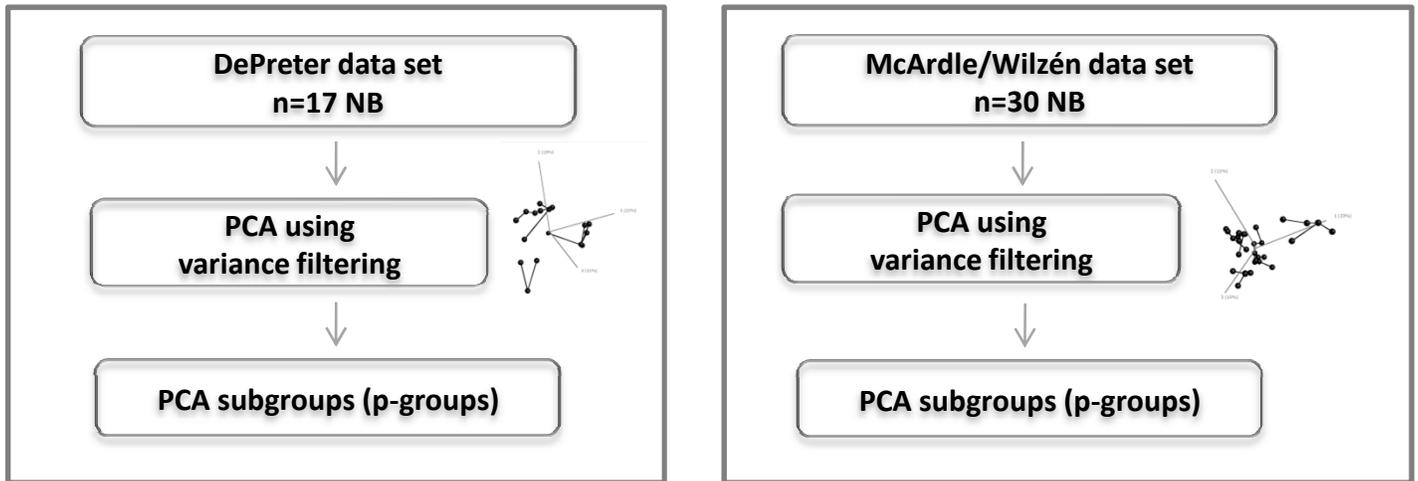


Step 1: Subgroup discovery by PCA (data sets 1 & 2)

Defining expression subgroups

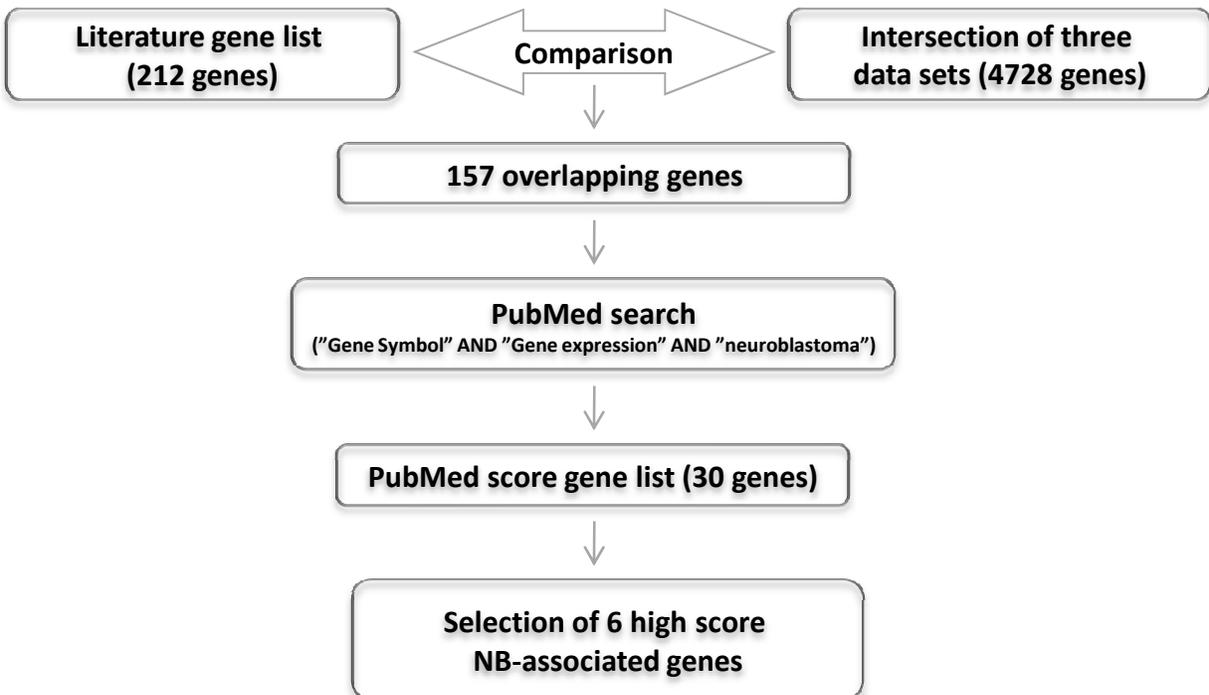
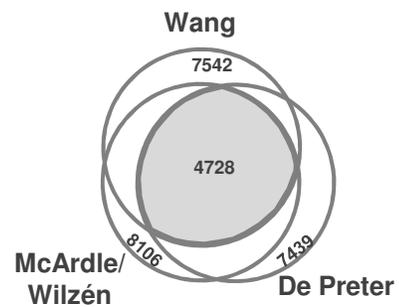


Data-mining of gene lists from literature



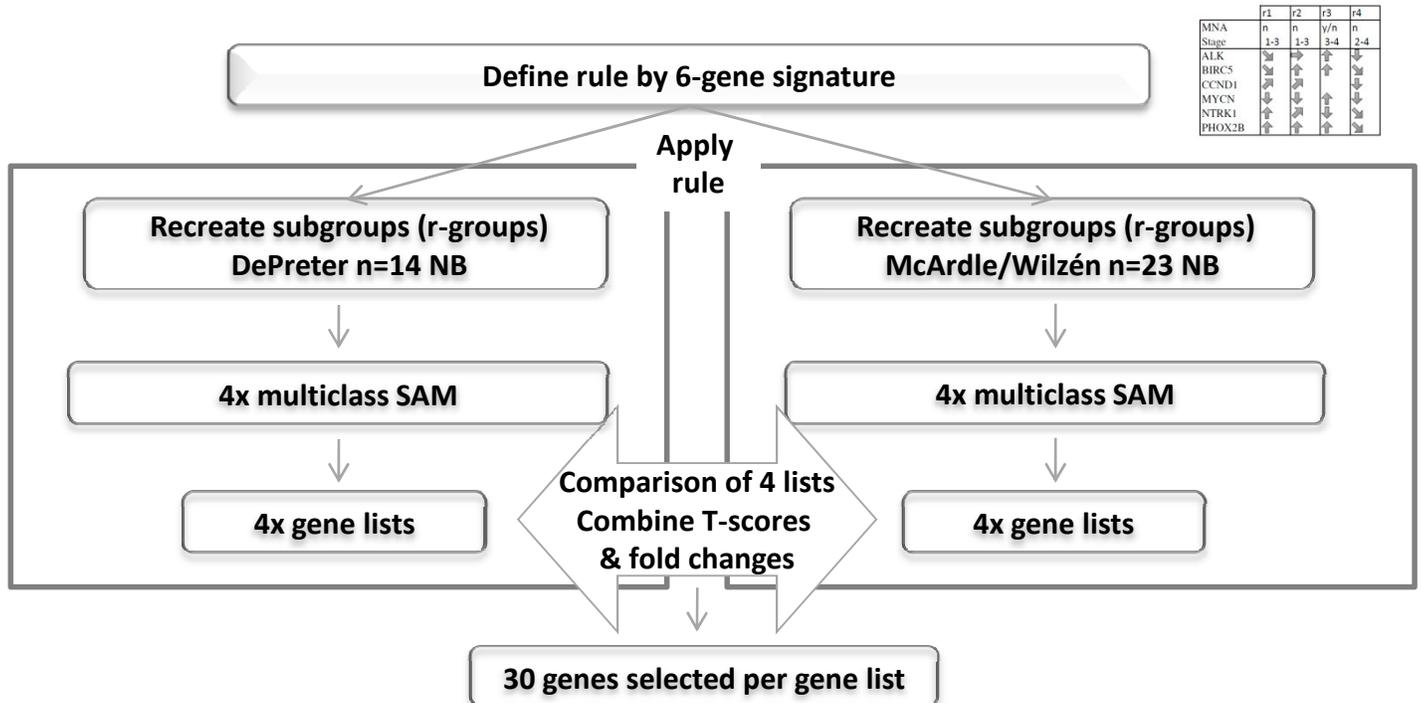
Gene lists from 15 expression studies

↳ 212 genes present in ≥ 2 studies



Step 2: Verification by hierarchical clustering & PCA (data set 3)

Defining a classifier gene set



Hierarchical clustering & PCA

