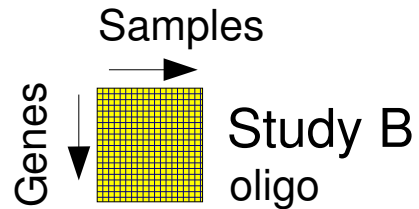
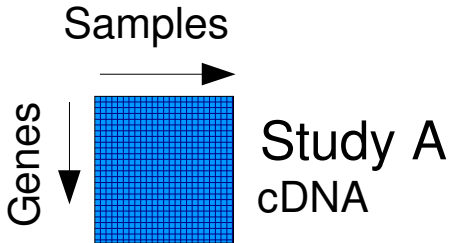
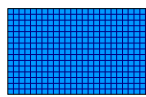


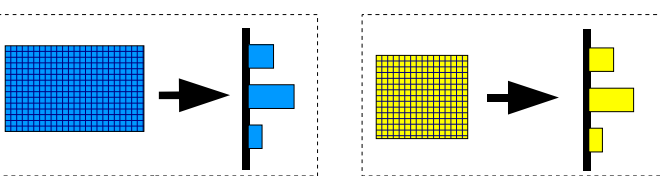
retrieve and preprocess
gene expression data



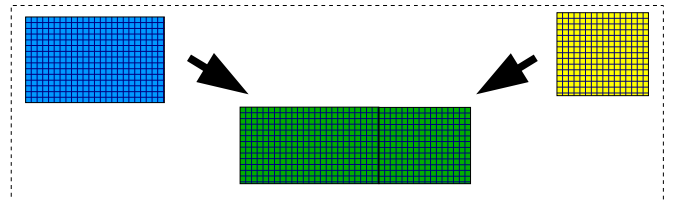
match to UniGene and
intersect



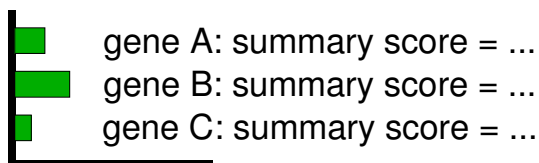
derive summary
information



data integration
with MRS or QD

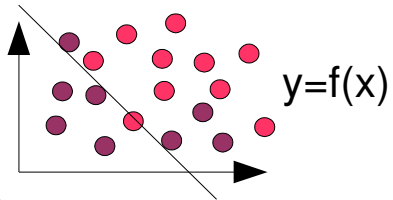


combine summary
information in
meta-analysis

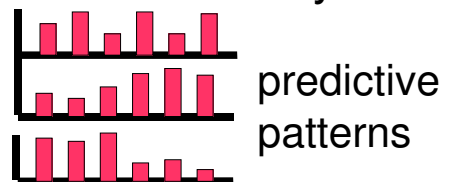


meta-analysis

classification analysis



gene selection by RFE



cross-platform classification