

Updates gene set  
annotations

Get experiment level  
estimates

Combine experiment level  
estimates in 2 parts

Aggregate results into  
a single matrix

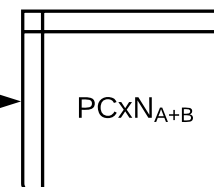
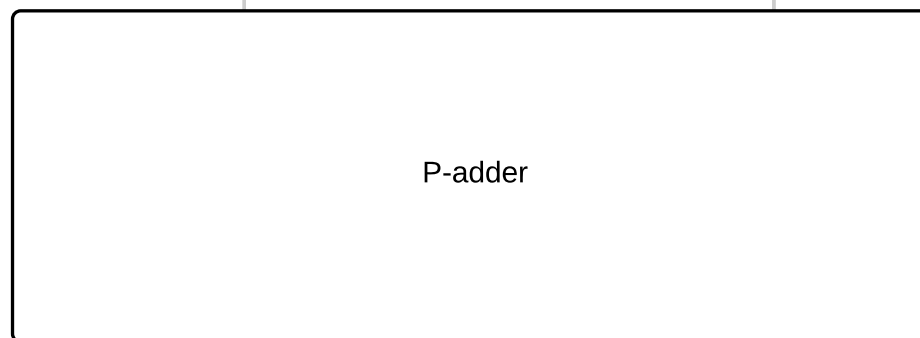
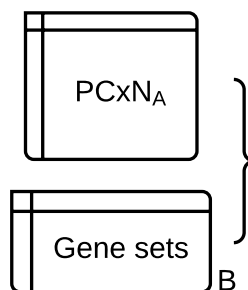
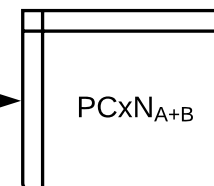
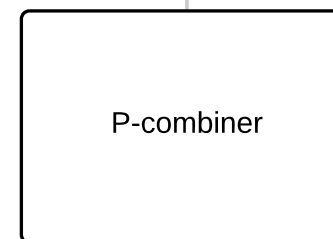
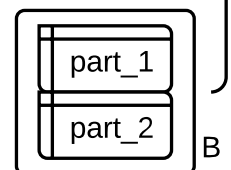
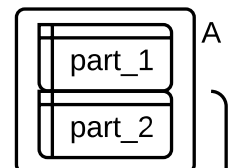
Estimates00

Estimates01

Estimates02

Estimates03

improved\_PCxN



**\*checks for identical gene  
members**

**\*Skips calculated pairs**