Drug repositioning based on <u>co</u>expressed <u>gene-set enrichment</u> <u>a</u>nalysis (cogena)

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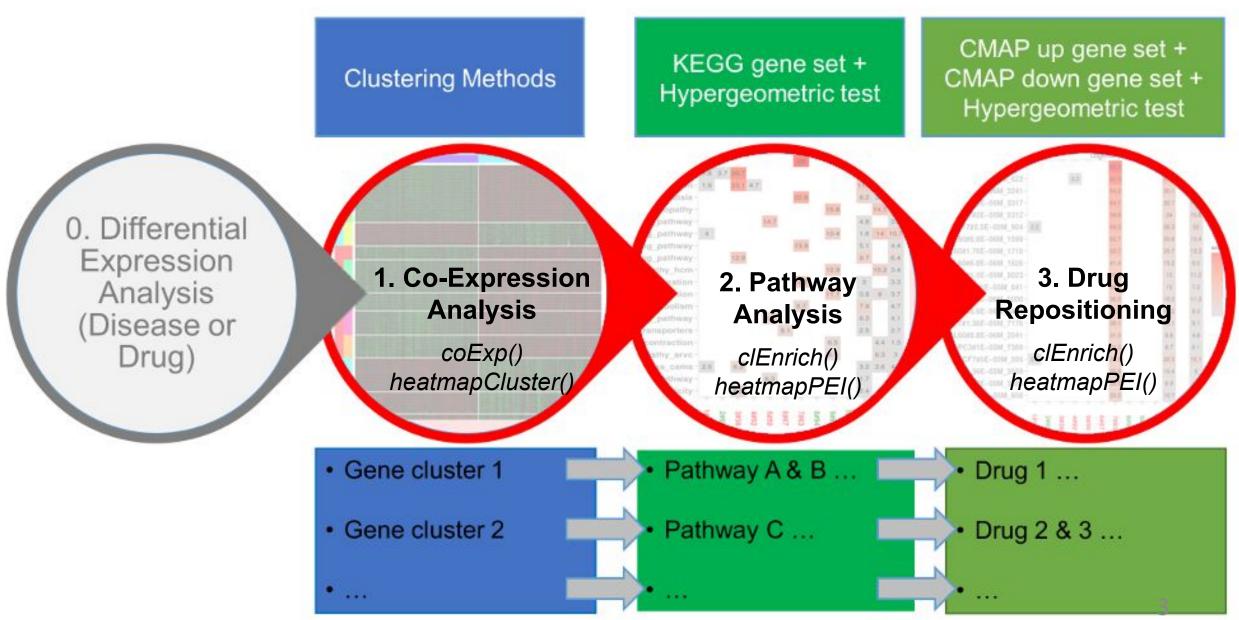
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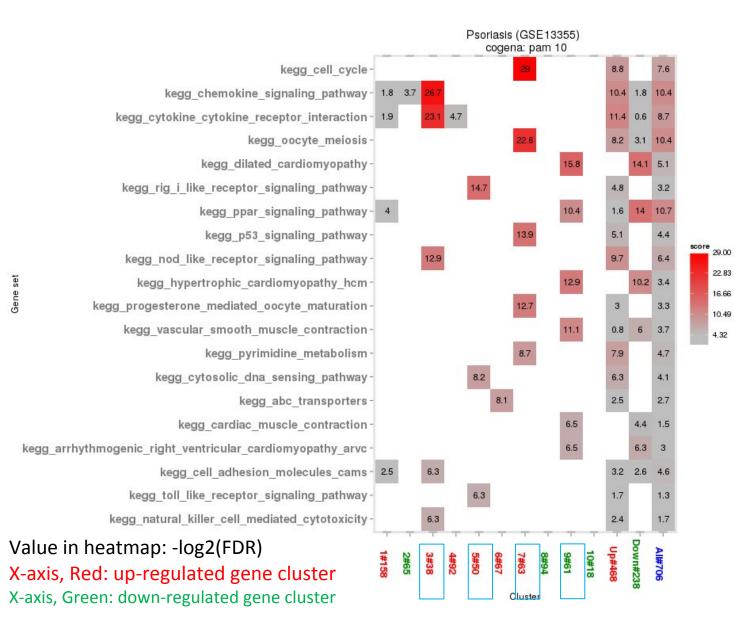
Cogena: a framework for co-expressed geneset enrichment analysis

- Co-expressed genes may be very biologically relevant
 - Co-expressed genes are more highly connected in pathways
 - May be mechanistically important in disease
 - May represent more robust drug targets
- Pathway analysis and drug repositioning analysis on co-expressed genes can be more informative than transcriptome wide analysis
- cogena can be applied to drug repositioning and mode of action discovery, illustrated by a psoriasis exemplar.
 - Code available at http://zhilongjia.github.io/psoriasis/

Drug repositioning pipeline based on cogena

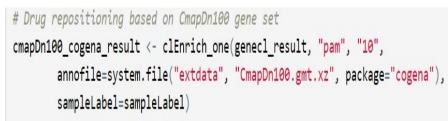


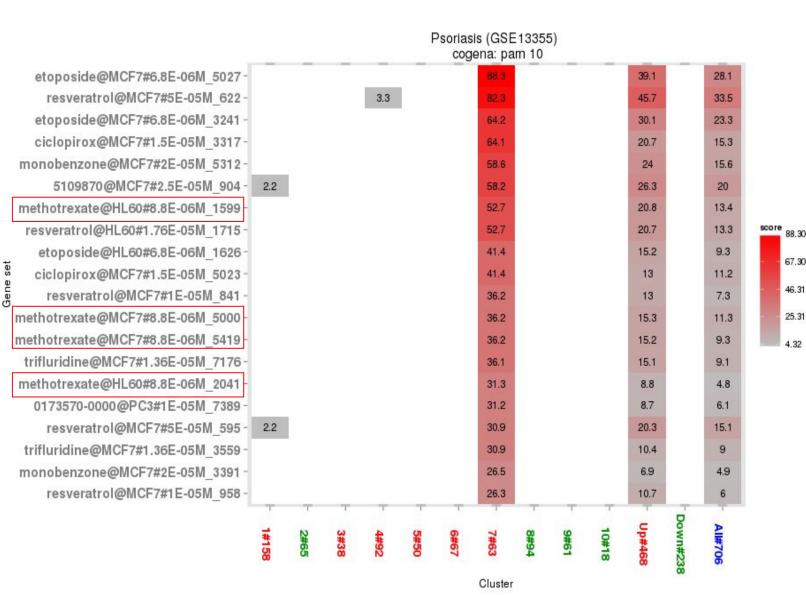
Pathway analysis for *psoriasis* based on cogena



- Immune-related pathways in cluster 3 and 5.
- Cell cycle and p53 signalling pathways in cluster 7.
- PPAR signalling pathways in cluster 9.

Drug repositioning for psoriasis based on cluster 7 (cell cycle & P53 pathway)





Methotrexate (ranked 7th) is a first-line drug for psoriasis and an inhibitor of the enzyme folic acid reductase, which leads to the inhibition of DNA synthesis and cellular replication.

Ciclosporin (ranked 9th), an FDAapproved drug for psoriasis, is a powerful immunosupressant. (Enriched in Cluster 5, including RIG-i like receptor signaling pathway; Data not shown)

Conclusions

- In a Psoriasis example, cogena retrieves disease relevant mechanisms and drugs
- Cogena is a powerful Bioconductor package for drug repositioning and drug mode of action discovery
- Cogena is available from Bioconductor or https://github.com/zhilongjia/cogena

• Reproduciable research: http://zhilongjia.github.io/psoriasis/