

Co-expressed gene-set enrichment analysis for drug repositioning with examples of psoriasis and periodontal diseases

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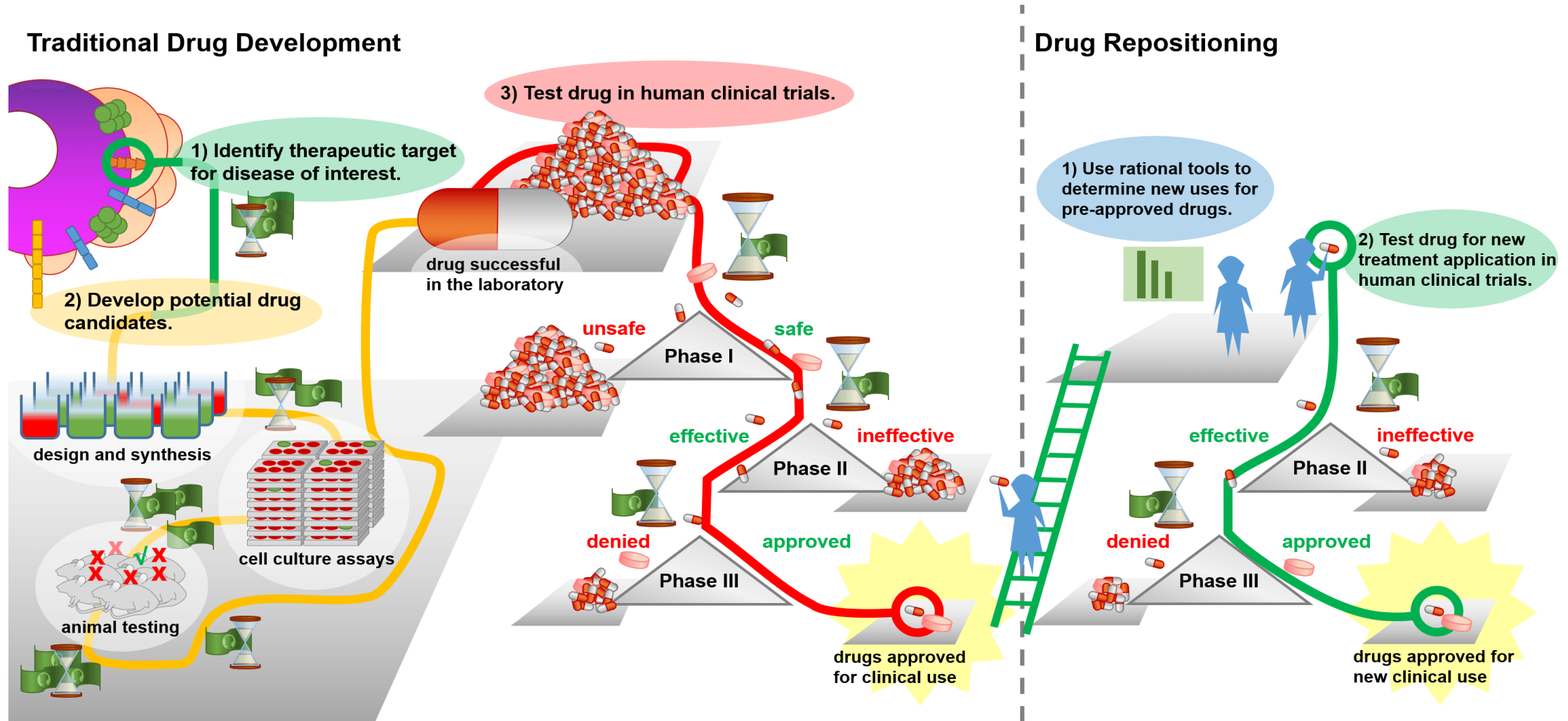
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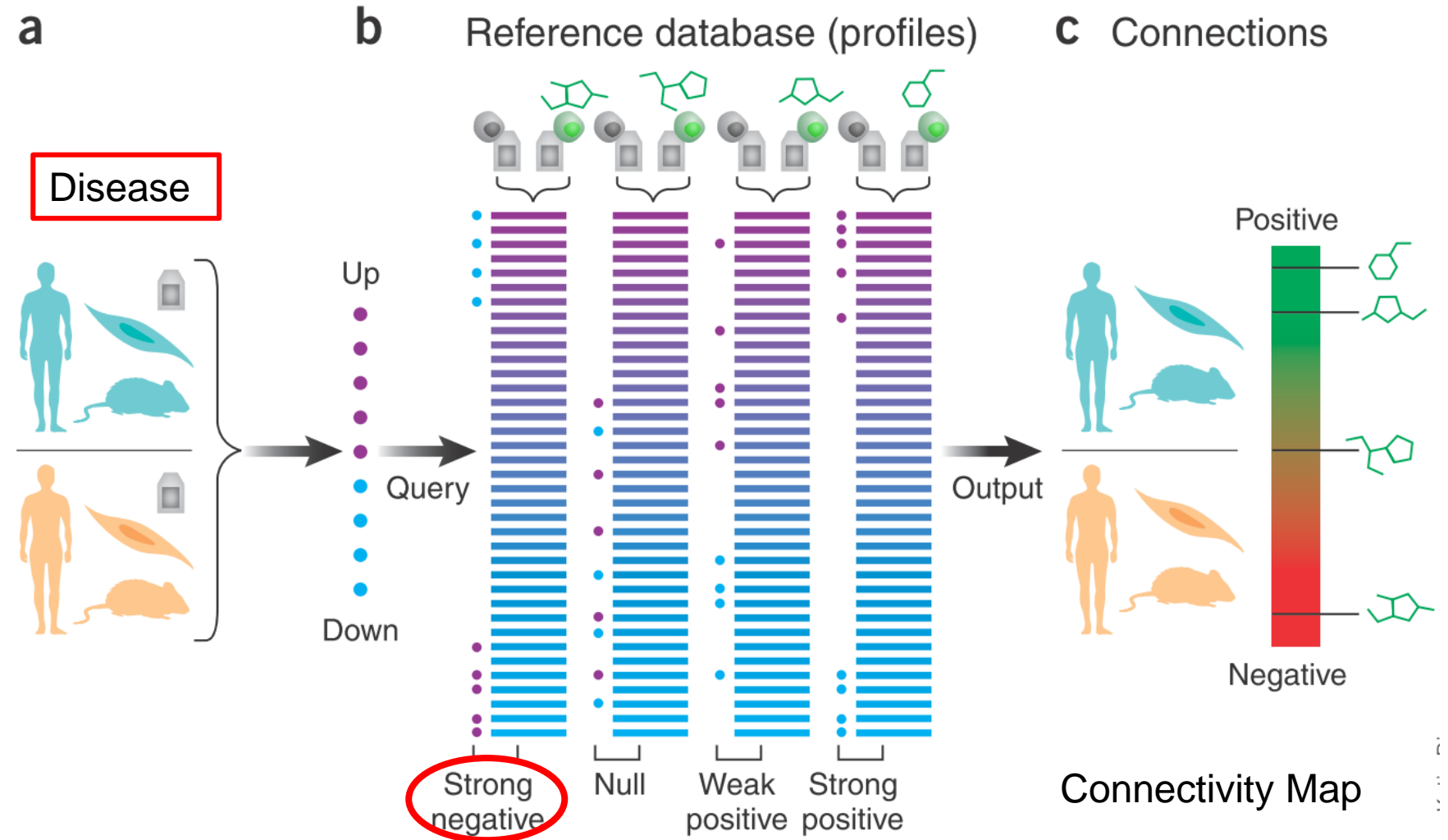
Drug repositioning: discovering new indications for approved drugs



- **Advantages of Drug repositioning:** Less steps, shorten period, more effective cost and lower risk!

Basic hypothesis of transcriptome-based drug repositioning

If a drug can recovery the changed genes caused by a disease, the drug probably can treat the disease.



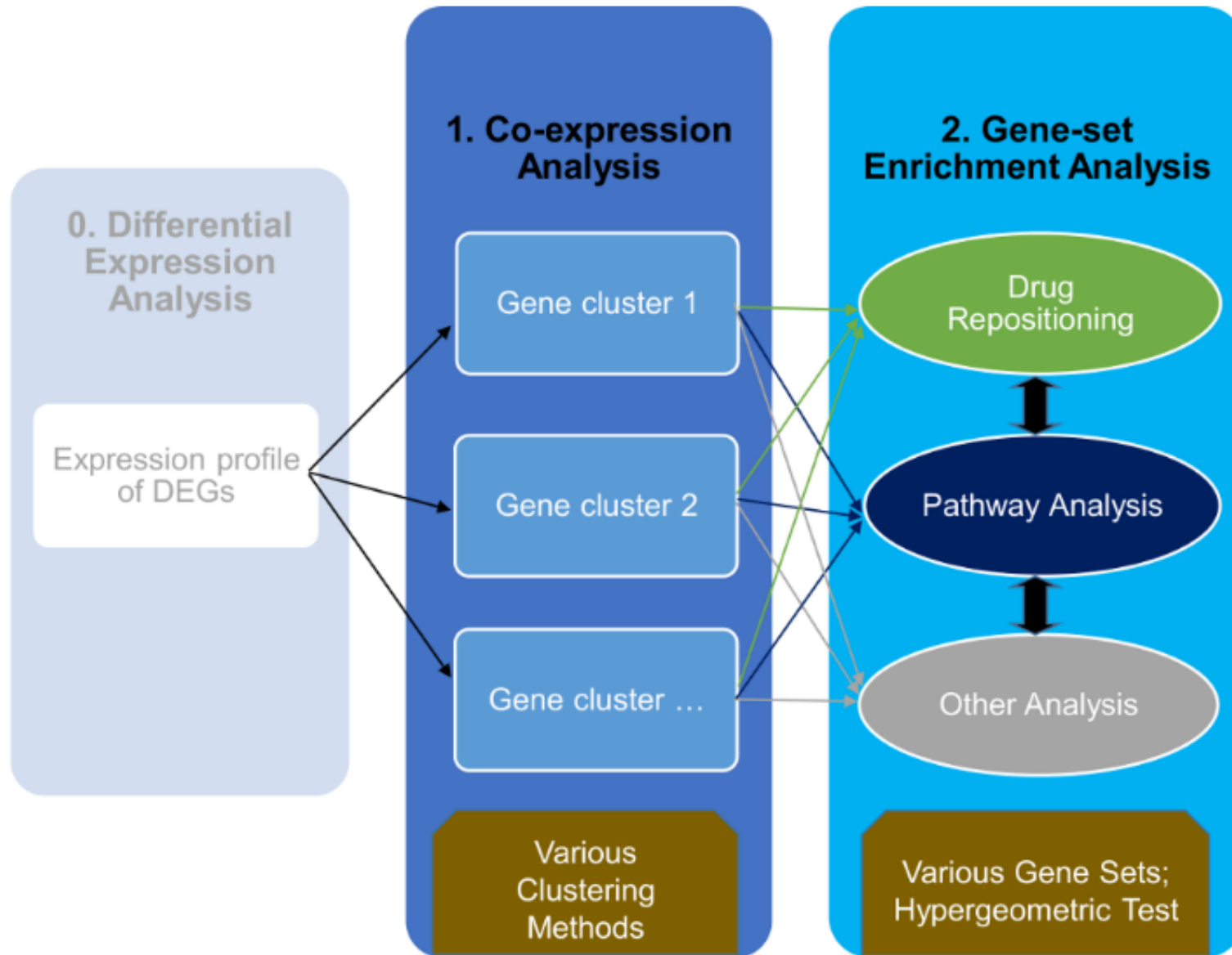
Lamb, Justin, et al. "The Connectivity Map: using gene-expression signatures to connect small molecules, genes, and disease." *science* 313.5795 (2006): 1929-1935.

Michnick, Stephen W. "The connectivity map." *Nature chemical biology* 2.12 (2006): 663-664.

Issues

- No versatile Co-expression analysis tool
 - In general, co-expressed genes functions collaboratively
 - They are probably involved in similar or same pathway or Gene Ontology
- No clue to the Drug Mode of Action (MoA) in the CMap-based drug repositioning pipeline
 - Pathway and GO are a clue to drug MoA

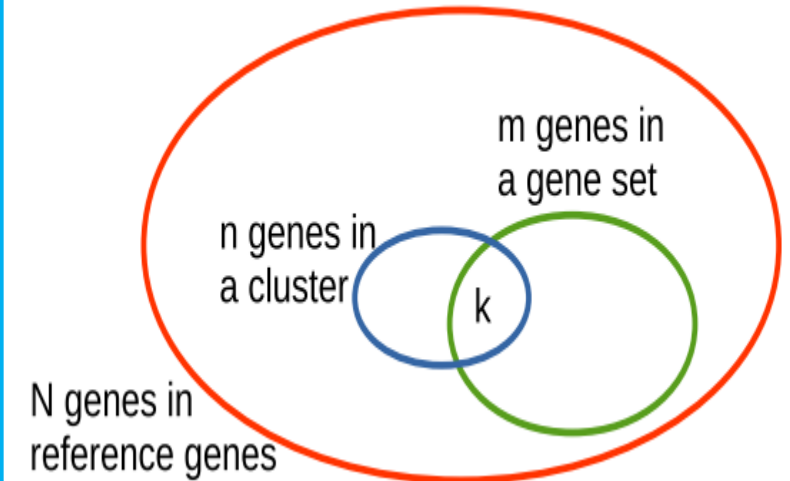
Cogena: Co-expressed gene-set enrichment analysis



Gene sets:



Hypergeometric test :

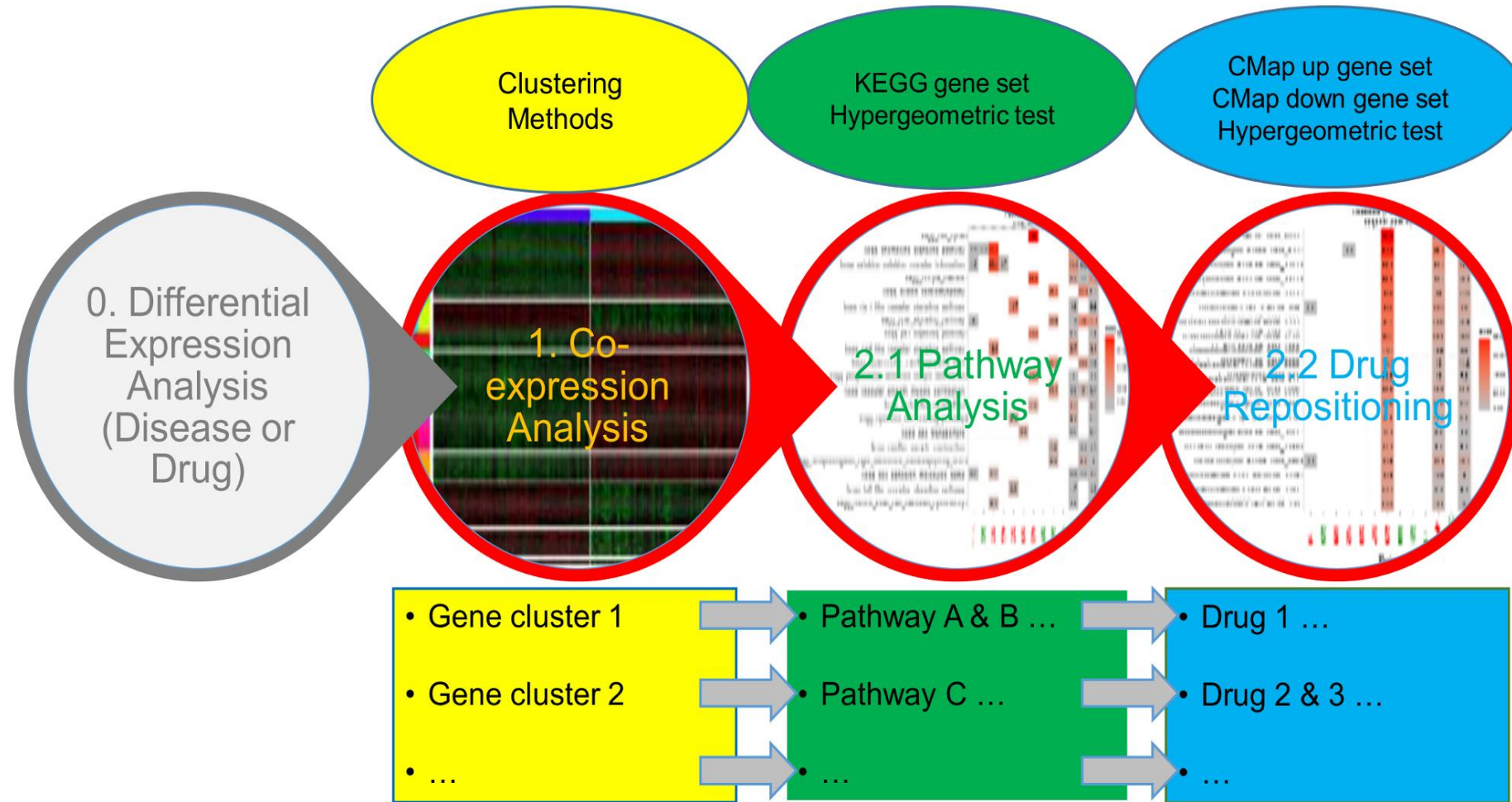


$$P\{x \geq k\} = \sum_{x=k}^{\infty} \binom{m}{x} \binom{N-m}{n-x} / \binom{N}{n}$$

$$ES = -\log_2(\text{FDR}) \quad 6$$

Cogena for drug repositioning

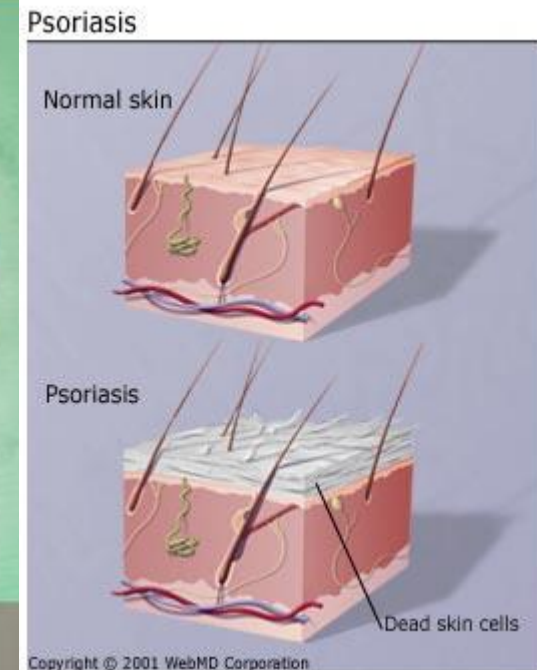
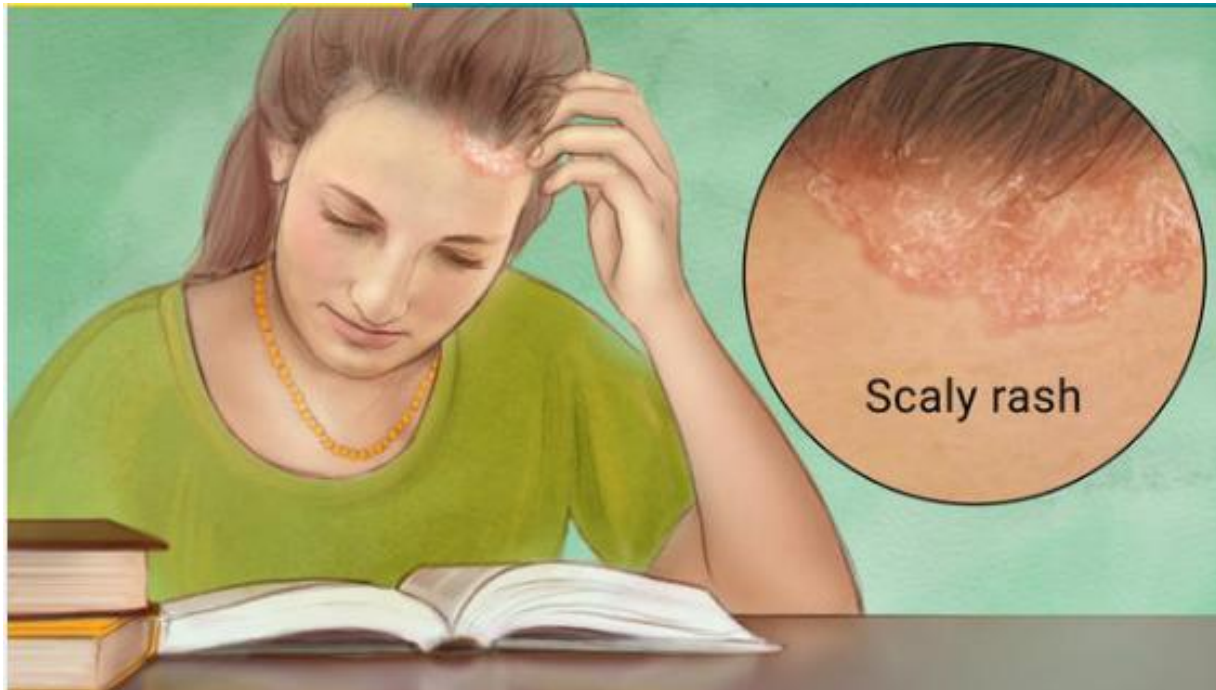
Targeting co-expressed genes could be more precise than all the DEGs.



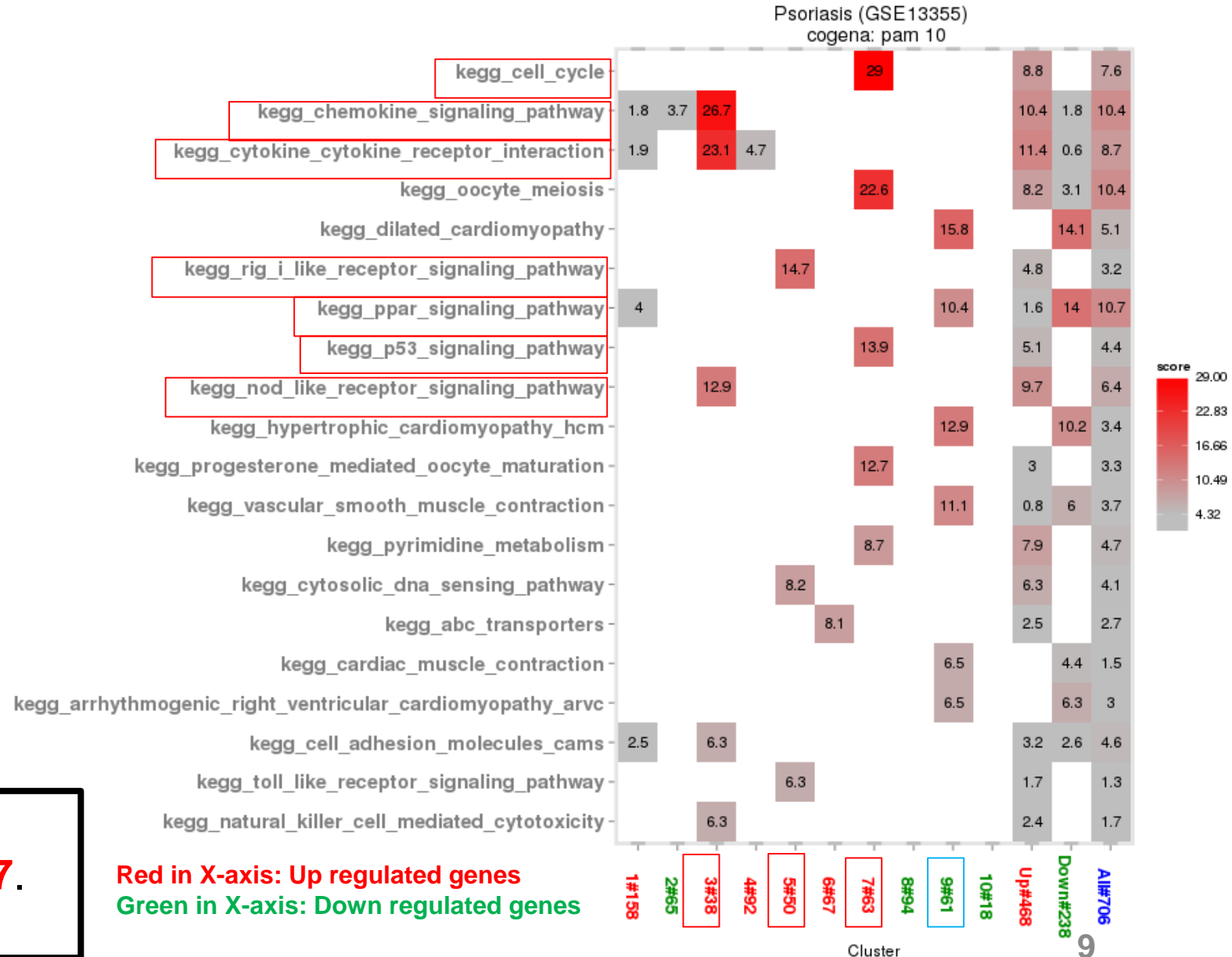
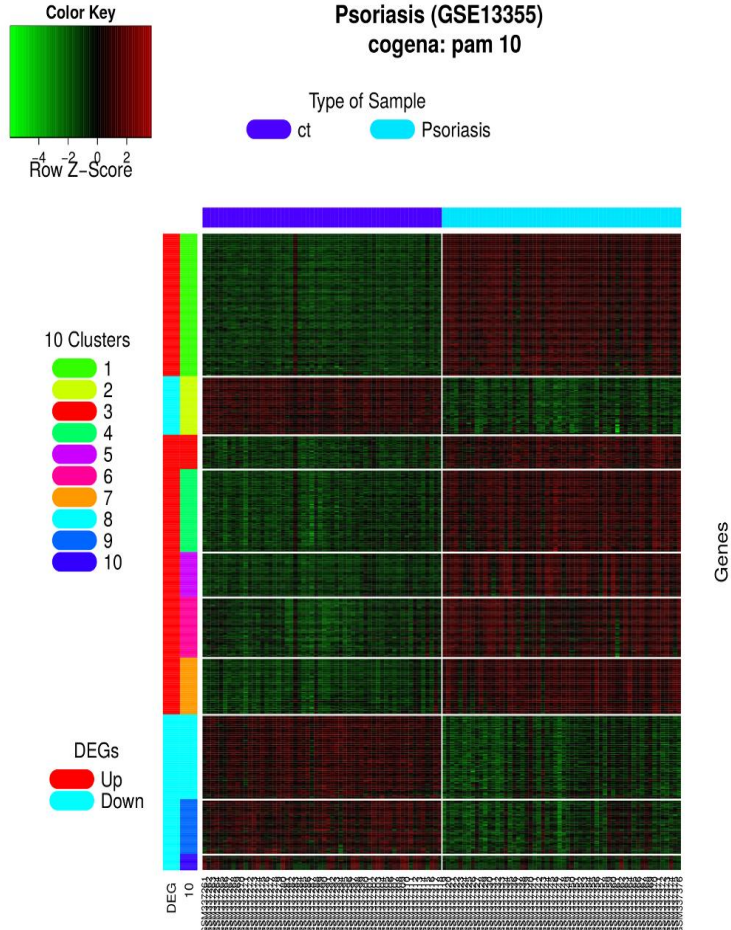
- Combination Pathway and Drug repositioning provides a clue to the MoA of Drug.

Cogena-based drug repositioning for psoriasis

- Psoriasis is a long-lasting autoimmune disease characterized by patches of abnormal skin.
- 2%-3% of people throughout the world.
- Treatments for moderate to severe psoriasis include Methotrexate and Biologics.



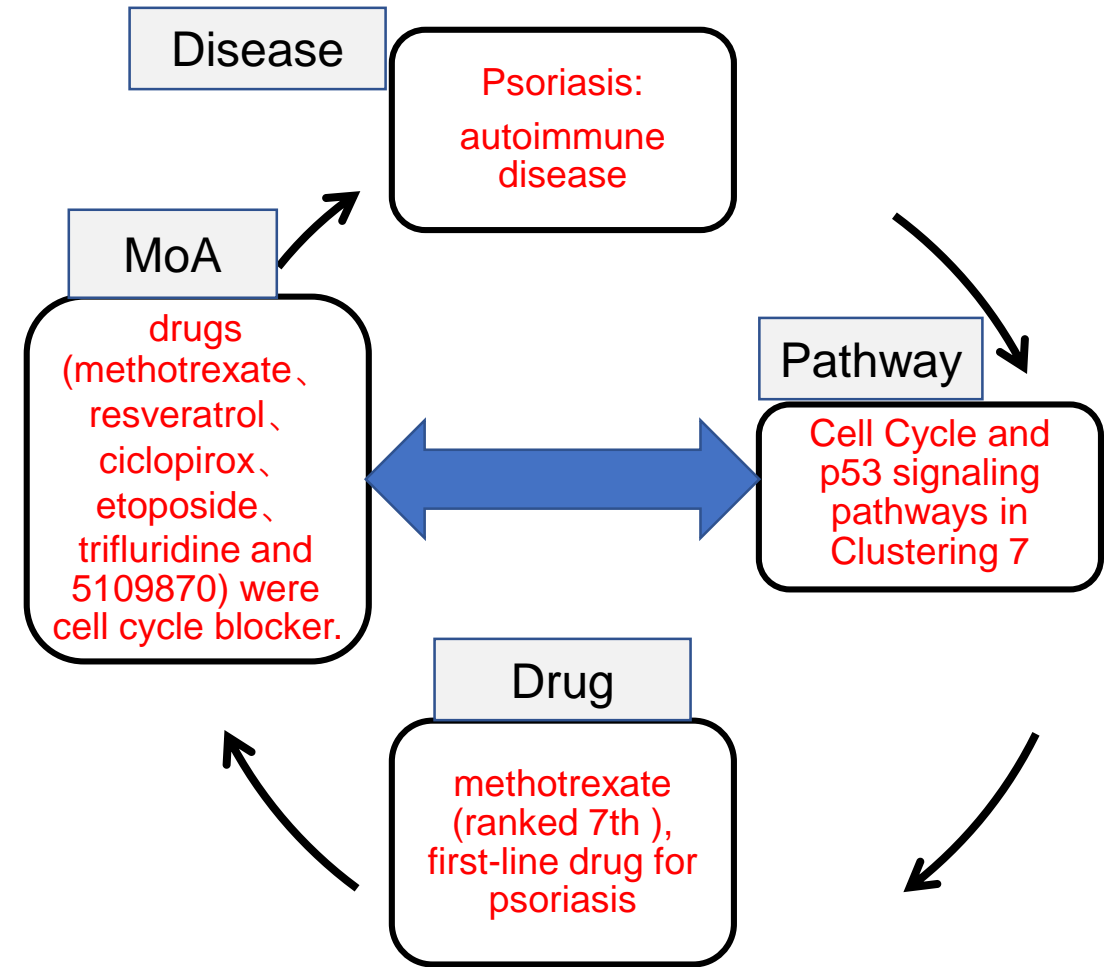
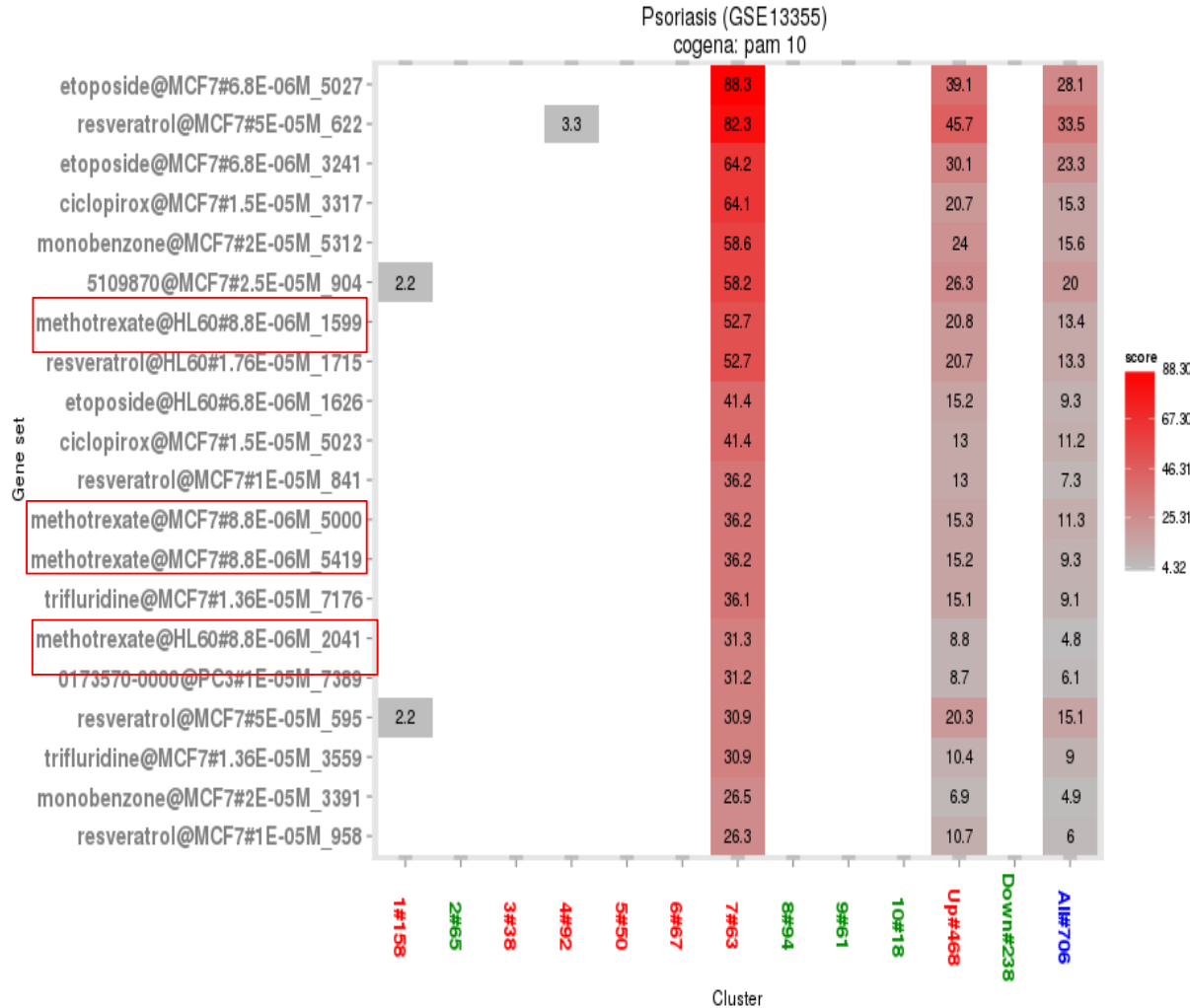
Cogena-based drug repositioning for psoriasis



Immune-related pathways in C3, C5.
Cell Cycle/p53 signaling pathways in C7.
PPAR signaling pathway in C9.

Red in X-axis: Up regulated genes
Green in X-axis: Down regulated genes

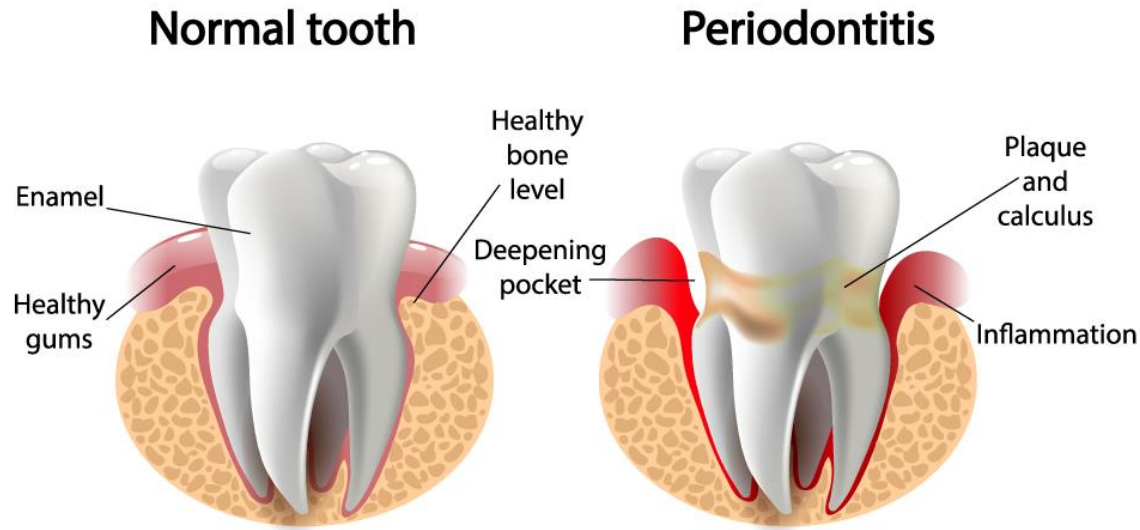
Cogena-based drug repositioning for psoriasis



Most drugs in cluster 7 are cell cycle blockers and probably could be promising candidate drugs for psoriasis and even more generally autoimmune diseases.

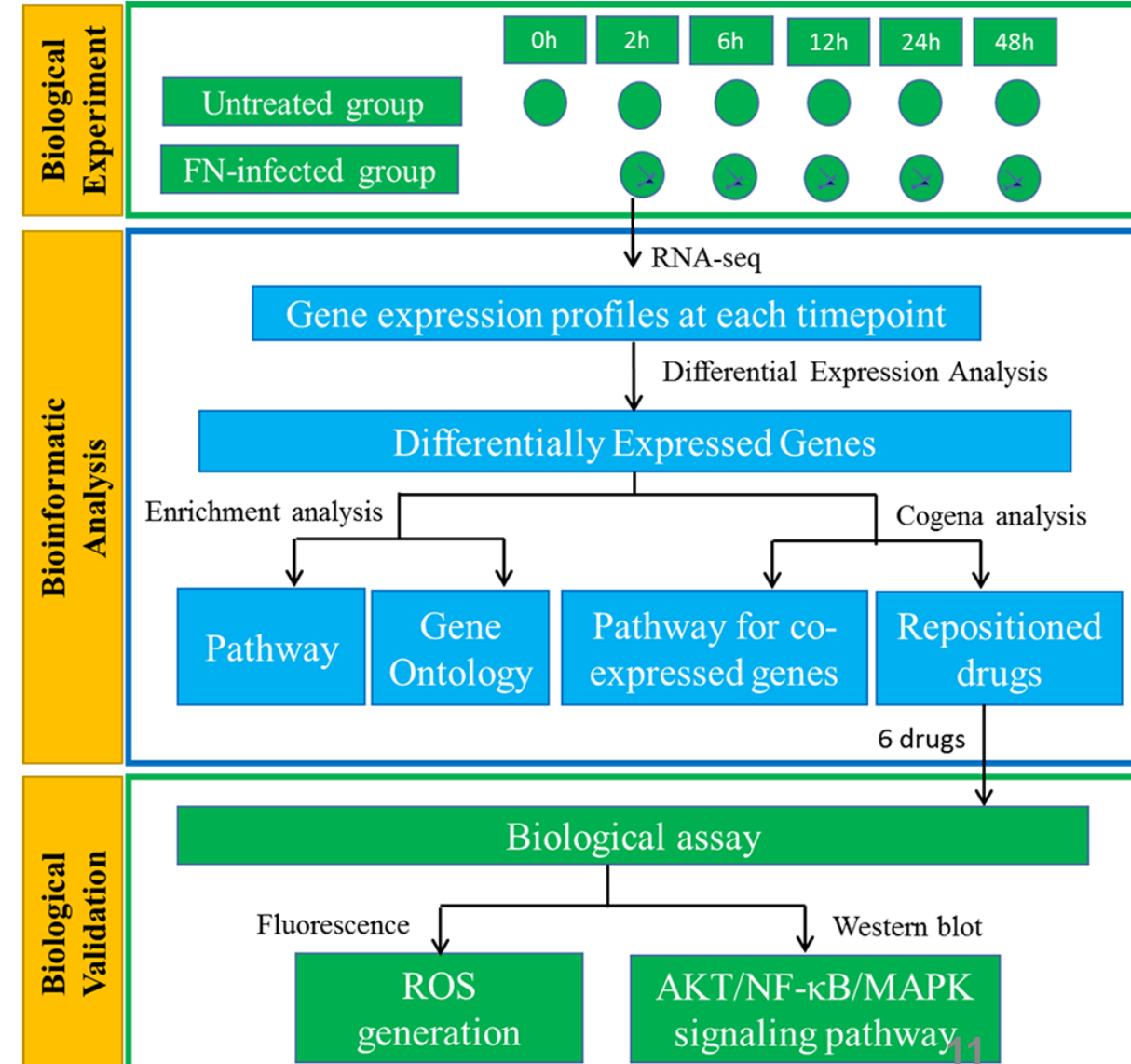
Cogena-based drug repositioning for periodontal diseases

- Periodontal disease is caused by the interaction of dental plaque biofilm and the host immune system.
- Fusobacterium nucleatum* (FN) is a high-frequency pathogen in periodontal disease.
- Gingival fibroblasts (GF) are the most abundant cell types in periodontal connective tissues

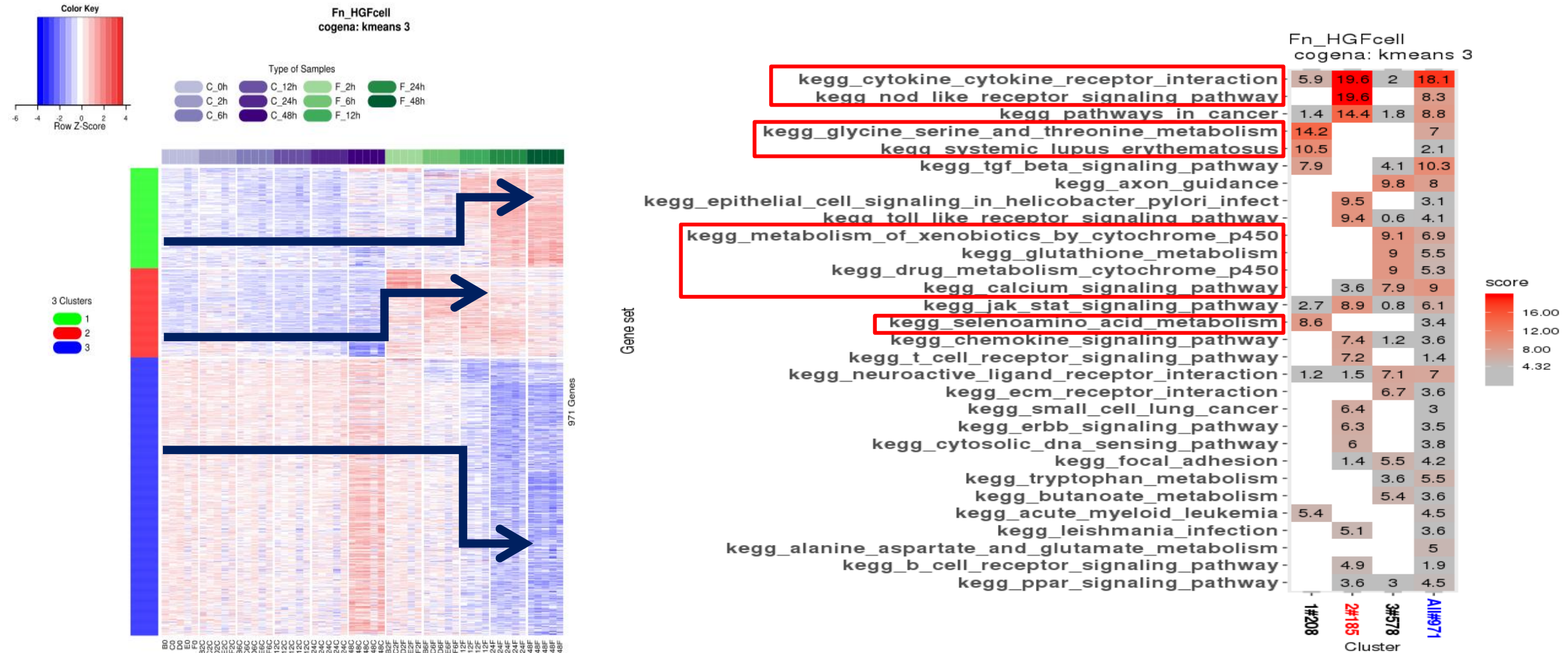


Ref (left pic): <https://www.saebo.com/the-hidden-connection-between-gum-disease-and-stroke/>

The entire experimental design



Cogena-based drug repositioning for periodontal diseases

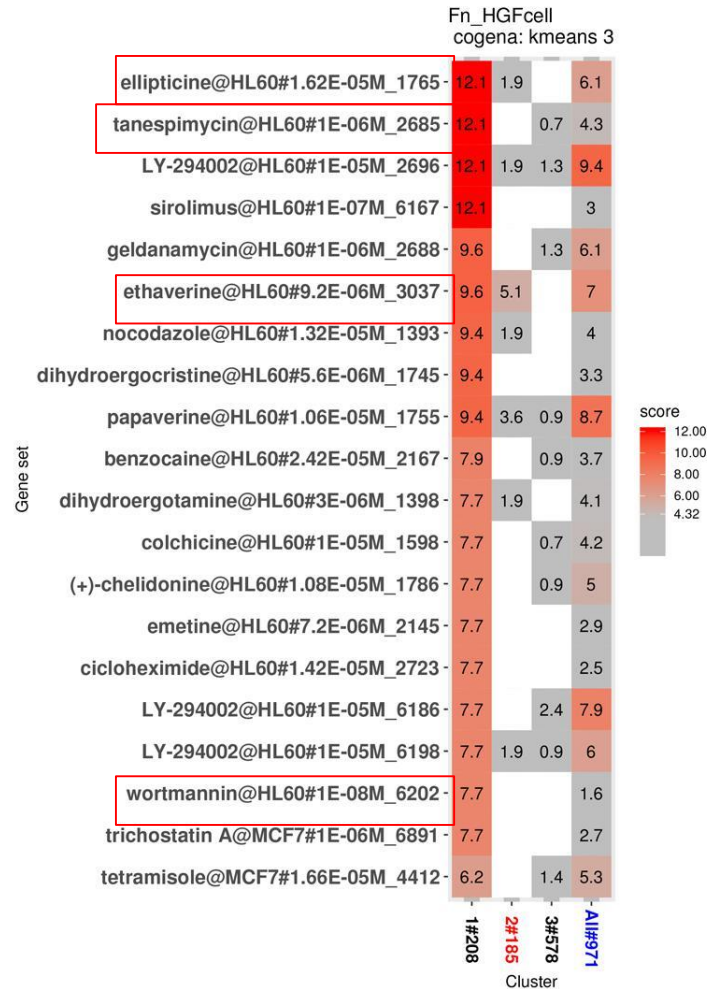


Cluster 1: up after 12h and Cluster 3: down after 12h -- mainly enriched in metabolism-related pathways
Cluster 2: up immediately-- highly enriched in the cytokine–cytokine receptor signaling pathways.

Cogena-based drug repositioning for periodontal diseases

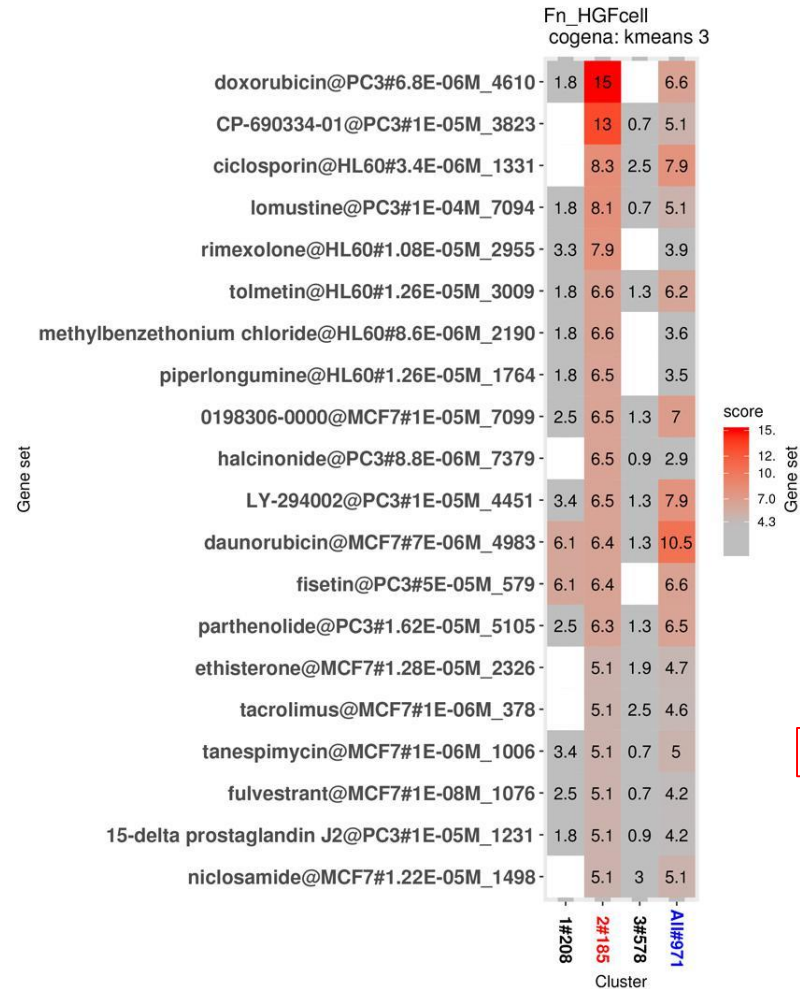
A

Drug candidates for cluster 1



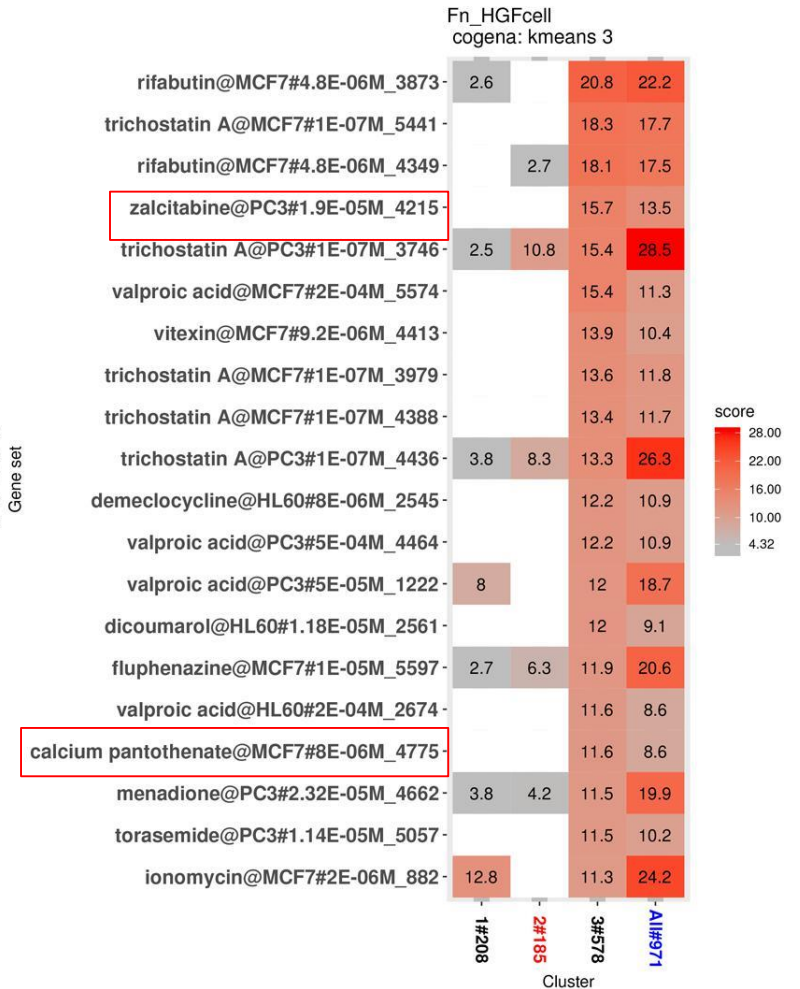
B

Drug candidates for cluster 2



C

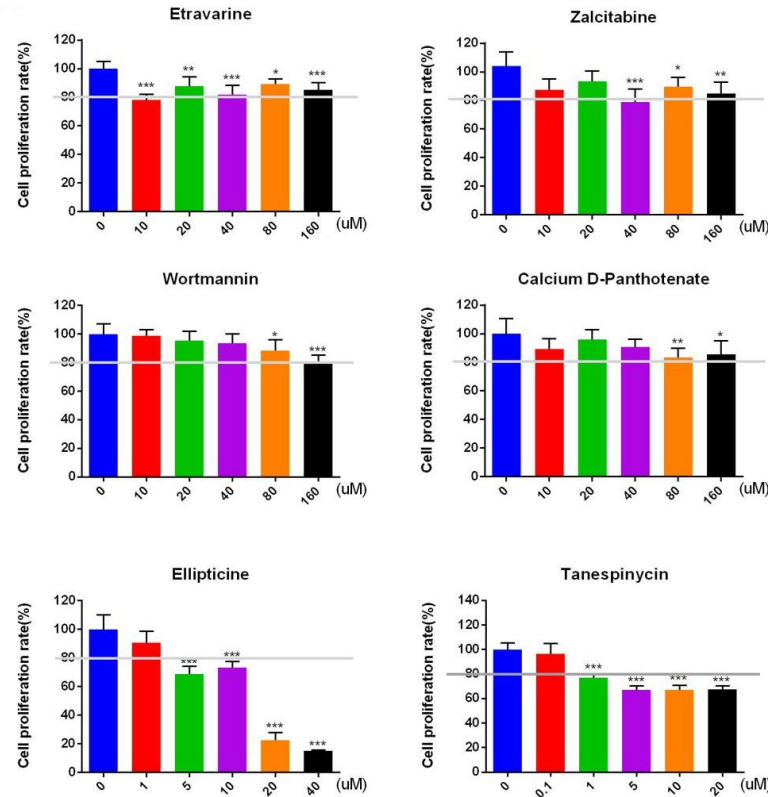
Drug candidates for cluster 3



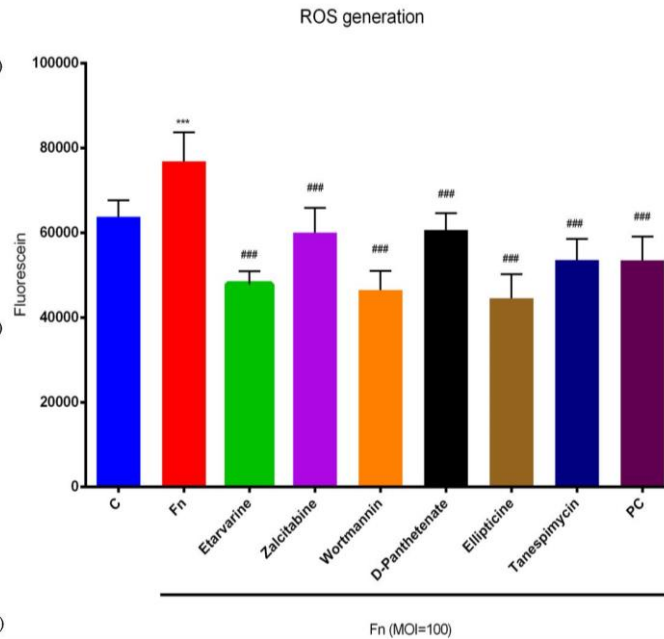
Six drugs (etravirine, zalcitabine, wortmannin, calcium D-pantothenate, ellipticine, and tanespimycin) were selected to be tested due to the scale of biological experiments and literature-based investigations.

Cogena-based drug repositioning for periodontal diseases

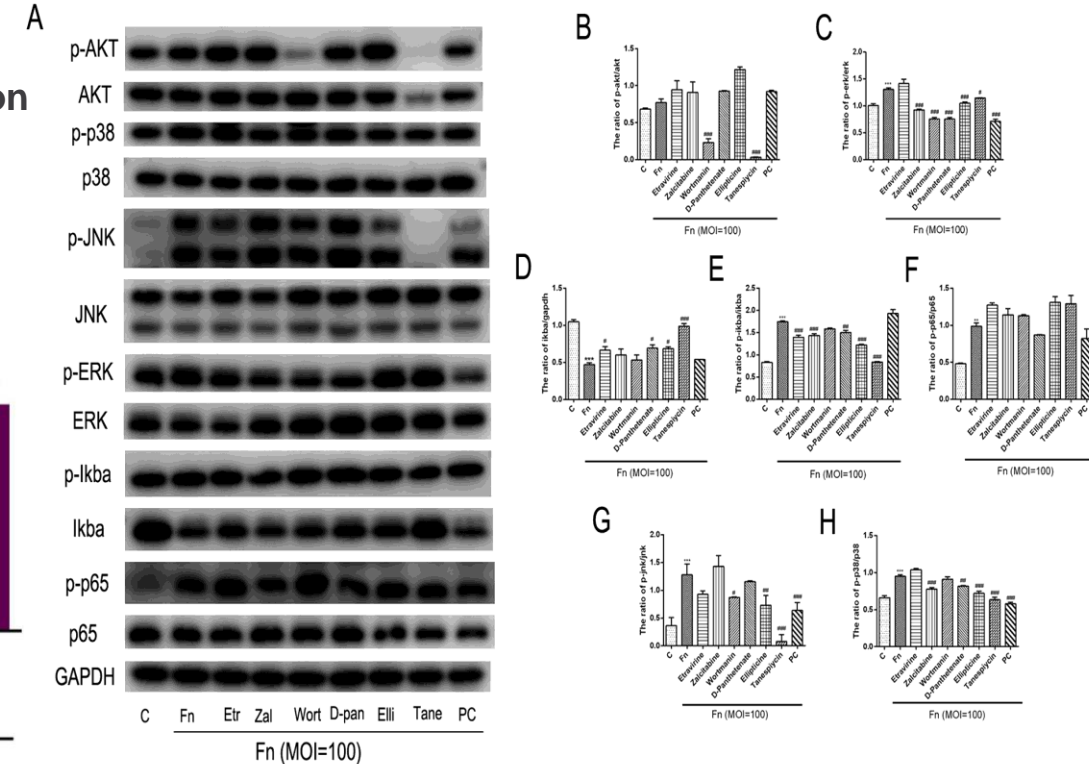
Effects of candidate drugs on GF proliferation



Effects of candidate drugs on F. nucleatum-induced ROS generation



Effects of candidate drugs on F. nucleatum-activated NF- κ B, MAPK, and AKT signaling pathways in GFs



Six drugs (etravirine, zalcitabine, wortmannin, calcium D-pantothenate, ellipticine, and tanespimycin) could significantly **decrease FN-induced ROS generation** and **block the PKB/AKT/MAPK signaling pathways**.

Kang, Wenyan, et al. "Time-course Transcriptome Analysis for Drug Repositioning in Fusobacterium nucleatum-infected Human Gingival Fibroblasts." *Frontiers in Cell and Developmental Biology* 7 (2019): 204.

Summary

- Cogenia is a tool of gene set enrichment analysis for co-expression genes.
- Cogenia can be used in drug repositioning and discovering the MoA of drugs, illustrated with two disease examples.
- Cogenia can be used to discovery similar drugs or treatable diseases (if the input of cogenia is drug signature)

Acknowledgements

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Reproducible Research

- **Cogena** Bioconductor Package:
 - <http://www.bioconductor.org/packages/cogena/>
 - <https://github.com/zhilongjia/cogena>
- **Psoriasis** project:
 - <https://github.com/zhilongjia/psoriasis>
- **Periodontal diseases** project:
 - <https://bigd.big.ac.cn/gsa/browse/CRA001739>
 - https://github.com/zhilongjia/Fn_HGFcell

Thank you
Q&A

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