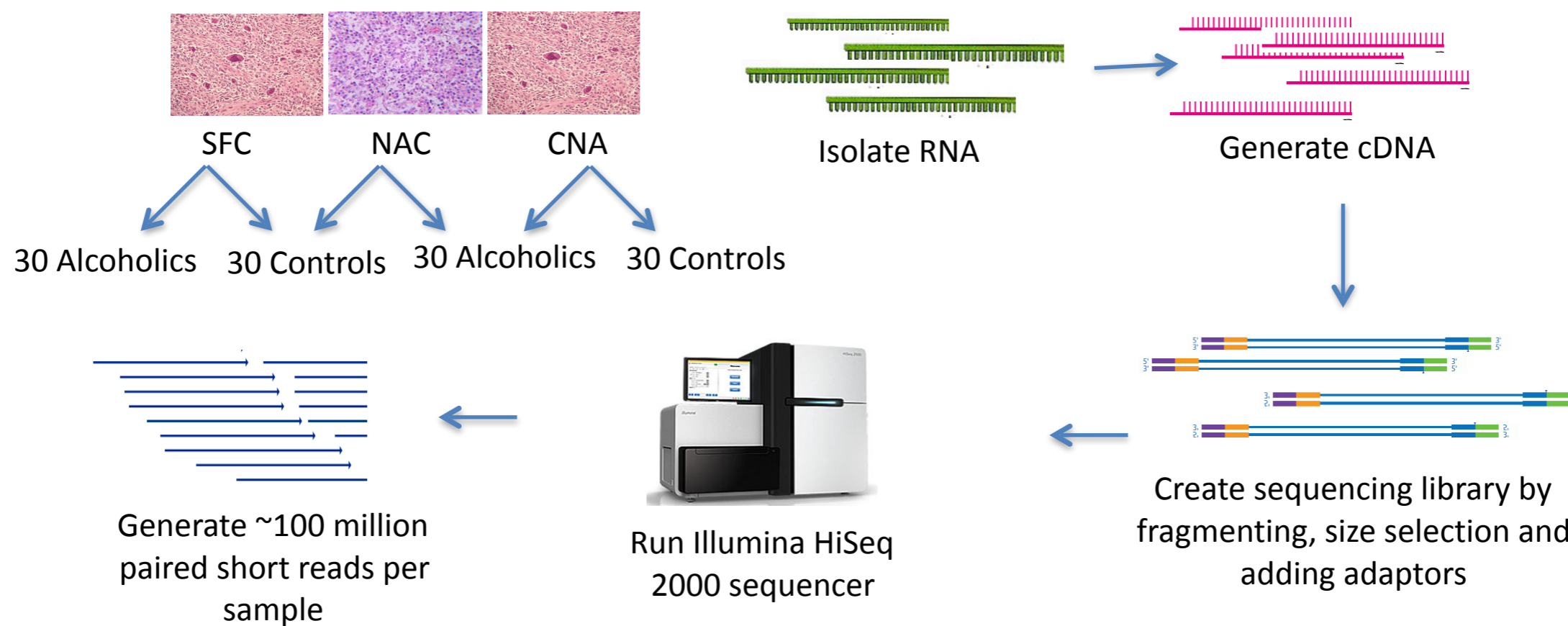


Downstream Analysis of RNA-Seq Data

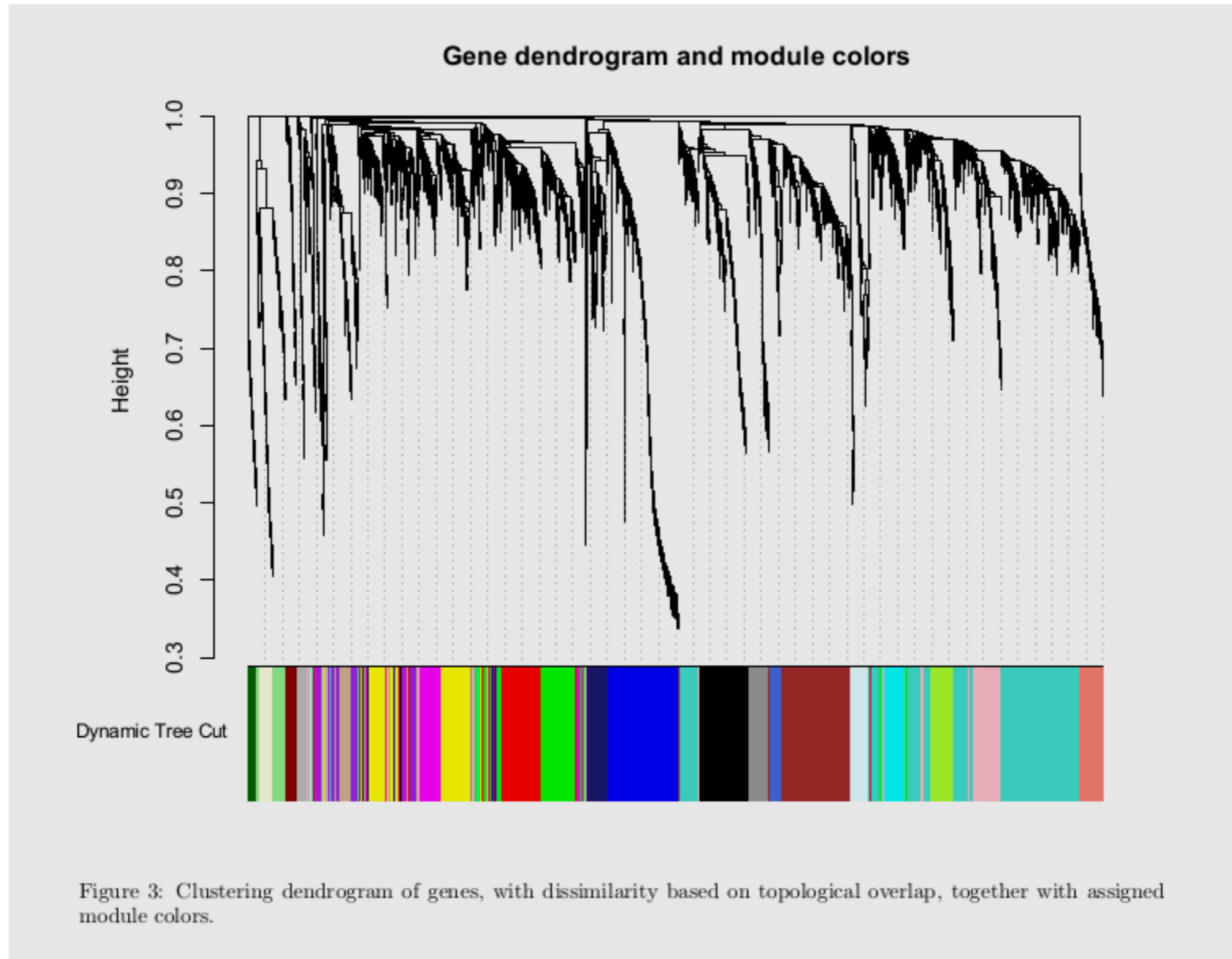
Example RNA-Seq Project

Transcriptome profiling of alcoholism

How does gene expression differ in Alcoholics vs. Healthy brains? How do these gene expression profiles vary across three different brain regions?

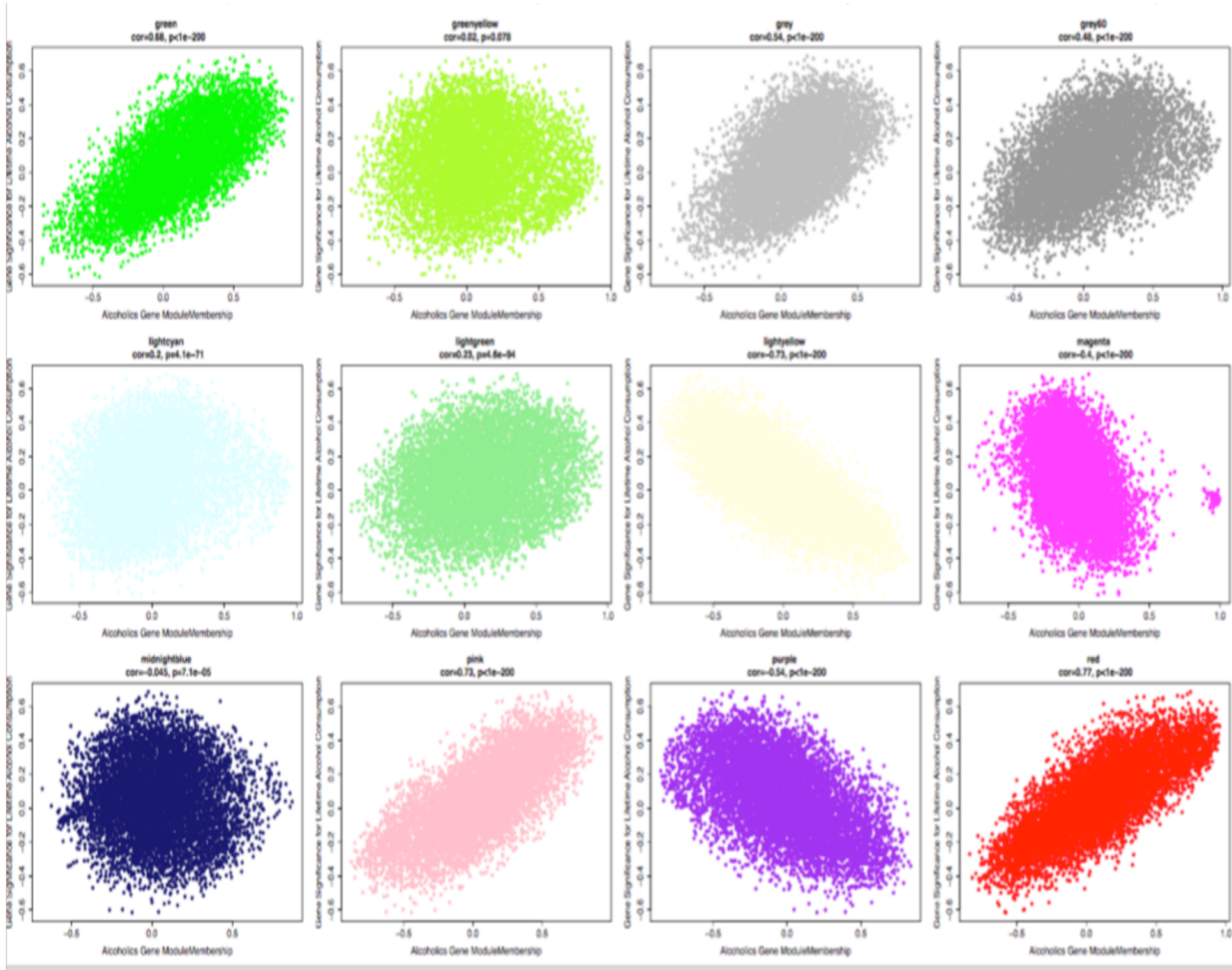


Identify groups of genes with correlated expressions



Identify groups of genes with correlated expressions

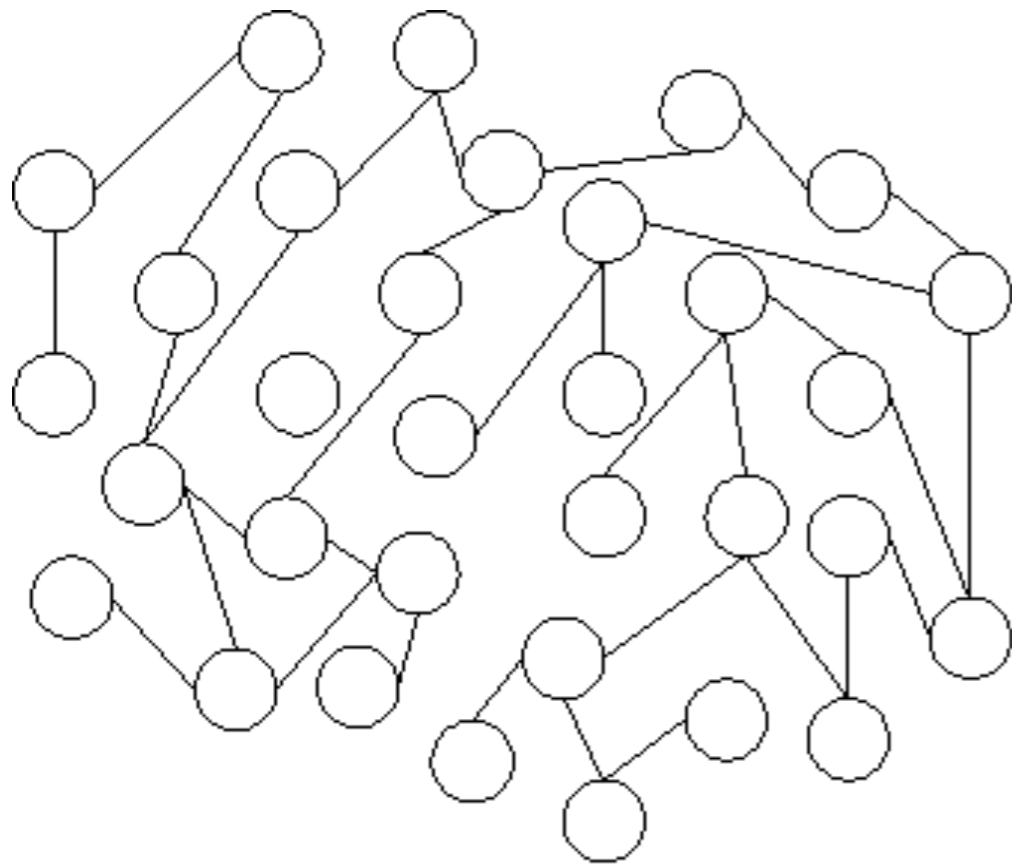
Module Correlation to Lifetime Alcohol Consumption in CNA Brain Region



COEXPRESSED
GENE
MODULES

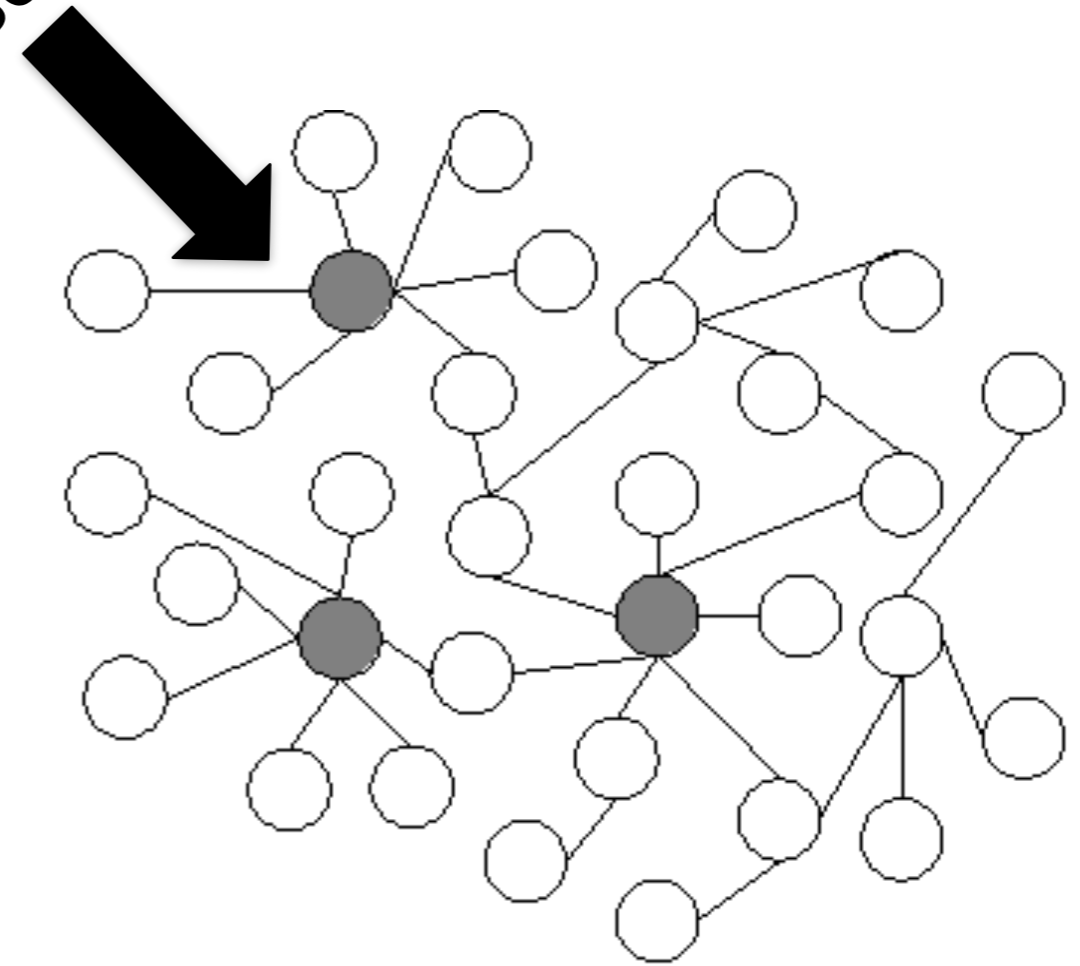
Identify groups of genes with correlated expressions

GENES DRIVING
CHANGE



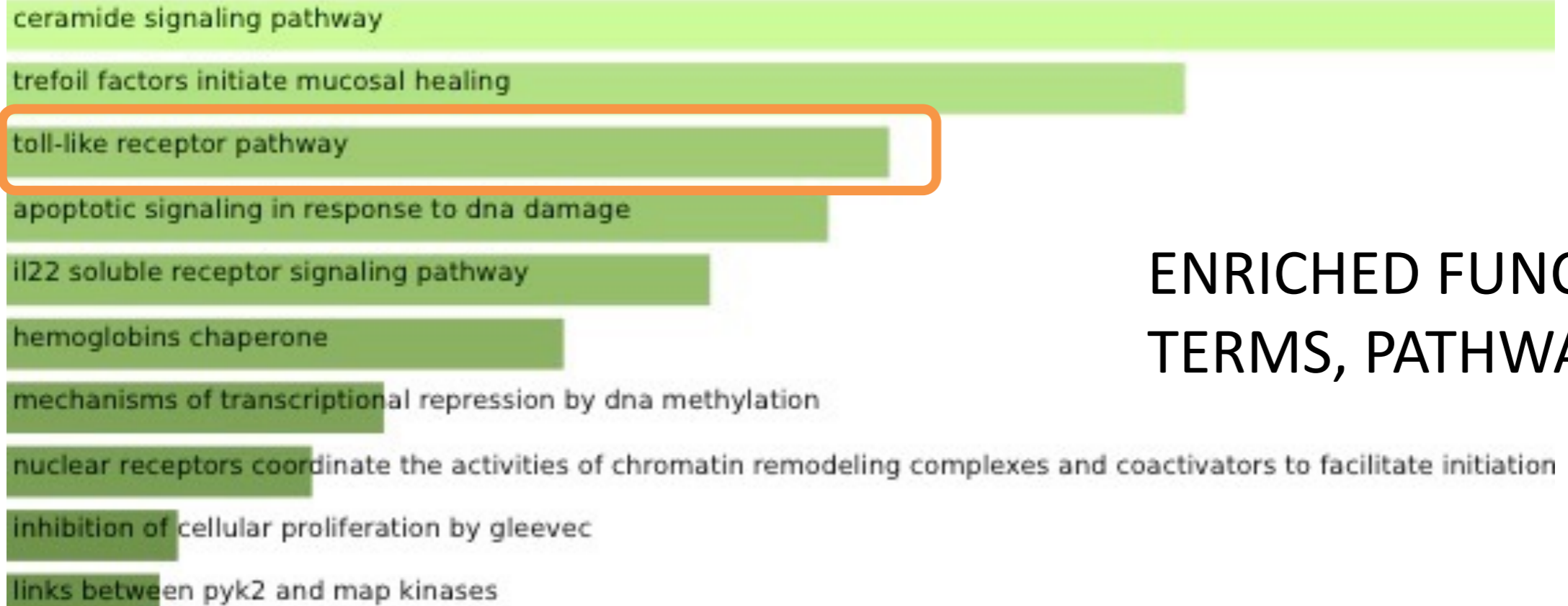
(a) Random network

transcription factor?

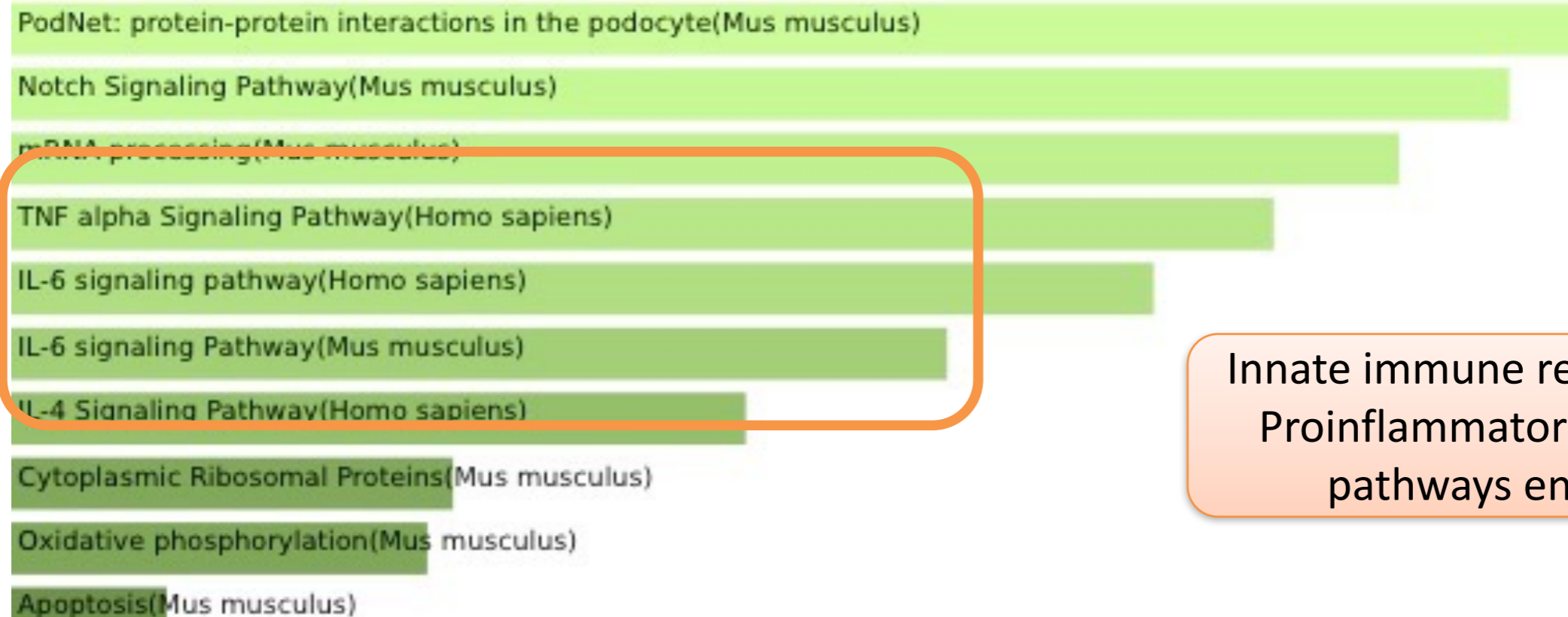


(b) Scale-free network

Pathway/Functional Group Analysis



ENRICHED FUNCTIONAL TERMS, PATHWAYS

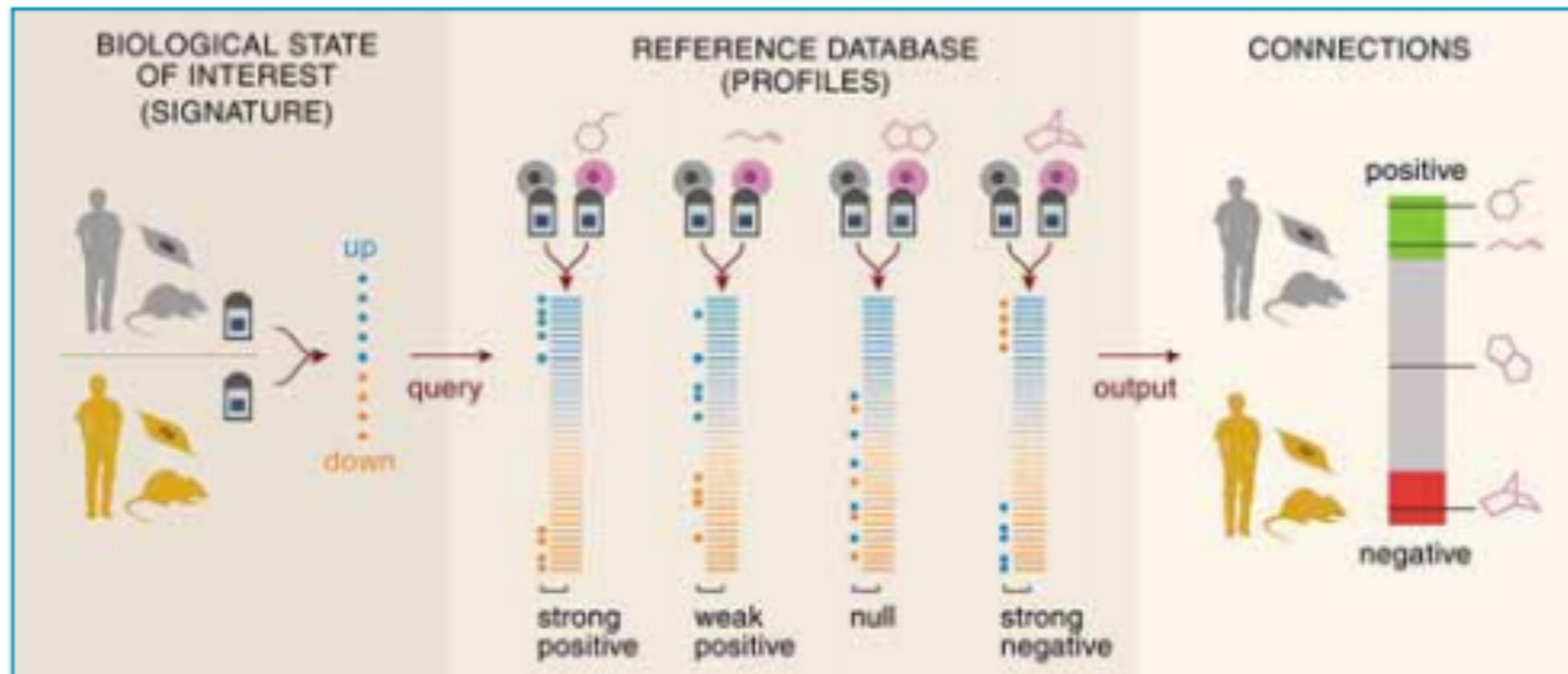


Innate immune response and Proinflammatory cytokine pathways enriched

Connecting Expression to Drug Compounds

DRUG COMPOUNDS WHICH CAN REGULATE GENE CHANGES

USING LINCS DATABASE (library of integrated cellular signatures)



Connecting Expression to Drug Compounds

DRUG COMPOUNDS WHICH CAN REGULATE GENE CHANGES

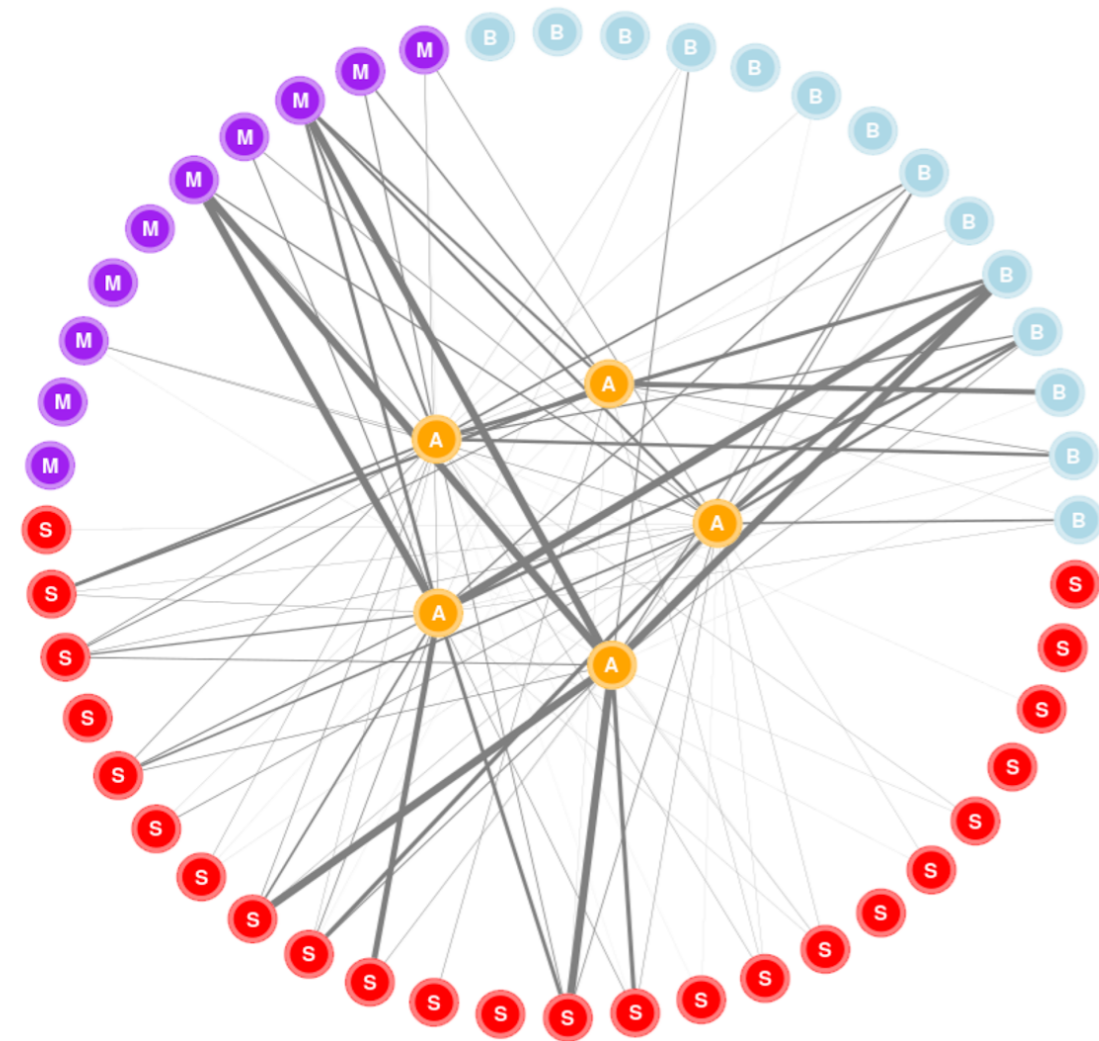
FDA Approved drugs in NAC Module

Drug Name	FDA Status	Connectivity Score
formestane	Approved, Investigational, Withdrawn	-95.11
metolazone	Approved	-96.17
Erlotinib	Approved	-96.38
timolol	Approved	-95.3
finasteride	Approved	-95.65
flunarizine	Approved	-98.77
alprazolam	Approved, Illicit, Investigational	-95.95
ipratropium bromide	Approved	-96.71
labetalol	Approved	-95.56
etodolac	Approved, Investigational, Vet Approved	-95.18
clofibrate	Approved	-95.91

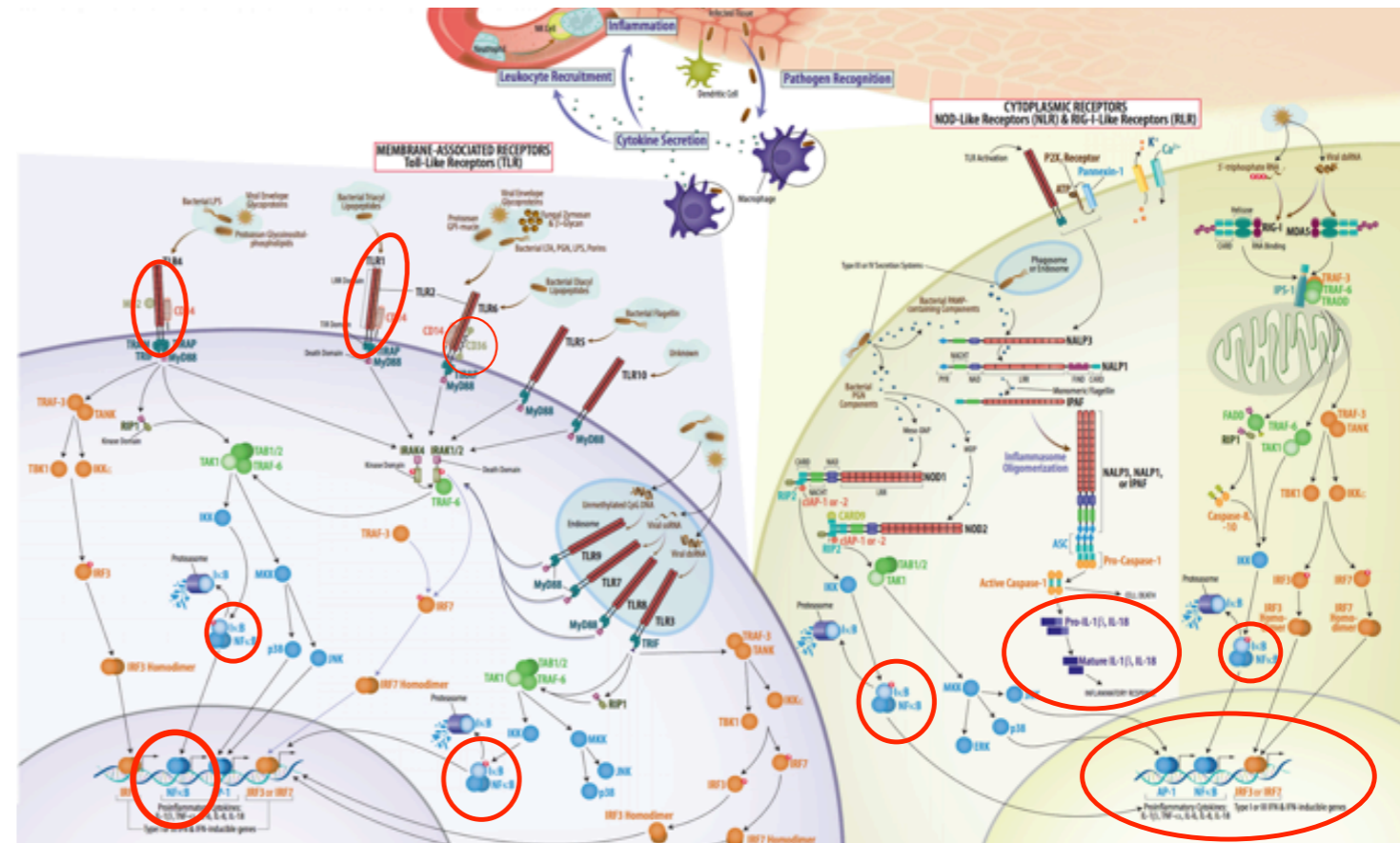
Contributed by Bo Miller

Comparing to Other Datasets

Comparison of genes co-expressed in alcohol abuse to other psychiatric disorders (Bipolar, depression, schizophrenia) using publicly available data



Modules
 ● BPD
 ● MDD
 ● SCH
 ● Alcoholism



Contributed by Khoi Le, Lee Rao