

Seminar

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1 End of MCnebula

2 Next Research

3 END

End of MCnebula

Pathway enrichment ACs

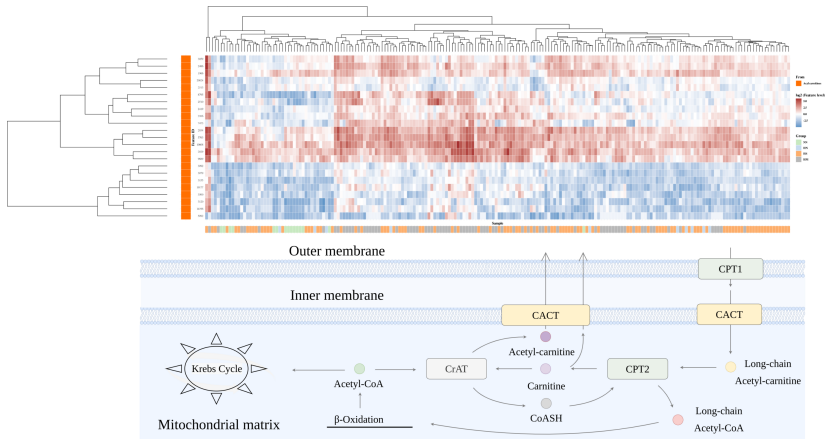


Figure 1: ACs

Pathway enrichment LPCs

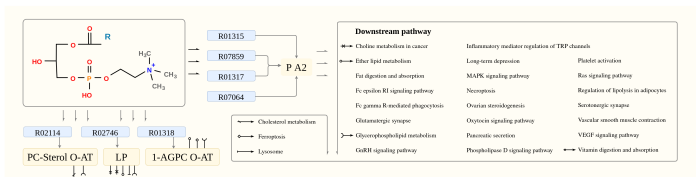
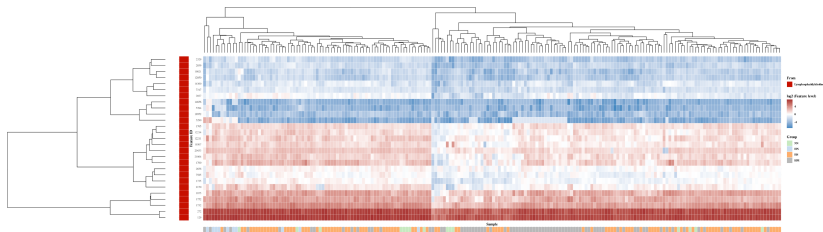


Figure 2: LPCs

Pathway enrichment BAs

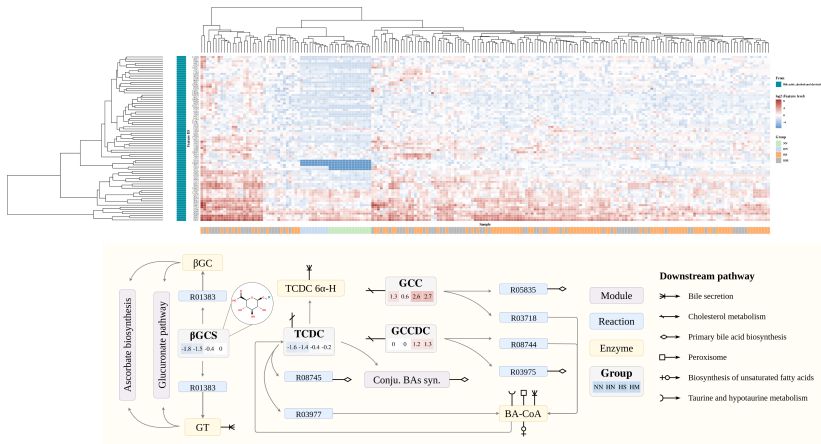


Figure 3: BAs

MCnebula in github

The screenshot displays the GitHub interface for the repository `Cao-lab-zcmu / MCnebula`. The repository is public and has 0 notifications, 0 forks, and 0 stars. The main navigation bar includes links for Code, Issues, Pull requests, Actions, Projects, Wiki, Security, and Insights. The repository structure is shown as follows:

File/Folder	Description	Last Commit
echo revise		aa7575b 4 days ago 56 commits
R	Merge branch 'ref'	6 days ago
data-raw	instance_data	2 months ago
data	instance_data	2 months ago
inst/extdata	instance_data	2 months ago
man	object documentation	6 days ago
tests	initial submit	3 months ago
vignettes	initial submit	3 months ago
.Rbuildignore	instance_data	2 months ago
.gitignore	initial submit	3 months ago

The right sidebar contains the 'About' section, which describes the repository as 'MCnebula algorithm integration in R'. It also includes links to the README, View license, and statistics: 0 stars, 1 watching, and 0 forks. Below this, the 'Releases' section shows 'No releases published', and the 'Packages' section shows 'No packages published'.

Figure 4: MCnebula

Next Research

Preliminary research route

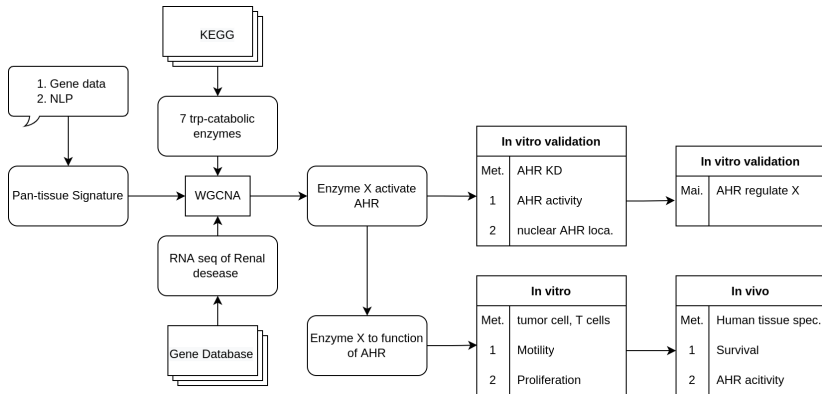


Figure 5: research route

Microarray and RNA-seq analysis

limma:

Linear Models for Microarray and RNA-Seq Data
User's Guide

Gordon K. Smyth, Matthew Ritchie, Natalie Thorne,
James Wettenhall, Wei Shi and Yifang Hu
Bioinformatics Division, The Walter and Eliza Hall Institute
of Medical Research, Melbourne, Australia

First edition 2 December 2002

Last revised 14 November 2021

Figure 6: limma

Experimental design

Table 1: In vivo experiment with rats

No.	group	drug	n	low	medium	high
1	control	-	15	-	-	-
2	model	-	15	-	-	-
3	raw.eu	raw.eu	15	5	10	20
4	pro.eu	pro.eu	15	5	10	20
5	positive	pos.drug	15	-	-	-
6	extra	MTA	15	-	-	-

Processing and extracting of *E. ulmoides*

Known Variables

$$D_{base} = 15 \text{ (g)}$$

$$w_{rat.weight} = 400 \text{ (g)}$$

$$w_{hum.weight} = 60,000 \text{ (g)}$$

$$C_{coefficient} = 6$$

$$T_{times} = 28 \text{ (day)}$$

$$n_{group.rat.number} = 15$$

$$\begin{aligned} M_{group.h.m.l} &= 2 \times (5 + 10 + 20) \\ &= 70 \text{ (multiple)} \end{aligned}$$

Require *E. ulmoides* (W_{eu}):

$$\begin{aligned} W_{eu} &= D \times \frac{w_{rat.}}{w_{hum.}} \times C \times T \times n \times M \\ &= 17640 \text{ (g)} \\ &= 17.640 \text{ (kg)} \end{aligned}$$

Figure 7: Material

END