





# Perseus-Powered Transcriptomics: Omics Insights Made Easy

This workshop will **focus** on topics ranging from **basic experimental design** to **advanced downstream analyses** of **RNA-seq** data as well as **tips to publish in high-impact journals**.

Upon the completion of this **3-day hybrid hands-on workshop**, participants will be **capable** of the following:

- Conceptualising a transcriptome experiment with different considerations
- **Understanding** a complete **workflow** of RNA-seq data acquisition, processing, assembly, functional annotation, analysis, and interpretation
- Aware of the different analysis workflows and software packages in RNA-seq analysis
- **Exposure** to different approaches for **integrating** transcriptomics with other **omics** towards **systems understanding**
- Network opportunity with other researchers in the field & sharing of knowledge via voluntary flash talks
- Hands-on in *de novo* transcriptomic analysis and mining data
- Hands-on in downstream **functional analyses**: WEGO, KEGG pathway, clustering, functional enrichment

#### Why should you attend

- ✓ You're a researcher, academician, or student interested in transcriptomics and its applications.
- ✓ You're interested in advanced skills in RNA-seq analysis.
- ✓ You want user-friendly platforms for the interpretation of RNA-seq data.
- ✓ You got the transcriptome **results** from the sequencing vendor but not sure how to **interpret** them.
- ✓ You wish to analyse the transcriptome data hands-on, especially as a biologist instead of a data scientist.
- ✓ You want to learn on how to mine for biological knowledge from transcriptome data without programming/coding.
- ✓ You work on a transcriptomic project and want to learn on how to **publish** your results as a **high-impact** journal article.

This workshop is opened to **local** & **international** participants, from **beginner** to **advanced researchers** in **molecular biology** / **omics**. The **content** of this workshop is **transferable** to other omics studies such as **proteomics**.

# 25-27 JUL 2023

[0900-1700 GMT+8]

**VENUE: HYBRID** 

### **TRAINER**

Assoc. Prof. Dr. Goh Hoe Han

FEE

**Online** 

**MYR300** 

**USD80** (International)

On-site INBIOSIS
MYR750

# **Register Now!**

\*Priority will be given to those who made payment before 14<sup>th</sup> Jul 2023

Organised by





Further details inbiosis.workshop@gmail.com +603 8921 4547/4558/4549

#### **TENTATIVE PROGRAMME**

#### Day 1: 25 Jul (Tue)

Buy 1. 20 out (10c)		
Time	Activity/Event	
0830	Registration	
0900	Introduction	
	Lecture 1: Principles of transcriptomics	
1030	Morning Break	
1045	Lecture 2: A crash course on RNA-seq analysis	
1230	Technical talk 1 – Current developments of NGS	
1300	Lunch Break	
1400	Practical 1: Perseus for data pre-processing/exploration	
1600	Afternoon Break	
1615	Participant flash talks, Summary, discussion & wrap-up	
1700	End of Day 1	

#### Day 2: 26 Jul (Wed)

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Time	Activity/Event	
0900	Introduction to Day 2 & Recap	
0910	Lecture 3: Exploring RNA-seq analysis for biological discovery	
	(Functional Annotation, DEG & Downstream Analyses)	
1030	Morning Break	
1045	Practical 2: Statistical analysis with Perseus	
1230	Technical talk 2 – Single-cell RNA-seq	
1300	Lunch Break	
1400	Practical 3: Advanced statistical analysis: Multiple testing correction and FDR	
1600	Afternoon Break	
1615	Participant flash talks, Summary, discussion & wrap-up	
1700	End of Day 2	

## Day 3: 27 Jul (Thu)

Time	Activity/Event
0900	Introduction to Day 3 & Recap
0910	Lecture 4: RNA-seq downstream analyses using Perseus
1030	Morning Break
1045	Practical 4: Cluster analysis and visualisation
1230	Technical talk 3 – Applications of RNA-seq
1300	Lunch Break
1400	Practical 5: Functional enrichment analysis
1600	Afternoon Break
1615	Summary, discussion & conclusion
1700	End of Workshop

# **LOCATION MAP**

