Introduction to Bulk RNA-seq Data Analysis in R

In this workshop, you will learn how to analyze bulk RNA-seq data in R.

Session 1	Introduction to R
Session 2	Analyze bulk RNA-seq data with DESeq2 package
Session 3	Pathway analysis in R

This workshop is for biologists interested in learning how to work with R to analyze their bulk RNA-seq data and perform pathway analysis.

You do not need any prior knowledge of R for this workshop.

This will be a code-along, in-person workshop, and each session will take 2 hours.

https://forms.gle/vwx1PSmT67MCviW16

Scan the QR code or click on the link to fill out the form to find the best time for holding the workshop.

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