

# Preparation

## ASSUMED BACKGROUND

If you are not familiar with any of these topics, or have forgotten about them, then please look them up on Wikipedia, for example, before the conference.

**Linear algebra:** vector space; linear map; matrix; kernel, image, cokernel, and rank of a linear map; rank-nullity theorem; row reduction of a matrix; (real) inner product space; (real) Hilbert space; the sequence space,  $\ell^2$ ; the space of square-integrable functions,  $L^2$

**Statistics:** statistical hypothesis testing; p value

**Topology:** simplicial complex; abstract simplicial complex; simplicial map; simplicial homology (with coefficients taken to be in a field instead of being integers so that one obtains vector spaces instead abelian groups)

**Category theory:** category; functor; natural transformation

## PREPARATION FOR THE LAB SESSIONS

**Install R and RStudio.** Download and install the following free software on your laptop. First R from <https://cran.rstudio.com/> and then RStudio Desktop from <https://www.rstudio.com/products/rstudio/download/>.

**Introduction to R.** Work through the following R introduction by typing (or copying and pasting) commands from <https://www.r-tutor.com/r-introduction> into the console in RStudio.

## SUGGESTED READING

To obtain a thorough background, it is recommended that you read (or skim) the beginning of one of the following two excellent recent books.

**For the mathematically inclined reader:** Chapters 1 and 2 in Topological Data Analysis for Genomics and Evolution, by Raúl Rabadán and Andrew Blumberg, Cambridge University Press, 2020.

**For the computationally inclined reader:** Chapters 1–3 in Computational Topology for Data Analysis, by Tamal Krishna Dey and Yusu Wang, Cambridge University Press, 2022. <https://www.cs.purdue.edu/homes/tamaldey/book/CTDAbook/CTDAbook.pdf>