

GENESTAT-B / NORBIS: Genetic epidemiology and genome-wide association analyses

30 May - 3 June 2022

Venue

At the University of Bergen. Rooms to be determined.

Course Home Page

https://folk.uib.no/gjessing/genetics/software/haplin/other/NORBIS_2022/intro/

Github Repository:

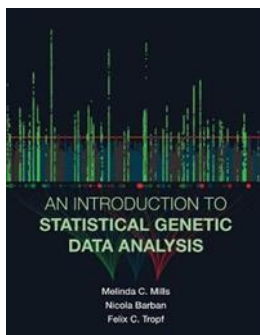
<https://github.com/jromanowska/Genestat2022>

For scripts, course data etc.

NORBIS course description

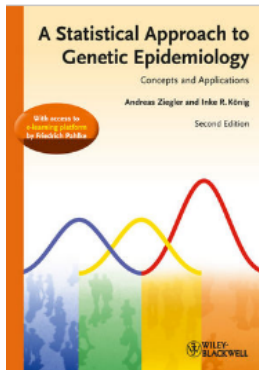
<https://norbis.w.uib.no/event/genetic-epidemiology-and-genome-wide-association-analyses-genestat-b/>

Course book



Melinda C. Mills: An Introduction to Statistical Genetic Data Analysis. *MIT press*, 2020. **Note:** The book is quite useful, but the lectures will not follow the book very closely. Available from, for instance, [Adlibris](#).

Another generally useful book is:



Andreas Ziegler, Inke R. König, Friedrich Pahlke: A Statistical Approach to Genetic Epidemiology: Concepts and Applications, 2nd Ed. Wiley-Blackwell, 2010

Lecturers

Håkon K. Gjessing (course coordinator)

Anil Jugessur

Rolv Terje Lie

Øystein Ariansen Haaland

Julia Romanowska

Jon Bohlin

Yunsung Lee

Ellisif Nygaard

Bjørn Olav Åsvold

Marc Vaudel

Gemma Sharp