PRE-CONGRESS WORKSHOP on December 5tth, 2019 San Pedro de Atacama, Chile II Congreso Chileno de Parasitología

Biostatistical Analysis of Genotype Data

Aims

After the course, participants will:

Recognize population genetics as background for genetic mapping studies Understand the principles of genetic association studies Be able to apply basic techniques to analyze genetic case-control data Be able to apply basic methods of risk calculation

Contents

The course will cover following topics: Study designs in genetic epidemiology Confounding in epidemiological and genetic association studies Power calculations and multiple testing Use of genotype data from public data repositories

The workshop is designed as a mixture of short lectures and practical exercises. Real data will be analyzed to illustrate theoretical concepts. Please bring your own laptop with you!

Instructor:

Prof. Dr. Justo Lorenzo Bermejo www.biometrie.uni-heidelberg.de/StatisticalGenetics Institute of Medical Biometry and Informatics University of Heidelberg, Germany

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Local coordinator:

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Workshop conducted within the research activities of the EU-LAC Health project "Towards Personalized Treatment of Chagas Disease by Molecular Profiling of Patients and Parasites"

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