

Oscar Rueda has a PhD in Mathematics (Statistics) from the University of Valladolid, Spain. He has previously conducted research at the Spanish National Cancer Centre (CNIO) as a graduate student working on Bayesian methods for detecting copy number aberrations in DNA and at CRUK (University of Cambridge) as a postdoc, in the Caldas Lab, working on METABRIC and in many other projects analysing breast cancer data from tumours and pre-clinical models. Since September of 2020 he leads a group at the MRC Biostatistics Unit (University of Cambridge) dedicated to the development of statistical models for the analysis of large genomic and transcriptomic datasets in order to identify biomarkers that can be used to stratify patients and to identify potential drug candidates for specific breast cancer subtypes.