



## **OMICs analysis in DataSHIELD workshop**

Given by: Juan Ramón González, ISGlobal

Open to: everyone in various projects (see target audience in the information below)

Video link: <a href="https://www.youtube.com/watch?v=X9cP2ztOurl">https://www.youtube.com/watch?v=X9cP2ztOurl</a>

Title	Omics analysis in DataSHIELD
Duration	2 hours
Target audience	Researchers interested in knowing how to analyse omic data in DataSHIELD
Brief description	This workshop will cover how to use perform state-of-the-art methods in omic data using non-disclosive approaches implemented in DataSHIELD including GWAS, gene expression and methylation data analysis
Topics covered	<ul> <li>GWAS using Bioconductor</li> <li>GWAS using PLINK</li> <li>Gene expression using limma, DESeq2 and edgeR</li> <li>Methylation using limma</li> </ul>
Pre-requisites	<ul> <li>Basic knowledge or R</li> <li>25<sup>th</sup> January 2021 DataSHIELD Beginners workshop on Youtube (link: <a href="https://www.youtube.com/watch?v=d-rqre00tlY">https://www.youtube.com/watch?v=d-rqre00tlY</a>)</li> <li>Resources in DataSHIELD course given at ATHLETE or in a previous DataSHIELD event (see <a href="https://www.datashield.ac.uk/events/">https://www.datashield.ac.uk/events/</a>)</li> </ul>
Before the workshop, you need to	<ul> <li>Download and Install R 4.0 or newer (<a href="https://www.r-project.org/">https://www.r-project.org/</a>)</li> <li>Download and Install RStudio         <a href="https://rstudio.com/products/rstudio/download/#download">https://rstudio.com/products/rstudio/download/#download</a>)</li> <li>Material and instructions:         <a href="https://github.com/isglobal-brge/dsOmics-workshop">https://github.com/isglobal-brge/dsOmics-workshop</a></li> </ul>