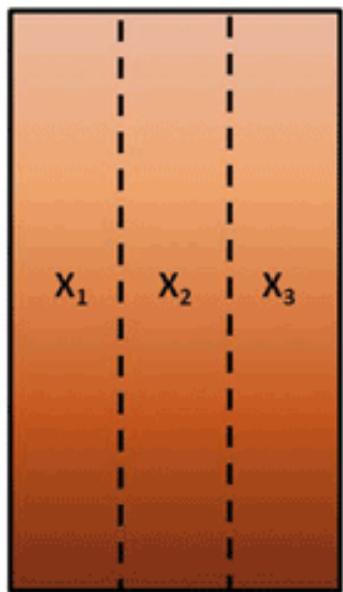


An overview of DataSHIELD

Dr Olly Butters, University of Liverpool

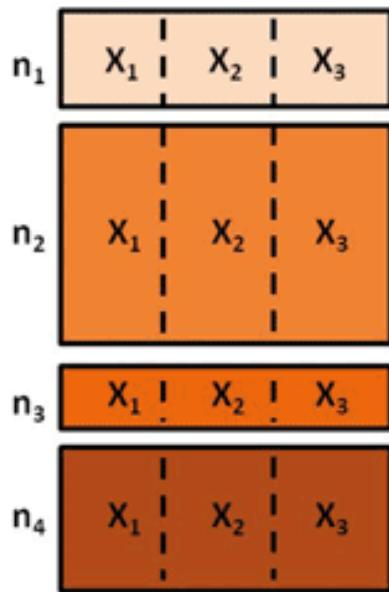
Data partitioning

← Variables →



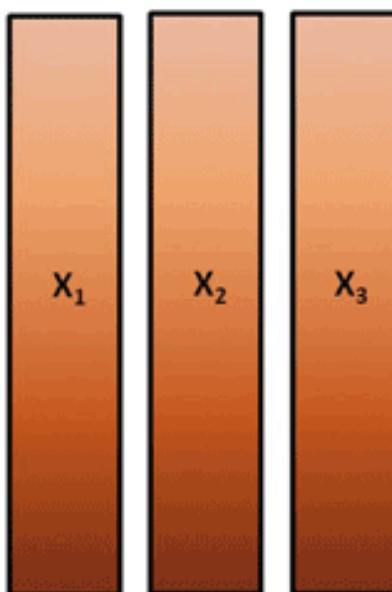
(a) Ideal:
All individual-level
data pooled
together in one file

← Variables →



(b) Horizontally partitioned:
individual-level data held in
4 separate data files, one for
each study

← Variables →



(c) Vertically partitioned :
individual-level data held in 3
datasets distinct data files, one
variable for each study

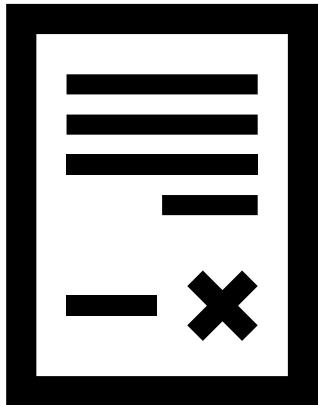
[Gaye et al, 2014: 10.1093/ije/dyu188](https://doi.org/10.1093/ije/dyu188)

Problems collaborating

Data sharing-access barriers result from a range of scenarios



Ethico-legal & governance



Control IP



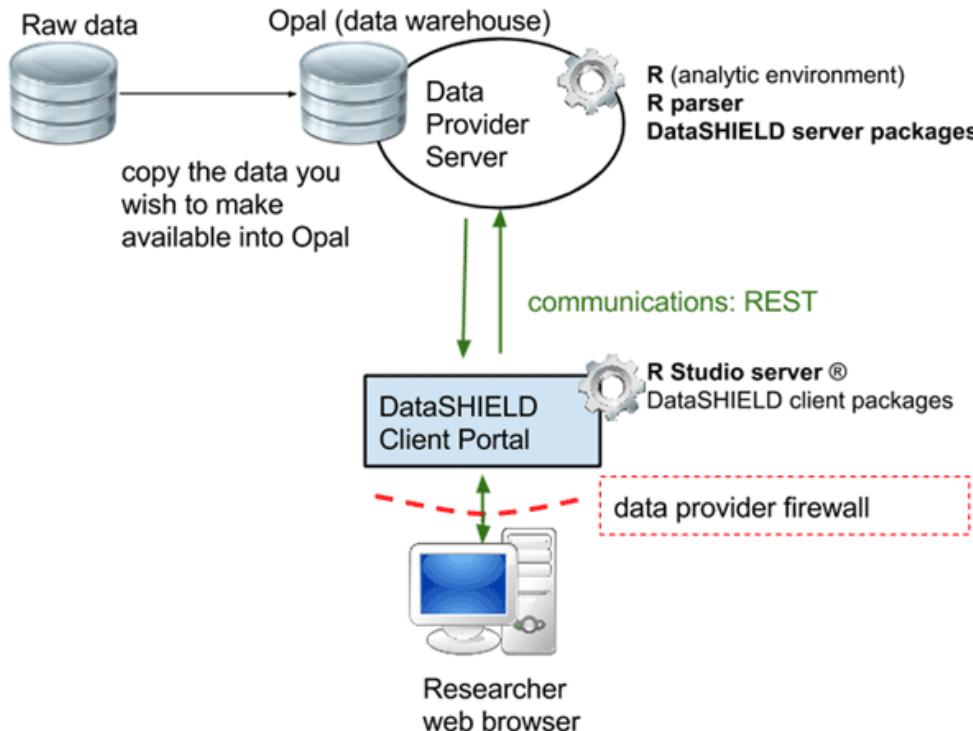
Physical size of data

- Methods to prevent disclosure of individual level data
- Typically in medical and health research:
 - Anonymisation
 - Pseudonymisation
 - Use of a data safe haven
 - Log onto a portal can view all the individual level data
 - Reliant on data governance measures
 - Sign a contract saying you won't misuse the data
 - Are any penalties enforced?
 - For very sensitive information human scrutiny of outputs
 - Financial and time cost

DataSHIELD: a solution

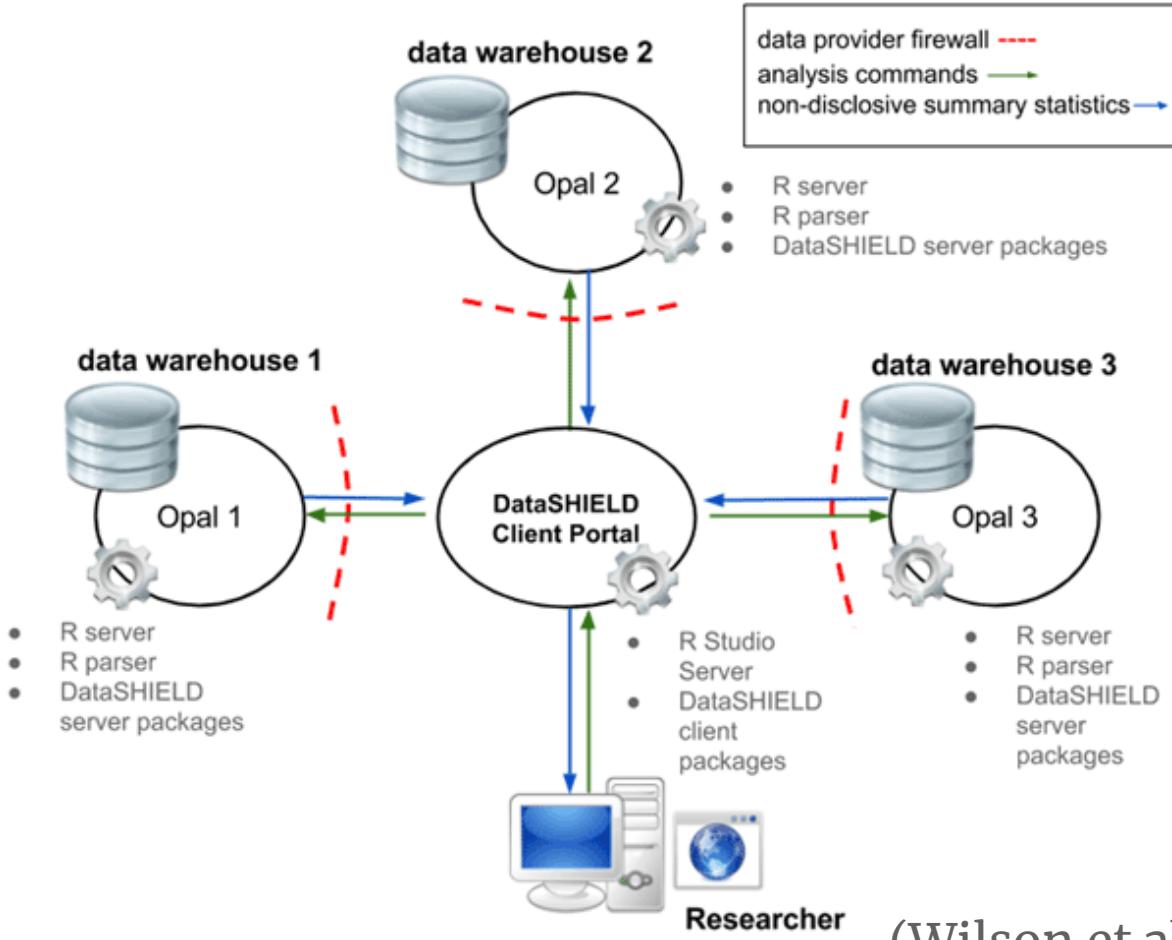
- Technological and methodological solution
 - Designed by epidemiologists and biostatisticians, community led
 - Individual level and raw data do not leave the data controller site
 - Simplifies and streamlines data governance process and decisions
 - Automated disclosure control built into the system
 - Addresses confidentiality restrictions
 - Only non-disclosive summary statistics returned to analyst
 - No delay waiting for human scrutiny of results
 - GDPR compliance
 - Analyse data from multiple studies (meta-analysis) or a single one
 - Researcher does their own analysis in real time
 - Open source – free at point of use (cost effective)
-

Single-site DataSHIELD



(Wilson et al., 2017: [10.5334/dsj-2017-021](https://doi.org/10.5334/dsj-2017-021))

Multi-site DataSHIELD



(Wilson et al., 2017: [10.5334/dsj-2017-021](https://doi.org/10.5334/dsj-2017-021))

DataSHIELD functions

Coercing functions

- `ds.asCharacter`
- `ds.asInteger`
- `ds.asMatrix`
- `ds.asList`
- `ds.asList`
- `ds.asNumeric`
- `ds.asFactor`
- `ds.asLogical`

Number manipulation functions

- `ds.make`
- `ds.Boole`
- `ds.list`
- `ds.abs`
- `ds.log`
- `ds.exp`
- `ds.sqrt`
- `ds.rep`
- `ds.seq`
- `ds.c`

Administrative functions

- `ds.listClientsideFunctions`
- `ds.listServersideFunctions`
- `ds.message`
- `ds.setSeed`
- `ds.listDisclosureSettings`
- `ds.ls`
- `ds.rm`
- `ds.testObjExists`
- `ds.look`
- `ds.listOpals`
- `ds.setDefaultOpals`

Data Frame/List manipulation functions

- `ds.completeCases`
- `ds.replaceNA`
- `ds.sample`
- `ds.merge`
- `ds.recodeValues`
- `ds.getWGSR`
- `ds.dataFrame`
- `ds.dataFrameSubset`
- `ds.dataFrameFill`
- `ds.dataFrameSort`
- `ds.tapply`
- `ds.tapply.assign`
- `ds.vectorCalc`
- `ds.list`
- `ds.unList`
- `ds.assign`
- `ds.cbind`
- `ds.rbind`

Factor manipulation functions

- `ds.changeRefGroup`
- `ds.recodeLevels`

Matrices Functions

- `ds.matrix`
- `ds.matrixDiag`
- `ds.matrixMult`
- `ds.matrixDet`
- `ds.matrixDimnames`
- `ds.matrixTranspose`
- `ds.matrixDet.report`
- `ds.matrixInvert`

Data structure queries

- `ds.levels`
- `ds.length`
- `ds.exists`
- `ds.colnames`
- `ds.names`
- `ds.dim`
- `ds.isValid`
- `ds.isNA`
- `ds.numNA`
- `ds.class`

Summary Statistics Functions

- `ds.mean`
- `ds.meanByClass`
- `ds.meanSdGp`
- `ds.quantileMean`
- `ds.rowColCalc`
- `ds.var`
- `ds.cov`
- `ds.cor`
- `ds.corTest`
- `ds.summary`
- `ds.skewness`
- `ds.kurtosis`

Table functions

- `ds.table`

Survival Analysis functions

- `ds.lexis`
- `ds.reshape`

Distribution Generating functions

- `ds.rNorm`
- `ds.rBinom`
- `ds.rPois`
- `ds.rUnif`

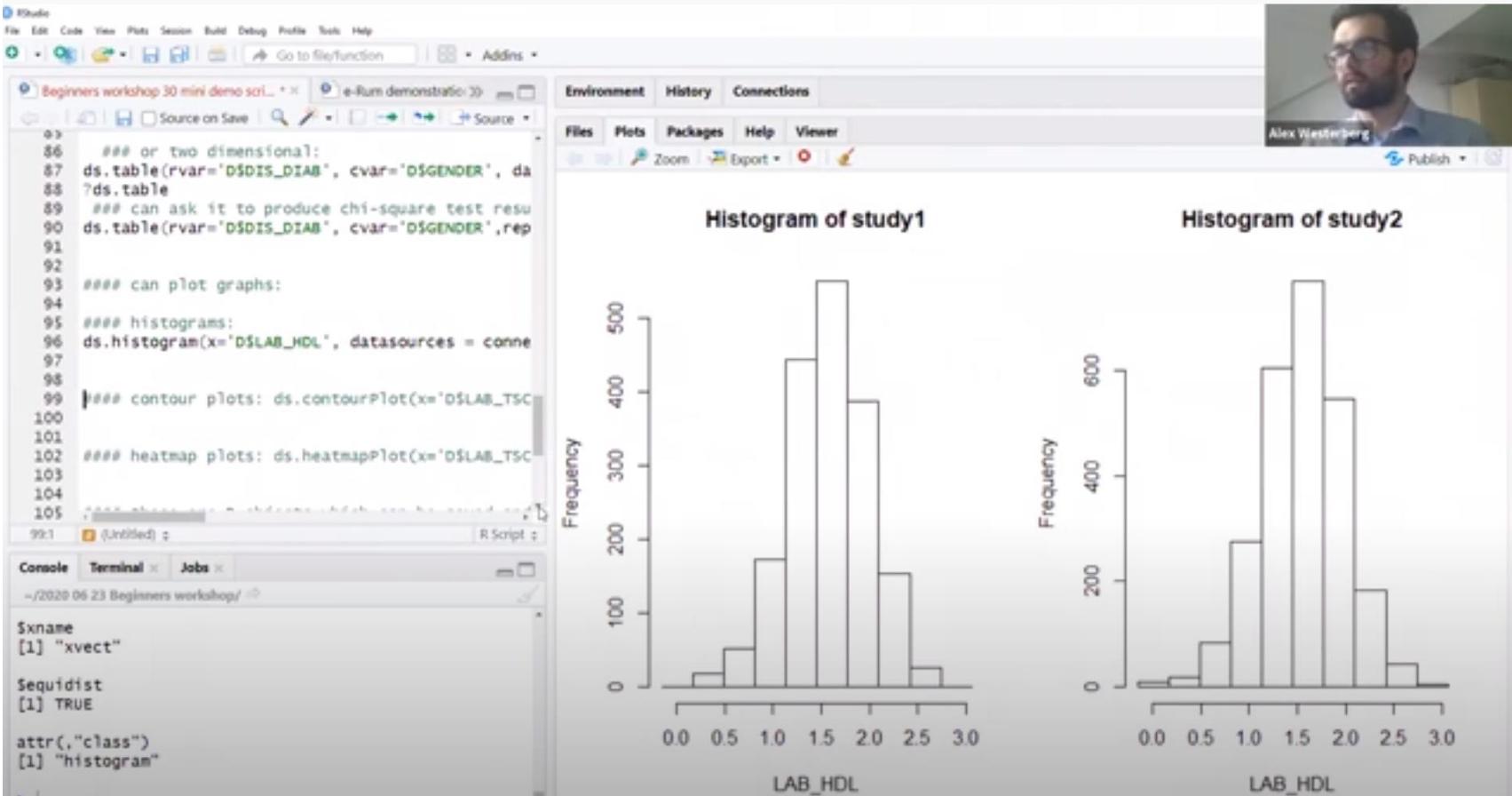
Modelling Functions

- `ds.glm`
- `ds.glmSummary`
- `ds.glmSLMA`
- `ds.glmPredict`
- `ds.lmerSLMA`
- `ds.glmerSLMA`

Plotting functions

- `ds.histogram`
- `ds.boxPlot`
- `ds.densityGrid`
- `ds.heatmapPlot`
- `ds.contourPlot`
- `ds.scatterPlot`

What's it like to use?

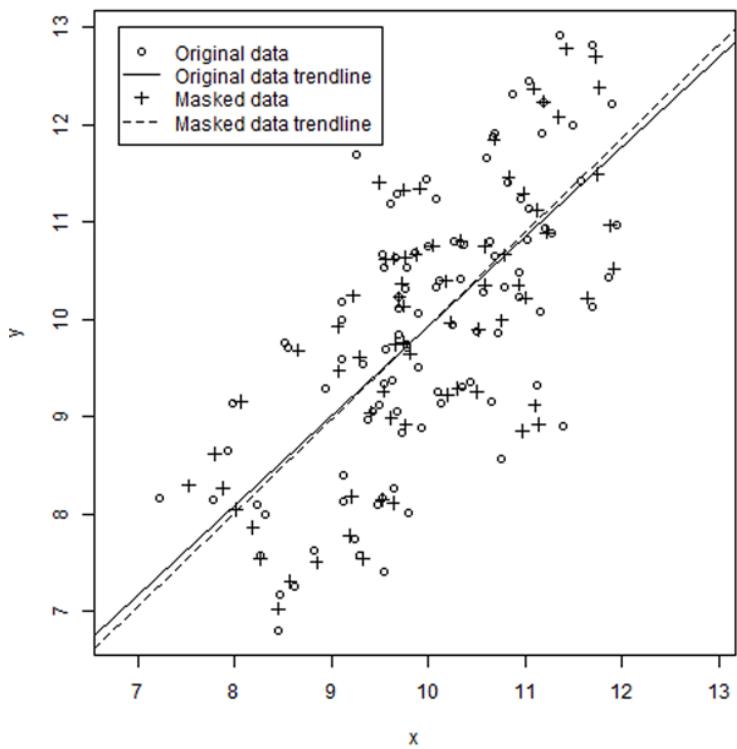


<https://youtu.be/f0NJNu--Oik>

Statistical Disclosure Control

- Systematically built into each function – technique depends on function
- No “print to screen” of microdata
- Cell suppression
 - Minimum cell count – set by each study (typically 5 or 3)
 - Prevent disclosure of individual level data in tables, histograms etc
- Subset suppression
- GLM number of parameters limit
- String length limit
- K-nearest neighbour K value
- Number of categories in categorical data
- Amount of noise to add
- Often just adapt a standard R function
 - Disclosive information blocked from function
 - ds.glm vs glm - residuals are blocked

Data visualization in DataSHIELD



- Scatter plots inherently disclosive
- Application of different methods in DataSHIELD
 - *k*-anonymization: preserves privacy by reducing the granularity of the data (suppression, generalization)
 - a deterministic approach: replaces individual-level observations with centroids of k nearest neighbours
 - a probabilistic procedure: perturbs individual attributes with addition of random stochastic noise
- Retain original data structure and features, no disclosure
- Now want to apply this to geospatial data – make use of full postcode/geolocation information, not aggregated areas

(Avraam et al., 2021:
[10.1140/epjds/s13688-020-00257-4](https://doi.org/10.1140/epjds/s13688-020-00257-4))

Other developments

- Other source data providers (Other DBs, flat files, connections etc)
- Genomics analysis (integration with BioConductor)
- Machine learning techniques
- Geospatial in the pipeline
- Exploring a spin out company to offer support for DataSHIELD
- About to write a grant application around disclosure control (interested? Let me know!)
- About to write a grant application about using DataSHIELD with TREs (interested? Let me know!)

- www.datashield.ac.uk
- DataSHIELD Conference (Online) 10-11 November 2021