### **DataSHIELD workshop**

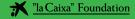
### Block 1 - Introduction to DataSHIELD

### Juan R Gonzalez Bionformatics Research Group in Epidemiology e-mail: juanr.gonzalez@isglobal.org





#### A partnership of:



















### **Biomedical Research Park of Barcelona**

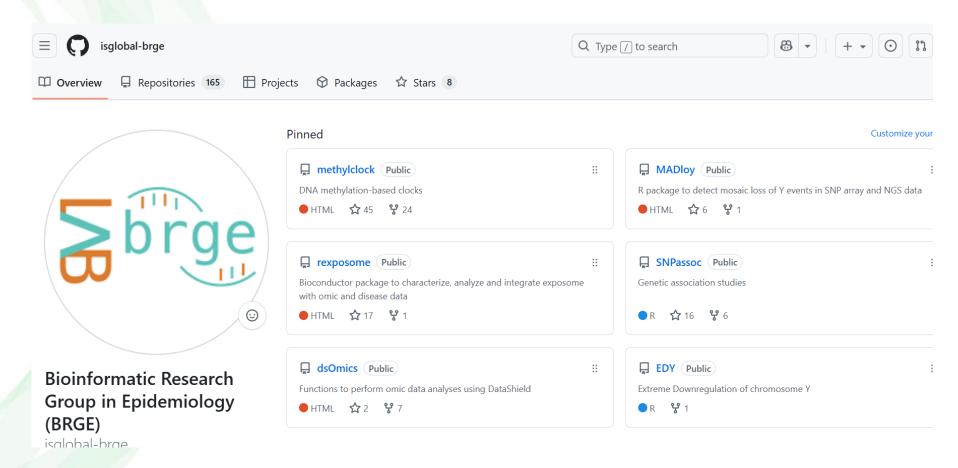


- Barcelona Institute for Global Health (ISGlobal)
- University Pompeu Fabra (UPF)
- □ Center for Genomic Regulation
- European Molecular Laboratory (EMBL)
- ☐ Institute of Medicine (Hospital)
- Evolutive Biology Laboratory

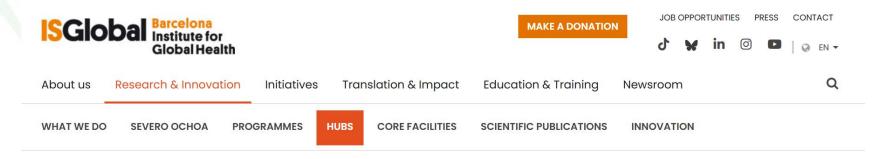


### **Bioinformatics Reserarch Group in Epidemiology (BRGE)**

### https://github.com/isglobal-brge



# Health Analytics Hub @ISGlobal



Research & Innovation > Hubs > Health Analytics Hub

RESEARCH

### Health Analytics Hub



### What is DataSHIELD

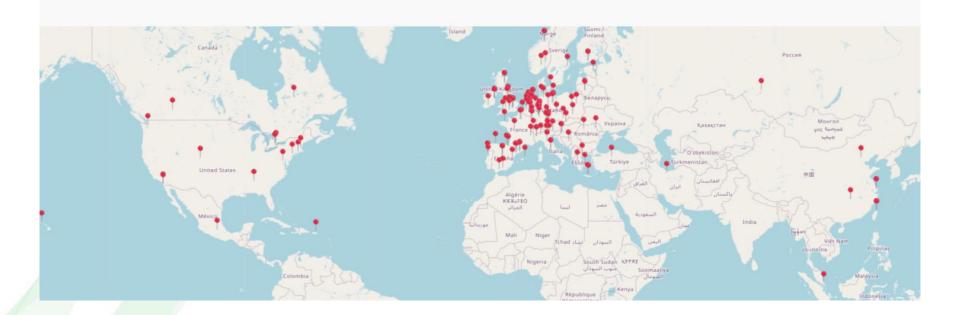
- ☐ A privacy-preserving, non-disclosive, federated analysis software
- What it is
  - way of analyzing sensitive data, at a individual level, without providing direct access
- What it is not
  - data harmonization platform
  - ☐ GUI based analysis tool (yet possible!)

# **What is DataSHIELD**

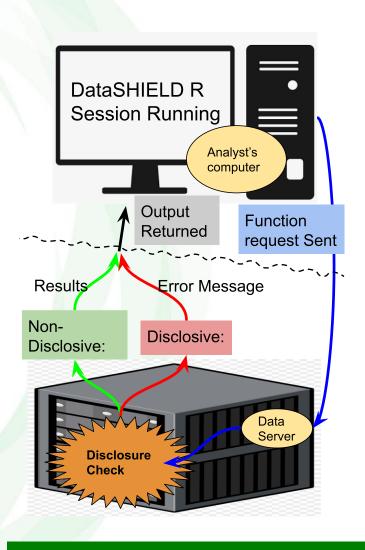
https://datashield.org/



About - For analysts For data custodians For developers News Suppor

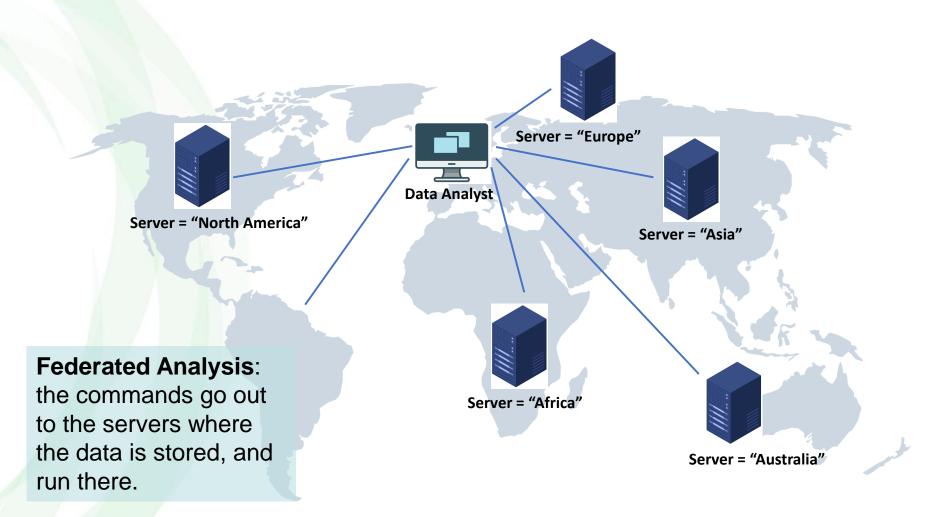


# **Privacy-preserving**

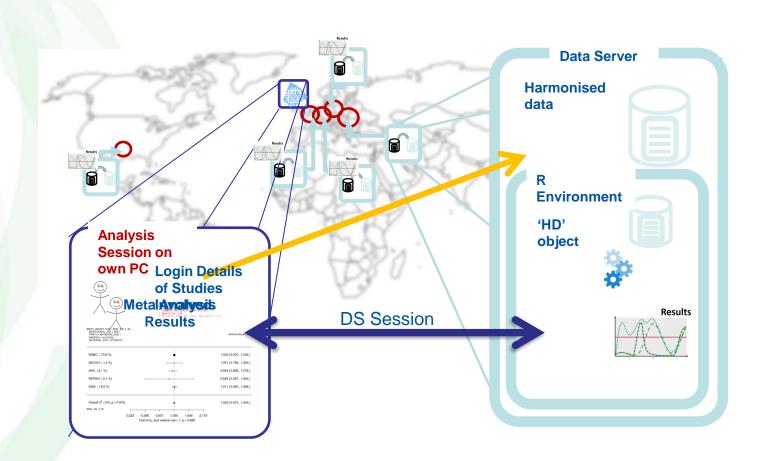


- ☐ Functions are sent to data server
- Server runs function
- Built into function is disclosure check
- ☐ If disclosive, check will discover and only return error message, not results.

# Federated analysis



# **Animation of a DataSHIELD Analysis**



# What kind of analysis can we do?

- ☐ There are 120 functions within the DataSHIELD clientside package to choose from, in version 6.3 (released November 2024)
- □A full list can be found on this wiki:
- https://data2knowledge.atlassian.net/wiki/spaces/DSDEV/pages/1184825438/List+of+all+DataSHIELD+functions+v6.3
- ☐ There are external developers creating new packages (ggplot, omics, mediation, tydiverse, ...)

# **Methodological papers**

#### JOURNAL ARTICLE

#### DataSHIELD: mitigating disclosure risk in a multisite federated analysis platform 3

Demetris Avraam 

, Rebecca C Wilson , Noemi Aguirre Chan , Soumya Banerjee ,
Tom R P Bishop , Olly Butters , Tim Cadman , Luise Cederkvist , Liesbeth Duijts ,
Xavier Escribà Montagut ... Show more

#### PLOS COMPUTATIONAL BIOLOGY

#### RESEARCH ARTICLE

Orchestrating privacy-protected big data analyses of data from different resources with R and DataSHIELD

Yannick Marcono<sup>1\*</sup>, Tom Bishopo<sup>2</sup>, Demetris Avraamo<sup>3</sup>, Xavier Escriba-Montagut<sup>4,5</sup>, Patricia Ryser-Welch<sup>3</sup>, Stuart Wheater<sup>6</sup>, Paul Burtono<sup>3</sup>, Juan R. Gonzálezo<sup>4,5,7,8</sup>\*

1 Epigeny, St Ouer, France, 2 MRC Epidemiclogy Unit, University of Cambridge, Cambridge, United Kingdom, 3 Population Health Science Institute, Newcrates University, Newcreatie, United Kingdom, 4 Barcelona Institute for Global Health (ISGlobal), Barcelona, Spain, 5 Universitat Promper Fabra (IJPF), Barcelona, Spain, 6 Apirar Technologies, Newcastel, United Kingdom, 7 Certhrol of Investigation Blomédica en Red on Epidemiclogia y Salud Pública (CIBERESP), Barcelona, Spain, 6 Dept. of Mathematics, Universitat Authorna de Barcelona (USB), Betalerra (Barcelona), Spain

\* yannick.marcon@obiba.org (YM); juanr.gonzalez@isglobal.org (JRG)

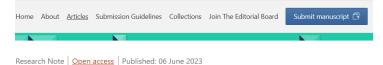
#### PLOS COMPUTATIONAL BIOLOGY

#### RESEARCH ARTICLE

Federated privacy-protected meta- and megaomics data analysis in multi-center studies with a fully open-source analytic platform

Xavier Escriba-Montagut<sup>1,2</sup>, Yannick Marcon<sup>3</sup>, Augusto Anguita-Ruiz<sub>©</sub><sup>1,2</sup>, Demetris Avraam<sup>4</sup>, Jose Urquiza<sup>1,2,5</sup>, Andrei S. Morgan<sub>©</sub><sup>6,7</sup>, Rebecca C. Wilson<sub>©</sub><sup>4</sup>, Paul Burton<sup>3</sup>, Juan R. Gonzalez<sub>©</sub><sup>1,2,4</sup>

#### **BMC Research Notes**



dsSurvival 2.0: privacy enhancing survival curves for survival models in the federated DataSHIELD analysis system

Soumya Banerjee <sup>™</sup> & Tom R. P. Bishop



#### JOURNAL ARTICLE

Software Application Profile: ShinyDataSHIELD—an R Shiny application to perform federated non-disclosive data analysis in multicohort studies 3

Xavier Escribà-Montagut, Yannick Marcon, Demetris Avraam, Soumya Banerjee,
Tom R P Bishop, Paul Burton, Juan R González ▼

### Use of DataSHIELD in science



Environment International



. . .

Green spaces and respiratory, cardiometabolic, and neurodevelopmental outcomes: An individual-participant data meta-analysis of >35.000 European children

Amanda Fernandes  $^{\circ}$  b  $^{\circ}$   $^{\circ}$   $^{\circ}$  Demetris Avraom  $^{\circ}$   $^{\circ}$  Im Cadman  $^{\circ}$  b  $^{\circ}$   $^{\circ}$  Payam Dadvand  $^{\circ}$  b  $^{\circ}$  Monica Guxens  $^{\circ}$  b  $^{\circ}$   $^{\circ}$  Anne-Cloire Binter  $^{\circ}$  b  $^{\circ}$  Angelo Pinot de Moiro  $^{\circ}$   $^{\circ}$  Moniteriore Mori Nieuwenhuijsen  $^{\circ}$  b  $^{\circ}$  Liebeth Duijts  $^{\circ}$   $^{\circ}$  1 John Julyez  $^{\circ}$  b  $^{\circ}$   $^{\circ}$  Moniterrat De Castro  $^{\circ}$  b  $^{\circ}$  Serena Fossati  $^{\circ}$  b  $^{\circ}$  Sandra Márquez  $^{\circ}$  b  $^{\circ}$  S, Tanja Vrijkotte  $^{\circ}$   $^{\circ}$  Ahmed Elhakeem  $^{\circ}$   $^{\circ}$  Rosemary McEochon  $^{\circ}$   $^{\circ}$  Iffany Yang  $^{\circ}$   $^{\circ}$  Marie Pedersen  $^{\circ}$  Johan Vinther  $^{\circ}$  , Johan Lapeule  $^{\circ}$   $^{\circ}$  Lapeule  $^{\circ}$   $^{\circ}$  Mortine Vrijkotte  $^{\circ}$  b  $^{\circ}$   $^{\circ}$ 

#### **PLOS MEDICINE**

#### nature communications

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### Identification of biomarkers for glycaemic deterioration in type 2 diabetes

Roderick C. Slieker, Louise A. Donnelly, Elina Akalestou, Livia Lopez-Noriega, Rana Melhem, Ayşim Güneş, Frederic Abou Azar, Alexander, Efanox, Eleni Georgiadou, Hermine Muniangi-Muhitu, Mahsa Sheikh, Giuseppe, N. Giordano, Mikael Akerlund, Emma Ahlqvist, Ashfaq Ali, Karina Banasik, Søren Brunak, Marko Bartovis, Gerard A. Bouland, Frédéric Burdet, Miskael Canouil, Julian Dragan, Petra J. M. Elders, Celine Fernandez, ... Guy A. Rutter



RESEARCH ARTICLE

Gestational age at birth and body size from infancy through adolescence: An individual participant data meta-analysis on 253,810 singletons in 16 birth cohort studies

Johan L. Vinthere]\*, Tim Cadmang<sup>2</sup>, Demetria Avraama<sup>3</sup>, Claus T. Ekstrema<sup>4</sup>, Thorkild I. A. Sørensen<sub>3</sub><sup>1,5</sup>, Ahmed Elhakeem<sup>2</sup>, Ana C. Santos<sub>6</sub><sup>5,7</sup>, Angela Pinot de Moira<sub>0</sub>, Barbara Heude<sub>9</sub>\*, Carmen Iñiguez<sub>2</sub><sup>6,10,11</sup>; Costanza Pizzi<sup>2</sup>, Elinor Simons<sub>6</sub><sup>5,7</sup>, H. Ellis Voerman <sup>5,16</sup>, Eva Corpelejin<sub>9</sub><sup>5,7</sup>, Faryal Zariouh<sup>18</sup>, Gilian Santorelli<sup>19</sup>, Hazel M. Inskip<sub>6</sub><sup>5,5,7</sup>, Henrique Barros<sup>6,7</sup>, Jennic Carson<sub>9</sub><sup>2,2,23</sup>, Jennifer R. Harris<sup>24</sup>, Johanna L. Nadere<sup>25</sup>, Justiina Ronkainen<sub>9</sub><sup>5,8</sup>, Katrine Strandberg-Larsen<sub>9</sub><sup>1</sup>, Loreto Santa-Marina<sub>9</sub><sup>5,7,26</sup>, Lucinda Calas<sub>9</sub><sup>6</sup>, Luise Cederkviste<sup>3</sup>, Maja Popovic<sub>9</sub><sup>1,2</sup>, Marie-Aline Charles<sub>9</sub><sup>1,6</sup>, Marie-Reiberg-Larsen<sub>9</sub><sup>1,6,7,6,7</sup>, Paul Burton<sub>9</sub><sup>8</sup>, Puishkumar J. Mandhane<sub>9</sub><sup>3,6</sup>, Rae-Chi Huang<sub>9</sub><sup>2,2,7</sup>, Rebecca C. Wilson<sub>9</sub><sup>3,6</sup>, Sido Haakma<sub>9</sub><sup>3,7</sup>, Silvia Fernández-Barrès<sub>9</sub><sup>2,5,9,6,9</sup>, Stuart Turvey<sup>4</sup>, Susana Santos<sub>9</sub><sup>1,6,7</sup>, Suzane C. Toughe<sup>4</sup>, Sylvian Sebertre<sup>6,7</sup>, The OJ. Moraes<sub>9</sub><sup>3,7</sup>, Theodosia Salika<sub>9</sub><sup>2,7</sup>, Vincent W. V. Jaddoe<sub>9</sub><sup>1,5,16</sup>, Deborah A. Lawlor<sup>2,4,3</sup>, Anne-Marie Nybo Aladersen<sub>9</sub><sup>1</sup>

#### Practice of Epidemiology

Associations of Maternal Educational Level, Proximity to Green Space During Pregnancy, and Gestational Diabetes With Body Mass Index From Infancy to Early Adulthood: A Proof-of-Concept Federated Analysis in 18 Birth Cohorts

Tim Cadman\*, Ahmed Elhakeem, Johan Lerbech Vinther, Demetris Avraam, Paula Carrasco, Lucinda Calas, Marloes Cardol, Marie-Aline Charles, Eva Corpeleijn, Sarah Crozier, Montserrat de Castro, Marisa Estarlich, Amanda Fernandes, Serena Fossatti, Dariusz Gruszfeld, Kathrin Guerlich, Veit Grote, Sido Haakma, Jennifer R. Harris, Barbara Heude, Rae-Chi Huang, Jesús Ibarluzea, Hazel Inskip, Vincent Jaddoe, Berthold Koletzko, Sandrine Lioret, Veronica Luque, Yannis Manios, Giovenale Moirano, George Moschonis, Johanna Nader, Mark Nieuwenhuijsen, Anne-Marie Nybo Andersen, Rosie McEachen, Angela Pinot de Moira, Maja Popovic, Theano Roumeliotaki, Theodosia Salika, Loreto Santa Marina, Susana Santos, Sylvain Serbert, Evangelia Tzorovili, Marina Vafeiadi, Elvira Verduci, Martine Vrijheid, T. G. M. Vrijkotte, Marieke Welten, John Wright, Tiffany C. Yang, Daniela Zugna, and Deborah Lawlor



The Lancet Regional Health - Europe



Articles

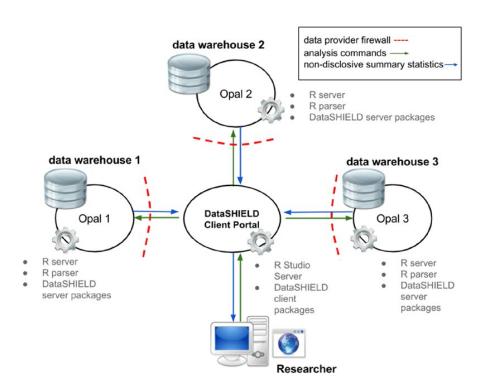
Early childcare arrangements and children's internalizing and externalizing symptoms: an individual participant data meta-analysis of six prospective birth cohorts in Europe

Katharine M. Barry <sup>a b</sup>, Demetris Avraam <sup>c</sup>, Tim Cadman <sup>c</sup>, Ahmed Elhakeem <sup>d</sup>,
Hanan El Marroun <sup>e f</sup>, Pauline W. Jansen <sup>e f</sup>, Anne-Marie Nybo-Andersen <sup>c</sup>,
Katrine Strandberg-Larsen <sup>c</sup>, Llúcia González Safont <sup>g f h</sup>, Raquel Soler-Blasso <sup>j j s</sup>,
Florencia Barreto-Zarza <sup>b k t</sup>, Jordi Julvez <sup>[u j</sup>, Martine Vrijheid <sup>[u j</sup>, Barbara Heude <sup>m n</sup>,
Marie-Aline Charles <sup>g p</sup>, Alexandre Ramchandar Gomajee <sup>a b q</sup>, Maria Melchior <sup>b</sup>  $\mathcal{R}$ 

# DataSHIELD standard platform

Preliminary: data located at Opal server

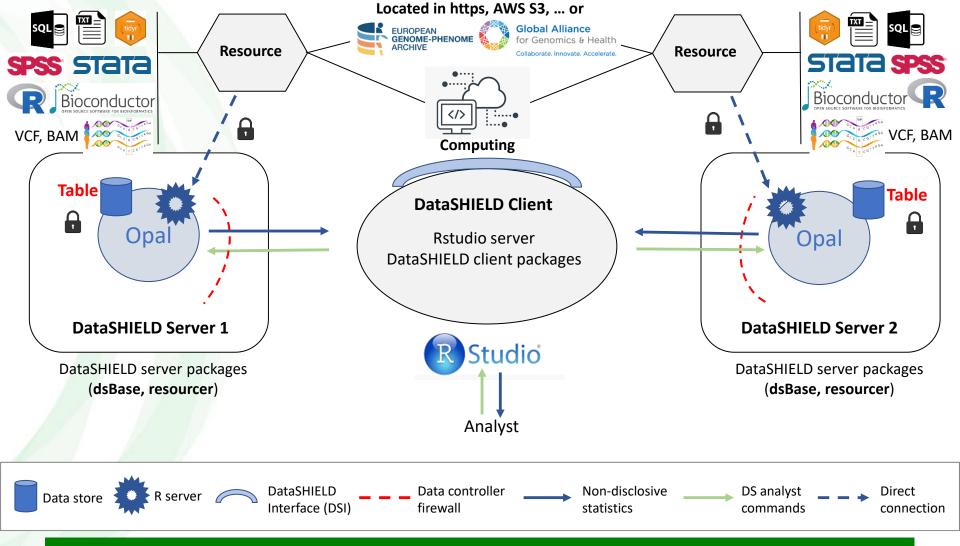
- Authenticate and authorize user
- ☐ Assign Opal table into the R server (data transfer)
- Execute DataSHIELDverified R commands in R server



# **DataSHIELD standard platform**

# Open Opal demo server

### The resources: idea



### The resources: rational

- use data at their original location (do not move/copy data when possible)
- use data in their original format (no preprocessing, no loss of information)
- use external computation facilities (no R limitations)

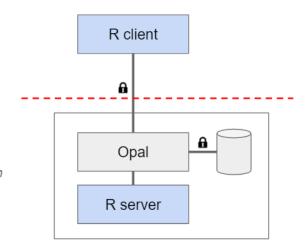
□=> use DataSHIELD with large/big datasets (omics etc.)

### The resources ...

### **Opal with tables**

DataSHIELD client

DataSHIELD server 1..n

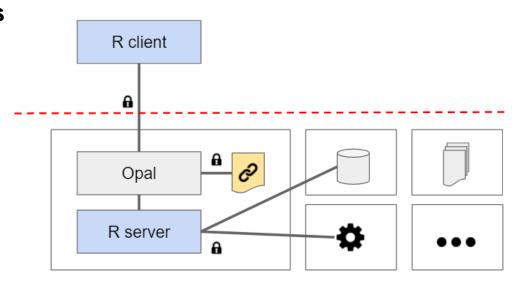


### **Opal with resources**

DataSHIELD client

DataSHIELD server 1..n

- Databases
- File repositories
- Big data analytics systems
- etc.



### Resources examples

CSV file (compressed or not) ■ SQL database table ☐ R object stored in a R data file ☐ HL7 FHIR dataset ☐ GA4GH server ■ AWS server (OMOP database) ☐ HPC server accessible through SSH Python script ☐ Big data analytics system (Apache Spark, Dremio, ...) Apps in docker images

# The resources ...

Property	Description	Examples
url	Location of the resource	https://example.org/some/file.rda file://path/to/file.csv ssh://example.org/work/dir?exec=plink mysql://dbhost:3306/mydb/mytable
format	Data format (optional)	SPSS ExpressionSet
credentials	Data access (optional)	token=Q3sDdsWq2dsx7 username=user1 password=xxxxxx

Not visible by DataSHIELD users

### **DataSHIELD disclosure controls**

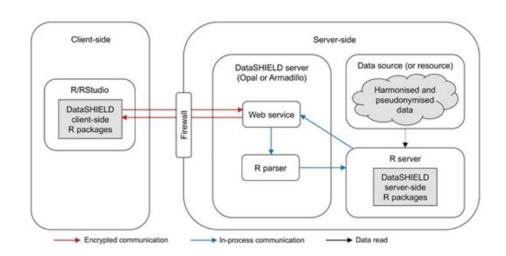
- Ensures no individuallevel data leaves the data holder's server.
- ☐ Analyses are performed remotely; only non-disclosive summaries are returned.
- ☐ Protects participant privacy while enabling collaborative research.
- ☐ Balances data utility with confidentiality.

#### JOURNAL ARTICLE

DataSHIELD: mitigating disclosure risk in a multi-site federated analysis platform 3

Demetris Avraam ★ , Rebecca C Wilson , Noemi Aguirre Chan , Soumya Banerjee , Tom R P Bishop , Olly Butters , Tim Cadman , Luise Cederkvist , Liesbeth Duijts , Xavier Escribà Montagut ... Show more

*Bioinformatics Advances*, Volume 5, Issue 1, 2025, vbaf046, https://doi.org/10.1093/bioadv/vbaf046



# DataSHIELD: mitigation disclosure risks

- System protection elements
- Analysis protection elements
- Governance protection elements
- ☐ Five Safe Framework

# **DataSHIELD:** system protection elements

■ Network security: ☐ Traffic encrypted. Connection requires SSL/TLS certificate. ☐ Firewall protection ☐ User authentication and certification: R server is only callable via middleware hosting DataSHIELD. R parser: Analyses are only able to use R server-side functions not native R Data management: Data comprise a snapshot not a live data (resources)

# DataSHIELD: analysis protection elements

Only use assign and aggregate functions: Only create objects in the server-side. Generate summarized data to the client-side. Implementation restriction: □ R functions are not implemented (*print*, *max*, ...) Disclose controls: ☐ Function to provide non-disclosive results (each package can have their own functions) Data level obfuscation: Anonymization, data synthesis techniques, ... ☐ DataSHIELD log files: log files visible for data custodians

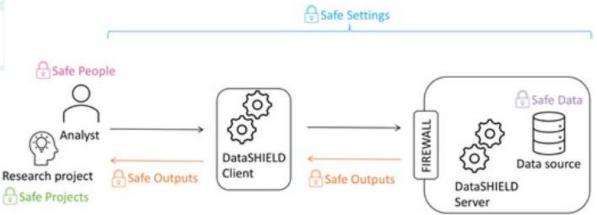
# DataSHIELD: examples of disclose controls

Name	Description
nfilter.tab	Prevents the return of a contingency table if any of its cells represents less than <i>nfilter.tab</i> observations. The value of <i>nfilter.tab</i> can be set to any non-negative integer. The default value is set to 3.
nfilter.subset	Prevents the creation of a dataset's subset if the subset has less than <i>nfilter.subset</i> rows. The value of <i>nfilter.subset</i> can be set to any positive integer. The default value is set to 3.
nfilter.glm	Prevents the fitting of a regression model that has more than <i>nfilter.glm x N</i> unknown parameters in a dataset with sample size <i>N</i> . The value of <i>nfilter.glm</i> can be set to any numeric value in the interval (0,1). The default value is set to 0.33.
nfilter.string, nfilter.stringShort	Blocks the evaluation of a string argument that passes from the client-side to the server-side, if it has a length greater than nfilter.string or nfilter.stringShort characters. The values of nfilter.string and nfilter.stringShort can be set to any positive integers. The default values are set to 80 and 20, respectively.
nfilter.levels.density	Prevents the return of the unique levels of a categorical variable if their length is more than <i>nfilter.levels.density x N</i> where <i>N</i> is the length of the vector of the categorical variable. The value of <i>nfilter.levels.density</i> can be set to any numeric value in the interval (0,1). The default value is set to 0.33.

# DataSHIELD: governance protection elements

■ Formal agreements: Data access, data analysis and approval of research results prior to publication. ☐ User can have access to: ■ Different tables or resources ■ Different variables from a table of resource Different data analysis packages ☐ It is controlled through the 'profiles'





Five Safes	Study mitigation	DataSHIELD mitigation
Safe People	<ul> <li>Formal data access request process</li> <li>Due diligence on prospective users         —'are they bona fide researchers?         Have they conducted mandatory         training/accreditation to work with data         safely?'</li> <li>Legal contracts, signed terms and         conditions of data access and use</li> <li>Sanctions policy</li> <li>DataSHIELD users are authorized for         data access by the study</li> </ul>	The authorization to access     DataSHIELD can be     delegated, under the principle     of subsidiarity to individual     studies

Five Study mitigation
Safes

Safe
projects

- Assess the requirement for access to the data—the context of the research project or the data being used

 Ensuring the data access/use does not contradict any necessary legal requirements, e.g. study consent

Five Safes	Study mitigation	DataSHIELD mitigation
Safe settings	<ul> <li>The operation and maintenance of robust computing infrastructure and hardware</li> <li>Following IT security best practice</li> </ul>	<ul> <li>DataSHIELD is a client-server architecture, the user does not connect directly to the study data</li> </ul>
	Log of registered users,     maintaining/blocking user access	<ul> <li>Analysis of individual level data occurs server-side (at study)</li> </ul>
	Preventative measures for unauthorized access	<ul> <li>Analysis environment server- side can only be called via authenticated users through</li> </ul>
	Deploy DataSHIELD to securely	Opal or Armadillo
	transfer information (analysis commands and outputs) via https	The server-side R Parser prevents invalid characters or
	Each study/consortium provides     unique authentication credentials for	non-approved functions from being run.
	users to log onto the DataSHIELD client and to connect to each study they are authorized for	Users can not directly view the individual-level data
		<ul> <li>User commands logged server-side, only accessible by the study. Can be manually scrutinized e.g. if data misused</li> </ul>

Five Safes	Study mitigation	DataSHIELD mitigation
Safe data	Data Protection Impact Assessment (risk assessment)	
	<ul> <li>Assessing the disclosure risk of the data</li> </ul>	
	Pseudonymized data used to lower the disclosure risk	

Five Safes	Study mitigation	DataSHIELD mitigation
Safe outputs	<ul> <li>Manual checking of analysis outputs for disclosure</li> <li>Legal contracts or terms and conditions making it mandatory for the user to check their own outputs for disclosure</li> </ul>	<ul> <li>DataSHIELD server-side functions prevent directly disclosive outputs being returned to the analyst</li> <li>DataSHIELD server-side</li> </ul>
	DataSHIELD disclosure setting	functions prevent viewing of individual-level data
	thresholds are set and maintained by the study, no one else can alter these	DataSHIELD has disclosure     settings based on established     statistical disclosure control     methods to conduct automated     checks for direct disclosure in     outputs