



Implementing disclosure controls in DataSHIELD demonstrated by the dsSurvival package

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Survival analysis

Rationale

- Survival analysis is widely used in medical sciences to analyze the expected duration of time until some event of interest occurs
- The most frequently used model is the cox proportional hazard model (Cox, 1972)
- Performing meta-analysis of survival models requires large amount of data from different sites
 - General Data Protection Regulation
 - Physical size of data

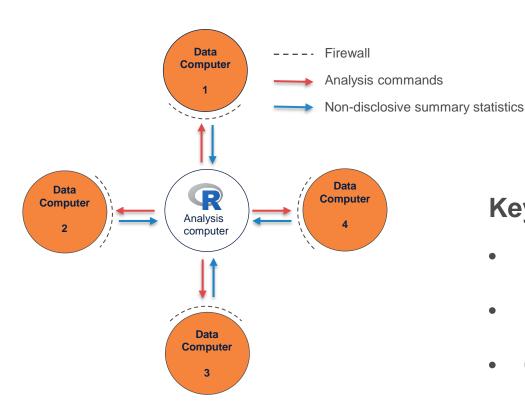
Alternative: DataSHIELD



The DataSHIELD approach

Take "analysis to data" not "data to analysis"

<u>Data Aggregation Through Anonymous Summary-statistics from Harmonized Individual-levEL Databases</u>



Key principles

- Enables federated analysis
- Uses client server architecture
- Controls disclosure risks

Gaye, Amadou, et al. "DataSHIELD: taking the analysis to the data, not the data to the analysis." International journal of epidemiology 43.6 (2014): 1929-1944.

https://www.datashield.org/

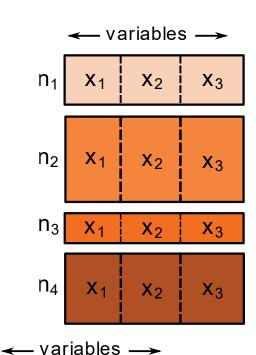


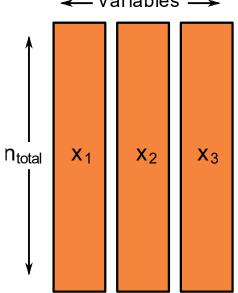
The DataSHIELD approach

Two classes of multi-score analysis

- Horizontal partitioning
 - meta-analysis setting

- Vertical partitioning
 - record linkage setting







dsSurvival

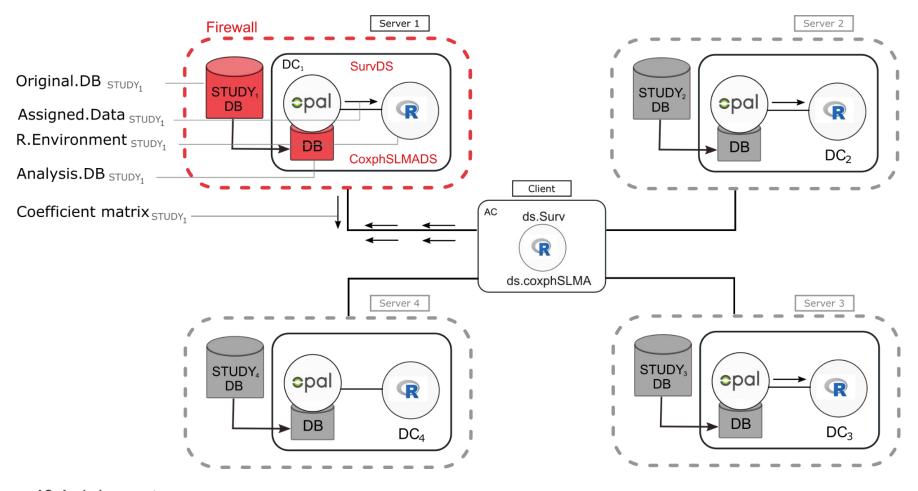
Privacy preserving fitting of Cox models

- Allow Cox models to be fitted at each study, and then meta analyse the results
- Implementation is restricted to being study-level meta-analysis (SLMA)
 rather than full likelihood
- Server-side package: <u>dsSurvival</u>
 - SurvDS(...)
 - coxphSLMADS(...)
- Client-side package: <u>dsSurvivalClient</u>
 - ds.Surv(...) → assign function
 - ds.coxphSLMA(…) → aggregate function



dsSurvival Framework

Privacy preserving fitting of Cox models



AC: Analysis computer DC: Data computer DB: Database



Disclosure risks

Survival analysis

- Controlling the risk that the data analyst can deliberately infer to the identity or to one of the key variables being analyzed.
- The results of a survival analysis are likely to be disclosive if:
 - Reveal identifying information, or exact values of variables, including <u>dates</u>, diagnoses, and comorbidities
 - Reveal status of observations

O'Keefe, Christine M., et al. "Confidentialising survival analysis output in a remote data access system." Journal of Privacy and Confidentiality 4.1 (2012).

Disclosure risks

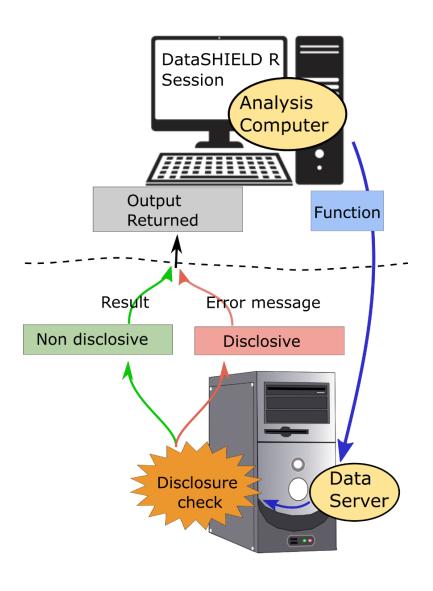
Cox proportional hazard models

$$h_i(t) = h_o(t) \exp\left(\sum_{j=1}^p \beta_j x_{ij}\right)$$
 Baseline hazard X Relative risk of covariates x_i

- Interested in the coefficient estimates β rather than the baseline hazard $h_o(t)$
- Do not release the values of the covariates x_{ij} for each participant
- Do not reveal the hazard function $h_i(t)$ (survival objects) for each participant



Disclosure control

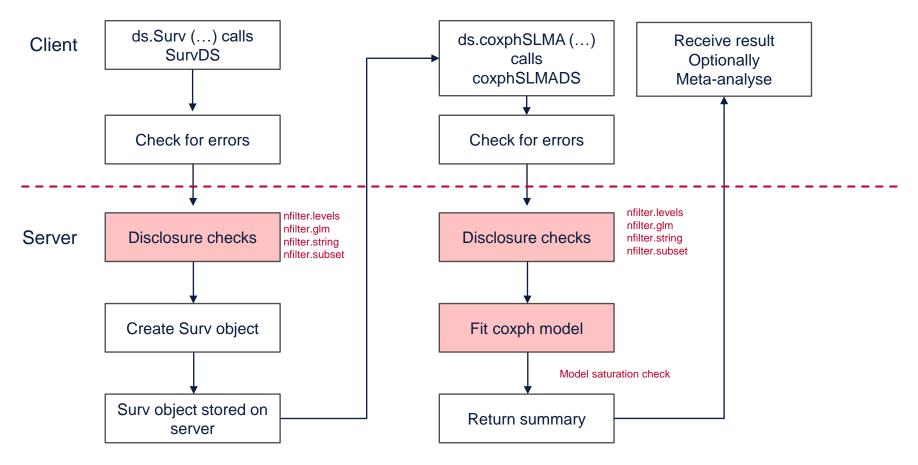


Disclosure checks

- nfilter.levels
- nfilter.tab
- nfilter.glm
- nfilter.string
- nfilter.subset



Disclosure checks



- Number of parameters in Cox model as a proportion of the sample size
- Default : 20% of sample size
- Prevents model oversaturation



Output presentation

\$study1

| | coef | exp(coef) | se(coef) | z | Pr(> z) |
|--------------------|---------|-----------|----------|-------|--------------|
| D\$age | 0.00815 | 1.008191 | 0.001248 | 6.535 | 6.35e-11 *** |
| D\$bmi | 0.00553 | 1.005551 | 0.030356 | 2.422 | 0.004245 ** |
| D\$factor(sex)male | 0.15224 | 1.164442 | 0.065621 | 0.215 | 0.000116 ** |
| | | | | | |

Signif. Codes: 0 '***' 0.001 '**' 0.05 '.' 0.1'' 1

\$study2

| | coef | exp(coef) | se(coef) | z | Pr(> z) |
|--------------------|----------|-----------|----------|--------|--------------|
| D\$age | 0.04067 | 1.04151 | 0.00416 | 9.776 | < 2e-16 *** |
| D\$bmi | -0.62756 | 0.53389 | 0.11767 | -5.333 | 9.66e-08 *** |
| D\$factor(sex)male | -0.66000 | 0.516850 | 0.099481 | -6.634 | 3.26e-11 *** |

Signif. Codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1' 1

\$study3

| | coef | exp(coef) | se(coef) | z | Pr(> z) |
|--------------------|-----------|-----------|----------|--------|------------|
| D\$age | 0.042145 | 1.043045 | 0.003086 | 13.655 | < 2e-16 * |
| D\$bmi | 0.006522 | 1.005551 | 0.03359 | 1.452 | 0.424513 * |
| D\$factor(sex)male | -0.599238 | 0.549230 | 0.084305 | -7.108 | 1.18e-12 * |

Signif. Codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1' 1



Metafor R package

- Meta-analysis of the hazard ratios
- Forest plots of estimates from RE model

| Study | logHR | SE | | | | | | 95%−CI | Weight (common) | Weight (random) |
|---|---|------|-------|------|-------|----------------------|--|--------|-------------------------|-------------------------|
| 1 2 3 | 1.0425 0.00 1.0415 0.00 1.0430 0.00 |)42 | | - | | — 1.041 . | 5 [1.0356; 5 [1.0334; 0 [1.0370; | 1.0497 | 33.4% 23.6% 42.9% | 33.4% 23.6% 42.9% |
| Common effect mode Random effects mode Heterogeneity: $I^2 = 0\%$, T | el | 1.03 | 1.035 | 1.04 | 1.045 | | 5 [1.0385; 5 [1.0385; | _ | 100.0% | 100.0% |

http://www.metafor-project.org



Summary

- DataSHIELD enables federated analysis and tailored disclosure controls
- dsSurvival is a DataSHIELD package for privacy preserving metaanalysis of survival data distributed across different sites
- A tutorial in bookdown format with code, diagnostics, plots and synthetic data is available here:
- https://neelsoumya.github.io/dsSurvivalbookdown/
- All code is available from the following repositories:
- https://github.com/neelsoumya/dsSurvivalClient/
- https://github.com/neelsoumya/dsSurvival/





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