



DEEPHEALTH

Hackathon - Course 1

Hackathon presentation

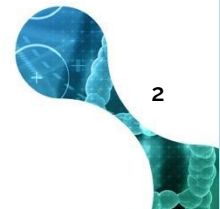


The project has received funding from the European Union's Horizon 2020 research and innovation programme under grant agreement No 825111.



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Introduction and Course Index

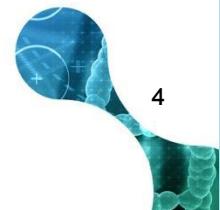


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Introduction

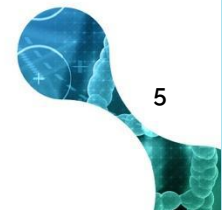
- How all the methodology associated with anonymisation methods and their implementation have been developed.
- Provide guidelines and best practices for the medical imaging community





Course Index

1. Hackathon presentation
2. De-identification of Radiological reports
3. De-identification of DICOM metadata
4. De-identification of Biomedical Images
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Pseudonymisation vs. Anonymisation



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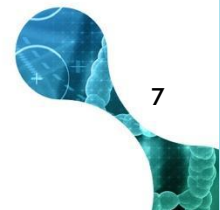


Definitions

De-identification is the process used to prevent someone's personal identity from being revealed.

The **anonymisation** of data is an irreversible process. Anonymised data cannot be linked to the original subject it identifies.

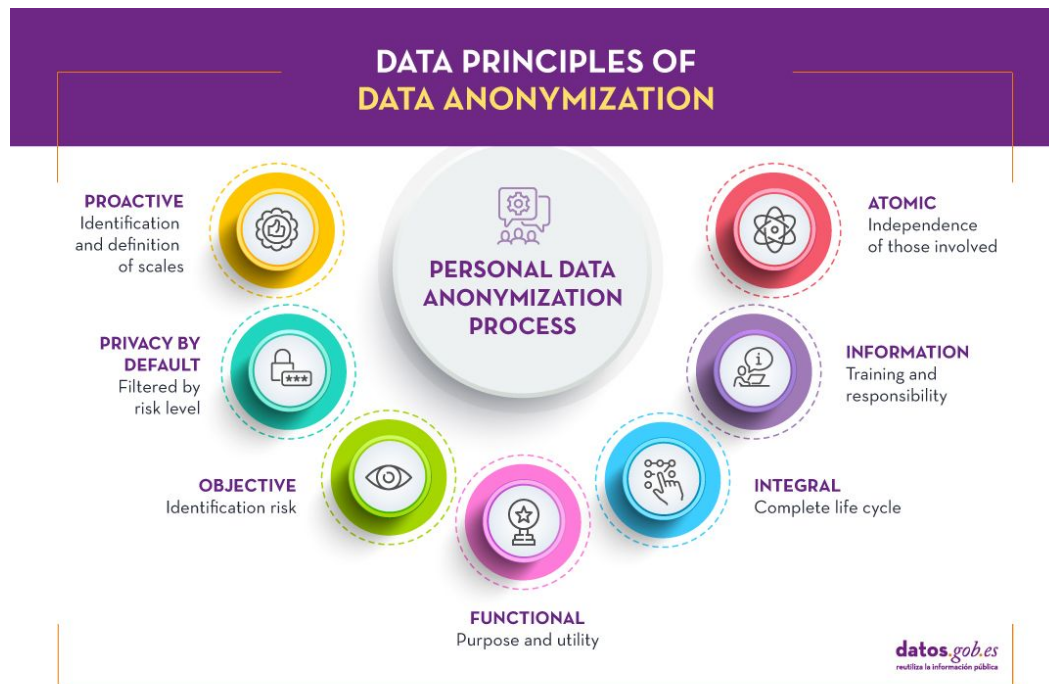
Pseudonymisation limits the traceability between the data and the original subject it identifies. It can be reversed.



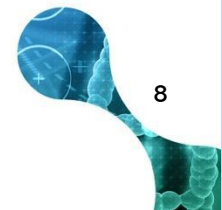


Basic principles

- Proactive
- Privacy by default
- Objective
- Functional
- Integral
- Information
- Atomic



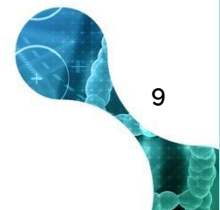
<https://datos.gob.es/en/blog/importance-anonymization-and-data-privacy>





Main risks

- **Singling out:** risk of extracting attributes that allow an individual to be identified.
- **Linkability:** risk of linking at least two attributes to the same individual or group, in one or more data sets.
- **Inference:** risk of deducing the value of a critical attribute from other attributes.

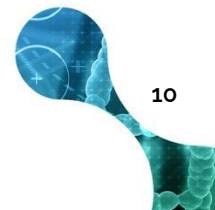




Anonymisation

Methods

- Randomisation
 - noise addition
 - permutation
 - differential privacy
- Generalisation
 - aggregation and anonymity "k"
 - diversity "l" / proximity "t"
- Encoding
 - Hash algorithms with secret key and key erasure
 - homomorphic encryption
 - time stamp



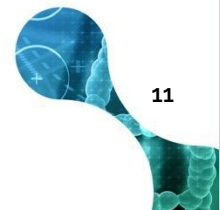


Pseudonymisation

Methods

The processing of personal data such that it can no longer be attributed to a specific data subject without using additional information.

- Hash algorithms
- Secret key encryption
- Token decomposition





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Clinical data and relevance of de-identification



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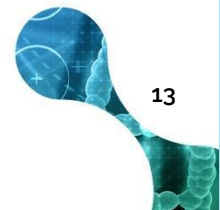
Medical Information Systems

HIPAA

The Health Insurance Portability and Accountability Act (**HIPAA**) proposes a set of good practices to protect:

- confidentiality
- integrity
- availability

of information in the health sector.



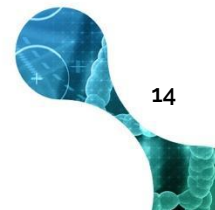


Medical Information Systems

Standards

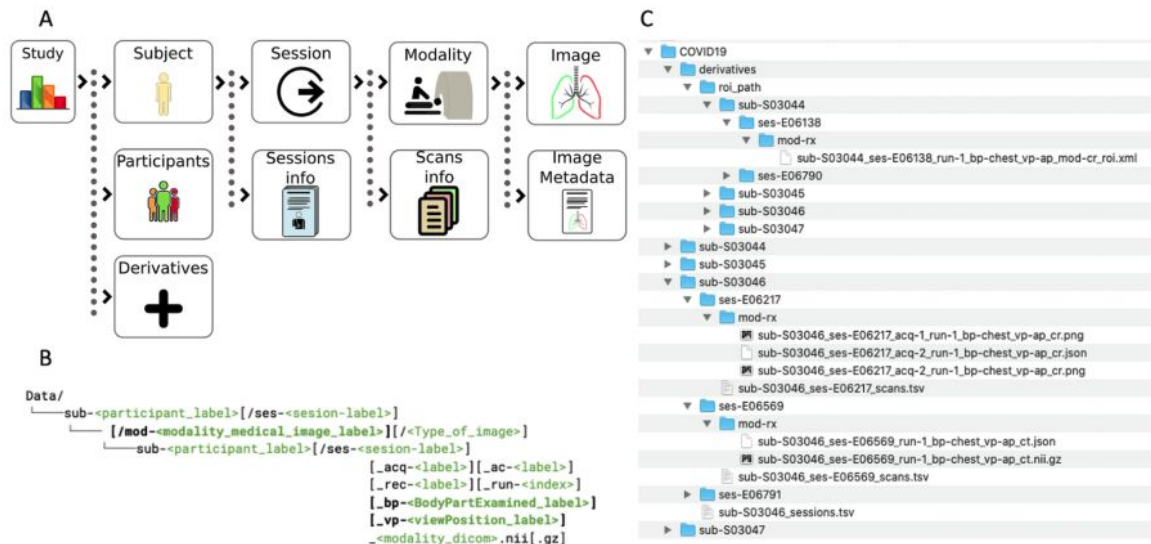
Standards for the development and management of hospital information:

- **HL7** (Health Level Seven Inc.) → standardise the exchange of medical information among applications
- **DICOM** (Digital Imaging and Communications in Medicine) → ensure the interoperability among heterogeneous medical imaging equipment and systems



MIDS

Medical Imaging Data Structure (**MIDS**) is the methodology proposed by FISABIO to standardise the organization and management of medical imaging data.





Types of anonymisable clinical data

Radiological reports

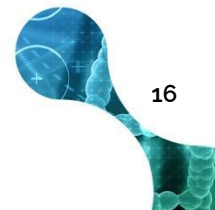
- Traditionally de-identified through pattern matching and regular expressions
- Natural Language Processing (NLP) de-identification as an alternative

DICOM meta information

- Data De-identification Guidelines and Protocol
- **Smart-Upload:** FISABIO's open-source DICOM de-identification software tool

Pixel data and graphic information

- Facial information
- Text annotations superimposed to the image





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GDPR on Clinical Data



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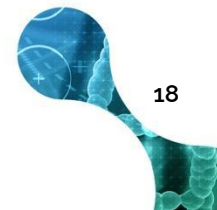
GDPR

General Data Protection Regulation

- It is applied to the processing of personal data
- It is applied in the form of national laws.
- Possibility of varying interpretations and legal implementations

Personal data must be collected, processed and shared under a lawful basis of the GDPR:

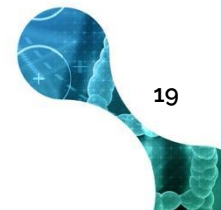
- Informed consent
- Public interest
- Legitimate interest





Data ownership

- The GDPR does not explicitly define or assign data ownership
- The subject has to give consent for:
 - collection
 - processing
 - sharingof their data
- This consent can be revoked at any time



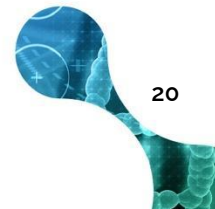
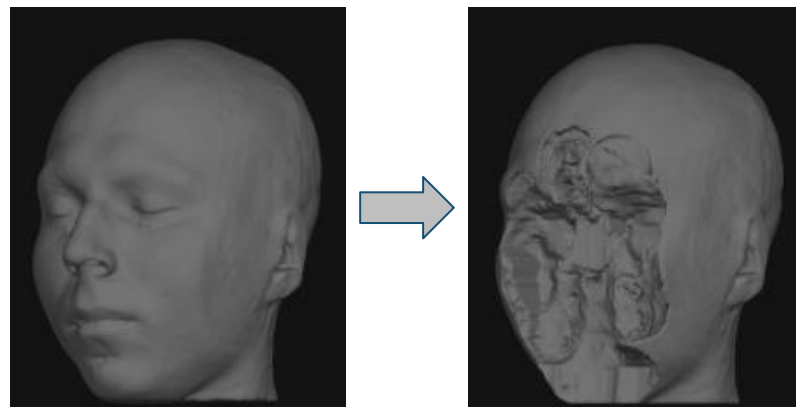


Clinical Data and GDPR

What constitutes anonymous data when it comes to clinical data?

e.g., Brain MRI

- Are they anonymous?
 - Facial features could be reconstructed
- How can they be anonymised?
 - Defacing





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BIMCV

Medical Imaging Data Bank of
Valencia Region



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BIMCV life cycle





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**Thank you for your
attention!**



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1.

Rocher, L., et al. Estimating the success of re-identifications in incomplete datasets using generative models. Nat. Commun.10, 1–9 (2019).

Yeh, F.-C. et al. Quantifying differences and similarities in whole-brain white matter architecture using local connectome fingerprints. PLoS Comput. Biol. 12, e1005203 (2016).

