

Data management plan checklist

<p>Table of content</p>	<ol style="list-style-type: none"> 1. Administrative Data 2. Data Collection and organization 3. Documentation and Metadata 4. Ethics and Legal Compliance 5. Storage and Backup 6. Long-term preservation 7. Data Availability 8. DMP-online
<p>1. Administrative Data (kort afsnit da alle disse oplysninger skal stå i protokollen)</p>	<p>-Name of the persons involved in the project</p> <p>-Project Description: short introduction and clearly defined objectives</p> <p>-Method: Design, participants, sites and dates, guidelines followed (e.g. CONSORT, STROBE, and TRIPOD)</p> <p>-Related policies: Study permissions and approvals (registration numbers), founding, study owners, data protection/security policy from the institution, and data management policy.</p> <p><i>Henvis evt. til protokollen og SAP</i></p>
<p>2. Data collection and organization</p>	<p>-Type and source of the data (questionnaire, interview, biological samples, extraction from medical record mv.)</p> <p>-Recommend a figure illustrating how data is to be combined e.g.</p> <div style="text-align: center;"> <pre> graph TD Population[Population] --> Interview[Interview] Population --> MedicalRecord[Medical record] Population --> BiologicalSample[Biological sample] Interview --> RedCap[RedCap or other] MedicalRecord --> RedCap BiologicalSample --> Microbiology[Microbiology data base] RedCap --> Sharepoint[Sharepoint] Microbiology --> Sharepoint </pre> </div> <p>-Names of folders and files. Do-files should include detailed information on data process and management, including labelling and preparation for the analysis. Results files should match precisely the tables and figures applied in the papers.</p> <div style="text-align: center;"> <pre> graph TD subgraph Data-file RD1[Raw data (Microbiology data)] --> CD1[clean data (Labeling/grouping)] RD2[Raw data (RedCap or other)] --> CD2[clean data (Labeling/grouping)] CD1 --> WDF[Work data file Merged.dta] CD2 --> WDF end subgraph Do-file DRD[Do-Raw data] --> MDL[Master Do-file logged] DPC[Do-preparation and cleaning] --> MDL DOA[Do-analysis -descriptive -outcome -sensitivities] --> MDL end subgraph Results-file R[Results -Tables -Figures] end </pre> </div>

	<p>-Include a system describing the version of the documents and files e.g. Nameyyyymmdd (year, month, day)</p> <p>-consider a quality assurance process (e.g. supervision, external data check)</p> <p>Henvis her til terminologilisten og evt. til SAP.</p>																																										
<p>3. Documentation and Metadata</p>	<p>-Interpretation of the data in the future, helping secondary users to understand and reuse it, including a codebook e.g. [henvis til terminology-listen]</p> <p>(Clinical studies)</p> <table border="1" data-bbox="466 631 1437 801"> <thead> <tr> <th>Variable</th> <th>Code</th> <th>Classification</th> <th>Data source</th> <th>Data measures</th> <th>Time of measurement</th> <th>consideration</th> </tr> </thead> <tbody> <tr> <td>gender</td> <td>Journal_sex</td> <td>Binary 1 male 2 female</td> <td>-at patient interview -extracted from the patient identification number</td> <td></td> <td>-At admission</td> <td>-reason of why this variable is of interest</td> </tr> <tr> <td>crp</td> <td>Blod_crp</td> <td>Continuous</td> <td>-patient medical record</td> <td>-Mg/L</td> <td>-At baseline, 3 and 6 months from baseline</td> <td>-predictor of pneumonia</td> </tr> </tbody> </table> <p>(Register studies)</p> <table border="1" data-bbox="466 891 1437 1061"> <thead> <tr> <th>Variable</th> <th>Code</th> <th>Classification</th> <th>Data source</th> <th>Data measures</th> <th>Time of measurement</th> <th>consideration</th> </tr> </thead> <tbody> <tr> <td>gender</td> <td>Journal_sex</td> <td>Binary 1 male 2 female</td> <td>-at patient interview -extracted from the patient identification number</td> <td></td> <td>-At admission</td> <td>-reason of why this variable is of interest</td> </tr> <tr> <td>crp</td> <td>Blod_crp</td> <td>Continuous</td> <td>-patient medical record</td> <td>-Mg/L</td> <td>-At baseline, 3 and 6 months from baseline</td> <td>-predictor of pneumonia</td> </tr> </tbody> </table> <p>- Data control for; requirements of the values for categorical variables, intervals allowed for numerical variables, assumptions for dates/times, and the lengths of time intervals.</p> <p>-[Clinical studies] Handling of missing data: description of the expected reasons for missing data, the expected type of missing (MCAR, MAR, MNAR), and how missing data in each relevant variable is handled (exclude observations, impute values, or apply other handling strategies). Henvis her evt. til SAP.</p> <p>-Timeline for data collection and follow-ups (example below for a clinical study). Henvis her evt. til protokollen eller SAP.</p> <p>Timeline for data collection of the INDEED study (Skjøt-Arkil H et al BMJ open 2021):</p>	Variable	Code	Classification	Data source	Data measures	Time of measurement	consideration	gender	Journal_sex	Binary 1 male 2 female	-at patient interview -extracted from the patient identification number		-At admission	-reason of why this variable is of interest	crp	Blod_crp	Continuous	-patient medical record	-Mg/L	-At baseline, 3 and 6 months from baseline	-predictor of pneumonia	Variable	Code	Classification	Data source	Data measures	Time of measurement	consideration	gender	Journal_sex	Binary 1 male 2 female	-at patient interview -extracted from the patient identification number		-At admission	-reason of why this variable is of interest	crp	Blod_crp	Continuous	-patient medical record	-Mg/L	-At baseline, 3 and 6 months from baseline	-predictor of pneumonia
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6. Long-term preservation	<p>-Time the data (reports, biological samples) will be available -Achievement of the data</p> <p>Example for biological samples preservation of the INDEED study (Skjøt-Arkil H et al BMJ open 2021):</p> <table border="1"> <thead> <tr> <th></th> <th>Blood for analysis of PCT and suPAR</th> <th>Blood for research biobank</th> </tr> </thead> <tbody> <tr> <td>Collection</td> <td>The blood will be collected in an EDTA plasma tube.</td> <td>Biobank blood is only collected for patient in track A and includes one tube of EDTA plasma and one tube of LiHeparin plasma.</td> </tr> <tr> <td>Storage</td> <td>At two of the sites, the analysis will be performed within is tested within two hours from the collection of the blood sample. At the third site, samples will be stored locally in a -80 °C freezer. The samples are pseudonymized with a code to protect the identity of the participants. All samples and code lists are stored safely and separately to prevent unauthorized access.</td> <td>All samples will be stored locally in a -80 °C freezer. The samples are pseudonymized with a code to protect the identity of the participants. All samples and code lists are stored safely and separately to prevent unauthorized access.</td> </tr> <tr> <td>Sample analysis</td> <td><i>Serum procalcitonin (PCT)</i> Serum PCT concentration is quantified with an automated sandwich immunoassay "ECLIA" (Elecsys®, BRAHMS PCT-analyses) on Cobas e801. Calibration is performed after Cobas e pack has been registered in the instrument and is standardized to the BRAHMS PCT LIA assay. The correlation of Elecsys BRAHMS PCT analyses has been compared to BRAHMS PCT LIA and to BRAHMS PCT sensitive KRYPTOR with similar results of r=0.981 and r=0.988 respectively.</td> <td>Molecular analysis for future use in ancillary studies will take place after all samples have been collected.</td> </tr> <tr> <td>Evaluation</td> <td>The results will be saved in a study database and not be visible for the physician in the medical journal.</td> <td>The results will be saved in a study database. The expiry date of the research biobank is expected to be October 2022. After expiry date, the remaining material in the research bank will be destroyed.</td> </tr> <tr> <td>Location</td> <td>Samples will be located at Bloodsamples, Biochemistry and Immunology, University Hospital of Southern Denmark, Aabenraa, Denmark</td> <td>Samples will be located at: - Bloodsamples, Biochemistry and Immunology, University Hospital of Southern Denmark, Aabenraa, Denmark</td> </tr> </tbody> </table> <p>[Link til hjemmeside, når anbefalingerne er lagt ud]</p>		Blood for analysis of PCT and suPAR	Blood for research biobank	Collection	The blood will be collected in an EDTA plasma tube.	Biobank blood is only collected for patient in track A and includes one tube of EDTA plasma and one tube of LiHeparin plasma.	Storage	At two of the sites, the analysis will be performed within is tested within two hours from the collection of the blood sample. At the third site, samples will be stored locally in a -80 °C freezer. The samples are pseudonymized with a code to protect the identity of the participants. All samples and code lists are stored safely and separately to prevent unauthorized access.	All samples will be stored locally in a -80 °C freezer. The samples are pseudonymized with a code to protect the identity of the participants. All samples and code lists are stored safely and separately to prevent unauthorized access.	Sample analysis	<i>Serum procalcitonin (PCT)</i> Serum PCT concentration is quantified with an automated sandwich immunoassay "ECLIA" (Elecsys®, BRAHMS PCT-analyses) on Cobas e801. Calibration is performed after Cobas e pack has been registered in the instrument and is standardized to the BRAHMS PCT LIA assay. The correlation of Elecsys BRAHMS PCT analyses has been compared to BRAHMS PCT LIA and to BRAHMS PCT sensitive KRYPTOR with similar results of r=0.981 and r=0.988 respectively.	Molecular analysis for future use in ancillary studies will take place after all samples have been collected.	Evaluation	The results will be saved in a study database and not be visible for the physician in the medical journal.	The results will be saved in a study database. The expiry date of the research biobank is expected to be October 2022. After expiry date, the remaining material in the research bank will be destroyed.	Location	Samples will be located at Bloodsamples, Biochemistry and Immunology, University Hospital of Southern Denmark, Aabenraa, Denmark	Samples will be located at: - Bloodsamples, Biochemistry and Immunology, University Hospital of Southern Denmark, Aabenraa, Denmark																																																																																																																																								
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<p>7. Data Availability</p>	<p>-Due to Danish laws on personal data, data cannot be shared publicly: Data protection laws https://www.datatilsynet.dk/english/legislation</p> <p>-describe how to request the data and from whom -Consider at what level, the data can be shared and what kind of data (common projects or international collaboration)</p> <p>Example: Availability of data and material: “Due to Danish laws on personal data, data cannot be shared publicly. Please contact the corresponding author XXXX to request this data. The person responsible for the research was the principal investigator and corresponding author (XXXXX), and together with the Department of XXXX and the University Hospital of Southern Denmark own the data and have access to the final data set.</p>
<p>8. DMP- online register</p>	<p>https://dmponline.deic.dk/</p> <p>Benefits: -More people can work on the document (e.g. supervisors) -DMP plan has the possibility of public sharing -The document can be adapted on an ongoing basis</p> <p>Limitations: -Not possible to attach figures and tables</p>