



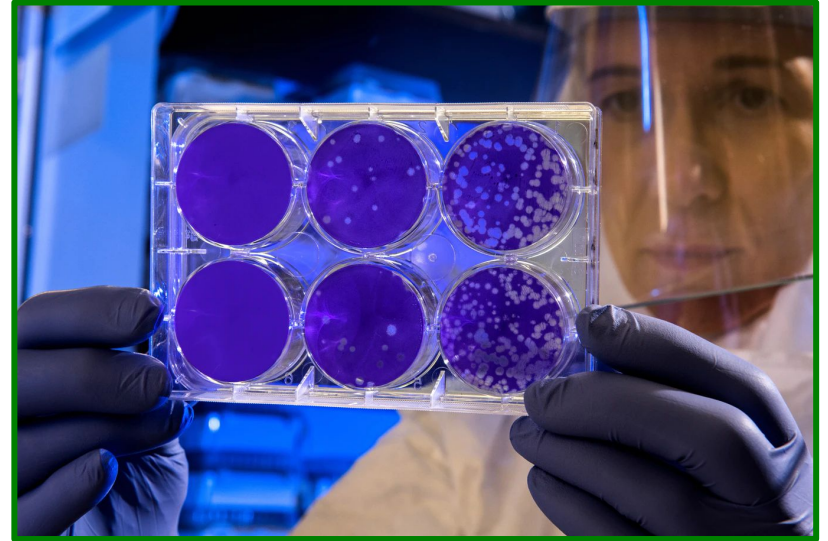
Ready for BioData Management?

How to Make a DMP

João Cardoso, Daniel Faria

Let Us Consider a Mock Project

- **Project X**
 - Unveiling the mechanisms of Disease X
 - BioData.pt is applying for funding from the Fundação para a Ciência e Tecnologia (FCT).



Let Us Consider a Mock Project

- **Motivation**

- The cause of **Disease X** has been recently discovered to be a **virus**, phage X.
- It **infects** normal **gut bacteria** and leads them to become virulent and cause **chronic intestinal infection**.
- This disease has been **spreading rapidly** in Europe, with **costs** in health-care reaching the **tens of millions of Euros**.



Let Us Consider a Mock Project

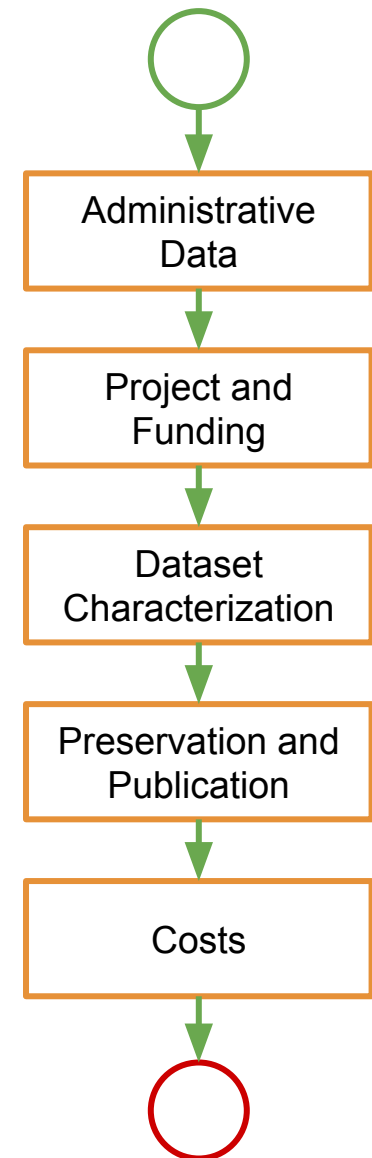
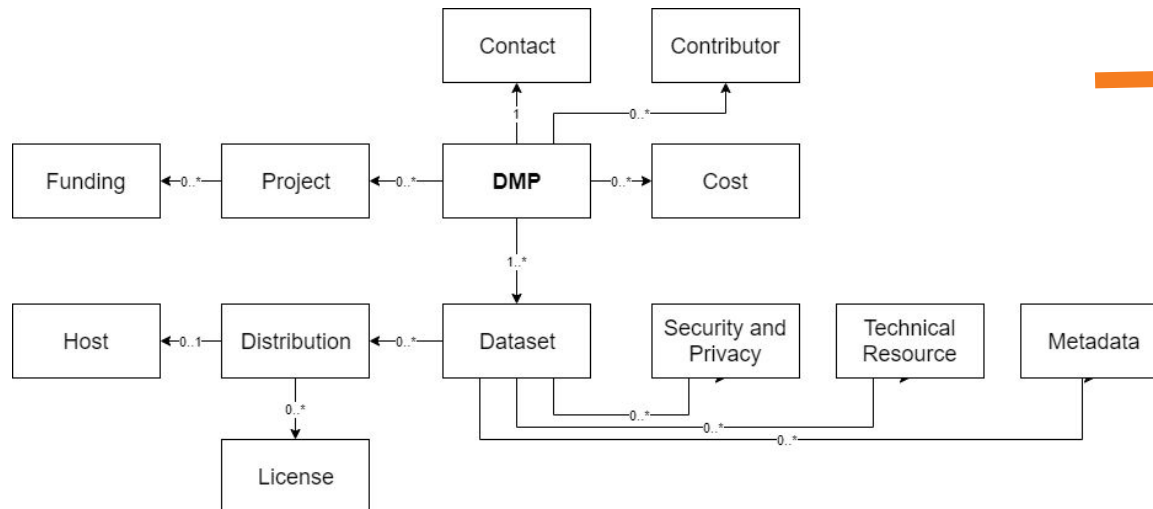
- **Objectives**

- **Uncover the mechanisms** of disease X by **sequencing phage X** and studying the **effects of its infection** in human gut microbiota at the **population and molecular level**.
- **Assess which bacterial taxa** are **infected** by phage X and what are the **effects on the relative abundance** of the various taxa.
- Study the **effects** on the **infected taxa** at the **gene expression level**.
- **Improving treatment** for Disease X, and potentially being able to **cure it**.



Creating a DMP

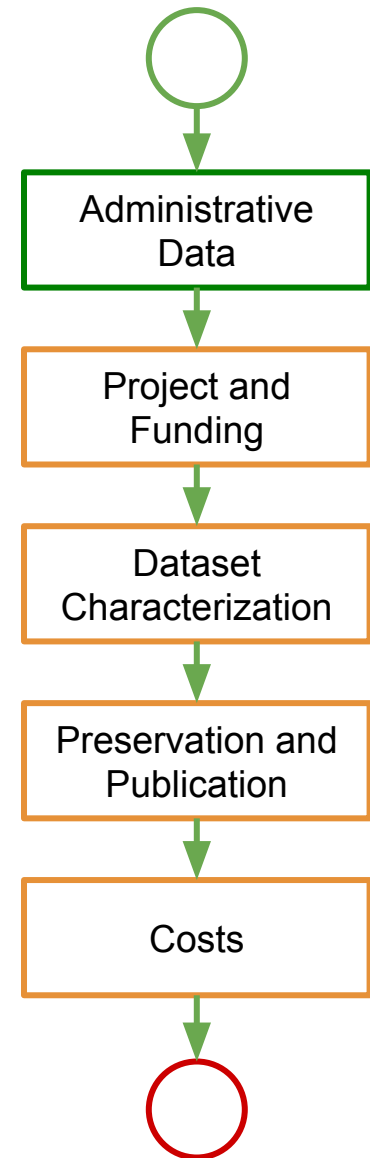
- The **DMP Creation Methodology** comprises 5 steps.
- Each step focuses on a **specific aspect** of the DMP.
- It is **based** on the **RDA's DMP Common Standards** metadata application standard.



Creating a DMP (Step 1)

- **Step 1 - Administrative Data**

- **Characterization** of the **DMP document**, and the **responsibilities** of all the **people** mentioned.
- The information is classified in three sections:
 - **General information** characterizing the **DMP document**.
 - **Contact** (person of institution) for the DMP.
 - A listing of all **collaborators** and their **roles** in the DMP.



Creating a DMP (Step 1)

In the project

Project X (Application to Fundação para a Ciência e Tecnologia)

Title of the project: Unveiling the mechanisms of Disease X

Participants:

- Prof. Coor Dinator (coor.dinator@biodata.pt) [PI & DMP Coordinator]
- Dr. Dat Manger (dat.manger@biodata.pt) [Data Manager]
- Dr. Col Hector (col.hector@biodata.pt) [Clinical Data & Sample Collector]
- Dr. R. Sercher (r.sercher@biodata.pt) [Researcher]
- Mrs. A. D'Min (a.dmin@biodata.pt) [Project Manager]

Host Institution: BioData.pt

Start date: January 1st, 2021

Duration: 36 months

Creating a DMP (Step 1)

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- Generic information on the DMP document:
 - Title, institution, start date, duration, etc.
 - Description from the abstract.

Creating a DMP (Step 1)

In the project

Abstract:

The cause of Disease X has been recently discovered to be a virus, phage X, which infects normal gut bacteria and leads them to become virulent and cause chronic intestinal infection. Although non-fatal, this disease has been spreading rapidly in Europe, with costs in health-care reaching the tens of millions of Euros.

This project aims to uncover the mechanisms of disease X by sequencing phage X and studying the effects of its infection in human gut microbiota at the population and molecular level. We will assess which bacterial taxa are infected by phage X and what effect the infection has on the relative abundance of the various taxa, as well as what effect the infection has on the abundance of the various taxa at the gene expression level.

The project will be a key step towards improving our understanding of the disease, potentially being able to cure it.

- Generic information on the DMP document:
 - Description from the abstract.

Creating a DMP (Step 1)

In the project

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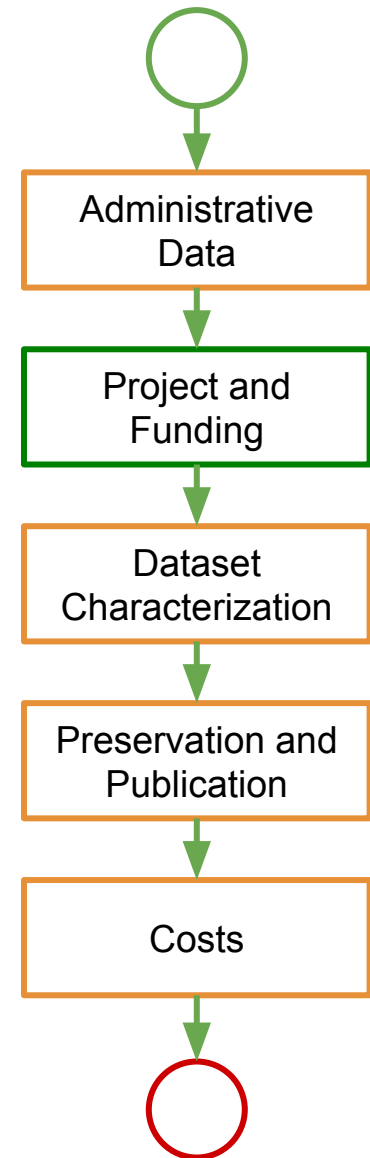
Duration: 36 months

- A listing of all **collaborators** and their **roles** in the DMP.

Creating a DMP (Step 2)

- **Step 2 - Project and Funding**

- **Characterization** of the **project(s)** and their sources of **funding**.
- The information is classified in two sections:
 - Information regarding the **project(s)** to which the **DMP** is **associated**.
 - Information pertaining the **funding** of a **particular project**.



Creating a DMP (Step 2)

In the project

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Title of the project: Unveiling the mechanisms of Disease X

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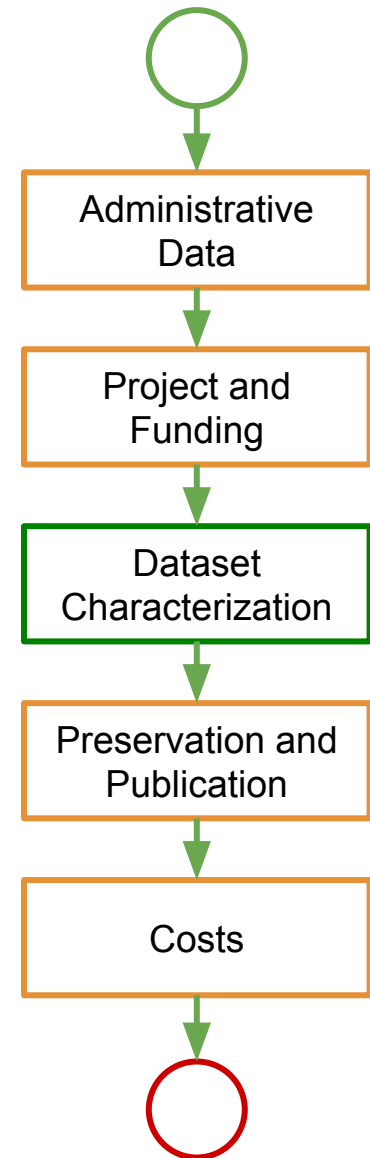
Duration: 36 months

- Information pertaining the **funding** of a **particular project**.

Creating a DMP (Step 3)

- **Step 3 - Dataset Characterization**

- **Characterization** of the **dataset(s)** that are encompassed by the DMP. Apart from **generic information** on the dataset, **additional** descriptions of **security and privacy** policies, **technical resources** and **metadata** standards can also be given.
- The information is classified in four sections:
 - **General information** about all **datasets**.
 - Any **security and privacy** policies associated with the datasets.
 - **Technical resources** associated with the datasets.
 - **Metadata** associated with the datasets.



Creating a DMP (Step 3)

In the project

Research plan and method summary:

The project will be divided into four activities:

1. Sample collection
2. Phage X sequencing
3. 16S sequencing
4. Metatranscriptomics

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In the project

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4. Metatranscriptomics

- **General information** about all datasets.

Creating a DMP (Step 3)

In the project

Sample Collection:

In the sample collection activity, we will define a study group of volunteer disease X patients, numbering no less than 20, and a control group comprising their close relatives, 1-2 per patient. We will collect stool samples from each of the volunteers.

Creating a DMP (Step 3)

In the project

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- Any security and privacy policies associated with the datasets.

Creating a DMP (Step 3)

In the project

Phage X sequencing:

In the Phage X sequencing activity, we will carry out DNA sequencing of the stool samples and assemble the genome of Phage X. In order to facilitate the assembly while enabling the reliable identification of sequence variants, we will combine the higher quality but short read sequencing technology of the Illumina NextSeq 500 sequencer with the long read but lower quality technology of the Nanopore MinION sequencer.

Creating a DMP (Step 3)

In the project

Phage X sequencing:

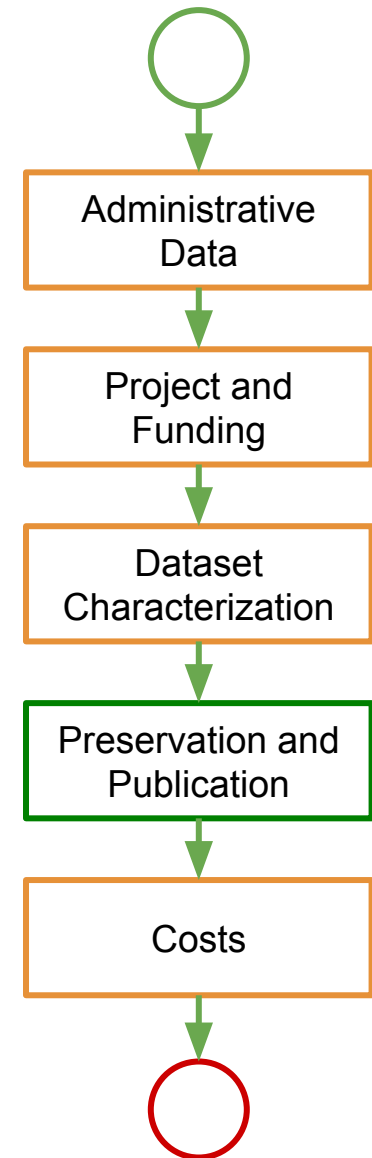
In the Phage X sequencing activity, we will carry out DNA sequencing of the stool samples and assemble the genome of Phage X. In order to facilitate the assembly while enabling the reliable identification of sequence variants, we will combine the higher quality but short read sequencing technology of the Illumina NextSeq 500 sequencer with the long read but lower quality technology of the Nanopore MinION sequencer.

- **Technical resources** associated with the datasets.
- **Metadata** associated with the datasets.

Creating a DMP (Step 4)

- **Step 4 - Preservation and Publication**

- **Characterization** of the **preservation** and **publication** policies **for each** of the identified **datasets**.
- The information is classified in three sections:
 - Information regarding the **policies** on how each dataset is **distributed**.
 - Information on the data **host for each** of the identified **distributions**.
 - Characterization of the **licenses** associated with each **distribution policies**.



Creating a DMP (Step 4)

In the project



Creating a DMP (Step 4)

In the project

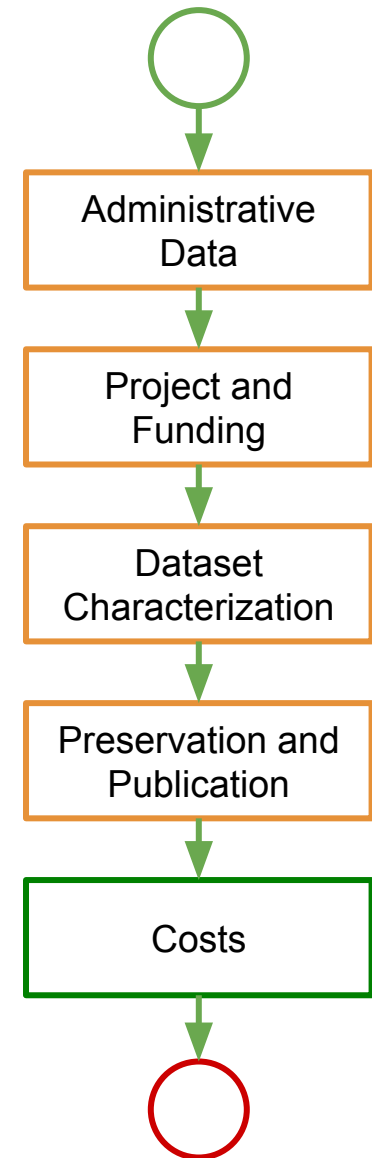
Why is this empty? Not usually covered in project descriptions.
What would you write?

- Information regarding the **policies** on how each dataset is **distributed**.
- Information on the data **host** **for each** of the identified **distributions**.
- Characterization of the **licenses** associated with each **distribution policies**.

Creating a DMP (Step 5)

- **Step 5 - Costs**

- **Characterization** of the **costs** associated with this DMP.
 - The numeric value associated with each cost can be a rough estimate.



Creating a DMP (Step 5)

In the project

Expected Data & Metadata Outputs:

1. Sample Collection:
 - Patient clinical data (< 1 MB)
 - Sample identification table (< 1 MB)
2. Phage X sequencing
 - Raw FASTQ sequencing data - NextSeq (60 MB)
 - Sample preparation & sequencing metadata - NextSeq (< 1 MB)
 - Raw FASTQ sequencing data - MinION (1 GB)
 - Sample preparation & sequencing metadata - MinION (< 1 MB)
 - Assembled Phage X genome (< 1 MB)
 - Assembly metadata (< 1 MB)
3. ...

Creating a DMP (Step 5)

In the project

Expected Data & Metadata Outputs:

1. Sample Collection:

- Patient clinical data (< 1 MB)
- Sample identification table (< 1 MB)

2. Phage X sequencing:

- Raw FASTQ sequencing data - NextSeq (60 MB)
- Sample preparation & sequencing metadata - NextSeq (< 1 MB)
- Raw FASTQ sequencing data - MinION (1 GB)
- Sample preparation & sequencing metadata - MinION (< 1 MB)
- Assembled Phage X genome (< 1 MB)
- Assembly metadata (< 1 MB)

What other costs would you consider?

- **Characterization** of the costs associated with this DMP.

Thank you!

BioData.pt

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