

FWO DMP Template - Flemish Standard Data Management Plan

Version KU Leuven

Project supervisors (from application round 2018 onwards) and fellows (from application round 2020 onwards) will, upon being awarded their project or fellowship, be invited to develop their answers to the data management related questions into a DMP. The FWO expects a **completed DMP no later than 6 months after the official start date** of the project or fellowship. The DMP should not be submitted to FWO but to the research co-ordination office of the host institute; FWO may request the DMP in a random check.

At the end of the project, the **final version of the DMP** has to be added to the final report of the project; this should be submitted to FWO by the supervisor-spokesperson through FWO's e-portal. This DMP may of course have been updated since its first version. The DMP is an element in the final evaluation of the project by the relevant expert panel. Both the DMP submitted within the first 6 months after the start date and the final DMP may use this template.

The DMP template used by the Research Foundation Flanders (FWO) corresponds with the Flemish Standard Data Management Plan. This Flemish Standard DMP was developed by the Flemish Research Data Network (FRDN) Task Force DMP which comprises representatives of all Flemish funders and research institutions. This is a standardized DMP template based on the previous FWO template that contains the core requirements for data management planning. To increase understanding and facilitate completion of the DMP, a standardized **glossary** of definitions and abbreviations is available via the following [link](#).

1. General Project Information	
Name Grant Holder & ORCID	Alexandra Bacquelaine Veloso, 0000-0001-5807-7578
Contributor name(s) (+ ORCID) & roles	
Project number ¹ & title	D-2024-2579
Funder(s) GrantID ²	12AZ324N
Affiliation(s)	<input checked="" type="checkbox"/> KU Leuven <input type="checkbox"/> Universiteit Antwerpen <input type="checkbox"/> Universiteit Gent <input type="checkbox"/> Universiteit Hasselt <input type="checkbox"/> Vrije Universiteit Brussel <input type="checkbox"/> Other: ROR identifier KU Leuven: 05f950310
Please provide a short project description	<p>The main consequence of these ectopic transcription factors, such as TLX1 and TLX3, is the deregulation of gene expression, which is one of the hallmarks of T-ALL development. Still, the mechanisms that link these two effects remains elusive. Thus, I am focused on characterising transcriptional deregulation caused by the ectopic expression of TLX1 and TLX3 transcription factors found in T-ALL. I hypothesise that TLX1 and TLX3 regulate gene expression with epigenetic and other transcription factors, to promote T-ALL development. To do so, I will first focus on identifying proteins that interact with TLX1 and TLX3 in T-ALL cells by using proteomics assays. Secondly, I will perform a CRISPR screen to determine transcriptional regulators and chromatin modifiers that together with TLX1 or TLX3 promote T-ALL proliferation. Finally, I will perform a drug screen against the most pertinent targets to obtain the most efficient treatment.</p> <p>Overall, this project will reveal the members within these ectopic transcriptional complexes, their role in T-ALL development, and the best targets to kill T-ALL cells from these two subsets.</p>

¹ "Project number" refers to the institutional project number. This question is optional. Applicants can only provide one project number.

² Funder(s) GrantID refers to the number of the DMP at the funder(s), here one can specify multiple GrantIDs if multiple funding sources were used.

2. Research Data Summary

List and describe all datasets or research materials that you plan to generate/collect or reuse during your research project. For each dataset or data type (observational, experimental etc.), provide a short name & description (sufficient for yourself to know what data it is about), indicate whether the data are newly generated/collected or reused, digital or physical, also indicate the type of the data (the kind of content), its technical format (file extension), and an estimate of the upper limit of the volume of the data ³.

Dataset Name	Description	New or Reused	Digital or Physical	ONLY FOR DIGITAL DATA	ONLY FOR DIGITAL DATA	ONLY FOR DIGITAL DATA	ONLY FOR PHYSICAL DATA
				Digital Data Type	Digital Data Format	Digital Data Volume (MB, GB, TB)	Physical Volume
Single cell RNA-seq CRISPR screen	Single-cell RNA-seq data from mouse pro-T and DP cells upon CRISPR inactivation of library of genes	<input checked="" type="checkbox"/> Generate new data <input type="checkbox"/> Reuse existing data	<input checked="" type="checkbox"/> Digital <input type="checkbox"/> Physical	Experimental	<i>nextgeneration sequence data (Illumina)</i>	<input type="checkbox"/> < 1 GB <input checked="" type="checkbox"/> < 100 GB <input type="checkbox"/> < 1 TB <input type="checkbox"/> < 5 TB <input type="checkbox"/> > 5 TB <input type="checkbox"/> NA	
Protein interaction data	<i>proteomics data from interaction partners of TLX transcription factors</i>	<input checked="" type="checkbox"/> Generate new data	<input checked="" type="checkbox"/> Digital	Experimental	<i>proteomics dataset (MassSpec analysis)</i>	<input checked="" type="checkbox"/> < 100 GB	

GUIDANCE:

The data description forms the basis of your entire DMP, so make sure it is detailed and complete. It includes digital and physical data and encompasses the whole spectrum ranging from raw data to processed and analysed data including analysis scripts and code. Physical data are all materials that need proper management because they are valuable, difficult to replace and/or ethical issues are associated. Materials that are not considered data in an RDM context include your own manuscripts, theses and presentations; documentation is an integral part of your datasets and should be described under documentation/metadata.

[RDM Guidance on data](#)

³ Add rows for each dataset you want to describe.

If you reuse existing data, please specify the source, preferably by using a persistent identifier (e.g. DOI, Handle, URL etc.) per dataset or data type.	NA
Are there any ethical issues concerning the creation and/or use of the data (e.g. experiments on humans or animals, dual use)? If so, refer to specific datasets or data types when appropriate and provide the relevant ethical approval number.	<input type="checkbox"/> Yes, human subject data; provide SMEC or EC approval number: <input checked="" type="checkbox"/> Yes, animal data; provide ECD reference number: 030/2023 <input type="checkbox"/> Yes, dual use; provide approval number: <input type="checkbox"/> No Additional information: <i>The experiments for this project will be conducted on primary mouse pro-T cells for which we only need very few animals. We will isolate bone marrow cells from the animals and culture the cells ex vivo. Thus, we only need post-mortem animals, there is no suffering, and there is no housing of the animals needed. Total number of animals (mice) will be below maximum 20 mice for the entire project.</i>
Will you process personal data ⁴ ? If so, please refer to specific datasets or data types when appropriate and provide the KU Leuven or UZ Leuven privacy register number (G or S number).	<input type="checkbox"/> Yes (provide PRET G-number or EC S-number below) <input checked="" type="checkbox"/> No Additional information:
Does your work have potential for commercial valorization (e.g. tech transfer, for example spin-offs, commercial exploitation, ...)? If so, please comment per dataset or data type where appropriate.	<input checked="" type="checkbox"/> Yes <input type="checkbox"/> No If yes, please comment: <i>Both the RNA-seq dataset and the proteomics dataset can lead to the identification of new targets for therapy. We will discuss the data and potential commercial valorization with the Tech transfer office at VIB and LRD at KU Leuven to determine the possibilities for tech transfer. We will work with them to determine a publication plan to ensure that publication does not affect the tech transfer possibilities.</i>
Do existing 3rd party agreements restrict exploitation or dissemination of the data you (re)use (e.g. Material/Data transfer agreements, research collaboration agreements)? If so, please explain to what data they relate and what restrictions are in place.	<input type="checkbox"/> Yes <input checked="" type="checkbox"/> No If yes, please explain:

⁴ See Glossary Flemish Standard Data Management Plan

<p>Are there any other legal issues, such as intellectual property rights and ownership, to be managed related to the data you (re)use? If so, please explain to what data they relate and which restrictions will be asserted.</p>	<p><input type="checkbox"/> Yes <input checked="" type="checkbox"/> No If yes, please explain:</p>
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3. Documentation and Metadata	
<p>Clearly describe what approach will be followed to capture the accompanying information necessary to keep data understandable and usable, for yourself and others, now and in the future (e.g. in terms of documentation levels and types required, procedures used, Electronic Lab Notebooks, README.txt files, Codebook.tsv etc. where this information is recorded).</p> <p>RDM guidance on documentation and metadata.</p>	<p><i>Each dataset will be accompanied by a detailed excel file and text file explaining how the experiment was performed (cells used, oncogenes used, cell culture conditions, amounts of cells used, RNA/protein isolation methods, purification methods, meaning of the different labels used in the dataset).</i></p> <p><i>For the RNA-seq data: this will be the result of a CRISPR screen, so each of the gRNA sequences used in the screen will be described.</i></p> <p><i>For the proteomics data: this will be the result of protein-protein interactions that are mapped, so the exact method to isolate the protein-protein interactions will be described.</i></p>

<p>Will a metadata standard be used to make it easier to find and reuse the data?</p> <p>If so, please specify which metadata standard will be used. If not, please specify which metadata will be created to make the data easier to find and reuse.</p> <p><i>REPOSITORIES COULD ASK TO DELIVER METADATA IN A CERTAIN FORMAT, WITH SPECIFIED ONTOLOGIES AND VOCABULARIES, I.E. STANDARD LISTS WITH UNIQUE IDENTIFIERS.</i></p>	<p><input checked="" type="checkbox"/> Yes <input type="checkbox"/> No</p> <p>If yes, please specify (where appropriate per dataset or data type) which metadata standard will be used: <i>RNA-seq data is next-generation sequencing data for which specific standards are used to deposit the data. We will follow the recommendations of the KU Leuven genomics core facility. MIAME guidelines will be followed: https://www.ncbi.nlm.nih.gov/geo/info/MIAME.html</i> <i>Proteomics data has specific standards for data deposits. We will follow the recommendations of the VIB proteomics facility. We will follow the guidelines of the PRIDE data repository: https://www.ebi.ac.uk/pride/markdownpage/specificsoftwareformats</i></p> <p>If no, please specify (where appropriate per dataset or data type) which metadata will be created:</p>
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4. Data Storage & Back-up during the Research Project	
<p>Where will the data be stored?</p> <p><i>Consult the interactive KU Leuven storage guide to find the most suitable storage solution for your data.</i></p>	<p><input checked="" type="checkbox"/> Shared network drive (J-drive) <input type="checkbox"/> Personal network drive (I-drive) <input checked="" type="checkbox"/> OneDrive (KU Leuven) <input type="checkbox"/> Sharepoint online <input type="checkbox"/> Sharepoint on-premis <input type="checkbox"/> Large Volume Storage <input type="checkbox"/> Digital Vault <input type="checkbox"/> Other: <i>KU Leuven network drives, external hard drive (as additional backup); VSC (Vlaams Supercomputer Center)</i></p>

<p>How will the data be backed up?</p> <p><i>WHAT STORAGE AND BACKUP PROCEDURES WILL BE IN PLACE TO PREVENT DATA LOSS?</i></p>	<p><input checked="" type="checkbox"/> Standard back-up provided by KU Leuven ICTS for my storage solution</p> <p><input checked="" type="checkbox"/> Personal back-ups I make (specify)</p> <p><input type="checkbox"/> Other (specify)</p> <p><i>external hard drive (as additional backup)</i></p>
<p>Is there currently sufficient storage & backup capacity during the project? If yes, specify concisely. If no or insufficient storage or backup capacities are available, then explain how this will be taken care of.</p>	<p><input checked="" type="checkbox"/> Yes</p> <p><input type="checkbox"/> No</p> <p><i>We pay yearly for storage space at VSC and KU Leuven.</i></p> <p>If no, please specify:</p>
<p>How will you ensure that the data are securely stored and not accessed or modified by unauthorized persons?</p> <p><i>CLEARLY DESCRIBE THE MEASURES (IN TERMS OF PHYSICAL SECURITY, NETWORK SECURITY, AND SECURITY OF COMPUTER SYSTEMS AND FILES) THAT WILL BE TAKEN TO ENSURE THAT STORED AND TRANSFERRED DATA ARE SAFE.</i></p> <p><u>Guidance on security for research data</u></p>	<p><i>secure login (2 factor authorization login)</i></p>
<p>What are the expected costs for data storage and backup during the research project? How will these costs be covered?</p>	<p><i>70 Euro per TB per year. These costs can be covered by our consumable costs.</i></p>

5. Data Preservation after the end of the Research Project

<p>Which data will be retained for at least five years (or longer, in agreement with other retention policies that are applicable) after the end of the project? In case some data cannot be preserved, clearly state the reasons for this (e.g. legal or contractual restrictions, storage/budget issues, institutional policies...).</p> <p>Guidance on data preservation</p>	<p><input checked="" type="checkbox"/> All data will be preserved for 10 years according to KU Leuven RDM policy</p> <p><input type="checkbox"/> All data will be preserved for 25 years according to CTC recommendations for clinical trials with medicinal products for human use and for clinical experiments on humans</p> <p><input type="checkbox"/> Certain data cannot be kept for 10 years (explain)</p> <p><i>Upon completion of the project, data is deposited at GEO (RNA-seq) or PRIDE (proteomics).</i></p>
<p>Where will these data be archived (stored and curated for the long-term)?</p> <p>Dedicated data repositories are often the best place to preserve your data. Data not suitable for preservation in a repository can be stored using a KU Leuven storage solution, consult the interactive KU Leuven storage guide.</p>	<p><input type="checkbox"/> KU Leuven RDR</p> <p><input type="checkbox"/> Large Volume Storage (longterm for large volumes)</p> <p><input type="checkbox"/> Shared network drive (J-drive)</p> <p><input type="checkbox"/> Other (specify):</p> <p><i>Upon completion of the project, data is deposited at GEO (RNA-seq) or PRIDE (proteomics).</i></p>
<p>What are the expected costs for data preservation during the expected retention period? How will these costs be covered?</p>	<p><i>Data storage at GEO and PRIDE is free.</i></p>

6. Data Sharing and Reuse

<p>Will the data (or part of the data) be made available for reuse after/during the project? Please explain per dataset or data type which data will be made available.</p> <p><i>NOTE THAT 'AVAILABLE' DOES NOT NECESSARILY MEAN THAT THE DATA SET BECOMES OPENLY AVAILABLE, CONDITIONS FOR ACCESS AND USE MAY APPLY. AVAILABILITY IN THIS QUESTION THUS ENTAILS BOTH OPEN & RESTRICTED ACCESS. FOR MORE INFORMATION:</i> https://wiki.surfnet.nl/display/STANDARDS/INFO-EU-REPO/#INFO-EURO-ACCESSRIGHTS</p>	<p><input checked="" type="checkbox"/> Yes, as open data <input type="checkbox"/> Yes, as embargoed data (temporary restriction) <input type="checkbox"/> Yes, as restricted data (upon approval, or institutional access only) <input type="checkbox"/> No (closed access) <input type="checkbox"/> Other, please specify:</p>
<p>If access is restricted, please specify who will be able to access the data and under what conditions.</p>	<p>NA</p>
<p>Are there any factors that restrict or prevent the sharing of (some of) the data (e.g. as defined in an agreement with a 3rd party, legal restrictions)? Please explain per dataset or data type where appropriate.</p>	<p><input type="checkbox"/> Yes, privacy aspects <input type="checkbox"/> Yes, intellectual property rights <input type="checkbox"/> Yes, ethical aspects <input type="checkbox"/> Yes, aspects of dual use <input type="checkbox"/> Yes, other <input checked="" type="checkbox"/> No</p> <p><i>We will make sure that IP rights are handled before the data is deposited.</i></p>
<p>Where will the data be made available? If already known, please provide a repository per dataset or data type.</p>	<p><input type="checkbox"/> KU Leuven RDR <input checked="" type="checkbox"/> Other data repository (specify) <input type="checkbox"/> Other (specify) <i>Upon completion of the project, data is deposited at GEO (RNA-seq) or PRIDE (proteomics).</i></p>

When will the data be made available?	<input checked="" type="checkbox"/> Upon publication of research results <input type="checkbox"/> Specific date (specify) <input type="checkbox"/> Other (specify) <i>Upon completion of the project, or earlier at each publication requiring the data deposit.</i>
Which data usage licenses are you going to provide? If none, please explain why. <i>A DATA USAGE LICENSE INDICATES WHETHER THE DATA CAN BE REUSED OR NOT AND UNDER WHAT CONDITIONS. IF NO LICENSE IS GRANTED, THE DATA ARE IN A GREY ZONE AND CANNOT BE LEGALLY REUSED. DO NOTE THAT YOU MAY ONLY RELEASE DATA UNDER A LICENCE CHOSEN BY YOURSELF IF IT DOES NOT ALREADY FALL UNDER ANOTHER LICENCE THAT MIGHT PROHIBIT THAT.</i> Check the RDR guidance on licences for data and software sources code or consult the License selector tool to help you choose.	<input type="checkbox"/> CC-BY 4.0 (data) <input type="checkbox"/> Data Transfer Agreement (restricted data) <input type="checkbox"/> MIT licence (code) <input type="checkbox"/> GNU GPL-3.0 (code) <input checked="" type="checkbox"/> Other (specify) <i>The dataset generated in this project are from mouse cells and will be made publicly available without license.</i>
Do you intend to add a PID/DOI/accession number to your dataset(s)? If already available, please provide it here. <i>INDICATE WHETHER YOU INTEND TO ADD A PERSISTENT AND UNIQUE IDENTIFIER IN ORDER TO IDENTIFY AND RETRIEVE THE DATA.</i>	<input checked="" type="checkbox"/> Yes, a PID will be added upon deposit in a data repository <input type="checkbox"/> My dataset already has a PID <input type="checkbox"/> No
What are the expected costs for data sharing? How will these costs be covered?	<i>No costs for datasharing.</i>

7. Responsibilities	
Who will manage data documentation and metadata during the research project?	Sofie Demeyer

Who will manage data storage and backup during the research project?	Sofie Demeyer
Who will manage data preservation and sharing?	Jan Cools and Alexandra Bacquelaine Veloso
Who will update and implement this DMP?	Alexandra Bacquelaine Veloso