

## **Materials Design Analysis Reporting (MDAR)**

### **Checklist for Authors**

The MDAR framework establishes a minimum set of requirements in transparent reporting applicable to studies in the life sciences (see Statement of Task: [doi:10.31222/osf.io/9sm4x](https://doi.org/10.31222/osf.io/9sm4x)). The MDAR checklist is a tool for authors, editors, and others seeking to adopt the MDAR framework for transparent reporting in manuscripts and other outputs. Please refer to the MDAR Elaboration Document for additional context for the MDAR framework.

**For all that apply, please note where in the manuscript the required information is provided.**

**Materials:**

<b>Newly created materials</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
The manuscript includes a dedicated "materials availability statement" providing transparent disclosure about availability of newly created materials including details on how materials can be accessed and describing any restrictions on access.	Information can be found in Section "Data and materials availability" in the main text.	
<b>Antibodies</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
For commercial reagents, provide supplier name, catalogue number and <a href="#">RRID</a> , if available.		n/a
<b>DNA and RNA sequences</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
<b>Short novel DNA or RNA including primers, probes:</b> Sequences should be included or deposited in a public repository.		n/a
<b>Cell materials</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
<b>Cell lines:</b> Provide species information, strain. Provide accession number in repository <b>OR</b> supplier name, catalog number, clone number, <b>OR</b> RRID.		n/a
<b>Primary cultures:</b> Provide species, strain, sex of origin, genetic modification status.		n/a
<b>Experimental animals</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
<b>Laboratory animals or Model organisms:</b> Provide species, strain, sex, age, genetic modification status. Provide accession number in repository <b>OR</b> supplier name, catalog number, clone number, <b>OR</b> RRID.		n/a
<b>Animal observed in or captured from the field:</b> Provide species, sex, and age where possible.		n/a
<b>Plants and microbes</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
<b>Plants:</b> provide species and strain, ecotype and cultivar where relevant, unique accession number if available, and source (including location for collected wild specimens).		n/a
<b>Microbes:</b> provide species and strain, unique accession number if available, and source.		n/a
<b>Human research participants</b>	<b>indicate where provided: section/legend; or state if these demographics were not collected</b>	<b>n/a</b>
If collected and within the bounds of privacy constraints report on age, sex and gender or ethnicity for all study participants.	Information can be found in Section "Study design" and Table 1 in the main text.	

**Design:**

<b>Study protocol</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
If study protocol has been pre-registered, provide DOI. For clinical trials, provide the trial registration number <b>OR</b> cite DOI.	Information can be found in Section “ <b>Ethics statement:</b> ” in the main text.	
<b>Laboratory protocol</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
Provide DOI <b>OR</b> other citation details if detailed step-by-step protocols are available.		n/a
<b>Experimental study design (statistics details)</b>		
<b>For in vivo studies:</b> State whether and how the following have been done	<b>indicate where provided: section/legend. If it could have been done, but was not, write not done</b>	<b>n/a</b>
Sample size determination		n/a
Randomisation		n/a
Blinding		n/a
Inclusion/exclusion criteria		n/a
<b>Sample definition and in-laboratory replication</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
State number of times the experiment was replicated in laboratory.		n/a
Define whether data describe technical or biological replicates.		n/a
<b>Ethics</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
<b>Studies involving human participants:</b> State details of authority granting ethics approval (IRB or equivalent committee(s), provide reference number for approval.	Information can be found in Section “ <b>Ethics statement:</b> ” in the main text.	
<b>Studies involving experimental animals:</b> State details of authority granting ethics approval (IRB or equivalent committee(s), provide reference number for approval.		n/a
<b>Studies involving specimen and field samples:</b> State if relevant permits obtained, provide details of authority approving study; if none were required, explain why.	Information can be found in Section “ <b>Ethics statement:</b> ” in the main text.	
<b>Dual Use Research of Concern (DURC)</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
If study is subject to dual use research of concern regulations, state the authority granting approval and reference number for the regulatory approval.		n/a

**Analysis:**

<b>Attrition</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
Describe whether exclusion criteria were preestablished. Report if sample or data points were omitted from analysis. If yes report if this was due to attrition or intentional exclusion and provide justification.	Information can be found in Section “ <b>2.Statistical analysis</b> ” in the supplementary Material.	
<b>Statistics</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
Describe statistical tests used and justify choice of tests.	Information can be found in Section “ <b>Statistical analysis</b> ” in the main text.	
<b>Data availability</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
For newly created and reused datasets, the manuscript includes a data availability statement that provides details for access or notes restrictions on access.	Information can be found in Section “ <b>Data and materials availability</b> ” in the main text.	
If newly created datasets are publicly available, provide accession number in repository <b>OR DOI OR URL</b> and licensing details where available.	Information can be found in Section “ <b>Data and materials availability</b> ” in the main text.	
If reused data is publicly available provide accession number in repository <b>OR DOI OR URL, OR citation.</b>	Information can be found in Section “ <b>Data and materials availability</b> ” in the main text.	
<b>Code availability</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
For all newly generated custom computer code/software/mathematical algorithm or re-used code essential for replicating the main findings of the study, the manuscript includes a data availability statement that provides details for access or notes restrictions.	Information can be found in Section “ <b>Data and materials availability</b> ” in the main text.	
If newly generated code is publicly available, provide accession number in repository, <b>OR DOI OR URL</b> and licensing details where available. State any restrictions on code availability or accessibility.	Information can be found in Section “ <b>Data and materials availability</b> ” in the main text.	
If reused code is publicly available provide accession number in repository <b>OR DOI OR URL, OR citation.</b>	Information can be found in Section “ <b>Data and materials availability</b> ” in the main text.	

## **Reporting**

MDAR framework recommends adoption of discipline-specific guidelines, established and endorsed through community initiatives. Journals have their own policy about requiring specific guidelines and recommendations to complement MDAR.

<b>Adherence to community standards</b>	<b>indicate where provided: section/legend</b>	<b>n/a</b>
State if relevant guidelines (e.g., ICMJE, MIBBI, ARRIVE) have been followed, and whether a checklist (e.g., CONSORT, PRISMA, ARRIVE) is provided with the manuscript.		n/a