# <u>Materials Design Analysis Reporting (MDAR)</u> 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.). 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:**

Newly created materials	indicate where provided: page no/section/legend)	n/a
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.	Yes; main text acknowledgements section "Data and materials availability"	
Antibodies	indicate where provided page to (section (legand)	- 1a
For commercial reagents, provide supplier name,	indicate where provided: page no/section/legend)	n/a
catalogue number and <u>RRID</u> , if available.		n/a
DNA and RNA sequences	indicate where provided: page no/section/legend)	n/a
Short novel DNA or RNA including primers, probes: Sequences should be included or deposited in a public repository.	Yes; main text acknowledgements section "Data and materials availability"	
Cell materials	indicate where provided: page no/section/legend	n/a
Cell lines: Provide species information, strain.	, , , , , , , , , , , , , , , , , , , ,	
Provide accession number in repository <b>OR</b> supplier		
name, catalog number, clone number, <b>OR</b> RRID.		n/a
Primary cultures: Provide species, strain, sex of		
origin, genetic modification status.		n/a
Experimental animals	indicate where provided: page no/section/legend)	n/a
Laboratory animals or Model organisms: Provide		
species, strain, sex, age, genetic modification status.		
Provide accession number in repository <b>OR</b> supplier		n/a
name, catalog number, clone number, <b>OR</b> RRID.		
Animal observed in or captured from the field:	Voc. first main tout costion (untitled) main tout costion	
Provide species, sex, and age where possible.	Yes; first main text section (untitled), main text section "Structure of the baboon hybrid zone", and table s1	
Plants and microbes	indicate where provided: page no/section/legend)	n/a
Plants: provide species and strain, ecotype and	- Francis Provided Page 110/ 2000/01/ 10Bella/	, ~
cultivar where relevant, unique accession number if		
available, and source (including location for collected		n/a
wild specimens).		
Microbes: provide species and strain, unique		n/2
accession number if available, and source.		n/a
II.	indicate where provided: page no/section/legend) or	,
Human research participants	state if these demographics were not collected	n/a
If collected and within the bounds of privacy		<u> </u>
constraints report on age, sex and gender or		n/a
ethnicity for all study participants.		

## Design:

Study protocol	indicate where provided: page no/section/legend)	n/a
If study protocol has been pre-registered, provide DOI. For clinical trials, provide the trial registration number <b>OR</b> cite DOI.		n/a

Laboratory protocol	indicate where provided: page no/section/legend)	n/a
Provide DOI <b>OR</b> other citation details if detailed step- by-step protocols are available.		n/a

Experimental study design (statistics details)		
For in vivo studies: State whether and how the	indicate where provided: page no/section/legend. If it	2/2
following have been done	could have been done, but was not, write not done	n/a
Sample size determination	Yes; Supplementary Materials Section 1.2 and 1.3	
Randomisation		n/a
Blinding		n/a
Inclusion/exclusion criteria	Yes; Supplementary Materials Section 1.2 and 1.3	

Sample definition and in-laboratory replication	indicate where provided: page no/section/legend	n/a
State number of times the experiment was		n/a
replicated in laboratory.		II/a
Define whether data describe technical or biological		n/a
replicates.		11/ d

Ethics	indicate where provided: page no/section/legend	n/a
Studies involving human participants: State details of authority granting ethics approval (IRB or equivalent committee(s), provide reference number for approval.		n/a
Studies involving experimental animals: State details of authority granting ethics approval (IRB or equivalent committee(s), provide reference number for approval.		n/a
Studies involving specimen and field samples: State if relevant permits obtained, provide details of authority approving study; if none were required, explain why.	Yes; main text acknowledgements (untitled first section) and Supplementary Materials Section 1.2	

Dual Use Research of Concern (DURC)	indicate where provided: page no/section/legend	n/a
If study is subject to dual use research of concern		
regulations, state the authority granting approval		n/a
and reference number for the regulatory approval.		

## **Analysis:**

Attrition	indicate where provided: page no/section/legend	n/a
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.	Yes; Supplementary Materials section 1.2, 5	

Statistics	indicate where provided: page no/section/legend	n/a
Describe statistical tests used and justify choice of tests.	Yes; main text all sections, with justification in the main text and supplementary materials	

Data availability	indicate where provided: page no/section/legend	n/a
For newly created and reused datasets, the manuscript includes a data availability statement that provides details for access or notes restrictions on access.	Yes; main text acknowledgements section "Data and materials availability"	
If newly created datasets are publicly available, provide accession number in repository <b>OR</b> DOI <b>OR</b> URL and licensing details where available.	Yes; main text acknowledgements section "Data and materials availability" and table s1 list accession numbers for genomic data; main text acknowledgements section "Data and materials availability" also provides Zenodo DOI for analyzed data	
If reused data is publicly available provide accession number in repository <b>OR</b> DOI <b>OR</b> URL, <b>OR</b> citation.	Yes; main text acknowledgements section "Data and materials availability" and table s1 list accession numbers and publicly available data is also cited in the References and Notes section	

Code availability	indicate where provided: page no/section/legend	n/a
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.	Yes; main text acknowledgements section "Data and materials availability" provides Zenodo DOI for custom scripts	
If newly generated code is publicly available, provide accession number in repository, <b>OR</b> DOI <b>OR</b> URL and licensing details where available. State any restrictions on code availability or accessibility.	Yes; main text acknowledgements section "Data and materials availability" provides Zenodo DOI for custom scripts	
If reused code is publicly available provide accession number in repository <b>OR</b> DOI <b>OR</b> URL, <b>OR</b> citation.	Yes; all Supplementary Materials sections contain citations to publicly available code	

## 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.

Adherence to community standards	indicate where provided: page no/section/legend	n/a
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.	Yes; this MDAR checklist	