<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.

Materials

| Antibodies | Yes (indicate where provided: section/paragraph) | n/a |
|--|--|-----|
| For commercial reagents, provide supplier | Materials and methods/paragraph10-13 | |
| name, catalogue number and RRID, if available. | | |
| | | |
| Cell materials | Yes (indicate where provided: section/paragraph) | n/a |
| Cell lines: Provide species information, strain. | Materials and methods/paragraph10 | |
| Provide accession number in repository OR | | |
| supplier name, catalog number, clone number, | | |
| OR RRID | | |
| Primary cultures: Provide species, strain, sex of | | N/A |
| origin, genetic modification status. | | |
| | | |
| Experimental animals | Yes (indicate where provided: section/paragraph) | n/a |
| Laboratory animals: Provide species, strain, sex, age, | | |
| genetic modification status. Provide accession | | N/A |
| number in repository OR supplier name, catalog | | |
| number, clone number, OR RRID | | |
| Animal observed in or captured from the | | |
| field: Provide species, sex and age where | | N/A |
| possible | | |
| Model organisms: Provide Accession number | | N/A |
| in repository (where relevant) OR RRID | | |
| | | |
| Plants and microbes | Yes (indicate where provided: section/paragraph) | n/a |
| Plants: provide species and strain, unique accession | | |
| number if available, and source (including location | | N/A |
| for collected wild specimens) | | |
| Microbes: provide species and strain, unique | | |
| accession number if available, and source | | N/A |
| | · | |
| Human research participants | Yes (indicate where provided: section/paragraph) | n/a |
| Identify authority granting ethics approval (IRB or | | N/A |
| equivalent committee(s), provide reference number | | |
| for approval. | | |
| Provide statement confirming informed consent | | N/A |
| obtained from study participants. | | |
| Report on age and sex for all study participants. | | N/A |

<u>Design</u>

| Study protocol | Yes (indicate where provided: section/paragraph) | n/a |
|--|---|------------|
| For clinical trials, provide the trial registration number OR cite DOI in manuscript. | | N/A |
| Laboratory protocol Provide DOI or other citation details if detailed step- | Yes (indicate where provided: section/paragraph) | n/a N/A |
| by-step protocols are available. | | |
| Experimental study design (statistics details) | Yes (indicate where provided: section/paragraph) | n/a |
| State whether and how the following have been done, or if they were not carried out. | Yes | |
| Sample size determination | Materials and methods/paragraph1 | |
| Randomisation | | N/A |
| Blinding | Matarials and matheds/paragraph1.14 | N/A |
| | Materials and methous/paragraph1,14 | |
| Sample definition and in-laboratory replication | Yes (indicate where provided: section/paragraph) | n/a |
| State number of times the experiment was replicated in laboratory | Materials and methods/paragraph11-13 | |
| Define whether data describe technical or biological replicates | Materials and methods/paragraph11-13 | |
| Ethics | Yes (indicate where provided: section/paragraph) | n/a |
| Studies involving human participants: State details of authority granting ethics approval (IRB or equivalent committee(s), provide reference number for approval. | First Affiliated Hospital, Hainan Medical College, Haikou 570216, China . Number for approval : 2023-KYS-161 | |
| 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. | | N/A |
| Dual Use Research of Concern (DURC) | Yes (indicate where provided: section/paragraph) | n/a |
| If study is subject to dual use research of concern, state the authority granting approval and reference number for the regulatory approval | | N/A |

Analysis

| Attrition | Yes (indicate where provided: section/paragraph) | n/a |
|---|--|-------|
| State if sample or data point from the analysis is excluded, and whether the criteria for exclusion were determined and specified in advance. | Materials and methods/paragraph14 | |
| | | |
| Statistics | Yes (indicate where provided: section/paragraph) | n/a |
| Describe statistical tests used and justify choice of tests. | Materials and methods/paragraph14 | |
| Data Availability | Yes (indicate where provided: section/paragraph) | n/a |
| State whether newly created datasets are available | res (maleate where provided, section, paragraph, | 11/ 4 |
| including protocols for access or restriction on | | NI/A |
| | | N/A |
| | | |
| number in repository or DOI or URL. | URL: https://xenabrowser.net/datapages/ | |
| If publicly available data are reused, provide | URL: https://xenabrowser.net/datapages/ | |
| accession number in repository or DOI or URL, where | | |
| possible. | | |
| | | |
| Code Availability | Yes (indicate where provided: section/paragraph) | n/a |
| For all newly generated code and software essential | Yes | |
| for replicating the main findings of the study: | | |
| State whether the code or software is available. | Yes | |
| If code is publicly available, provide accession | DOI: 10.1093/bioinformatics/btab561 | |
| number in repository, or DOI or URL. | | |

Reporting

| Adherence to community standards | Yes (indicate where provided: section/paragraph) | n/a |
|---|--|-----|
| MDAR framework recommends adoption of discipline-specific guidelines, established and | Yes | |
| endorsed through community initiatives. Journals | | |
| have their own policy about requiring specific | | |
| guidelines and recommendations to complement | | |
| MDAR. | | |
| State if relevant guidelines (eg., ICMJE, MIBBI, | ICMJE guidelines were followed, as the journal follows | |
| ARRIVE) have been followed, and whether a checklist | ICMJE recommendations for publication. | |
| (eg., CONSORT, PRISMA, ARRIVE) is provided with | | |
| the manuscript. | | |

Article information: https://dx.doi.org/10.21037/tcr-23-1744

The REMARK checklist

| | | Reported on Page | Reported on |
|------------------------------|--|-----------------------|-------------------|
| Item to be reported | | Number/Line Number | Section/Paragraph |
| INTR | ODUCTION | | |
| 1 | State the marker examined, the study objectives, and any pre-specified hypotheses. | | |
| MAT | ERIALS AND METHODS | | |
| Patie | nts | | |
| 2 | Describe the characteristics (e.g., disease stage or co-morbidities) of the study patients, including their source and inclusion and exclusion criteria. | | |
| 3 | Describe treatments received and how chosen (e.g., randomized or rule-based). | | |
| Spec | imen characteristics | | |
| 4 | Describe type of biological material used (including control samples) and methods of preservation and storage. | | |
| Assa | y methods | | |
| 5 | Specify the assay method used and provide (or reference) a detailed protocol, including specific reagents or kits used, quality control procedures, reproducibility assessments, quantitation methods, and scoring and reporting protocols. Specify whether and how assays were performed blinded to the study endpoint. | | |
| Study | y design | | |
| 6 | State the method of case selection, including whether prospective or retrospective and whether stratification or matching (e.g., by stage of disease or age) was used. Specify the time period from which cases were taken, the end of the follow-up period, and the median follow-up time. | | |
| 7 | Precisely define all clinical endpoints examined. | | |
| 8 | List all candidate variables initially examined or considered for inclusion in models. | | |
| 9 | Give rationale for sample size; if the study was designed to detect a specified effect size, give the target power and effect size. | | |
| Statistical analysis methods | | | |
| 10 | Specify all statistical methods, including details of any variable selection procedures and other model-building issues, how model assumptions were verified, and how missing data were handled. | | |
| 11 | Clarify how marker values were handled in the analyses; if relevant, describe methods used for cutpoint determination. | | |

| RESULTS | | | |
|---------------------------|---|--|--|
| Data | Data | | |
| 12 | Describe the flow of patients through the study, including the number of patients included in each stage of the analysis (a diagram may be helpful) and reasons for dropout. Specifically, both overall and for each subgroup extensively examined report the numbers of patients and the number of events. | | |
| 13 | Report distributions of basic demographic characteristics (at least age and sex), standard (disease-specific) prognostic variables, and tumor marker, including numbers of missing values. | | |
| Analysis and presentation | | | |
| 14 | Show the relation of the marker to standard prognostic variables. | | |
| 15 | Present univariable analyses showing the relation between the marker and outcome, with the estimated effect (e.g., hazard ratio and survival probability). Preferably provide similar analyses for all other variables being analyzed. For the effect of a tumor marker on a time-to-event outcome, a Kaplan-Meier plot is recommended. | | |
| 16 | For key multivariable analyses, report estimated effects (e.g., hazard ratio) with confidence intervals for the marker and, at least for the final model, all other variables in the model. | | |
| 17 | Among reported results, provide estimated effects with confidence intervals from an analysis in which the marker and standard prognostic variables are included, regardless of their statistical significance. | | |
| 18 | If done, report results of further investigations, such as checking assumptions, sensitivity analyses, and internal validation. | | |
| DISCUSSION | | | |
| 19 | Interpret the results in the context of the pre-specified hypotheses and other relevant studies; include a discussion of limitations of the study. | | |
| 20 | Discuss implications for future research and clinical value. | | |

From: McShane LM, Altman DG, Sauerbrei W, Taube SE, Gion M, Clark GM: Reporting recommendations for tumor marker prognostic studies (REMARK). J Natl Cancer Inst 2005; 97: 1180-1184.