Corresponding Author Name:_____ Manuscript Number:_____



Reporting Checklist

This checklist is used to ensure good reporting standards and to improve the reproducibility of published results. **Please respond completely to all questions relevant to your manuscript.** For more information, please read the journal's Guide to Authors.

□ Check here to confirm that the following information is available in the Material & Methods section:

- the exact sample size (n) for each experimental group/condition, given as a number, not a range;
- a description of the sample collection allowing the reader to understand whether the samples represent technical or biological replicates (including how many animals, litters, culture, etc.);
- a statement of how many times the experiment shown was replicated in the laboratory;
- **definitions of statistical methods and measures**: (For small sample sizes (n<5) descriptive statistics are not appropriate, instead plot individual data points)
 - \circ very common tests, such as *t*-test, simple χ^2 tests, Wilcoxon and Mann-Whitney tests, can be unambiguously identified by name only, but more complex techniques should be described in the methods section;
 - are tests one-sided or two-sided?
 - are there adjustments for multiple comparisons?
 - statistical test results, e.g., P values;
 - definition of 'center values' as median or mean;
 - o definition of error bars as s.d. or s.e.m. or c.i.

Please ensure that the answers to the following questions are reported **in the manuscript itself.** We encourage you to include a specific subsection in the methods section for statistics, reagents and animal models. Below, provide the page number or section and paragraph number.

Statistics and general methods

- How was the sample size chosen to ensure adequate power to detect a pre-specified effect size? (Give section/paragraph or page #)
- For animal studies, include a statement about sample size estimate even if no statistical methods were used.
- Describe inclusion/exclusion criteria if samples or animals were excluded from the analysis. Were the criteria pre-established? (Give section/paragraph or page #)
- If a method of randomization was used to determine how samples/animals were allocated to experimental groups and processed, describe it. (Give section/paragraph or page #)
- For animal studies, include a statement about randomization even if no randomization was used.



- If the investigator was blinded to the group allocation during the experiment and/or when assessing the outcome, state the extent of blinding. (Give section/paragraph or page #)
- For animal studies, include a statement about blinding even if no blinding was done.
- 5. For every figure, are statistical tests justified as appropriate?
- Do the data meet the assumptions of the tests (e.g., normal distribution)?
- Is there an estimate of variation within each group of data?
- Is the variance similar between the groups that are being statistically compared? (Give section/paragraph or page #)

Reagents

- 6. Report the source of antibodies (vendor and catalog number)
- Identify the source of cell lines and report if they were recently authenticated (e.g., by STR profiling) and tested for mycoplasma contamination

Animal Models

- 8. Report species, strain, sex and age of animals
- For experiments involving live vertebrates, include a statement of compliance with ethical regulations and identify the committee(s) approving the experiments.
- 10. We recommend consulting the ARRIVE guidelines (*PLoS Biol.* **8**(6), e1000412,2010) to ensure that other relevant aspects of animal studies are adequately reported.

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Human subjects

- 11. Identify the committee(s) approving the study protocol.
- 12. Include a statement confirming that informed consent was obtained from all subjects.
- 13. For publication of patient photos, include a statement confirming that consent to publish was obtained.
- 14. Report the clinical trial registration number (at <u>ClinicalTrials.gov</u> or equivalent).
- 15. For phase II and III randomized controlled trials, please refer to the <u>CONSORT statement</u> and submit the CONSORT checklist with your submission.
- 16. For tumor marker prognostic studies, we recommend that you follow the **REMARK reporting guidelines**.

Data deposition

- Provide accession codes for deposited data. Data deposition in a public repository is mandatory for:
 - a. Protein, DNA and RNA sequences
 - b. Macromolecular structures
 - c. Crystallographic data for small molecules
 - d. Microarray data

Reported in section/paragraph or page

Deposition is strongly recommended for many other datasets for which structured public repositories exist; more details on our data policy are available in the Guide to Authors. We encourage the provision of other source data in supplementary information or in unstructured repositories such as <u>Figshare</u> and <u>Dryad</u>. We encourage publication of Data Descriptors (see <u>Scientific Data</u>) to maximize data reuse.

18. If computer code was used to generate results that are central to the paper's conclusions, include a statement in the Methods section under "Code availability" to indicate whether and how the code can be accessed. Include version information as necessary and any restrictions on availability.



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