nature portfolio

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Reporting Summary

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

Statistics

For	all st	atistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Cor	firmed
	×	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	X	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	×	The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.
	×	A description of all covariates tested
	X	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	×	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)
	×	For null hypothesis testing, the test statistic (e.g. F, t, r) with confidence intervals, effect sizes, degrees of freedom and P value noted Give P values as exact values whenever suitable.
X		For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
×		For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
X		Estimates of effect sizes (e.g. Cohen's d, Pearson's r), indicating how they were calculated
		Our web collection on <u>statistics for biologists</u> contains articles on many of the points above.

Software and code

Policy information about availability of computer code					
Data collection	N/A				
Data analysis	N/A				

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

The authors declare that all data supporting the findings of this study are available in the article, its Extended Data, its Source Data or from the corresponding authors upon request.

Research involving human participants, their data, or biological material

Policy information about studies with <u>human participants or human data</u>. See also policy information about <u>sex, gender (identity/presentation)</u>, <u>and sexual orientation</u> and <u>race, ethnicity and racism</u>.

Reporting on sex and gender	(N/A
Reporting on race, ethnicity, or other socially relevant groupings	N/A
Population characteristics	N/A
Recruitment	N/A, the data from a previous clinical study was re-used in the present study.
Ethios oversight	The multi-contex study of SCAD was approved by the Duilin Llespital Ethics Committee normit number 2017, 190 with
Ethics oversight	The multi-center study of SCAP was approved by the Ruijin Hospital Ethics Committee permit number 2017-186 with patients' consent. All experiments were performed and information released according to the instructions of this permit.

Note that full information on the approval of the study protocol must also be provided in the manuscript.

Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

▼ Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size	For biological replicates, three or more independent experiments were performed.
Data exclusions	N/A
Replication	All replications were successful.
Randomization	N/A
Blinding	(N/A

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

Materials & experimental systems			Methods	
n/a	Involved in the study	n/a	Involved in the study	
	X Antibodies	×	ChIP-seq	
	X Eukaryotic cell lines	×	Flow cytometry	
×	Palaeontology and archaeology	×	MRI-based neuroimaging	
×	Animals and other organisms			
	X Clinical data			
×	Dual use research of concern			
×	Plants			

Antibodies

Antibodies used Anti-myc (CST, Cat. No. 2272S), anti-HA (CST, Cat. No. 3724S), anti-beta-actin (CST, Cat. No. 5125S), Anti-ALCAM (SinoBiological, Cat. No. 80221-RP02), Anti-CAR (Abcam, Cat. No. ab272711), Anti-DSG-2(CST, Cat. No. 88970S)

Validation

All antibodies have been validated for intended use (western blotting)

Eukaryotic cell lines

Policy information about cell lines and Sex and Gender in Research

Cell line source(s)	HEK-293A (Procell, No. CL-0003), HEK-293T (SIBS, Cat. No. SCSP-502), CHO-K1 cells (SIBS, Cat. No. MD12), HeLa (ATCC, CRL-1958), U-87MG (ATCC, HTB-14), A549 (ATCC, CCL-185)
Authentication	All cells used in the present study were validated by STR methods
Mycoplasma contamination	The cells used in the present study were free of mycoplasma contamination
Commonly misidentified lines (See <u>ICLAC</u> register)	N/A

Clinical data

Policy information about <u>clinical studies</u>

All manuscripts should comply with the ICMJEguidelines for publication of clinical research and a completed CONSORT checklist must be included with all submissions.			
Clinical trial registration	N/A		
Study protocol	N/A		
Data collection	N/A		
Outcomes	N/A		

Plants

Seed stocks	N/A
Novel plant genotypes	N/A
Authentication	N/A