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Reporting Summary

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When statistical analyses are reported, confirm that the following items are present in the relevant location (e.g. figure legend, table legend, main

Statistical parameters

text, or Methods section).			
n/a	n/a Confirmed		
	The exact sample size (n) for each expe	rimental group/condition, given as a discrete number and unit of measurement	
	An indication of whether measurement	s were taken from distinct samples or whether the same sample was measured repeatedly	
	The statistical test(s) used AND whethe Only common tests should be described sole.	r they are one- or two-sided ly by name; describe more complex techniques in the Methods section.	
X	A description of all covariates tested		
X	A description of any assumptions or con	rections, such as tests of normality and adjustment for multiple comparisons	
\boxtimes	V	ing <u>central tendency</u> (e.g. means) or other basic estimates (e.g. regression coefficient) AND sociated <u>estimates of uncertainty</u> (e.g. confidence intervals)	
	For null hypothesis testing, the test state Give P values as exact values whenever suita	cistic (e.g. F , t , r) with confidence intervals, effect sizes, degrees of freedom and P value noted ble.	
\boxtimes	For Bayesian analysis, information on the	ne choice of priors and Markov chain Monte Carlo settings	
X	For hierarchical and complex designs, id	dentification of the appropriate level for tests and full reporting of outcomes	
\boxtimes	Estimates of effect sizes (e.g. Cohen's d	, Pearson's \emph{r}), indicating how they were calculated	
	Clearly defined error bars State explicitly what error bars represent (e.s.	g. SD, SE, CI)	
Our web collection on <u>statistics for biologists</u> may be useful.			

Software and code

Policy information about $\underline{availability}$ of $\underline{computer}$ \underline{code}

Data collection

Custom written LabView 2015 microscope control software

Data analysis

Custom written Matlab 2018a,b scripts. Code is available from the corresponding authors on request. Statistical analysis was performed using the software Origin (version 8.5.6; OriginLab)

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/reviewers upon request. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.

Data

Policy information about availability of data

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

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

The datasets generated and analysed in the current study are available from the corresponding authors upon reasonable request. Raw data for Fig. 3c and 4c are presented in Supplementary Data 1 and 2, respectively.

Field-specific reporting			
Please select the be	est 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 t	the document with all sections, see nature.com/authors/policies/ReportingSummary-flat.pdf		
Life scier	nces study design		
All studies must disclose on these points even when the disclosure is negative.			
Sample size	No particular statistical method was used to define sample size. Sample size was determined based on previous studies in the field. A minimum of 3 independent experiments were carried out.		
Data exclusions	Images with a too low signal to noise ratio of the Brillouin shift and width were excluded. Representative data sets were chosen for the Figures.		
Replication	Experiments were reliably reproduced. At least 3 independent experiments were carried out. Experiments were performed on different days with different batches of ovaries.		
Randomization	No randomization was applied.		
Blinding	Authors were blinded to any group allocation since the mice were chosen randomly for the study.		
Materials & experimental systems Methods N/a Involved in the study N/a Involved in th			
Antibodies used	Primary pAb Anti-Collagen IV (Millipore, AB756P) used at 1:100. Secondary antibodies targeting rabbit immunoglobulin-coupled Alexa Fluor 546 (Invitrogen, A10040) was used at 1:200.		
Validation	C Anderson, et al. 'Sonic hedgehog-dependent synthesis of laminin alpha1 controls basement membrane assembly in the myotome', Development 136(20):3495-504, 2009		
Animals and	other organisms		
Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research			
Laboratory anima	Embryonic and postnatal mice ovaries were obtained in the animal facility at the European Molecular Biology Laboratory, with permission from the institutional veterinarian overseeing the operation (ARC number 2020-01-06RP). The animal facilities are operated according to international animal welfare rules (Federation for Laboratory Animal Science Associations guidelines and recommendations).		

The study did not involve wild animals.

The study did not involve field-collected samples.

Wild animals

Field-collected samples