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Reporting Summary

Nature Research 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 Research policies, see Authors & Referees and the Editorial Policy Checklist.

Statistics			
For all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
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The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
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			
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. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>			
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			
Estimates of effect sizes (e.g. Cohen's <i>d</i> , Pearson's <i>r</i>), 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 <u>availability of computer code</u>			
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/reviewers. 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 data availability statement. 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 that support the findings of this study are available in GEO Database [https://www.ncbi.nlm.nih.gov/geo/query/acc.cgi?acc=GSE124106]. They have been assigned GEO accession numbers GSE 124105 and GSE 124020. Further data that support the findings of this study are available within the article and Supplementary Files. All other data, including raw data used in each figure will be provided upon reasonable request to the corresponding author

Field-spe	cific reporting			
Please select the or	e 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 t	ne document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf			
Life scier	ces study design			
All studies must dis	close on these points even when the disclosure is negative.			
Sample size	All experiments were performed at least in triplicates in order to perform statistical tests on the experiments.			
Data exclusions	n.a.			
Replication	All experiments were performed at least in triplicates in order to perform statistical tests on the experiments.			
Randomization	n.a.			
Blinding	n.a.			
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Human rese	arch participants			
Policy information a	about studies involving human research participants			
Population chara	eteristics n.a.			

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

n.a.

n.a.

Recruitment

Ethics oversight

Flow Cytometry

Plots

Confirm that:

\times	The axis	labels state	the marker	and fluorochrome	used (e.g.	CD4-FITC)
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The axis scales are clearly visible. Include numbers along axes only for bottom left plot of group (a 'group' is an analysis of identical markers).

 ${\begin{tabular}{|c|c|c|c|c|c|}}$ All plots are contour plots with outliers or pseudocolor plots.

🔀 A numerical value for number of cells or percentage (with statistics) is provided.

Methodology

Sample preparation	detail within manuscript
Instrument	detail within manuscript
Software	detail within manuscript
Cell population abundance	detail within manuscript
Gating strategy	detail within manuscript

Tick this box to confirm that a figure exemplifying the gating strategy is provided in the Supplementary Information.