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>.

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St	at	isti	CS

For	all statistical ar	nalyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.	
n/a	Confirmed		
	The exact	sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement	
	A stateme	ent on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly	
	The statis Only comn	tical test(s) used AND whether they are one- or two-sided non tests should be described solely by name; describe more complex techniques in the Methods section.	
\boxtimes	A descrip	tion of all covariates tested	
	A descrip	tion 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)		
\boxtimes	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 Give <i>P</i> values as exact values whenever suitable.		
\boxtimes	For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings		
\boxtimes	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes		
\square Estimates of effect sizes (e.g. Cohen's d , Pearson's r), indicating how they were calculated			
	'	Our web collection on statistics for biologists contains articles on many of the points above.	
So	ftware an	d code	
Poli	cy information	about <u>availability of computer code</u>	
Da	ata collection	No Software was used.	
Da	ata analysis	Statistical tests were performed using GraphPad Prism version 10 or higher.	
		g 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 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

No data with mandated deposition was generated as a result of this study. Other data included in this study will be made available upon request to amcguire@fredhutch.org.

Research in	volving hu	man participants, their data, or biological material
,		with

Flow cytometry

MRI-based neuroimaging

Eukaryotic cell lines Palaeontology and archaeology

Animals and other organisms

Clinical data
Dual use research of concern

| | Plants

Antibodies

Antibodies used

anti-mouse PE secondary antibody (BioLegend Cat. #405307), rabbit anti-His tag antibody (SigmaAldrich Cat. #SAB5600227), anti-mouse IgG-HRP (SouthernBiotech Cat. #2010-05), goat anti-human IgG-HRP (Jackson ImmunoResearch Cat. # 109-035-088), anti-CD3 (ThermoFisher Cat. #14-0037-82), anti-CD28 (ThermoFisher Cat. #14-0281-82), CD45 BUV805 (BD Bioscience Cat. #568336), CD3

BUV395 (BD Bioscience Cat. #740268), CD8 BUV737 (BD Bioscience Cat. #612759), and CD4 PerCPCy5.5 (Thermofisher Cat. #45-0042-80), anti-mouse IFN-γ AF488 (Biolegend Cat. #505815), hCD45 FITC (eBioscience Cat. #5010066), mCD45 APC (eBioscience Cat. #17-0451-82), h, hCD33 PE (BD Bioscience Cat. #555450), hCD19 BV711 (Biolegend Cat.# 302246), hCD4 AF700 (eBioscience Cat. #56-0048-82), hCD8 BV421 (BD Bioscience Cat. #562429).

Validation

Antibodies were used in accordance with the manufactuer's website, assuming species and target specificity was validated by manufacturer.

Animals and other research organisms

Policy information about <u>studies involving animals</u>; <u>ARRIVE guidelines</u> recommended for reporting animal research, and <u>Sex and Gender in Research</u>

Comparative immunogenicity studies were performed in groups of 4 or 5 C57BL/6 mice between 7 and 10 weeks of age. Mice were purchased from the Jackson Laboratory. Six week old NSG female mice were used for humanized mouse studies.

Wild animals

The study did not use wild animals.

Equal numbers of male and female mice were used for comparative immunogenicity studies. Female NSG mice were used for huCD34 engraftment. Human serum was randomly selected for assays with no consideration for sex. Data was not analyzed with considerations for sex.

Field-collected samples The study did not use field collected samples.

Ethics oversight Fred Hutch Cancer Center Institutional Animal Care and Use Committee .

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

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

Flow Cytometry

Plots

Confirm that:

The axis labels state the marker and fluorochrome used (e.g. CD4-FITC).

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).

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	Cultured cells were transfected and stained with flourescent antibodies or infected with fluorescent reporter virus.
Instrument	BD FACSCelesta Special Order Research Product, Luminex Guava HT
Software	FlowJo v10.8 Software (BD Life Sciences).
Cell population abundance	No sorting was performed.
Gating strategy	Gating strategy is provided in supplementary information

DRMC from whole blood collected from humanized mice. Colonecutes were markenically disrupted from CE7/DIC mice

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