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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 our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

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For	all statistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.
n/a	Confirmed
	\square 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
\boxtimes	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.
\boxtimes	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)
\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 <i>Give P values as exact values whenever suitable.</i>
\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
\boxtimes	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 statistics for biologists contains articles on many of the points above.

Software and code

Policy information about availability of computer code

Data collection

We used J-Link RTT Viewer (V6.62a) for collecting the raw magnetic field values from the iMAG pills and the nRF receiver. We used MATLAB (R2020a, Mathworks) for position decoding using the field values. We used Arduino Mega 2560 and the Arduino IDE for FOV characterization and LUT creation. All custom codes will be available upon request to authors.

Data analysis

We used MATLAB (R2020a, Mathworks) and GraphPad Prism (9.4.1) for data analysis and plotting. Illustrations were made in Adobe Illustrator (26.4.1) and Microsoft PowerPoint.

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 Research guidelines for submitting code & software for further information.

Data

Policy information about <u>availability of data</u>

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

All data supporting the findings of this study are available within the manuscript and its supplementary material. Raw data values for any of the experiments reported in the manuscript are available upon reasonable request from the corresponding authors.

Field-spe	ecific reporting				
Please select the or	ne below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.				
Life sciences	Life sciences Behavioural & social sciences Ecological, evolutionary & environmental sciences				
For a reference copy of t	the document with all sections, see <u>nature.com/documents/nr-reporting-summary-flat.pdf</u>				
Life scier	nces study design				
All studies must dis	sclose on these points even when the disclosure is negative.				
Sample size	Sample sizes were chosen based on preliminary experiments and our prior work with ingestible devices, so as to provide sufficient power for statistical comparison (where appropriate).				
Data exclusions	o data values were excluded from the analyses.				
Replication	All replicates reported in the manuscript. Multiple tests were taken on different days and using different animals to ensure that the data was reproducible.				
Randomization	randomization of animals. Not applicable to these studies.				
Blinding	Blinding was not applicable to our study as no control vs. experimental groups were used for these studies. All data collection, processing, and analysis methods were quantitative and identical across experimental groups.				
Reportin	g for specific materials, systems and methods				
'	on from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each materia ted 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 & exp	perimental systems Methods				
n/a Involved in th	n/a Involved in the study n/a Involved in the study				
Antibodies ChIP-seq					
Eukaryotic cell lines Flow cytometry					
Palaeontology and archaeology MRI-based neuroimaging					
Clinical dat					
Dual use research of concern					
Animals and	other organisms				
Policy information	about studies involving animals; ARRIVE guidelines recommended for reporting animal research				
Laboratory anima	boratory animals Female Yorkshire Swine aged 3-5 months (35-70 kg).				

Wild animals

No wild animals were used.

Field-collected samples No field-collected samples were used.

Ethics oversight Committee for Animal Care of the Massachusetts Institute of Technology (MIT) for animal experiments.

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