

Life Sciences Reporting Summary

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► Experimental design

1. Sample size

Describe how sample size was determined.

The appropriate sample size of the behavioural and vibration experiments was determined from the results of pilot experiments.

2. Data exclusions

Describe any data exclusions.

No data was excluded.

3. Replication

Describe whether the experimental findings were reliably reproduced.

The behavioural results were replicated within the paper and have since been replicated in several independent experiments (unpublished). The vibration results were replicated between the individual participants in the paper.

4. Randomization

Describe how samples/organisms/participants were allocated into experimental groups.

The order of sensing modes in Behavioural Experiment 1 was randomly assigned to each participant.

5. Blinding

Describe whether the investigators were blinded to group allocation during data collection and/or analysis.

Blinding to experimental condition was not possible when running the behavioural experiments. However, as the experimenter was blind to the judgments made by participants, this would not influence the results.

Note: all studies involving animals and/or human research participants must disclose whether blinding and randomization were used.

6. Statistical parameters

For all figures and tables that use statistical methods, confirm that the following items are present in relevant figure legends (or in the Methods section if additional space is needed).

n/a Confirmed

- ☐ ☒ The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement (animals, litters, cultures, etc.)
- ☐ ☒ A description of how samples were collected, noting whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
- ☐ ☒ A statement indicating how many times each experiment was replicated
- ☐ ☒ The statistical test(s) used and whether they are one- or two-sided (note: only common tests should be described solely by name; more complex techniques should be described in the Methods section)
- ☐ ☒ A description of any assumptions or corrections, such as an adjustment for multiple comparisons
- ☐ ☒ The test results (e.g. P values) given as exact values whenever possible and with confidence intervals noted
- ☐ ☒ A clear description of statistics including central tendency (e.g. median, mean) and variation (e.g. standard deviation, interquartile range)
- ☐ ☒ Clearly defined error bars

See the web collection on [statistics for biologists](#) for further resources and guidance.

► Software

Policy information about [availability of computer code](#)

7. Software

Describe the software used to analyze the data in this study.

Behavioural data was analysed using Matlab 2016b. Accelerometer data was preprocessed (e.g., filtered, aligned, etc.) in Matlab 2016b. The computational model was implemented in Matlab 2016b. Classification analysis, model fitting, and model comparison was implemented in R version 3.2.3.

For manuscripts utilizing custom algorithms or software that are central to the paper but not yet described in the published literature, software must be made available to editors and reviewers upon request. We strongly encourage code deposition in a community repository (e.g. GitHub). *Nature Methods* [guidance for providing algorithms and software for publication](#) provides further information on this topic.

► Materials and reagents

Policy information about [availability of materials](#)

8. Materials availability

Indicate whether there are restrictions on availability of unique materials or if these materials are only available for distribution by a for-profit company.

No unique materials were used.

9. Antibodies

Describe the antibodies used and how they were validated for use in the system under study (i.e. assay and species).

No antibodies were used.

10. Eukaryotic cell lines

a. State the source of each eukaryotic cell line used.

No eukaryotic cell lines were used.

b. Describe the method of cell line authentication used.

No eukaryotic cell lines were used.

c. Report whether the cell lines were tested for mycoplasma contamination.

No eukaryotic cell lines were used.

d. If any of the cell lines used are listed in the database of commonly misidentified cell lines maintained by [ICLAC](#), provide a scientific rationale for their use.

No commonly misidentified cell lines were used.

► Animals and human research participants

Policy information about [studies involving animals](#); when reporting animal research, follow the [ARRIVE guidelines](#)

11. Description of research animals

Provide details on animals and/or animal-derived materials used in the study.

No non-human animals were used.

Policy information about [studies involving human research participants](#)

12. Description of human research participants

Describe the covariate-relevant population characteristics of the human research participants.

Sixty-three participants in total completed our behavioural experiments: Ten in Experiment 1 (9 right-handed, 4 males, mean of 27.2 years of age), ten in Experiment 2 (10 right-handed, 5 males, mean of 27.8 years of age), ten in Experiment 3 (9 right-handed, 4 males, mean of 26.5 years of age), twenty in Experiment 4 (all right-handed, 12 males, mean of 22.6 years of age), and ten in Experiment 5 (9 right-handed, 7 males, mean of 29.1 years of age); these same ten participants also completed Experiment 6 (order counterbalanced). A further three right-handed participants completed a more in-depth psychophysical experiment: LO (female; 24 years of age), AY (female; 25 years of age), and EA (male; 20 years of age). All participants had normal or corrected-to-normal vision and no history of neurological impairment.