Reporting Checklist for Nature Neuroscience

This checklist is used to ensure good reporting standards and to improve the reproducibility of published results. For more information, please read Reporting Life Sciences Research.

Please note that in the event of publication, it is mandatory that authors include all relevant methodological and statistical information in the manuscript.

### Statistics reporting, by figure

- Please specify the following information for each panel reporting quantitative data, and where each item is reported (section, e.g. Results, & paragraph number).
- Each figure legend should ideally contain an exact sample size (n) for each experimental group/condition, where n is an exact number and not a range, a clear definition of how n is defined (for example x cells from x slices from x animals from x litters, collected over x days), a description of the statistical test used, the results of the tests, any descriptive statistics and clearly defined error bars if applicable.
- For any experiments using custom statistics, please indicate the test used and stats obtained for each experiment.
- Each figure legend should include a statement of how many times the experiment shown was replicated in the lab; the details of sample collection should be sufficiently clear so that the replicability of the experiment is obvious to the reader.
- For experiments reported in the text but not in the figures, please use the paragraph number instead of the figure number.

Note: Mean and standard deviation are not appropriate on small samples, and plotting independent data points is usually more informative. When technical replicates are reported, error and significance measures reflect the experimental variability and not the variability of the biological process; it is misleading not to state this clearly.

<table>
<thead>
<tr>
<th>TEST USED</th>
<th>n</th>
<th>DESCRIPTIVE STATS (AVERAGE, VARIANCE)</th>
<th>P VALUE</th>
<th>DEGREES OF FREEDOM &amp; F/T/Z/R/ETC VALUE</th>
</tr>
</thead>
<tbody>
<tr>
<td>one-way ANOVA</td>
<td>9, 9, 10, 15 mice from at least 3 litters/group</td>
<td>error bars are mean +/- SEM</td>
<td>p = 0.044</td>
<td>F(3, 36) = 2.97</td>
</tr>
<tr>
<td>unpaired t-test</td>
<td>15 slices from 10 mice</td>
<td>error bars are mean +/- SEM</td>
<td>p = 0.006</td>
<td>t(28) = 2.808</td>
</tr>
<tr>
<td>binomial test</td>
<td>226 Neurons combined in monkey O and U</td>
<td></td>
<td>p = 1.90e-4</td>
<td></td>
</tr>
</tbody>
</table>

Nature Neuroscience: doi:10.1038/nn.3918
<table>
<thead>
<tr>
<th>FIGURE NUMBER</th>
<th>WHICH TEST?</th>
<th>SECTION &amp; PARAGRAPH #</th>
<th>EXACT VALUE</th>
<th>DEFINED?</th>
<th>SECTION &amp; PARAGRAPH #</th>
<th>REPORTED?</th>
<th>DESCRIPTIVE STATS (AVERAGE, VARIANCE)</th>
<th>P VALUE</th>
<th>SECTION &amp; PARAGRAPH #</th>
<th>REPORTED?</th>
<th>EXACT VALUE</th>
<th>SECTION &amp; PARAGRAPH #</th>
<th>VALUE</th>
</tr>
</thead>
<tbody>
<tr>
<td>2a</td>
<td>2 proportion z-test</td>
<td>Results, para 4</td>
<td>226</td>
<td>Neurons combined in monkey O and U</td>
<td>Results, para 4</td>
<td>p = 1.01e-6</td>
<td>Results, para 4</td>
<td>(AVERAGE, VARIANCE)</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>3b</td>
<td>paired t-test</td>
<td>Results, para 5</td>
<td>395</td>
<td>trials</td>
<td>not stated</td>
<td>shaded region is mean +/- SEM</td>
<td>figure legend</td>
<td>p = 1.40e-10</td>
<td>Results, para 5</td>
<td>t(393) = 6.592</td>
<td>not stated</td>
<td></td>
<td></td>
</tr>
<tr>
<td>4a</td>
<td>binomial test</td>
<td>Results, para 6</td>
<td>226</td>
<td>Neurons combined in monkeys O and U</td>
<td>online methods, Statistical Analysis</td>
<td>p = 2.12e-13</td>
<td>Results, para 6</td>
<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td></td>
<td>chi-squared test</td>
<td>Results, para 9</td>
<td></td>
<td>fraction of neurons encoding both HVL and choice</td>
<td>Results, para 9</td>
<td>p = 3.14e-4</td>
<td>only p-value reported : Results, para 9</td>
<td>x2(1) = 12.987</td>
<td>not stated</td>
<td></td>
<td></td>
<td></td>
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</tr>
<tr>
<td>5a</td>
<td>correlation coefficient</td>
<td>Results, para 9</td>
<td>226</td>
<td>Neurons combined in monkeys O and U</td>
<td>Results, para 9</td>
<td>p = 2.25e-19</td>
<td>Results, para 9</td>
<td>224</td>
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<tr>
<td></td>
<td>chi-squared test</td>
<td>Results, para 10</td>
<td></td>
<td>fraction of neurons encoding PRL and choice</td>
<td>Results, para 10</td>
<td>p = 0.007</td>
<td>Results, para 10</td>
<td>x2(1) = 7.344</td>
<td>not stated</td>
<td></td>
<td></td>
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<td></td>
</tr>
<tr>
<td>5b</td>
<td>correlation coefficient</td>
<td>Results, para 10</td>
<td>226</td>
<td>Neurons combined in monkeys O and U</td>
<td>Results, para 10</td>
<td>p = 0.215</td>
<td>Results, para 10</td>
<td>224</td>
<td></td>
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</tr>
<tr>
<td>5c</td>
<td>permutation test</td>
<td>Results, para 11</td>
<td>10000 shuffles</td>
<td>HVL</td>
<td>Online methods, para 12</td>
<td>p = 0.0001</td>
<td>Results, p.7, paragraph 2</td>
<td></td>
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<td></td>
<td></td>
</tr>
<tr>
<td>5c</td>
<td>permutation test</td>
<td>Results, para 11</td>
<td>10000 shuffles</td>
<td>PRL</td>
<td>Online methods, para 12</td>
<td>p = 0.100</td>
<td>Results, para 11</td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>5d</td>
<td>permutation test</td>
<td>Results, para 11</td>
<td>10000 shuffles</td>
<td>For both HVL and PRL</td>
<td>Online methods, para 12</td>
<td>p = 0.0001</td>
<td>Results, para 11</td>
<td></td>
<td></td>
<td></td>
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<td></td>
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</tr>
<tr>
<td>6b</td>
<td>paired t-test (reward over 50%)</td>
<td>Results para 13</td>
<td>216</td>
<td>Neurons in which number of trials were sufficient for decoding (&gt;=5 trials per category)</td>
<td>Online methods, statistical analyses</td>
<td>shaded region is mean +/- SEM</td>
<td>figure legend</td>
<td>p = 7.32e-13</td>
<td>Results, para 13</td>
<td>t(215) = 7.635</td>
<td>not stated</td>
<td></td>
<td></td>
</tr>
<tr>
<td>6b</td>
<td>paired t-test (unrewarded over 50%)</td>
<td>Results para 13</td>
<td>216</td>
<td>Neurons in which number of trials were sufficient for decoding (&gt;=5 trials per category)</td>
<td>Online methods, statistical analyses</td>
<td>shaded region is mean +/- SEM</td>
<td>figure legend</td>
<td>p = 0.517</td>
<td>Results, para 13</td>
<td>t(215) = 0.650</td>
<td>not stated</td>
<td></td>
<td></td>
</tr>
<tr>
<td>6b</td>
<td>paired t-test: last chosen location (rewarded vs unrewarded)</td>
<td>Results, para 13</td>
<td>216</td>
<td>Neurons in which number of trials were sufficient for decoding (&gt;=5 trials per category)</td>
<td>Online methods, statistical analyses</td>
<td>shaded region is mean +/- SEM</td>
<td>figure legend</td>
<td>p = 0.816</td>
<td>Results, para 13</td>
<td>t(215) = 0.233</td>
<td>not stated</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>
### Representative figures

1. Are any representative images shown (including Western blots and immunohistochemistry/staining) in the paper?
   - If so, what figure(s)?
   - N/A

2. For each representative image, is there a clear statement of how many times this experiment was successfully repeated and a discussion of any limitations in repeatability?
   - If so, where is this reported (section, paragraph #)?
   - N/A

### Statistics and general methods

1. Is there a justification of the sample size?
   - If so, how was it justified?
   - Where (section, paragraph #)?
   - Behavioral data reported in this study are over-powered, as reflected in unusually small p-values, since they were concurrently collected during the neurophysiological recording experiments.
   - Even if no sample size calculation was performed, authors should report why the sample size is adequate to measure their effect size.
   - We analyzed the activity recorded from 226 neurons in the DLPFC, which is larger than or comparable to most neurophysiological studies in the primate DLPFC.
   - N/A

2. Are statistical tests justified as appropriate for every figure?
   - Where (section, paragraph #)?
   - No.
   - a. If there is a section summarizing the statistical methods in the methods, is the statistical test for each experiment clearly defined?
   - Yes.
   - b. Do the data meet the assumptions of the specific statistical test you chose (e.g. normality for a parametric test)?
   - No.
<table>
<thead>
<tr>
<th>Question</th>
<th>Answer</th>
</tr>
</thead>
<tbody>
<tr>
<td>c. Is there any estimate of variance within each group of data?</td>
<td>N/A</td>
</tr>
<tr>
<td>Is the variance similar between groups that are being statistically compared?</td>
<td></td>
</tr>
<tr>
<td>Where is this described (section, paragraph #)?</td>
<td></td>
</tr>
<tr>
<td>d. Are tests specified as one- or two-sided?</td>
<td>All tests were two-tailed.</td>
</tr>
<tr>
<td>e. Are there adjustments for multiple comparisons?</td>
<td>N/A.</td>
</tr>
<tr>
<td>3. Are criteria for excluding data points reported?</td>
<td>Any cell that was recorded for fewer than 320 trials was not included. This criterion was established prior to data collection (see Online Methods).</td>
</tr>
<tr>
<td>Was this criterion established prior to data collection?</td>
<td></td>
</tr>
<tr>
<td>Where is this described (section, paragraph #)?</td>
<td></td>
</tr>
<tr>
<td>4. Define the method of randomization used to assign subjects (or samples) to the experimental groups and to collect and process data. If no randomization was used, state so. Where does this appear (section, paragraph #)?</td>
<td>We recorded from all neurons in which we were able to keep for at least 320 trials (see Online Methods)</td>
</tr>
<tr>
<td>5. Is a statement of the extent to which investigator knew the group allocation during the experiment and in assessing outcome included? If no blinding was done, state so. Where (section, paragraph #)?</td>
<td>No blinding was necessary for these experiments.</td>
</tr>
<tr>
<td>6. For experiments in live vertebrates, is a statement of compliance with ethical guidelines/regulations included? Where (section, paragraph #)?</td>
<td>Yes, see Online Methods (Animal preparation).</td>
</tr>
<tr>
<td>7. Is the species of the animals used reported?</td>
<td>Yes, see Online Methods (Animal preparation).</td>
</tr>
<tr>
<td>Where (section, paragraph #)?</td>
<td></td>
</tr>
<tr>
<td>8. Is the strain of the animals (including background strains of KO/transgenic animals used) reported? Where (section, paragraph #)?</td>
<td>N/A.</td>
</tr>
<tr>
<td>9. Is the sex of the animals/subjects used reported?</td>
<td>Yes, see Online Methods (Animal preparation).</td>
</tr>
<tr>
<td>Where (section, paragraph #)?</td>
<td></td>
</tr>
<tr>
<td>10. Is the age of the animals/subjects reported?</td>
<td>No.</td>
</tr>
<tr>
<td>Where (section, paragraph #)?</td>
<td></td>
</tr>
<tr>
<td>11. For animals housed in a vivarium, is the light/dark cycle reported?</td>
<td>No.</td>
</tr>
<tr>
<td>Where (section, paragraph #)?</td>
<td></td>
</tr>
</tbody>
</table>
12. For animals housed in a vivarium, is the housing group (i.e. number of animals per cage) reported?
   Where (section, paragraph #)?
   No, but both were pair-housed.

13. For behavioral experiments, is the time of day reported (e.g. light or dark cycle)?
   Where (section, paragraph #)?
   No. All recording was on Monday-Friday between 9am and 6pm.

14. Is the previous history of the animals/subjects (e.g. prior drug administration, surgery, behavioral testing) reported?
   Where (section, paragraph #)?
   No. One monkey was naive and the other had been previously trained on a joystick task.

   a. If multiple behavioral tests were conducted in the same group of animals, is this reported?
      Where (section, paragraph #)?
      N/A

15. If any animals/subjects were excluded from analysis, is this reported?
   Where (section, paragraph #)?
   No animals were excluded from the analysis.

   a. How were the criteria for exclusion defined?
      Where is this described (section, paragraph #)?
      N/A

   b. Specify reasons for any discrepancy between the number of animals at the beginning and end of the study.
      Where is this described (section, paragraph #)?
      N/A

Reagents

1. Have antibodies been validated for use in the system under study (assay and species)?
   N/A

   a. Is antibody catalog number given?
      Where does this appear (section, paragraph #)?
      N/A

   b. Where were the validation data reported (citation, supplementary information, Antibodypedia)?
      Where does this appear (section, paragraph #)?
      N/A

2. If cell lines were used to reflect the properties of a particular tissue or disease state, is their source identified?
   Where (section, paragraph #)?
   N/A
Data deposition

Data deposition in a public repository is mandatory for:

a. Protein, DNA and RNA sequences
b. Macromolecular structures
c. Crystallographic data for small molecules
d. Microarray data

Deposition is strongly recommended for many other datasets for which structured public repositories exist; more details on our data policy are available here. We encourage the provision of other source data in supplementary information or in unstructured repositories such as Figshare and Dryad.

1. Are accession codes for deposit dates provided?

   Where (section, paragraph #)?

   No.

Computer code/software

Any custom algorithm/software that is central to the methods must be supplied by the authors in a usable and readable form for readers at the time of publication. However, referees may ask for this information at any time during the review process.

1. Identify all custom software or scripts that were required to conduct the study and where in the procedures each was used.

   Matlab was used to analyze the data.

2. Is computer source code/software provided with the paper or deposited in a public repository? Indicate in what form this is provided or how it can be obtained.

   It is not publicly available, but .mat files can be provided upon request.

Human subjects

1. Which IRB approved the protocol?

   Where is this stated (section, paragraph #)?

2. Is demographic information on all subjects provided?

   Where (section, paragraph #)?

3. Is the number of human subjects, their age and sex clearly defined?

   Where (section, paragraph #)?

4. Are the inclusion and exclusion criteria (if any) clearly specified?

   Where (section, paragraph #)?
5. How well were the groups matched?
   Where is this information described (section, paragraph #)?

6. Is a statement included confirming that informed consent was obtained from all subjects?
   Where (section, paragraph #)?

7. For publication of patient photos, is a statement included confirming that consent to publish was obtained?
   Where (section, paragraph #)?

### fMRI studies

For papers reporting functional imaging (fMRI) results please ensure that these minimal reporting guidelines are met and that all this information is clearly provided in the methods:

1. Were any subjects scanned but then rejected for the analysis after the data was collected?
   a. If yes, is the number rejected and reasons for rejection described?
      Where (section, paragraph #)?

2. Is the number of blocks, trials or experimental units per session and/or subjects specified?
   Where (section, paragraph #)?

3. Is the length of each trial and interval between trials specified?

4. Is a blocked, event-related, or mixed design being used? If applicable, please specify the block length or how the event-related or mixed design was optimized.

5. Is the task design clearly described?
   Where (section, paragraph #)?

6. How was behavioral performance measured?

7. Is an ANOVA or factorial design being used?

8. For data acquisition, is a whole brain scan used?
   a. How was this region determined?
9. Is the field strength (in Tesla) of the MRI system stated?
   
   a. Is the pulse sequence type (gradient/spin echo, EPI/spiral) stated?
   
   b. Are the field-of-view, matrix size, slice thickness, and TE/TR/flip angle clearly stated?
   
10. Are the software and specific parameters (model/functions, smoothing kernel size if applicable, etc.) used for data processing and pre-processing clearly stated?

11. Is the coordinate space for the anatomical/functional imaging data clearly defined as subject/native space or standardized stereotaxic space, e.g., original Talairach, MNI305, ICBM152, etc? Where (section, paragraph #)?

12. If there was data normalization/standardization to a specific space template, are the type of transformation (linear vs. nonlinear) used and image types being transformed clearly described? Where (section, paragraph #)?

13. How were anatomical locations determined, e.g., via an automated labeling algorithm (AAL), standardized coordinate database (Talairach daemon), probabilistic atlases, etc.?

14. Were any additional regressors (behavioral covariates, motion etc) used?

15. Is the contrast construction clearly defined?

16. Is a mixed/random effects or fixed inference used?
   
   a. If fixed effects inference used, is this justified?

17. Were repeated measures used (multiple measurements per subject)?
   
   a. If so, are the method to account for within subject correlation and the assumptions made about variance clearly stated?

18. If the threshold used for inference and visualization in figures varies, is this clearly stated?

19. Are statistical inferences corrected for multiple comparisons?
   
   a. If not, is this labeled as uncorrected?
20. Are the results based on an ROI (region of interest) analysis?
   a. If so, is the rationale clearly described?
   b. How were the ROI’s defined (functional vs anatomical localization)?

21. Is there correction for multiple comparisons within each voxel?

22. For cluster-wise significance, is the cluster-defining threshold and the corrected significance level defined?

» Additional comments

Additional Comments