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seq<sub>i</sub>nact.pdf[Neuronalpopulationdynamics and inactivation experiments] Neuronal population dynamics and inactivation experiments of the sequence of the seque
   seq_i nactc). These results were obtained days or weeks after the mouse achieved plateau behavioral performance, suggesting that Foundation and the sequence of the sequence
   learning phase. These results were in agreement with earlier work that used pharmacological methods to inactivate the PPC and other phases. The series of the property of th
   PPC activity could be involved in the transformation of the sensor yinformation into a behavioral action planor in some aspect of various and the property of the property o
                                      _{a}ll.pdf[Reorganization of activity within a trial across days.] Reorganization of activity within a trial across days.
   70 cmshift) on a subsequent day. Shading indicates mean sem (n = 1)
  5mice; some large interval data points had fewer than 5mice; see Supplemental Figure S1A). The gray shaded are a indicates 95 configure S1A and S1A are supplemental from the supplemental Figure S1A. The gray shaded are a indicates 95 configure S1A and S1A are supplemental from the su
   10^-8, ANOVA.c, Fraction of cells that had a significant peak on the noted day. <math>Fraction vs. time:
   0.85, ANOVA. In panels C-
   D, errorbarsindicate mean sem, n =
   5, 5, 4mice for the time intervals shown. \mathbf{d}, Left:
  \label{local-control} For cells with a significant peak on dayn and dayn+\\ x, the fraction of peak sthat shifted by greater than 35 cm, 50 cm and 1 m. Fraction moved 35 cm vs. time:
   0.019, ANOVA. Center:
   For cells with a highly significant peak on dayn, the fraction of cells that did not have a significant peak on dayn+
   x.Fraction lost vs.time:
   10^-9, ANOVA. Right:
   For cells without a significant peak of activity on dayn, the fraction of cells that had a highly significant peak on dayn <math>+
   x.Fraction gained vs.time:
 p = 0.96ANOVA.
                                      _{a}ll.pdf [Reorganization of information about trial-\\
   typeacrossdays | Reorganization of information about trial-type across days a, Decoding accuracy for trial type based of the second sec
   11, ANOVA. {f e}, On a given day, the cells with the top 20 and bottom 20 of decoding accuracies were identified. The distributions of decoding accuracies were identified. The distributions of decoding accuracies were identified and the distributions of the distribution and the distribution and the distribution and the distribution are decoding accuracies were identified and the distribution and the distribution are decoding accuracies were identified and the distribution and the distribution are decoding accuracies were identified and the distribution and the distribution are decoding accuracies were identified and the distribution and the distribution are decoding accuracies were identified and the distribution are decoding accuracies were identified and the distribution are decoding accuracies were identified and the distribution and the distribution are decoding accuracies were identified and the distribution are decoding accuracies were identified and the distribution are decoding accuracies were identified and the distribution and the distribution are decoding accuracies and accuracies are dec
   right turn trials. A model weight was determined at each spatial bin in the maze, and the mean weight was calculated for each cell <math>{f g}, {f E}
   typeinformation.Top:
   mean fluorescence image of the cellbody. Bottom:
   mean activity of the cell on correct white cue-
   leftturn(blue) and black cue-
   rightturn(red)trials. {f h}, On a given day, the cells with the top 20 largest weights for white cue-plane is a superior of the contraction of 
   right turn trials were identified. The distributions of trial-
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type weights are shown in comparison to the distribution for all cells after intervals of 1, 10, and 20 days.

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