

Experience-Driven Rate Modulation is Reinstated During Hippocampal Replay

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Replay, the sequential reactivation of a neuronal ensemble, is thought to play a central role in the hippocampus during the consolidation of a recent experience into a long-term memory. Following a contextual change (e.g. entering a novel environment), hippocampal place cells typically modulate their in-field firing rate and shift the position of their place field, providing a rate and place representation for the behavioral episode, respectively. However, replay has been largely defined by only the latter- based on the fidelity of sequential activity across neighboring place fields. Here we show that dorsal CA1 place cells in rats can modulate their firing rate between the replay of two different contexts, mirroring the same pattern of rate modulation observed during behavior. This context-driven rate modulation within replay events was experience-dependent, observable during both behavioral episodes and throughout the subsequent rest period, but not prior to experience. Furthermore, we demonstrate that both the temporal order and firing rate of place cells can independently be used to decode contextual information within a replay event, revealing the existence of two separable but complementary neural representations available for memory consolidation processes.

A key function of the hippocampus is the initial encoding and subsequent consolidation of episodic memories. Hippocampal place cells are activated in sequential patterns during behavioral episodes (e.g. a rodent running towards a goal), with each neuron's activity modulated by the animal's position within an environment (i.e. a place field) (1, 2). During offline periods such as sleep, place cells spontaneously reactivate, replaying the same sequential pattern previously activated during the behavioral episode (3–6). This phenomenon of neural replay is postulated to drive systems-level memory consolidation (7–11). Hippocampal replay has largely been defined by the sequential patterns of place cells (12, 13), which depends on the location of each neuron's place field in the environment (here referred to as place representation). Yet, place cells also carry additional information in the magnitude of their place field's activity (here referred to as rate representation). The magnitude of a place field's activity (e.g. peak in-field firing rate) is modulated by both local and global contextual cues (14–17), as well as behavioral variables (e.g. animal's acceleration during locomotion) (18, 19), and cognitive events (e.g. animal's attention and perception) (20–23). However, we do not know whether the hippocampus is capable of reinstating this rate modulation in place cells during offline replay.

To address this question, we recorded extracellularly from the dorsal CA1 of rats ($n = 3$, male Lister-hooded), trained to run back and forth along two novel linear tracks (2m) to collect a liquid reward at each end (Fig. 1A, Fig. S1A, Table S1, see Methods). A view-obstructing divider was placed between tracks and distinguishing visual cues were present within each environment to facilitate contextual discrimination (and in turn, hippocampal remapping). Additionally, rats rested in a quiet remote location, both before (PRE) and after (POST) the exploration of the two tracks.

We observed global remapping of place cells between the two linear tracks [Pearson correlation coefficient (r) between tracks population vectors, $r = 0.04 \pm 0.11$], such that the majority of place fields shifted their location along the track and/or modulated their firing rate (i.e. changed to higher or lower peak in-field firing rate) (Fig. S1B). Both the context (which linear track) and current position (location of the animal on the linear track) could be inferred from place cell activity with high accuracy using a naïve Bayes decoder (Fig. 1B, Fig.S1C). Track detection accuracy exceeded 89% during RUN periods on both tracks [all sessions: $89 \pm 5\%$] with an average median error of 5cm [all sessions: 5.17 ± 0.69 cm], while a population vector analysis showed high correlation between the ratemaps of the first and second half of running within each track [Pearson correlation coefficient (r) within track population vectors, $r = 0.83 \pm 0.04$]. We also used Bayesian decoding to detect replay sequences of each linear track (Fig. 1B), with statistically significant replay trajectories identified by comparing the weighted correlation score obtained from the posterior probabilities of each decoded event to three types of event shuffles [Total replay events = 2805, PRE: Track1= 35 ± 20 , Track2= 37 ± 15 , POST: Track1= 138 ± 61 , Track2= 156 ± 34 , RUN: Track1= 98 ± 20 , Track2= 95 ± 31 , Fig. S2, see Methods].

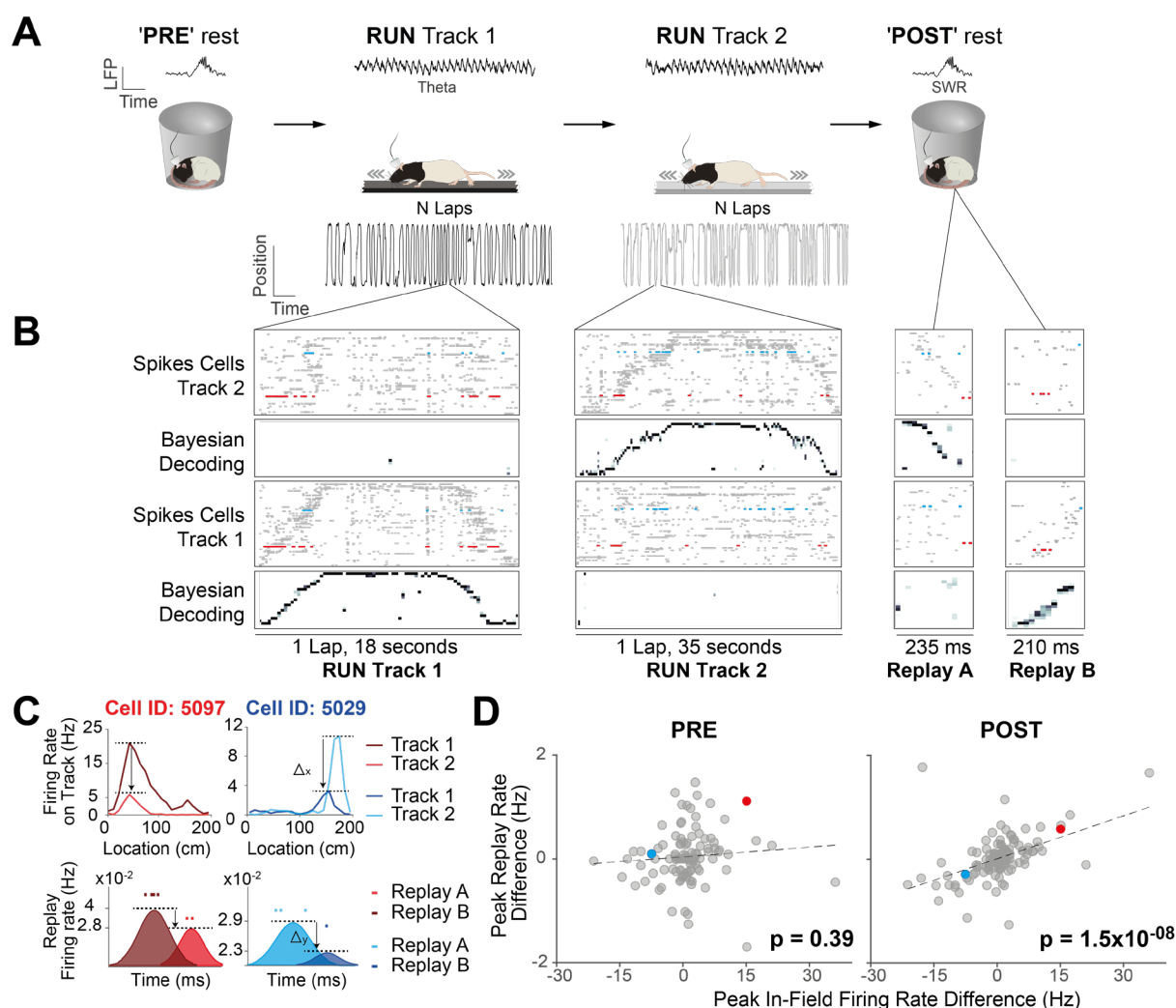


Figure 1. Rate coding in wake and rest. (A) Experimental design. Each recording session, the rats rested (PRE), ran back and forth on two different linear tracks (RUN), and then rested again (POST) (left to right). An example of LFP activity in each behavioral state is displayed [RUN-theta, PRE/POST-sharp wave ripples (SWR)] (B) Raster plots of spiking activity of cells sorted by their peak firing rate on the track (top: Track 2, bottom: Track 1), and the associated decoded positions for a single exemplar lap (first two columns) and replay event (last two columns), for each track (top: Track 2; bottom: Track 1). Example from Rat 3, session 2. (C) Two example neurons (highlighted in B) that changed their peak instantaneous firing rate (bottom) during two example replay events (displayed in B) in the same direction as their peak in-field firing rate difference (top). The activity and properties of the example neurons are color coded throughout all the panels. (D) The peak in-field firing rate difference (Track1-Track2) significantly predicts the average peak instantaneous replay rate difference (Track1-Track2) in POST rest, but not PRE rest. Each data point is a neuron active on both tracks (PRE: $n = 106$ cells, POST: $n = 123$ cells).

First, we sought to investigate whether the contextually-driven place fields' rate modulation between tracks is conserved during offline replay. Using all place cells active on both tracks and stable across the whole run (Fig.S1B, see Methods), we regressed the place fields' peak

in-field firing rate differences between tracks (Track1-Track2) against the difference (Track1-Track2) in average peak instantaneous rate of those same cells during significant replay events (Fig. 1C). For replay events during POST (but not PRE), the difference in peak in-field firing rate between tracks was predictive of the difference in average peak instantaneous rates during replay events from both tracks [PRE: $\beta = .06$, $F(1,104) = 0.75$, $P = 0.39$, $R^2 = .007$; POST: $\beta = .04$, $F(1,121) = 36.95$, $P < 0.001$, $R^2 = .23$; Fig. 1D]. In other words, if a place field had a higher firing rate on Track 2 (compared to Track 1), it also tended to have a higher mean firing rate across all Track 2 replay events (compared to Track 1 events) during POST, but not during PRE. This suggests that any contextually-driven rate modulation between two behavioral episodes is maintained during offline replay. This observation was consistent when using a more selective criteria of place field activity (e.g. place fields with higher minimum peak firing rate or place cells with a single field) (Fig. S3, S4) as well as for alternative metrics for both replay firing rate (e.g. mean number of spikes per replay event and median replay rate) (Fig. S4) and place fields firing rate (e.g. place field's area under the curve) (Fig. S4). Furthermore, as expected, selecting only cells that were classified as significantly rate modulated led to an even stronger effect (Fig. S4).

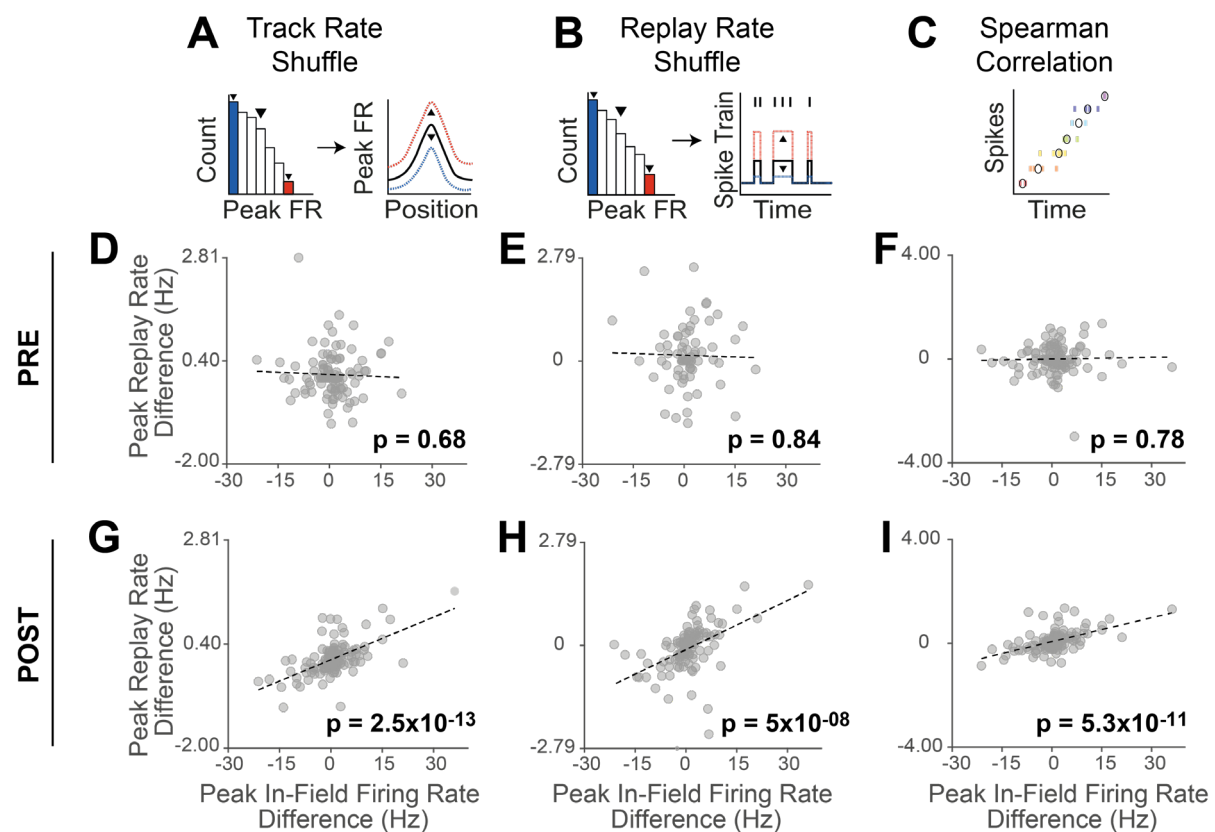


Figure 2. The reinstatement of rate modulation during rest replay is observed using replay detection methods insensitive to firing rate. (A-C) Three methods of replay detection that are insensitive to firing rate: **(A)** Track rate shuffle (PRE: $n = 106$ cells, POST: $n = 123$ cells), **(B)** Replay rate shuffle (PRE: $n = 69$ cells, POST: $n = 117$ cells), **(C)** Spearman correlation (PRE: $n = 111$ cells, POST: $n = 123$ cells). **(A-B)** Example of the original rate (in black) randomly scaled up (in red) or

down (in blue). **(D-F)** Regression of peak-in-field firing rate vs. peak replay rate difference (Track 1-Track 2) for PRE replay events with modified detection method: **(D)** Track rate shuffle, **(E)** Replay rate shuffle, **(F)** Spearman correlation. **(G-I)** Regression of peak-in-field firing rate vs. peak replay rate difference (Track 1-Track 2) for POST replay events with modified detection method **(G)** Track rate shuffle, **(H)** Replay rate shuffle, **(I)** Spearman correlation.

Replay detection methods generally detect the statistical significance of cells sequences based on templates derived from the behavioral episode, which are sensitive to the order of place field activation (during RUN and replay). However, because Bayesian decoding can also be influenced by firing rate, we next tested whether firing rate differences between tracks may bias the selection of the replay events used in the regression analysis. To remove the influence of firing rate in the replay event selection, we employed two independent types of shuffles before repeating the detection of replay events: 1) randomizing the overall firing rate of place fields within a track (Track rate shuffle, Fig.2A) and 2) randomizing the cell's firing rate during replay events (Replay rate shuffle) (Fig. 2B). For both shuffles, each place cell's firing rate was scaled based on a value randomly drawn from the corresponding rate distribution obtained from all cells. These significant replay events were then used to repeat the regression analysis using their original content (i.e. the original replay spike train, instead of the shuffled one) against the original differences in the place fields' in-peak firing rate. For both shuffles, we observed a significant contextual rate modulation of replay, where track differences in the place fields' in-peak firing rate remained predictive of track differences in the average peak instantaneous rates during replay events in POST, but not PRE [PRE: Track rate shuffle $P = .68$, Replay rate shuffle: $P = .84$, POST: Track rate shuffle $P < .001$, Replay rate shuffle: $P < .001$; Fig. 2D,E,G,H]. As an additional control, we detected replay events using a rank order correlation method independent of Bayesian decoding, which relies only on comparing the median spike times (across place cells) within a replay event compared to the sequential order of place field activity along the track (Fig. 2C). Repeating the regression analysis based on the track differences in the place fields' in-peak firing rate against the replay events detected using a Spearman correlation still resulted in a significant contextual rate modulation of the replay events rate in POST, but not PRE [PRE: $\beta = .002$, $F(1,111) = .08$, $P = .78$, $R^2 = .0007$, POST: $\beta = .03$, $F(1,121) = 51.99$, $P < .001$, $R^2 = 0.3$; Fig. 2F,I]. Overall, these results indicate that the contextual rate modulation observed during replay is not a result of a bias within the replay detection analysis (namely, Bayesian decoding). Importantly, we did not observe any significant track differences in place field or replay properties (Fig. S5, including the place fields' peak in-field firing rate [two-sided two-sample KS test, $P = .62$] and replay average peak instantaneous rate [two-sided two-sample KS test, $P = .79$], and confirmed that our regression analysis was not influenced by any intrinsic biases between tracks by remeasuring the statistical significance relative to a shuffled distribution (Fig. S6).

The phenomenon of replay has also been reported during awake quiescent states (e.g. while pausing on the track), and it is thought to serve different functional roles, ranging from planning to memory consolidation (24–27). We observed that during the behavioral episode on the

tracks, contextual rate modulation was expressed across local awake replay events [RUN: $\beta = .05$, $F(1,120) = 197.02$, $P < .001$, $R^2 = .61$; Fig. 3A]. The rate modulation during RUN was stronger than the one observed in rest replay events during POST [$\beta = .04$, $F(1,121) = 36.95$, $P < .001$, $R^2 = .23$; Fig. 1D]. Place cells during RUN already showed evidence of contextual rate modulation during awake local replay events within the first 2 laps of experience [2 laps: $\beta = .06$, $F(1,8) = 5.91$, $P = .04$, $R^2 = .42$; Fig. 3B]. Furthermore, place cells expressing rate modulation during RUN replay events were also likely to show a similar direction of contextual rate modulation (e.g. Track 1 > Track 2) during POST replay events [$\beta = .78$, $F(1,168) = 95.36$, $P < .001$, $R^2 = .36$, Fig. 3C], but not PRE [$\beta = .02$, $F(1,143) = 1.25$, $P = .2$, $R^2 = .008$, Fig S7]. Contextual rate modulation during replay was significantly expressed throughout the entire rest period during POST (at least 1 hour cumulatively, limited only by the maximum time period tested), with the strongest effect observed in the earliest portion of POST rest (Fig. 3D).

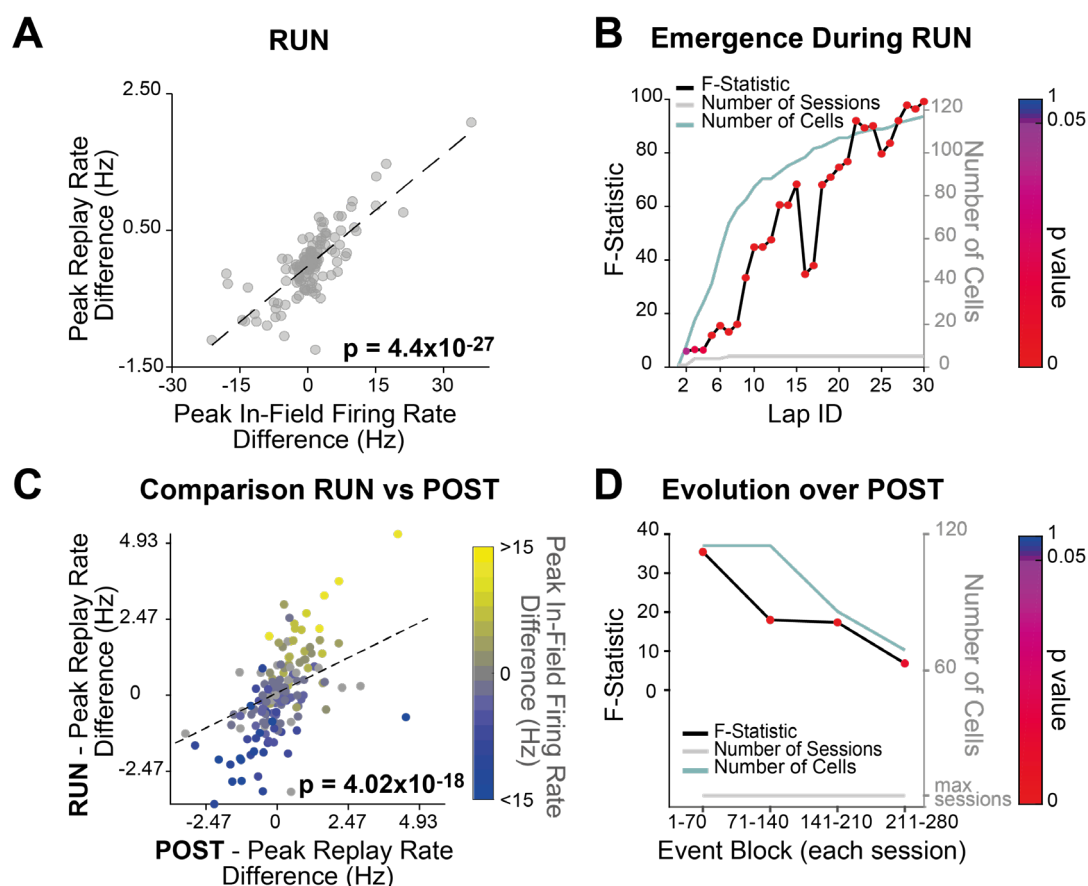


Figure 3. Rate modulation during replay is observed from the beginning of RUN and continues throughout the entire RUN and POST epochs. (A) Regression of peak in-field firing rates against peak replay rates (track differences) for all local RUN replay events. (B) Lap by lap emergence of rate modulation for RUN local replay events. F-statistic of regression - peak in-field firing rate vs. peak replay rate (track difference) - for replay events occurring both prior to and including the specified lap (lap ID). The p-value of each regression is indicated by color. The number of cells and number of data sessions contributing in each regression are indicated with the green and grey line, respectively. (C) Peak replay rate difference for replay events (RUN vs. POST), color coded by peak in-field rate

differences. **(D)** Temporal dynamics of rate modulation during POST. The regression's F statistic (y-axis) and the corresponding p-value (marker color), analyzed in non-overlapping blocks of 70 consecutive replay events (per session).

Our data suggest that both place (i.e. sequences) and rate information is embedded within hippocampal replay events. If this is the case, then the spatial context (i.e. track identity) associated with a replay event should be identifiable using either the place or rate representations alone. To address this question, we modified the current Bayesian framework such that it would rely on either or both place and rate information. For each decoded replay event, we measured the probabilistic bias towards one of the two tracks using the z-scored log odd(28), $(\log \frac{\sum prob(Track_1)}{\sum prob(Track_2)})$. This method summed the decoded posterior probabilities for each track within a replay event, quantified the log of the between track ratio, and then z-scored this ratio relative to a shuffled distribution (Ratemap track label shuffle, see Methods). For example, a more positive z-scored log odd would indicate a greater probabilistic bias towards Track 1. Therefore, if the content of a replay event was detected for Track 1 (based on a statistically significant sequence), the z-scored log odd for this event replay event would reflect how well a place and/or rate representation could be used to correctly classify this replay event as Track 1 (rather than Track 2).

In particular, we used two measurements to assess the replay track identity classification performance when using either the rate and/or place representation: 1) the log odd difference between two tracks compared to a Replay track identity shuffled distribution (see Methods) and 2) the binary discriminability of a replay event's track identity using Receiver Operator Characteristic (ROC) curves, in which the area under the ROC curve (AUC) of 0.5 indicates chance discriminability (and an AUC of 1 indicates perfect discriminability). For both measurements, only replay events with a statistically significant replay sequence were used. When both place and rate information were unaltered, the z-scored log odd difference and the track discriminability were significantly positive and highest for RUN [$\Delta(\text{z-scored log odd}) = 2.77, P < .001, \text{AUC} = .98$], significant but marginally lower for POST [$\Delta(\text{z-scored log odd}) = 1.29, P < .001, \text{AUC} = .80$], but indistinguishable from the null distribution and at chance levels for PRE [$\Delta(\text{z-scored log odd}) = .08, P = .22, \text{AUC} = 0.52$] (Fig. 4A, Table S2).

We then proceeded to selectively remove either the place or firing rate information available to the Bayesian decoder and assess its ability to determine the track being replayed. First, we removed the firing rate information by setting each place field's in-field peak firing rate to its average across both tracks, and by rescaling each cell's replay spike trains to its average firing rate across all replay events (Fix rate manipulation). We found that place information alone was sufficient to correctly classify replay events to each track, although both the z-scored log odd difference and the AUC were slightly lower than when both place and firing rate information were available [AUC difference < -0.1 compared to the original for all task periods, Fig. 4B,F-H, Table S2].

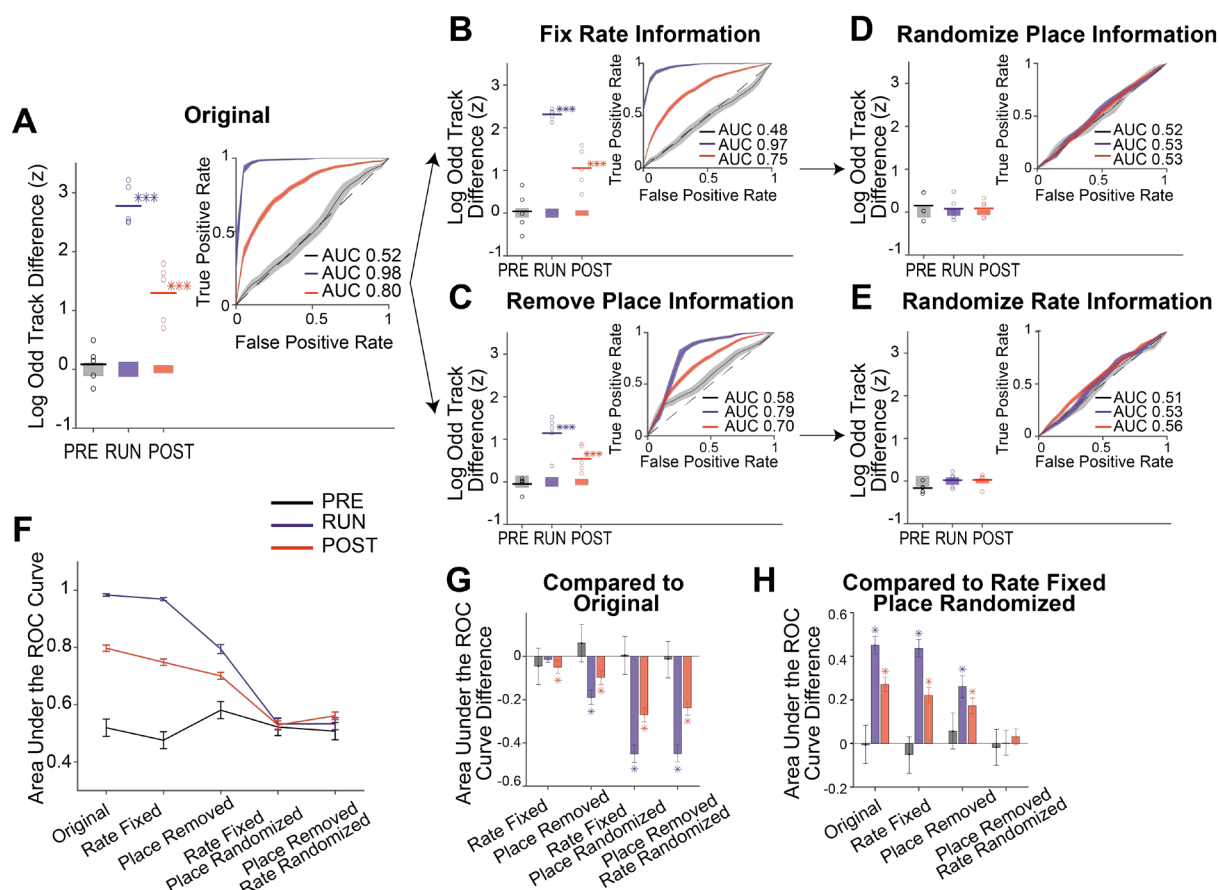


Figure 4. Contextual information can be decoded from either place or rate representations during replay. (A-E) The decoding performance for context (Track 1 or Track 2) for each manipulation, restricting the information (rate and/or place) available to the decoder. (Main plots) Mean z-scored log odd difference between tracks for replay during PRE (black), RUN (blue) and POST (red). Each data point is the mean of an individual session, while the horizontal line indicates the mean across all sessions (* $P < 0.05$, ** $P < 0.01$, *** $P < 0.001$, one-tailed). The filled box indicates the SEM from the Replay track identity shuffles. (Inset) ROC curves of Track1/Track2 binary discrimination for replay events during PRE, RUN and POST. The dashed line along the diagonal indicates chance performance (AUC = 0.5). The shaded region indicates the bootstrapped SE of the ROC curves. (A) Original (*rate and place information intact*) (B) Fix rate manipulation (*only place information available*) (C) Remove place manipulation (*only rate information available*) (D) Fix rate manipulation with place information additionally randomized (*no rate or place information available*). (E) Remove place manipulation with rate information additionally randomized (*no rate or place information available*). (F) Summary of mean AUC under each condition, for PRE, RUN, and POST (G) Mean AUC difference between original and each decoding manipulation. (H) Mean AUC difference between *negative control* and other conditions For G and H, * indicates where the 98.75% confidence interval of the bootstrapped AUC difference distribution does not overlap with 0 (see Table S3, S4). Similar results were obtained using an alternative approach (Delong method) for measuring significance (Table S5, S6, see Methods).

Next, we selectively removed place information without altering rate information, by expanding the size of our position and time bin for decoding: a single position bin spanning the

entire track and a single time bin spanning the entire replay event (Remove place manipulation, Fig. 4C, F-H). This manipulation led to a significant log odd difference towards the correct track and a high track discriminability for RUN [$\Delta(\text{z-scored log odds}) = 1.14$, $P < .001$, $\text{AUC} = .79$], POST [$\Delta(\text{z-scored log odds}) = .53$, $P < .001$, $\text{AUC} = .70$], but not PRE ($\Delta(\text{z-scored log odds}) = -.04$, $P = 0.63$, $\text{AUC} = .58$], supporting the finding that rate modulation is reinstated during replay and is experience-dependent. Finally, as a negative control for both manipulations, where rate and place representations were disrupted, each remaining non-manipulated representation (place or rate) was then randomized (Rate fixed place randomized manipulation and Place removed rate randomized manipulation, respectively). Removing both place and rate representations resulted in chance level discriminability for both conditions (Fig. 4D-H, Table S2).

Here we have shown that contextually driven place and rate representations used by the hippocampus during behavior are reinstated during replay, with either representation being sufficient to successfully decode the context being replayed. This phenomenon was experience-dependent, as rate modulation within replay was only observed during and after a behavioral episode, but not prior to it. While place cell sequences are generally stable across a behavioral episode within a single environment, in-field firing rates can still increase (or decrease) with minor contextual changes (29) a phenomenon referred to as rate remapping (19, 22, 30). Our results suggest that during replay, firing rate modulation could allow the hippocampus to discriminate between two marginally similar contexts even if they share an identical sequential pattern of place cell activity.

Computational models of replay have commonly used modified synfire chains, sequentially connected neurons that are often embedded within distinct recurrent networks (31–33). While these models aim to replicate the sequential nature of replay, it remains to be seen whether they can also reinstate experience-dependent rate modulation, or if additional mechanisms may need to be considered. Compelling evidence suggests a key role for neocortical activity preceding a hippocampal replay event, in determining its content (10, 34, 35). If cortical inputs to the hippocampus during behavior underlie rate modulation (36), a reinstatement of these same inputs prior to replay may be necessary for the subsequent rate modulation of hippocampal place fields during offline states.

AUTHOR CONTRIBUTIONS

M.T.¹, M.H.G. and D.B. designed the experiment, M.T.¹ and M.H.G collected the data, M.T.¹, M.H.G. and L.K. contributed to methodological development, M.T.¹, M.H.G, and M.T.² analyzed the data, M.T.¹, M.H.G, M.T.², and D.B. wrote and revised the manuscript.

Order of co-first authorship was determined by coin-flip.

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Rat schematic in Fig. 1A adapted with permission from SciDraw.io (doi.org/10.5281/zenodo.3926277, doi.org/10.5281/zenodo.3926237, <https://creativecommons.org/licenses/by/4.0/>)

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SUPPLEMENTARY FIGURES

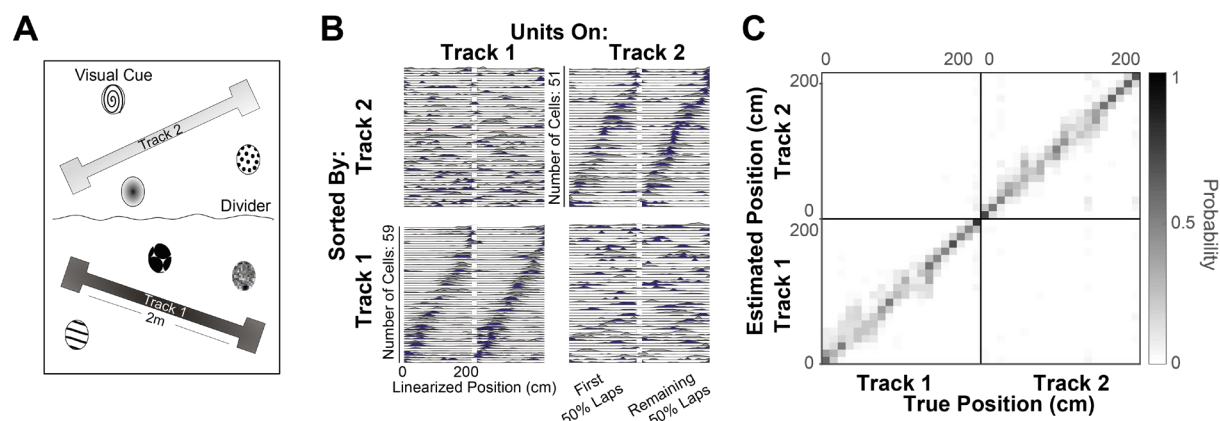


Figure S1. experimental setup, remapping and field stability. (A) Layout of the recording room - two linear tracks, each surrounded by visual cues and visually occluded from each other. (B) Ratemaps for Track 1 (left column) and Track 2 (right column) in an example session (rat 3, session 2) computed for the first 50% of laps (left half) and last 50% of laps (right half) and sorted by the ordered position of place fields on Track 1 (bottom) and Track 2 (top). (C) Confusion matrix for decoded position (posterior probability normalized over both tracks) for an example session (rat 3, session 2). Distribution of peak posterior probabilities (y-axis) is plotted at each True position during the behavioral episode (x-axis).

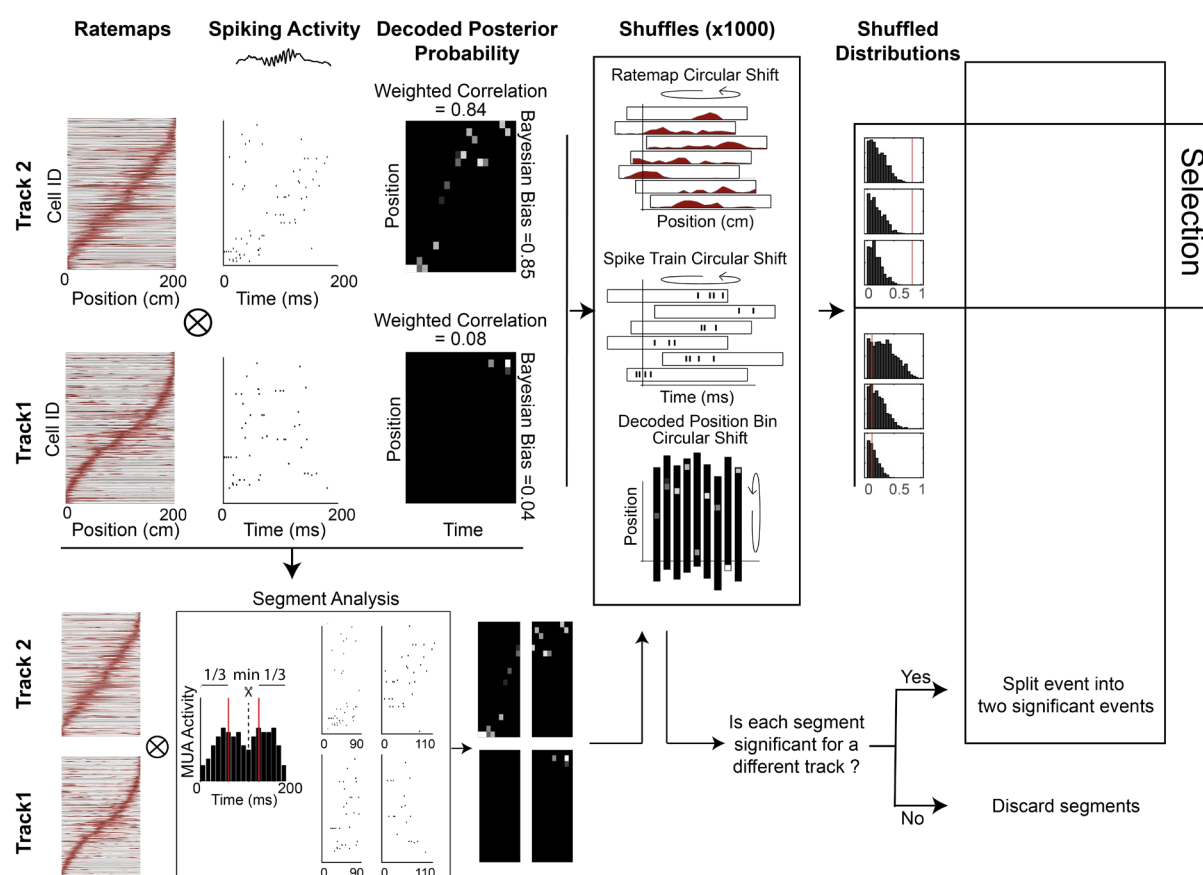


Figure S2. Analysis pipeline for detection of significant replay events. Detected replay spike trains were decoded using a Bayesian framework. Each track's ratemaps obtained during RUN were used to calculate the decoded posterior probabilities for each replay sequence (top left). In addition, all replay events were split into two shorter events based on the minimum in the MUA during the middle third of the replay event (bottom left, Segment analysis). For each replay event, weighted correlations were calculated on decoded posterior probability from both the whole replay sequence and each segment, and the three scores were compared to three different shuffles (ratemap circular shift, spike train circular shift, and decoded position bin circular shift, 1000 shuffles performed for each type). Only events with weighted correlation scores higher than 95% for every shuffle distribution (97.5% for split replay events) were considered statistically significant.

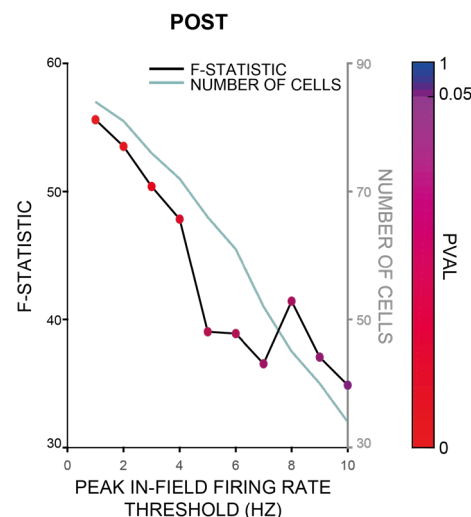


Figure S3. Significance of regression analysis using a stricter criterion of place cell activity. F-statistic of regression - peak in-field firing rate vs. peak replay rate (track difference) - for replay events occurring during POST rest, as the criterion for place cell activity (peak in-field firing rate) is progressively increased from 1 Hz. All regressions are statistically significant (up to and including using a threshold of 10 spk/s) The p-value of each regression is indicated by color. The number of cells used in each regression is indicated by the green line.

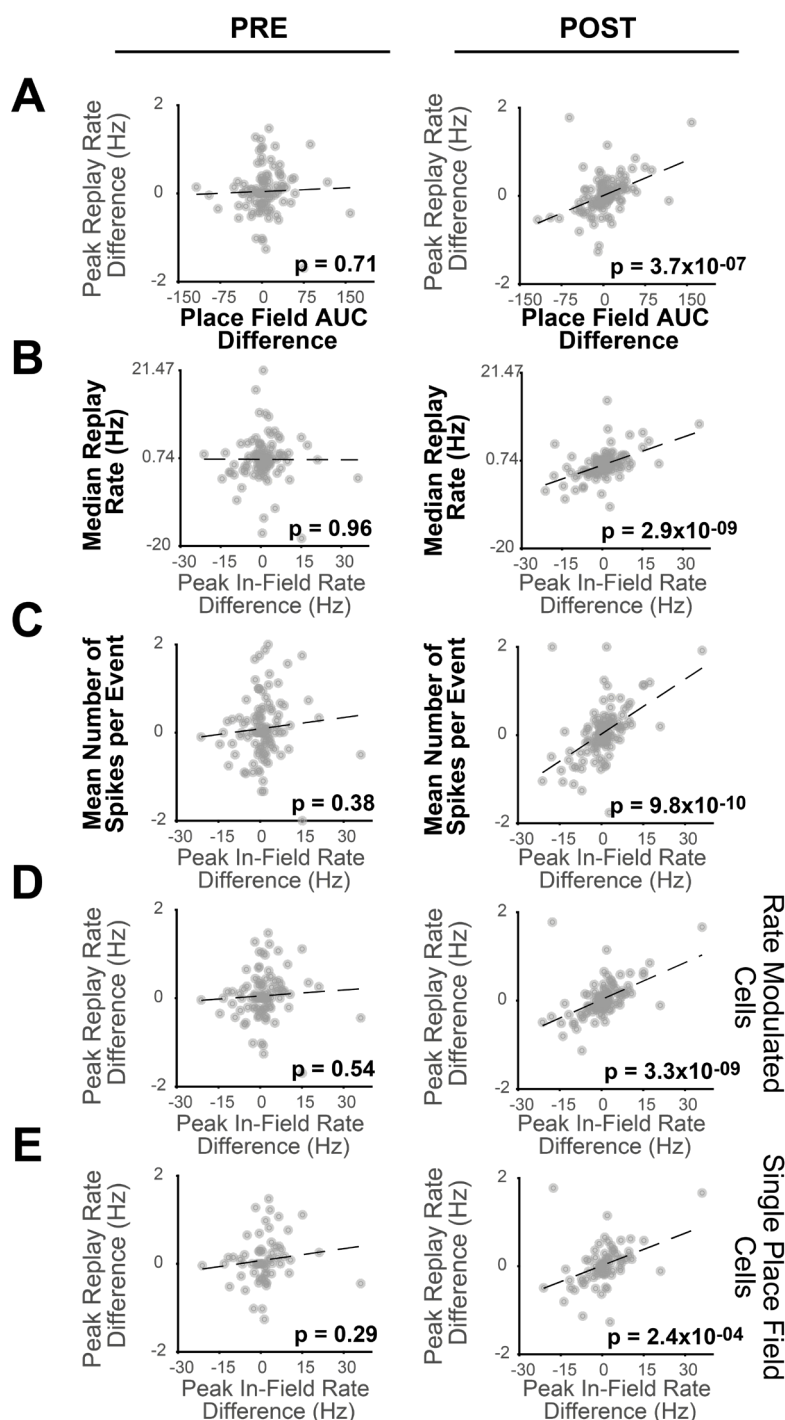


Figure S4. Significance of regression analysis with alternative place field and replay rate metrics, and stricter criteria for cell selection. F-statistic of regression - peak in-field firing rate vs. peak replay rate (track difference) - for replay events occurring during PRE (left column) and POST (right column) using alternative metrics and cell selection criteria. (A) Place field area under the curve (new x-axis, PRE: n = 106 cells, POST: n = 123 cells), (B) Median replay rate (new y-axis, PRE: n = 106 cells, POST: n = 123 cells), (C) Mean number of spike per event (new y-axis, PRE: n = 106 cells, POST: n = 123 cells), (D) Rate modulated cells (new criteria for cell selection, PRE: n = 97 cells, POST: n = 112 cells), (E) Single place field cells (new criteria for cell selection, PRE: n = 64 cells, POST: n = 77 cells).

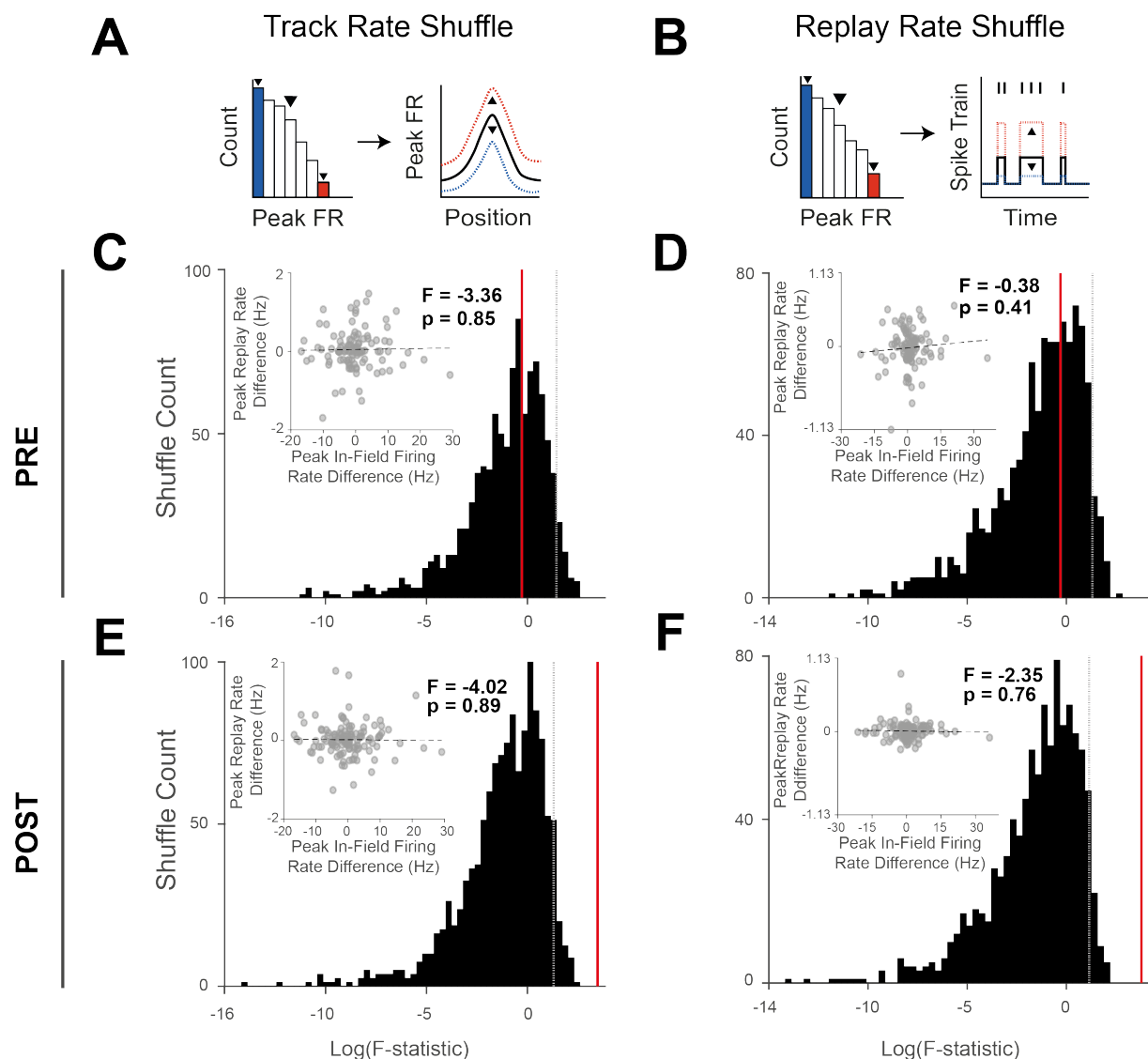


Figure S6. Re-computing the statistical significance of regression analysis relative to a shuffled distribution. A Track rate shuffle (A) and Replay rate shuffle (B) were used to randomize the x-axis and y-axis respectively and create F-Statistic distributions for measuring significance of the regression. (C-F) Distribution of shuffle, dashed gray line: $P < 0.05$, red line: F-Statistic of original data (main plot). One example of a regression plot within the shuffle distribution (inset plot). (C) PRE: Track rate shuffle. (D) PRE: Replay rate shuffle. (E) POST: Track rate shuffle. (F) POST: Replay rate shuffle.

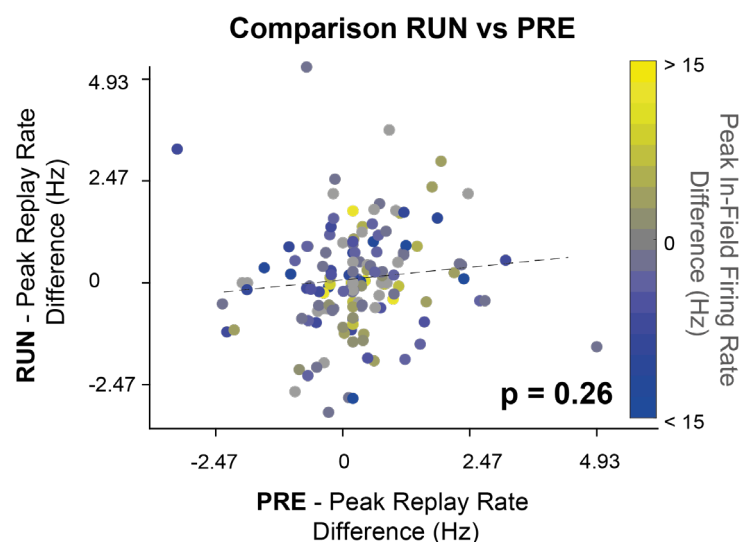


Figure S7. Between track differences in replay rate during PRE is not predictive of replay rate differences observed during RUN local replay events. Peak replay rate difference for replay events (RUN vs. PRE), color coded by peak in-field rate differences. A linear regression was not statistically significant ($P > 0.05$).

460 **SUPPLEMENTARY TABLES**

461 **Table S1: Overview of behavior, number of recorded cells and decoding accuracy for all sessions and rats.**

SESSION	TRACK	NUMBER OF LAPS	TIME RESTING (min)		CELLS			DECODING QUALITY (%)	
			PRE	POST	Total number of cells on track	Number of cells common to T1 and T2	Included cells in replay analysis	Classification accuracy	Local decoding accuracy
Rat 1	1	30	72	76	42	27	27	91	81
Rat 1	2	20	72	76	44	27	27	97	82
Rat 2, session 1	1	15	59	125	70	40	40	87	86
Rat 2, session 1	2	59	59	125	63	40	40	80	89
Rat 2, session 2	1	22	54	104	49	28	28	92	82
Rat 2, session 2	2	33	54	104	43	28	28	90	79
Rat 3, session 1	1	34	49	123	59	33	33	94	85
Rat 3, session 1	2	27	49	123	51	33	33	94	80
Rat 3, session 2	1	36	69	137	72	46	46	94	85
Rat 3, session 2	2	38	69	137	57	46	46	97	86

Table S2: Summary statistics of the z-scored log odd difference per session and across sessions for PRE, RUN and POST

	Session	Original			Rate fixed			Place removed			Rate fixed, Place randomized			Place removed, Rate randomized		
		Mean	SE	P value	Mean	SE	P value	Mean	SE	P value	Mean	SE	P value	Mean	SE	P value
PRE	1	-0.11		0.73	-0.02		0.56	-0.02		0.57	0.45		0.05	-0.30		0.92
	2	0.21		0.15	0.32		0.07	0.03		0.39	-0.21		0.80	-0.16		0.77
	3	-0.33		0.88	-0.54		0.96	-0.35		0.96	0.03		0.46	0.02		0.46
	4	0.15		0.32	-0.21		0.82	0.08		0.24	0.03		0.45	-0.15		0.73
	5	0.50		0.07	0.65		0.02	0.01		0.49	0.46		0.12	-0.25		0.73
	Mean	0.08	0.14	0.22	0.04	0.21	0.37	-0.05	0.08	0.64	0.15	0.13	0.11	-0.17	0.05	0.91
RUN	1	2.51		7.57×10^{-27}	2.13		9.24×10^{-26}	1.15		6.41×10^{-19}	0.01		0.48	-0.19		0.87
	2	3.22		9.61×10^{-36}	2.37		9.01×10^{-28}	1.39		2.05×10^{-28}	0.48		0.01	0.22		0.10
	3	2.55		5.93×10^{-23}	2.37		2.93×10^{-22}	1.27		5.08×10^{-18}	-0.19		0.78	0.12		0.22
	4	2.50		1.14×10^{-22}	2.44		1.38×10^{-22}	0.37		6.25×10^{-06}	-0.11		0.76	-0.15		0.79
	5	3.09		1.31×10^{-29}	2.26		4.38×10^{-25}	1.52		2.17×10^{-20}	0.20		0.16	0.09		0.29
	Mean	2.78	0.16	1.08×10^{-121}	2.31	0.06	1.03×10^{-101}	1.14	0.20	9.23×10^{-29}	0.08	0.12	0.18	0.02	0.08	0.43
POST	1	0.84		5.39×10^{-11}	0.43		1.09×10^{-05}	0.32		2.44×10^{-05}	0.17		0.09	0.15		0.18
	2	1.53		9.90×10^{-35}	1.03		1.96×10^{-24}	0.45		7.34×10^{-19}	0.33		0.003	0.10		0.15
	3	1.63		3.82×10^{-19}	1.44		3.30×10^{-16}	0.84		4.52×10^{-18}	-0.14		0.74	-0.25		0.98
	4	0.70		3.47×10^{-08}	0.78		5.46×10^{-11}	0.20		6.32×10^{-06}	0.20		0.03	0.09		0.20
	5	1.79		1.22×10^{-34}	1.59		9.58×10^{-30}	0.89		1.15×10^{-21}	-0.12		0.87	0.06		0.30
	Mean	1.30	0.22	1.03×10^{-91}	1.05	0.21	1.98×10^{-74}	0.54	0.14	8.12×10^{-15}	0.09	0.09	0.06	0.03	0.07	0.30

***Bold** values statistically significant*

466 Table S3: Summary statistics of the 98.75% confidence interval for the AUC difference between the original and the manipulated
 467 condition

		VS Original 98.75% confidence interval <i>BOLD values statistically significant from zero</i>		
		PRE	RUN	POST
Original VS rate fixed	Mean AUC Diff	-0.04	-0.02	-0.05
	SEM AUC Diff	0.04	0.01	0.02
	Lower CI	-0.15	-0.03	-0.09
	Upper CI	0.06	0.001	-0.01
Original VS Place removed	Mean AUC Diff	0.06	-0.19	-0.10
	SEM AUC Diff	0.04	0.02	0.02
	Lower CI	-0.04	-0.23	-0.14
	Upper CI	0.17	-0.15	-0.06
Original VS Rate fixed Place randomized	Mean AUC Diff	0.003	-0.45	-0.27
	SEM AUC Diff	0.04	0.02	0.02
	Lower CI	-0.10	-0.51	-0.31
	Upper CI	0.11	-0.40	-0.23
Original VS Place removed rate randomized	Mean AUC Diff	-0.01	-0.45	-0.24
	SEM AUC Diff	0.04	0.02	0.02
	Lower CI	-0.12	-0.50	-0.28
	Upper CI	0.09	-0.40	-0.19

468

469

470 Table S4: Summary statistics of the 98.75% confidence interval for the AUC difference between the negative control (Rate fixed place
471 randomized manipulation) and the other conditions

		VS Negative Control 98.75% confidence interval <i>BOLD values statistically significant from zero</i>		
		PRE	RUN	POST
Rate fixed VS Rate fixed Place randomized	Mean AUC Diff	-0.003	0.45	0.27
	SEM AUC Diff	0.04	0.02	0.02
	Lower CI	-0.11	0.40	0.23
	Upper CI	0.10	0.51	0.31
Place removed VS Rate fixed Place randomized	Mean AUC Diff	-0.05	0.44	0.22
	SEM AUC Diff	0.04	0.02	0.01
	Lower CI	-0.15	0.38	0.17
	Upper CI	0.06	0.49	0.27
Place removed VS Rate fixed Place randomized	Mean AUC Diff	0.06	0.26	0.17
	SEM AUC Diff	0.04	0.02	0.02
	Lower CI	-0.05	0.20	0.12
	Upper CI	0.17	0.33	0.22
Rate fixed, Place randomized VS Place removed rate randomized	Mean AUC Diff	-0.02	0.001	0.03
	SEM AUC Diff	0.04	0.03	0.02
	Lower CI	-0.11	-0.08	-0.02
	Upper CI	0.09	0.08	0.08

472

473

474 **Table S5: Summary statistics of the DeLong Test comparing the ROC curves between the original and the manipulated condition**

475

	DeLong Test versus Original (significance = 0.05/4 = 0.0125) (Bonferroni corrected) <i>BOLD values statistically significant</i>			
	Original VS Rate fixed	Original VS Place removed	Original VS Rate fixed place randomized	Original VS Place removed rate randomized
PRE	0.15	0.15	0.91	0.78
RUN	0.007	5.56×10⁻³⁴	2.28×10⁻⁹⁸	6.92×10⁻¹⁰⁰
POST	8.15×10⁻⁰⁸	6.33×10⁻¹²	2.65×10⁻⁵⁵	4.39×10⁻⁴³

476

477

478 **Table S6: Summary statistics of the DeLong Test comparing the ROC curves between the original and the manipulated condition**

479

	DeLong Test versus Negative control (significance = 0.05/4 = 0.0125) (Bonferroni corrected) <i>BOLD values statistically significant</i>			
	Rate information fixed VS Rate fixed place randomized	Place removed VS Rate fixed place randomized	Place information removed VS Rate fixed place randomized	Rate fixed, Place randomized VS Place removed rate randomized
PRE	0.91	0.23	0.18	0.70
RUN	2.28×10⁻⁹⁸	7.14×10⁻⁸⁸	1.17×10⁻²³	0.96
POST	2.65×10⁻⁵⁵	1.15×10⁻³⁵	2.03×10⁻²⁰	0.09

480

METHODS

Animals

Three male Lister-Hooded rats (350-450g) were implanted with a microdrive with 24 independently moveable tetrodes. Prior to surgery, rats were kept at 90% of their free-feeding weight and housed in pairs on a 12-hour light/dark cycle, with 1 hour of simulated dusk/dawn. All experimental procedures and postoperative care were approved and carried out in accordance with the UK Home Office, subject to the restrictions and provisions contained within the Animal (Scientific Procedures) Act of 1986.

Surgery

Animals were deeply anaesthetized under isoflurane anesthesia (1.5-3% at 2L/min) and implanted with a custom-made microdrive array carrying 24 independently moveable tetrodes (modified from microdrive first published by (12)). Each tetrode consisted of a twisted bundle of four tungsten microwires (12µm diameter, Tungsten 99.95% CS, California Fine Wire), gold-plated to reduce impedance to < 200kΩ. One rat was implanted with a dual-hippocampal microdrive targeting both dorsal hippocampal CA1 areas (AP: -3.48mm, ML: +/-2.4mm from Bregma), each output carrying 12 tetrodes. The two remaining rats were implanted with a microdrive targeting the right dorsal hippocampal CA1 area (AP: 3.72mm, ML: 2.5mm from Bregma) and the left primary visual cortex (AP: -5.76mm, ML: -3.8mm from Bregma), using 16 and 8 tetrodes respectively. After surgery, animals were housed individually and allowed to recover with food and water *ad libitum* for a week before returning to being kept at 90% of their free-feeding weight.

Experimental design

A given recording session started with a 1-hour rest period in which the rats were placed in a quiet, remote location (rest pot), to which they had been previously habituated. The rest pot consisted of a black circular enclosure of 20cm of diameter, surrounded by a 50cm tall black plastic sheet that isolated them from the surroundings. Following the rest period, the rats were exposed to two novel 2m linear tracks in which they were allowed to run back and forth for 15 min, except for one session in which the animal ran for 30 min in the second track. Liquid reward was dispensed at each end of the track (0.1mL chocolate flavored soy milk) to encourage the animals to traverse the entirety of the track. In all except one session, the exposures to the two tracks were separated by a 10min rest period in the rest pot. The recording session finished with a final 2-hours rest period inside the rest pot.

To simulate novel environments, the shape of the tracks was changed between recording sessions and their surfaces covered with different textured fabrics. In each session, the room was surrounded by black curtains with different high contrast visual cues. The tracks were separated using view-obstructing dividers.

519

520 Spike detection and unit isolation

521 Spiking data was sorted using the semi-automatic clustering software KlustaKwik 2.0
522 (K.Harris, <http://klustakwik.sourceforge.net/>) and then manually curated with either Phy-GUI
523 (<https://github.com/kwikteam/phy>) or Klustaviewa (<https://github.com/klusta-team/klustaviewa>).
524 Putative single units were discriminated based on the spike waveform, a clean inter-spike interval,
525 and their stability across the recording session. The rest of the clustered activity was classified in
526 either multi-unit activity or noise.

527

528 LFP analysis

529 The power spectral density (PSD) of the LFP was calculated using Welch's method (*pwelch*,
530 MATLAB) to identify the channels with higher power for theta (4-12Hz) and ripple (125-
531 300Hz) oscillations, as well as the channel with the largest difference in normalized theta to
532 ripple power. The LFP of selected channels was next downsampled from 30 kHz to 1 kHz and
533 band-passed filtered (MATLAB command *filtfilt*). The instantaneous phases were estimated
534 using Hilbert transform.

535

536 Criteria for Place cell selection

537 Putative principal cells were identified by selecting units with a Half Width Half Max
538 (HWHM) larger than 500 μ s and mean firing rate <5Hz across the entire recording session. For
539 place cell classification, spike trains were speed-filtered to only include the spiking activity
540 between 4cm/s and 50cm/s. A principal cell was classified as a place cell according to the
541 following criteria: (1) the minimum peak firing rate was >1Hz in the unsmoothed ratemap and
542 (2) a stable spiking activity across the first half and second half of the exploration on the track.
543 Only place cells satisfying these two criteria for each linear track independently, were included
544 in the regression analysis (Fig. 1-3) and log odd analysis (Fig. 4).

545 To generate firing ratemaps (the spike histogram divided by the total dwell time at each position
546 bin), the position data was discretized in 2cm bins for visualization and plotting, and 10 cm
547 bins for Bayesian decoding. Only raw (unsmoothed) ratemaps were used for all Bayesian
548 decoding analyses.

549

550 Population vector analysis (PPV)

551 A population vector analysis was used to assess the ratemap stability for each track as well as
552 the degree of between track remapping (29). For each session, the linearized ratemaps of all
553 recorded pyramidal cells with a firing rate > 1Hz were stacked into a 20 position bins x N cells
554 matrix for each track. To assess remapping between tracks, the ratemaps computed from the
555 entire track experience were used, while to assess ratemap stability within a track, the middle

lap was used as a reference to compute separate ratemaps for the first and second half of the behavioral episode. The population vector of rates at each position bin was then correlated with its counterpart vector: either on the other track or during the other half of the experience. The mean Pearson correlation value was first averaged across position bins and then across sessions.

Bayesian Decoding

A naïve Bayesian decoding algorithm was applied to reconstruct the animal's spatial trajectory during both behavior and replay events from CA1 hippocampal spiking activity (Zhang et al., 1998):

$$P(x|n) = CP(x) \left(\prod_{i=1}^N f_i(x)^{n_i} \right) \exp \left(-\tau \sum_{i=1}^N f_i(x) \right)$$

where P is the probability of the animal being at a specific position given the observed spiking activity, C is a normalization constant, x is the subject's position, $f_i(x)$ is the firing rate of the i^{th} place field at a given location x, and n is the number of spikes in the time window τ . The normalization constant was re-defined as the summed posterior probabilities across both tracks. The maximum probability of the subject's position was decoded using 250ms and 20ms time bin during behavior and replay, respectively.

The decoding error was defined as the differences between the real location of the animal and the estimated position with maximum-likelihood.

Candidate replay event detection

Candidate replay events were detected based on MUA and ripple power. MUA was first smoothed with a Gaussian Kernel (sigma = 5 ms) and binned into 1 ms steps. Only MUA bursts with a maximum duration of 300 ms and z-scored activity over 3 were selected. Ripples LFP signal was smoothed with a 0.1 s moving average filter and a threshold over 3 was set for the z-scored ripple power.

Candidate replay events passing both thresholds were next speed-filtered (above 5 cm/s), and discarded if their sequence had less than 5 different units active or if their duration was below 100 ms or over 750 ms (thus, candidate events had at least 5 consecutive 20 ms bins). Events detected within 50 ms were combined. In an effort to optimize detection of replay events and avoid a minority of events that were discarded due to noisy probability decoding at the beginning or the end of the event, candidate replay events were split in half. To do so, the minimum MUA activity in the middle third of the candidate replay event was used to determine a natural midpoint to split the event in two segments. Both segments were decoded and tested for significance independently following the same procedure as the 'intact' candidate replay

events (i.e., same criteria including minimum duration, etc.), for the exception of an adjusted p-value threshold ($P < 0.025$).

Replay events were classified as rest or awake local replay. Awake replay was defined as replay events occurring while the animals were running on either of the tracks and classified as local if the content of the replay event reflected the current track on which the animal was running. Replay events detected during sleep or rest periods within the rest pot were classified as rest replay events.

Replay events scoring and significance

Replay events were scored using weighted correlation between decoded posterior probabilities across position and time.

Weighted mean:

$$m(x; prob) = \frac{\sum_{i=1}^M \sum_{j=1}^N prob_{ij} x_i}{\sum_{i=1}^M \sum_{j=1}^N prob_{ij}}$$

Weighted covariance:

$$cov(x, t; prob) = \frac{\sum_{i=1}^M \sum_{j=1}^N prob_{ij} (x_i - m(x; prob))(t_j - m(y; prob))}{\sum_{i=1}^M \sum_{j=1}^N prob_{ij}}$$

Weighted correlation:

$$corr(x, t; prob) = \frac{cov(x, t; prob)}{\sqrt{cov(x, x; prob) cov(t, t; prob)}}$$

Where x_i is the i^{th} position bin, t_j is the j^{th} time bin and $prob_{ij}$ is the probability at the position bin i and time bin j .

The statistical significance of the weighted correlation for each candidate replay event was assessed by comparison with three different 1000 shuffle distributions:

1) Spike train circular shift, in which the spike count vectors for each cell were independently circularly shifted in time within each replay event, prior to decoding.

2) Place field shift, in which each ratemap was circularly shifted in space by a random amount of position bins prior to decoding.

3) Circular shift of position, in which posterior probability vectors for each time bin were independently circularly shifted by a random amount.

If the score of the candidate event was greater than the 95th percentile of the distribution for all three shuffles then the event was considered to be significant.

Reinstatement of rate modulation analysis

Linear regression was used to measure the reinstatement of rate modulation between place fields on the track and replay events (MATLAB function *fitlm*). For each place cell active in both tracks, rate modulation on the track was measured by calculating (1) the difference in peak in-field rate across tracks, and (2) the difference in the area under the curve (AUC; MATLAB function *trapz*). Place fields with a peak in-field rate < 1 Hz on either track were excluded. As a control, a variation of the analysis was done by excluding all cells with multiple place fields on the same track. Multiple peaks for a single cell were considered to be different place fields when their peak firing rate > 1 Hz, each place field width was over 2 spatial bins (20 cm) and the distance between the peaks was of 4 spatial bins (40 cm).

Rate modulation during replay events was calculated as the change in each cell's firing activity between the replay events encoding for each track. Included cells had to participate in at least one replay event for each track. The firing rate change was measured as (1) the difference in the average peak instantaneous rate across tracks, (2) the difference in the median rate across replay events, and (3) the difference in the mean number of spikes across replay events.

To obtain the peak instantaneous rate of each event, each cell's spike train was binned into 1ms steps, filtered with a 100ms long Gaussian window with $\sigma=20\text{ms}$ (MATLAB command *filter*), and the peak amplitude of the resulting signal measured.

Controls for replay detection analysis

Two types of shuffles were applied to the data before repeating the detection of replay events using Bayesian decoding: a Track rate shuffle and Replay rate shuffle. The Track rate shuffle consisted in the randomization of the firing rate of the place fields within a track, while the Replay rate shuffle randomized the cells' firing rate during the replay events. For both shuffles, each place cell's firing rate was scaled based on a value randomly drawn from a distribution of firing rates obtained from the analyzed place cells on the track (for the Track rate shuffle) or during the replay events (for the Replay rate shuffle). The shuffled data was then used to repeat the detection of replay events, and the newly identified significant replay events were used in a linear regression comparing their original spike content (instead of the shuffled one) against the original differences in the peak in-field rate.

Remapping classification

A bootstrapping procedure was used to identify the rate and place modulation properties of each cell. Modulation was defined when the track difference in a place field's specific parameter (e.g. peak in-field firing rate or peak location) exceeded the intrinsic variability of such parameter on a single track. The distribution of each parameter was computed for both tracks by calculating firing ratemaps from a randomly sampled N out of N laps (1000 iterations,

with replacement). The median value of a track distribution was then compared to the 5th/95th percentiles of the other track, and vice versa. If either median exceeded either percentile, the cell was classified as being modulated between tracks for that parameter.

Sequenceless Bayesian Decoding

The sequenceless Bayesian decoding was modified from Carey et al., (2019). Only cells with stable place fields on the track in the first and second half of the behavioral episode (peak rate > 1Hz) were included in our sequenceless decoding analysis.

Before Bayesian decoding, ratemaps for Track 1 and Track 2 were concatenated as a single matrix [nCell x (nPosition(T1) + nPositions(T2))]. Unless otherwise mentioned, 20ms time bins and 10cm position bins were used for decoding. After decoding, the posterior probabilities for each time bin were normalized across all position bins on both tracks to sum to 1. Then, the summed posterior probability across all time bins within each replay event was computed, and the probabilistic bias towards Track 1 or Track 2 for each replay event was quantified as $\log\left(\frac{\sum Prob(Track_1)}{\sum Prob(Track_2)}\right)$ (log odds), where $\sum Prob(Track_1)$ and $\sum Prob(Track_2)$ referred to the summed posterior probabilities for Track 1 and Track 2 respectively. To remove any intrinsic probabilistic bias, each raw log odd calculation was then z-scored relative to a shuffled log odd distribution (Ratemap track label shuffle), obtained by randomly permuting each cell's Track 1 and Track 2 ratemaps (either keeping a cell's ratemap unaltered or performing a between track swap) 1000 times for each replay event.

To test decoding using place or rate representation alone, the original data were manipulated as the following:

- Rate fixed manipulation: Rate information was fixed by rescaling each cell's peak in-field firing rate to the mean value across both tracks, and its spike count in each replay event to the average firing rate across all replay events.

- Place removed manipulation: The place (and sequence) information was removed by decoding with only a single position bin, which encompassed the entire track (mean firing rate on each track when speed is above 5cm/s) and a single time bin, which spanned the entire replay event.

Two negative controls were included to disrupt both rate and place information:

- Rate fixed place randomized manipulation: After fixing rate information, place information was randomized using cell ID shuffle in which all cells were randomly reassigned to a different cell's ratemap for each replay event.

- Place removed rate randomized manipulation: After removing place information, rate information was randomized by scaling the firing rate on each track based on a value randomly drawn from a distribution of firing rates obtained from the analyzed place cells on the track.

ROC analysis

The ‘ground truth’ track identity for each replay event was assigned based on the sequenceness of the replay determined by weighted correlation and compare to three shuffled distributions. The binary discriminability of a replay event’s identity (Track1 or Track 2) during PRE, POST or RUN was quantified using the Receiver Operative Characteristic (ROC) curve. By systematically shifting the discrimination threshold along the z-scored log odd distributions for Track 1 replay events and Track 2 replay events, a range of true positive rates and corresponding false positive rates at each threshold was used to construct the ROC curve. A trapezoidal approximation was used to estimate the area under each ROC curve (MATLAB function *perfcurve* for both constructing ROC curve and estimating area under the curve). The bootstrapped distributions of ROC curves and the area under the corresponding ROC curves were produced by resampling with replacement 1000 times.

Statistics

Z-scored Log Odd significance

The statistical significance of the z-scored log odd difference between Track 1 and Track 2 was determined by comparison with a shuffled distribution in which replay events were resampled with replacement 1000 times and their track identities were randomly assigned. The single z-scores were computed relative to the shuffled distributions, after which the associated one tailed p-value was calculated. The log odd difference was considered significantly different from the shuffled distribution at the significance level of 0.05.

Bootstrapping for ROC significance

To determine if rate and/or place representation manipulations significantly changed the area under the curve (AUC) of the ROC curves, we calculated the confidence interval for the difference between each condition’s bootstrapped AUC distribution and that of the original data. This was repeated later replacing the original data by one of the negative controls (rate fixed or place randomized manipulation). The 98.75% confidence interval was used to account for four comparisons ($1-(\alpha/n)$, where $\alpha=0.05$ and $n = 4$) within the same behavioral epochs. The mean difference between two bootstrapped AUC distributions were only considered statistically significant when the confidence interval did not overlap with 0.

Delong Test for ROC

To determine if rate and/or place manipulations statistically significantly changed the AUC of ROC curves within the same behavioral epochs, the AUC of each condition was compared with the AUC of the original data or the negative control (rate fixed, place randomized). The DeLong Test (37) was employed to perform pair-wise comparisons between two ROC curves, a non-parametric test that accounted for the correlated nature of the data from the same epoch. The DeLong variance-covariance matrix for two ROC curves was computed using MATLAB-

739 based algorithm [<https://github.com/PamixSun/DeLongUI>] developed by Sun and Xu (2014)
740 (36).

741 A z-score was calculated based on the following equation:

$$742 \quad z = \frac{|AUC_A - AUC_B|}{\sqrt{var(A) + var(B) - 2cov(A, B)}}$$

743 The p value threshold for the statistical significance of the z-score was Bonferroni corrected to
744 account for multiple comparisons.

745

746 Code Availability

747 All analysis was carried out using MATLAB (Mathworks, Natick MA) and all code is available
748 from the authors upon reasonable request.