Non-feature-specific elevated responses and feature-specific backward replay in human brain induced by visual sequence exposure

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ABSTRACT

The ability of cortical circuits to adapt in response to experience is a fundamental property of the brain. After exposure to a moving dot sequence, flashing a dot as a cue at the starting point of the sequence can elicit successive elevated responses even in the absence of the sequence. These cue-triggered elevated responses have been shown to play a crucial role in predicting future events in dynamic environments. However, temporal sequences we are exposed typically contain rich feature information. It remains unknown whether the elevated responses are feature specific and, more crucially, how the brain organizes sequence information after exposure. To address these questions, participants were exposed to a predefined sequence of four motion directions for about 30 min, followed by the presentation of the start or end motion direction of the sequence as a cue. Surprisingly, we found that cue-triggered elevated responses were not specific to any motion direction. Interestingly, motion direction information was spontaneously reactivated, and the motion sequence was backward replayed in a time-compressed manner. These effects were observed even after brief exposure. Notably, no replay events were observed when the second or third motion direction of the sequence served as a cue. Further analyses revealed that activity in the medial temporal lobe (MTL) preceded the ripple power increase in visual cortex at the onset of replay, implying a coordinated relationship between the activities in the MTL and visual cortex. Together, these findings demonstrate that visual sequence exposure induces two-fold brain plasticity that may simultaneously serve for different functional purposes. The non-feature-specific elevated responses may facilitate general

48   processing of upcoming stimuli, whereas the feature-specific backward replay may

49   underpin passive learning of visual sequence.

INTRODUCTION

The capacity of cortical circuits to undergo plasticity in response to experience is a fundamental feature of the brain (Buonomano and Merzenich, 1998; Costandi, 2016; Li, 2016). This plasticity can be induced not only by active, task-dependent training, such as visual perceptual learning (Watanabe and Sasaki, 2015; Chen et al., 2016; Lu and Dosher, 2022), but also by passive, repetitive exposure (Gutnisky et al., 2009; Sasaki et al., 2010; Ekman et al., 2017). For instance, after repeated exposure to a moving dot sequence, flashing a dot (i.e., cue) at the starting point of the sequence elicits successive elevated neural responses in visual cortex, akin to those induced by the actual moving dot sequence in both humans (Ekman et al., 2017, 2023; Lu et al., 2020) and animals (Eagleman and Dragoi, 2012; Xu et al., 2012). This cue-triggered reactivation is thought to be driven by expectations and is prediction-related (Ekman et al., 2017, 2023), as it facilitates the prediction of upcoming stimuli and influences the perception of sensory information (Gutnisky et al., 2009; Baker et al., 2014; Pojoga et al., 2020).

Unlike the simple white dot sequences used in previous studies (Xu et al., 2012; Ekman et al., 2017, 2023; Lu et al., 2020), temporal sequences to which we are exposed in daily life usually contain rich feature information. However, it remains unknown whether cue-triggered elevated responses are feature-specific. On the one hand, if these responses are not feature-specific, they may reflect a general state of cortical readiness for any upcoming stimuli. On the other hand, if these responses are indeed specific to a particular feature in the sequence, they would selectively facilitate

the processing of specific future events. This aligns with the view that expectation

sharpens neural tuning and enhances the processing of expected stimuli (Kok et al.,

2012). For instance, expectation has been shown to elevate the prestimulus baseline

activity of sensory neurons tuned to expected stimuli (Wyart et al., 2012; Kok et al.,

2014) and to preactivate stimulus templates in both the visual (Kok et al., 2017) and

auditory (Demarchi et al., 2019) cortices. Notably, however, all these prediction-related,

feature-specific activities were observed in static contexts. Whether prediction-related

responses in dynamic temporal contexts (e.g., visual sequences) are feature-specific

remains unclear.

After exposure to a feature-contained temporal sequence, another important

question is how the feature information in the sequence is encoded and organized in

neural activity following the cue. Previous studies have demonstrated that memory

consolidation (Carr et al., 2011; Gridchyn et al., 2020; Gillespie et al., 2021) or learning

process (Igata et al., 2021; Liu et al., 2021b, 2022) following training with a temporal

sequence is associated with a neural phenomenon known as replay. Replay refers to

the sequential reactivation of neural activity patterns associated with the trained

sequence in both sleep (Lee and Wilson, 2002) and awake (Foster and Wilson, 2006)

states. During replay, the neural representation of the trained sequence is temporally

compressed and can occur in either a forward or backward direction (Carr et al., 2011;

Joo and Frank, 2018; Nour et al., 2021; Liu et al., 2022; McFadyen et al., 2023). These

replay events frequently coincide with sharp-wave ripples (SWRs), which are high-

frequency (150–220 Hz) oscillations (O'Keefe and Nadel, 1978; Bush et al., 2022)

detected in hippocampal local field potentials (Buzsáki, 1986, 2015). To date, replay has

been observed in tasks involving sequences, such as nonlocal reinforcement learning

(Liu et al., 2021b) and episodic memory retrieval (Wimmer et al., 2020). It remains

unclear whether simple visual exposure to temporal sequences could also trigger replay

events in humans.

In the current study, we used magnetoencephalography (MEG) to investigate

whether cue-triggered elevated brain responses following visual sequence exposure are

feature-specific and, more importantly, to determine how feature information in the

exposed sequence is encoded and organized in the brain after exposure. To address

these questions, participants were initially exposed to a predetermined motion

sequence. We then decoded motion direction information during the blank period

following the presentation of either the first or last motion direction in the sequence as a

cue. Surprisingly, we found that cue-triggered elevated responses were not specific to

any motion direction. Interestingly, motion direction information was spontaneously

reactivated during the blank period, and the motion sequence was backward replayed in

a time-compressed manner. This backward replay was identified even after brief

exposure. However, neither forward nor backward replay was detected when an

intermediate motion direction in the sequence was presented as a cue. Lastly, MEG

source reconstruction analysis revealed that medial temporal lobe (MTL) activation

preceded visual cortical activation, implying that the replay events observed in visual

cortex may be triggered by activities in the MTL.

RESULTS

The visual stimuli used in the study were four random-dot kinematograms

(RDKs). All dots in an RDK moved in a single direction (i.e., 0°, 90°, 180°, or 270°) with

100% coherence. In Experiment 1, we included three trial conditions: full sequence

trials, start-only trials, and end-only trials (**Figure 1A**). In a full sequence trial,

participants were exposed to a predefined sequence of the four RDKs presented either

clockwise (i.e., 0° → 90° → 180° → 270°) or counterclockwise at the center of the

screen, with an inter-stimulus interval (ISI) of 300 ms between every two RDKs. In a

start- or end-only trial, the first or last RDK of the sequence was presented at the

beginning of the trial, followed by a blank period of 2.4 s. Participants were instructed to

complete three successive phases: functional localizer phase, exposure phase, and

main phase (**Figure 1B**). The functional localizer data were used to train models to

decode each motion direction in the sequence. During this phase, one of the four RDKs

was randomly presented for 1 s in each trial. During the exposure phase, participants

were exposed only to full sequence trials for about 30 min. In the main phase, 50%,

25%, and 25% of all trials were full sequence, start- and end-only trials, respectively.

The full sequence trials served as a topping-up exposure to maintain the exposure

effect.

Cue-triggered elevated responses are not feature-specific

We first measured the event-related field (ERF) activity evoked by RDKs in the

three trial conditions using all occipital gradiometer sensors (see Methods). **Figure 1C**

shows the evoked responses for full sequence trials (left panel, cluster-based

permutation test with a cluster forming threshold of t > 3.1 and 5000 permutations).

Remarkably, start- and end-only trials elicited similar wave-like responses as the full

sequence trials, despite the absence of stimuli following the first RDK (**Figure 1C**,

middle and right panels, cluster-based permutation test with a cluster forming threshold

of t > 3.1 and 5000 permutations). To further quantify the ERF peak amplitudes in the

three conditions while mitigating baseline confounds, we calculated each peak

amplitude using the 300 ms blank period just preceding the onset of the corresponding

RDK as the baseline (see Methods). We found that the four successive RDKs in the full

sequence trials evoked four comparable peaks after stimulus onset (**Figure 1C**, inset

figure in the left panel; two-tailed t-test; all $ts_{(17)} > 6.8072$, all $ps < 10^{-5}$). Interestingly, in

start- and end-only trials, significant peaks were still observed during the periods

corresponding to the intervals of the second and third RDKs in the full sequence trials

(**Figure 1C**, inset figures in the middle and right panels; start-only condition, two-tailed t-

test; all $ts_{(17)} > 4.5790$, all $ps < 10^{-3}$; end-only condition, two-tailed t-test; all $ts_{(17)} >$

6.0140, all $ps < 10^{-4}$). However, no significant peak was observed during the period

corresponding to the interval of the last RDK (start-only condition, two-tailed t-test; $t_{(17)} =$

1.3221, $p = 0.2036$; end-only condition, two-tailed t-test; $t_{(17)} = 0.2473$, $p = 0.8076$).

These results demonstrate cue-triggered elevated neural responses following visual

exposure to the RDK sequences, resembling previous studies using simple white dot

sequences (Ekman et al., 2017; Lu et al., 2020).

Figure 1. Stimuli, experimental procedure and evoked responses in Experiment 1

(A) Participants were presented with RDKs in three conditions. In the full sequence trial condition, four successive RDKs were presented. In the start- or end-only condition, only the first or last RDK in the sequence was presented at the beginning of the trial. **(B)** During MEG scanning, the participants were presented with two functional localizer runs (i.e., functional localizer phase) before providing any sequence information. Next, they were exposed with four full sequence runs (i.e., exposure phase). Finally, in the main

phase, full sequence, start- and end-only trials were presented in a pseudo-randomized

order. **(C)** Evoked neural responses as a function of time relative to trial onset in the full

sequence, start- and end-only trial conditions, respectively. Bold black lines at the

bottom indicate temporal clusters in which they reached significance when compared to

the pre-stimulus baseline. Inset figures at the top-right corner show the peak amplitudes

during the four corresponding RDK intervals after baseline correction.

Given that we observed elevated responses even in the absence of stimuli

following the cue, we next examined whether these responses were specific to a

particular feature (i.e., motion direction). Specifically, we asked whether motion

directions could be successfully decoded in start- and end-only trials, particularly during

the blank periods corresponding to the intervals of the second and third RDKs in full

sequence trials. To this end, we applied a time-resolved decoding analysis. For each

participant, we trained a one-versus-rest Lasso logistic regression model using the

functional localizer data to classify the neural activity pattern elicited by each motion

direction in the main phase. MEG signals from 72 occipital sensors were selected as

features for the training model.

To validate the reliability of our model, we first used a leave-one-out cross-

validation scheme on the localizer data to independently estimate decoding accuracy at

each time point. At the group level, decoding accuracies peaked at 411 ms, 464 ms, 444

ms, and 434 ms after stimulus onset for motion directions 0° (55.37% ± 1.21), 90°

(58.10% ± 1.14), 180° (54.07% ± 1.47), and 270° (53.64% ± 1.34), respectively. There

were no significant differences among the four motion directions in terms of either

latency (F(3, 51) = 0.375, p = 0.7716, η_p^2 = 0.0159) or decoding accuracy (F(3, 51) =

2.757, p = 0.0517, η_p^2 = 0.091) at the peak time point. Next, we trained a model using

the localizer data averaged between 100 ms and 500 ms after stimulus onset. The

trained model was then applied to the MEG signals recorded in the three trial conditions

in the main phase. Finally, we calculated the decoding probability (Liu et al., 2019; Nour

et al., 2021; Turner et al., 2023) for each motion direction at each time point at the group

level (**Figure 2**). Here, decoding probability for each motion direction reflects the

likelihood that the decoded stimulus had a specific motion direction at a given time point

(see Methods). Therefore, it provides a time-resolved decoding preference for each

motion direction, rather than only a single decoded label (e.g., 0°, 90°, 180°, or 270°).

In full sequence trials, the motion direction information could be successfully

decoded, as evidenced by the highest decoding probability during the interval

corresponding to the presentation of the respective RDK (**Figure 2A and 2B**, left

panels). In start- and end-only trials, we could also reliably decode the motion direction

of the first RDK (i.e., the cue) after stimulus onset (**Figure 2A and 2B**, middle and right

panels, only the decoding probability of the first RDK surpass the peak-level

significance threshold obtained from a nonparametric permutation test, FWE corrected

across time). Surprisingly, however, subsequent motion direction information was

absent at the group level during the post-cue blank period, where the cue-triggered

elevated responses were previously observed (i.e., 0.4 – 2.8 s after stimulus onset).

Together, these results demonstrate that the cue-triggered elevated response induced

by visual sequence exposure was not consistently specific to a particular feature across

participants and trials.

Figure 2. Time-resolved decoding probability for each motion direction

(A) Visualization of the time-resolved decoding probability in the three trial conditions. Each row shows the decoding probabilities for the four motion directions at that time point and each column indicates one of the four motion directions. **(B)** The line plots of the time-resolved decoding probability for the three trial conditions. Each colored line shows the time course of the decoding probability for each motion direction. For the start- and end-only conditions, we calculated the permutation threshold estimated by randomly shuffling of the labels and re-decoding, only the decoding probability of the cue surpassed the threshold.

Time-compressed backward replay of exposed motion sequence

How is the motion direction information encoded and organized in the brain

during the post-cue blank period? Clearly, the motion direction representation is not

time-locked to the onset of the cue in either start- or end-only trials. However, it is

possible that individual motion directions are encoded in the MEG signals in a

spontaneous way but are sequentially organized (Liu et al., 2019).

To test this hypothesis, we trained four decoding models using the localizer data

to capture the neural features of the four motion directions. We employed four decoding

models because our aim was to build feature-specific classifiers, each sensitive to only

one motion direction. These classifiers were designed to quantify the evidence of

feature-specific sequence in subsequent analyses. The models used a one-versus-rest

Lasso logistic regression algorithm, with MEG signals from 72 occipital sensors as

features (**Figure 3A**). For each participant and motion direction, we then selected the

time point with the highest decoding accuracy, which was estimated during the

validation of the functional localizer data, as the optimal time point. These optimal time

points were chosen because they are believed to carry the richest feature information

(Mo et al., 2019). For subsequent analyses, we obtained four classifiers that were

trained on the localizer data at their respective optimal time points. Each classifier

yielded significant decoding probability only when the stimulus matched the motion

direction it was trained to detect (**Figure 3B**). We did not find any spatial correlation

between any two trained classifiers (highest correlation $r < 0.12$, Figure S1A). Since the

optimal time point varied across participants and motion directions, we circularly shifted

the decoding probabilities over time and aligned them to a common time point

(arbitrarily set to 200 ms after stimulus onset) for visualization at the group level (**Figure**

3B).

Having established the classifiers for each participant and motion direction, we

then applied the classifiers to the MEG signals during the post-cue blank period in start-

and end-only conditions to estimate the reactivation probability (i.e., decoding

probability) for each motion direction at each time point (**Figure 3C;** Liu et al., 2019;

Wimmer et al., 2020; Nour et al., 2021). Example trials for the start- and end-only

conditions are shown in Figure S2. The results revealed that motion direction

reactivations occurred sparsely during the post-cue blank period, rather than crowded

within intervals corresponding to the RDK presentations in the full sequence condition,

suggesting that motion direction information might be reactivated in a spontaneous way.

Next, we applied temporally delayed linear modeling (TDLM) to quantify whether and

how these spontaneous reactivations followed the order of the exposed sequence

(**Figure 3D**). This algorithm includes two-level regression analyses; a first-level

regression that quantify evidence for each pairwise transition (e.g., $0° \rightarrow 90°$), and a

second-level regression that evaluates the extent to which reactivation patterns align

with a particular order of interest. Finally, we defined "sequenceness" as a metric of

forward (i.e., $0° \rightarrow 90° \rightarrow 180° \rightarrow 270°$) or backward (i.e., $270° \rightarrow 180° \rightarrow 90° \rightarrow 0°$)

replay (Kurth-Nelson et al., 2016; Liu et al., 2019, 2021a; Nour et al., 2021).

Figure 3. Illustration of replay analysis pipeline: Temporally delayed linear modeling (TDLM)

(A) A Lasso logistic regression model was trained for each participant and motion direction using MEG signals from the functional localizer data. **(B)** The four feature-specific models that were trained using functional localizer data. The decoding probabilities were aligned at 200 ms post-stimulus onset according to their corresponding optimal time point per participant and motion direction. Each colored line indicates the decoding results of a model applied to a motion direction dataset. The

dashed horizontal line indicates the permutation threshold estimated by random

shuffling of the labels and re-decoding. **(C)** The four models were next applied to MEG

signals during the post-cue blank period to derive a decoded reactivation matrix [time x

motion direction]. An illustration of backward sequential reactivations of motion direction

is shown on the left. Reactivation probabilities correspond to the decoding probabilities

derived from the four models. **(D)** Using TDLM, we quantified the evidence for

sequential replay of the motion sequence during the post-cue blank period in start- and

end-only conditions. We initially performed a time-lagged regression to create a [4 x 4]

regression coefficient matrix for each time lag by regressing each lagged predictor

matrix $X(\Delta t)$ onto the original reactivation matrix, Y (i.e., 1st level GLM analysis). The

degree to which the reactivations systematically follow a transition matrix of interest was

measured by "sequenceness" (i.e., 2nd level GLM analysis). We tested the magnitude of

this sequenceness at each time lag independently for all transition lags up to 600 ms.

The dashed line represents the corrected nonparametric statistical significance

threshold (see Methods). The green area indicates the lags when the evidence of

sequenceness in the backward direction exceeded the permutation threshold.

As shown in **Figure 4A** and **4B**, in the start-only condition, we found evidence of

backward replay of the exposed motion sequence (i.e., the replay sequence was 270°

→ 180° → 90° → 0° when the exposed motion sequence was 0° → 90° → 180° →

270°) during the post-cue blank period, with a peak transition lag at 28–40 ms (maximal

effect at 32 ms-lag: $\beta = -0.0202 \pm 0.002$, $p < 1/24 \approx 0.042$, peak-level significance

threshold derived from a nonparametric permutation test, FWE corrected across lags,

Figure 4B). For visualization, an example of backward replay of the motion sequence is

illustrated in **Figure 4A**. This effect was observed in most participants (Figure S3A). In

the end-only condition, we also found evidence of backward replay during the post-cue

blank period, with a peak transition lag at 28–36 ms (maximal effect at 32 ms-lag: β =

−0.0145 ± 0.0016, p < 1/24 ≈ 0.042, peak-level significance threshold obtained from a

nonparametric permutation test, FWE corrected across lags; **Figure 4E**). **Figure 4D**

shows an example of backward replay of the motion sequence in this condition, found in

most participants (Figure S3A). Note that the time lag in the horizontal axis in **Figure 4B**

and **4E** indicates the interval between the onsets of every two items in the replayed

motion sequence. We found that the replayed sequence was approximately 10 times

faster than the evoked activity sequence in the full sequence condition.

To further examine the period during which the backward replay occurred most

frequently, we divided the blank period (2.4 s) into 4 stages, each lasting 600 ms. In

start-only trials, we found that the backward replay predominantly appeared within the

third stage of the blank period (1.2 – 1.8 s after the onset of the blank period, Wilcoxon

signed-rank test, p = 0.0108), but there was no significant sequenceness in the other

three stages (first stage, p = 0.7112; second stage, p = 0.5566; fourth stage, p = 0.6791;

Figure 4C). In end-only trials, the backward replay was more likely to occur during the

second and third stages of the blank period, although the results approached, but did

not reach, significance (Wilcoxon signed-rank test, second stage, p = 0.0936; third

stage, p = 0.0642; **Figure 4F**), In contrast, the replay was not frequently observed

during the first and last stages (Wilcoxon signed-rank test, first stage, p = 0.9479; fourth

stage, p = 0.4997). Finally, in a control analysis, we conducted a comprehensive

examination of all 24 potential sequences. Only the backward replay was detected in

both start- and end-only conditions (Figure S4).

Figure 4. Backward replay in both start- and end-only conditions

(A and **D)** Examples of backward sequential reactivation in start- (A) and end- (D) only

conditions from a representative participant. Each row represents the reactivation

probabilities for the four motion directions at that time point and each column indicates

one of the four motion directions. **(B and E)** Backward replay of the exposed motion

sequence with peak transition lags at 28–40 ms in the start-only condition (B) and at 28

– 36 ms in the end-only condition (E). Horizontal dashed lines represent corrected

significance levels from a nonparametric permutation test at the 2nd-level GLM analysis

of TDLM. **(C and F)** Backward replay of the exposed motion sequence predominantly

appeared at 1.2 – 1.8 s in the start-only condition (C) and 0.6 – 1.8 s in the end-only

condition (F) after the onset of the blank period.

Backward replay is cue-dependent and depends on the amount of exposure

So far, we have demonstrated that the cue-triggered elevated responses induced

by the motion sequence exposure were not motion direction specific. However, motion

information was spontaneously reactivated and the motion sequence was backward

replayed in a time-compressed manner. It remains unknown whether the observed

backward replay of the motion sequence was cue-dependent. In other words, does the

replay of the motion sequence occur irrespective of which item of the sequence is

presented as a cue? To address this question, we conducted Experiment 2, mirroring

the design of Experiment 1 but with a different cue. Instead of flashing the first or last

RDK, we presented the second or third RDK as a cue (2nd-only or 3rd-only condition, see

Figure S5A and Methods) to examine whether these two non-terminal cues could

induce replay during the post-cue blank period. We found no evidence of either forward

or backward replay in either condition (**Figure 5A**; maximal nonsignificant effect at 32

ms-lag: 2nd-only condition, $\beta = -0.0024 \pm 0.0014$; 3rd-only condition, $\beta = -0.0018 \pm$

0.0015).

Figure 5. Backward replay is cue-dependent and depends on the amount of exposure

(A) No evidence for replay was found in either the 2nd-only (left) or 3rd-only (right) condition in Experiment 2. Horizontal dashed lines have the same meaning as those in Figure 4B and 4E. **(B)** In Experiment 3, immediately after the functional localizer phase, participants entered the main phase without the exposure phase. In the end-only condition, backward replay was observed; however, in the start-only condition, no such replay was observed.

Another interesting question is how varying the amount of exposure would affect

replay events during the blank period. To explore this issue, we conducted Experiment

3, also similar to Experiment 1 except for the removal of the exposure phase.

Accordingly, only full sequence trials (50% trials) in the main phase served for exposure

(see Figure S5B and Methods). In the start-only condition, we found a numerical trend

of backward replay at transition lags at 20–40 ms, although it did not reach statistical

significance (**Figure 5B**, left panel; maximum effect at 28 ms-lag: β = −0.0139 ±

0.0021). In the end-only condition, the evidence for backward replay just reached the

significance level at transition lags at 20 – 40 ms (**Figure 5B**, right panel; maximal effect

at 24 ms-lag: β = −0.0147 ± 0.0019, $p < 1/24 \approx 0.042$, using the peak-level significance

threshold from a nonparametric permutation test, FWE-corrected across lags).

Power increase in replay-associated sharp-wave ripple frequencies

Previous studies on rodents and humans showed that replay events are

associated with increased high-frequency ripple power (Buzsáki, 2015; Liu et al., 2019).

To investigate whether such replay-associated power increase could be observed in our

study, we performed a time-frequency analysis using combined data from the start- and

end-only conditions in Experiment 1. We first identified putative replay onsets during the

post-cue blank period in each trial. Replay onsets were determined by choosing time

points with a high (>95th percentile) probability for backward replay with a 32 ms-lag

transition (maximal replay effect at the group level for both start- and end-only

conditions, Figure 4B and 4E; see Methods for details). Using the MEG signals

recorded from whole brain sensors, we found a transient ripple power increase at 120–

180 Hz at the onset of replay events, compared to the baseline period of 50–100 ms

prior to replay onset (**Figure 6A**; cluster-based permutation test with cluster forming

threshold t > 3.1 and 5000 permutations).

Figure 6. Replay events align with ripple power, with source activation in the MTL

preceding activation in visual cortex

(A) Top: In Experiment 1, a time-frequency decomposition of sensor-level data revealing

a brief increase in high-frequency oscillatory power at replay onset. Bottom: a cluster-

based permutation test (cluster forming threshold, t > 3.1; number of permutations =

5000) could identify a significant cluster around 140 Hz. **(B)** Source localization of

ripple-band power 30 ms before replay onset showing significant activation in the MTL

(peak Montreal Neurological Institute [MNI] coordinates: X = 21, Y = 21, Z = 13.

Neurological orientation.) **(C)** Source localization of ripple-band power at replay onset

showing significant activation in visual cortex (peak MNI coordinates: X = 20, Y = 8, Z =

17). **(D)** The activation time course of the MTL at its peak MNI coordinate is shown in

red, whereas that of visual cortex at its peak MNI coordinate is displayed in green. The

MTL reached its peak activation before visual cortex.

Visual cortical activation lags MTL activation

Since replay events were obtained based on decoding probabilities of motion

directions from occipital sensors only, we next examined whether the observed replay

onsets were related to power increase within visual cortex. Using a linearly constrained

minimum variance (LCMV) beamforming algorithm (Van Veen et al., 1997), we first

epoched the data using replay onsets combined from both start- and end-only

conditions and then beamformed the broadband power into the source space (see

Methods). We found that the power increase at replay onset was localized to sources in

bilateral visual cortex (**Figure 6C**). Moreover, the right MTL, including the hippocampus

and amygdala, was associated with power increase prior to the onset of replay events

(**Figure 6B**). Specifically, activation (i.e., power increase) in the MTL occurred 30 ms

earlier than that in visual cortex ($p < 0.05$, corrected, whole brain nonparametric

permutation test). For display purposes, we extracted activations from the 10 most

activated voxels within the MTL and visual cortex, respectively, and plotted the time

courses of their broadband power. Peak activation in visual cortex at replay onset was

preceded by peak activation in the MTL (**Figure 6D**), implying an information flow from

the MTL to visual cortex.

DISCUSSION

We found the co-existence of time-locked, non-feature-specific elevated

responses and non-time-locked, feature-specific backward replay after visual sequence

exposure in human brain. The feature-specific backward replay occurred in a time-

compressed manner and could be triggered only by the first or last stimulus of the

sequence. Interestingly, even brief exposure to the sequence could still induce a trend

for the backward replay. Finally, we observed that MTL activity preceded the ripple

power increase in visual cortex at replay onset.

Our study provides two new findings in the fields of vision and learning. First,

different from many previous studies showing that expectation-based responses in

static contexts are feature-specific (Wyart et al., 2012; Kok et al., 2014, 2017), we show

here that prediction-related elevated responses in the dynamic temporal context are not.

This non-feature-specific elevated responses potentially facilitate the general

processing of any upcoming stimuli, rather than stimuli with a specific feature. Second,

contrary to the prevailing notion that replay in learning and memory requires lengthy

training with a task, our study shows that even brief exposure to a visual sequence is

sufficient to induce replay. The replay we unveiled here may play a critical role in

memorizing the visual sequence.

Regarding the two-fold neural consequences following the visual sequence

exposure, the non-feature-specific elevated responses could be induced by rhythmic

entrainment. Substantial evidence supports that rhythmic stimulation can entrain neural

oscillations, which in turn facilitates predictions about future inputs and enhances the

brain's readiness for incoming stimuli (Lakatos et al., 2008, 2013; Herrmann et al., 2016;

Barne et al., 2022). For instance, recent findings demonstrate that even task-irrelevant

information could be more effectively decoded when presented at moments close to a

highly probable target presentation (Auksztulewicz et al., 2019). In our study, the

rhythmic presentation of the motion sequence may have entrained oscillatory activity in

the brain, leading to periodic activation of sensory cortices. This rhythmic entrainment

likely serves as a possible mechanism supporting the general facilitation of neural

processing for any upcoming stimuli, independent of specific stimulus features.

In contrast, feature-specific replay may operate through a different mechanism

driven by intrinsic neural processes rather than direct external stimuli. Therefore, this

intrinsic activity is not time-locked to stimulus onset, but instead manifests in a

spontaneous way. Furthermore, since replay often occurs during offline periods (e.g.,

rest or blank period), it allows the neural system to reactivate and consolidate past

experiences without interference from ongoing external inputs. As a result, feature-

specific replay may facilitate the integration of fragmented experiences into coherent

representations, thereby subserving visual sequence learning and memory.

Another interesting question regarding the two-fold neural responses is whether

the elevated responses and the backward replay share the same neural origin, for

instance, originating from the hippocampus or other brain areas. For the elevated

responses, the cue serves to provide temporal information for upcoming stimuli but

without detailed feature information, thereby priming the cortices for activation and

facilitating the general processing of future events. Previous animal studies (Xu et al.,

2012; Gavornik and Bear, 2014) showed that this process can be implemented through

a simple local synaptic mechanism in visual cortex (Xu et al., 2012) without top-down

guidance. Moreover, Ekman et al. (2023) recently found no functional relationship

between activities in V1 and the hippocampus when exposing human participants with a

white dot sequence, further suggesting that the elevated responses may indeed

originate in visual cortex.

Different from the elevated responses, the replay events are likely initiated by the

hippocampus. This speculation is supported by two key findings. First, the replay of the

motion direction information manifested in a backward direction and a time-compressed

manner. Second, the replay is observed regardless of whether the cue is the first or last

RDK in the sequence. Both of these two properties cannot be explained by a simple

local synaptic mechanism (Xu et al., 2012) or a pattern completion-like mechanism

(Hindy et al., 2016; Ekman et al., 2017; Kok and Turk-Browne, 2018) within visual cortex

alone. Instead, the replay entails reorganizing the motion direction sequence in the

brain. Therefore, we propose the involvement of active communication between the

hippocampus and visual cortex in the occurrence of replay events, consistent with the

view that the hippocampus encodes relationships among stimuli whereas visual cortex

primarily acts as a platform for the manifestation of replay events (Whittington et al.,

2020). Previous studies on memory consolidation considered the exchange of

information between these two regions critical for facilitating replay events through

hippocampal-neocortical circuits (Buzsáki, 1996; Ji and Wilson, 2007; Carr et al., 2011;

Ólafsdóttir et al., 2016; Buch et al., 2021). Functionally, replay events offer a mechanism

for transferring recent experience from the hippocampus to the cortex, enabling the

encoding of stimulus relationships in the cortex (Marr and Brindley, 1971; Alvarez and

Squire, 1994; Redish and Touretzky, 1998; Dimakopoulos et al., 2022).

A natural behavioral consequence of visual sequence exposure is visual

sequence learning (Baker et al., 2014; Finnie et al., 2021; Ekman et al., 2023). How is

the learning implemented through hippocampus-dependent replay in visual cortex?

Three key processes are considered here. First, the hippocampus is involved in

encoding relationships among stimuli (Staresina and Davachi, 2009; Turk-Browne et al.,

2009; Hsieh et al., 2014; Garvert et al., 2017); in fact, the ability to encode such

relationships drastically decreases when the hippocampus is damaged (Chun and

Phelps, 1999; Hannula et al., 2006; Konkel et al., 2008; Schapiro et al., 2014; Finnie et

al., 2021). Second, visual cortical areas act as a "cognitive blackboard" where task-

relevant features are highlighted through feedback connections (Roelfsema and de

Lange, 2016). Such a blackboard can be flexibly written or edited. Third, there are

strong bidirectional connections between the hippocampus and sensory cortices

(Eichenbaum et al., 2007; Henke, 2010). As proposed by the hippocampal-cortical

backward projection model (Rolls, 2000), sequential reactivations of feature information

initially generated in the hippocampus can be quickly and accurately sent back to the

sensory cortices, consistent with the findings of Ji and Wilson, (2007). A recent study

also provided direct evidence that visual sequence plasticity is impaired when the

hippocampus is damaged (Finnie et al., 2021), supporting the hypothesis of functional

feedback information flow.

Why does the replay manifest in a reverse order? To date, there is no consensus

on this issue. In rodents, a seminal study of replay during the awake state showed that

when animals stopped at the end of a rewarded pathway, place cells were reactivated in

the reverse order of the previously experienced direction (Foster and Wilson, 2006).

However, a later study revealed that awake replay could occur in either a forward or

backward direction relative to behavioral experience (Diba and Buzsáki, 2007).

Similarly, in humans, both directions have been observed in different nonspatial

cognitive tasks (Liu et al., 2019, 2021b; Wimmer et al., 2020). Forward replay may be

associated with planning (Ólafsdóttir et al., 2015; Gillespie et al., 2021), providing

information pertaining to the assessment of future pathways (Diba and Buzsáki, 2007).

Backward replay may be more related to experience, as often observed at the end of

runs when animals consume a reward (Foster and Wilson, 2006; Ambrose et al., 2016).

Nevertheless, the exact function of the replay direction remains mysterious, as both

forward and backward replays are modulated by task demands (Ólafsdóttir et al., 2017).

Thus, the underlying neural mechanisms of backward replay in visual cortex remain to

be investigated.

Finally, we also found that replay occurrence is modulated by the cue. This result

highlights the importance of the start and end points of the sequence in the replay. One

fascinating proposal is that the replay event is sensitive to sequence boundaries, as

indicated by the role of the start or end point of the sequence as salient boundaries that

anchor place cell firing (Rivard et al., 2004). Accordingly, previous studies have shown

that when rats are trained to run along a linear track starting at different points, place

fields tend to be anchored to either the start or end of the journey (Gothard et al., 1996;

Redish et al., 2000), suggesting that a boundary is essential to sequence integrity and

may play a pivotal role in triggering replay onset. An alternative explanation to these

findings posits that the onset of the second or third stimulus in the sequence reinstate

the neural representations of a partial visual sequence (Ekman et al., 2023). For

example, for a given sequence, A $\rightarrow$ B $\rightarrow$ C $\rightarrow$ D, flashing the third stimulus (i.e., C)

only trigger a backward reactivation of the sequence giving C $\rightarrow$ B $\rightarrow$ A during the blank

period.

Taken together, we found that simple visual sequence exposure could

concurrently induce two-fold brain plasticity, that is, non-feature-specific elevated

responses and feature-specific backward replay in the human visual cortex. We

speculate that the non-feature-specific elevated responses may enhance general

processing of upcoming visual stimuli, whereas the feature-specific backward replay

may subserve visual sequence learning and memory. These findings significantly

advance our understanding of the task-independence and the multifaceted nature of

brain plasticity in response to visual experience.

METHODS

Participants

A total of 59 healthy participants (29 females) were involved in the three

experiments (Experiment 1: n = 21, 11 females, 22.1 $\pm$ 2.61 years; Experiment 2: n =

18, 10 females, 23.56 $\pm$ 3.29 years; Experiment 3: n = 20, 10 females, 20.65 $\pm$ 2.62

years). In Experiment 1, data from three participants were excluded before analyses, as

two showed large head motion (>20 mm) and the behavioral performance of one was at

chance level. In Experiment 3, data from two participants were excluded before

analyses because of large head motion (>20 mm). All participants were recruited in

exchange for monetary compensation (100 RMB/h). They reported normal or corrected-

to-normal vision and had no history of neurological disorders. They were naive to the

purposes of the study. The experiments reported here were carried out in accordance

with the guidelines expressed in the Declaration of Helsinki. All participants provided

written informed consent in accordance with the procedures and protocols approved by

the Human Subject Review Committee of Peking University.

Task

Experiment 1

Visual stimuli were RDKs with 100% coherence. All dots in an RDK moved in the

same direction (luminance: 2.86 cd/ m^2; diameter: 0.1°; speed: 8°/s) and were

presented against a gray background (luminance: 16.7 cd/m^2). At any one moment, 400

dots were visible within a 9° circular aperture and moved in one of the four directions: 0°

(right), 90° (up), 180° (left), and 270° (down). Each participant completed three phases

successively in the MEG scanner, namely, the functional localizer phase, the exposure

phase, and the main phase (Figure 1C). In the functional localizer phase, each trial

started with the presentation of an RDK for 1 s followed by a 1–1.5 s intertrial interval

(ITI). Participants did not need to perform any task in this phase. The motion direction of

the RDK was randomly chosen from the four directions. Each participant performed two

functional localizer runs and, each run comprised 100 trials, resulting in a total of 50

trials per motion direction. The localizer data were used to train motion direction

classifiers. This phase took approximately 10 min.

In the exposure phase, we showed participants the four RDKs in either clockwise

or counterclockwise order (e.g., $0° \rightarrow 90° \rightarrow 180° \rightarrow 270°$, Figure 1A and 1C); each was

displayed for 400 ms, followed by 300 ms of a blank screen with a fixation only.

Therefore, the full sequence lasted for 2.8 s. The order was counterbalanced among

participants, but once decided, it was fixed for each participant. Participants were

instructed to detect an oddball RDK by pressing a button, i.e., in 20% trials, dots in one

of the four RDKs traveled at a faster speed (9°/s). Each participant completed four runs

of 50 trials.

In the main phase, participants were presented with trials in three different

conditions: full sequence condition (50% trials), start-only condition (25% trials), and

end-only condition (25% trials). Trials in the full sequence condition were identical to

those in the exposure phase which served as "topping-up" exposure to maintain the

exposure effect, similar to "topping-up" adaptation in visual adaptation studies (Fang et

al., 2005). In the start- and end-only conditions; however, we only presented the first

RDK (start-only condition) or the last RDK (end-only condition) of the full sequence. For

a given run, the order of the three conditions was pseudorandomized with the restriction

that the start- and end-only trials were always preceded or followed by a full sequence

trial. Participants performed an identical oddball detection task as during the exposure

phase. The oddball occurred only in 10% of full sequence trials. Finally, each participant

completed four runs of 48 trials, yielding a total of 96 full sequence trials, 48 start-only

trials, and 48 end-only trials.

Experiments 2 and 3

In Experiment 2, we only presented the second or third RDK as a cue at the start

of the trial, referred to as the 2nd-only and 3rd-only conditions, respectively. The

procedure of Experiment 2 was similar to that of Experiment 1 except that the start- and

end-only conditions were replaced with the 2nd-only and 3rd-only conditions. Experiment

3 followed the same procedure as Experiment 1 except that the exposure phase was

removed.

QUANTIFICATION AND STATISTICAL ANALYSIS

MEG acquisition and preprocessing

Neuromagnetic signals were recorded continuously at 1000 Hz with a 306-

channel (204 planar gradiometers; 102 magnetometers), whole-head MEG system

(Elekta Neuromag TRIUX) in a magnetically shielded room. Before scanning, four-

headed position indicator coils attached to the scalp determined the head position

with respect to the sensor array. Coil location was digitized with respect to three

anatomical landmarks (nasion and preauricular points) with a 3D digitizer

(Polhemus Isotrak system). Participants sat upright inside the scanner, while the

stimuli were projected onto a screen suspended in front of them. Participants

responded using a MEG-compatible button box held in their right hand.

To reduce noise from external environment, the temporal extension of signal-

space separation (tSSS) method was applied at the preprocessing stage using the

Elekta Neuromag MaxFilter software (Taulu and Simola, 2006). MEG signals were

high-pass filtered at 0.5 Hz using a first-order IIR filter to remove slow-drifts. Data

were then downsampled from 1000 Hz to 250 Hz for sequenceness analysis or 500

Hz for time-frequency analysis. Excessively noisy segments and sensors were

automatically removed before independent component analysis (FastICA,

http://research.ics.aalto.fi/ica/fastica) and performed to remove artifacts including

cardiac signals, eye movements and blinks, and environmental noise. Artifact

components were removed by visual inspection of spatial topography, time course,

kurtosis of the time course, and frequency spectrum of all components. Only 72

sensors (including both gradiometers and magnetometers) covering the occipital

lobe, labeled as "Occipital" in the MEG data acquisition system were used for MEG

data analyses, except for source localization. The sensor selection was primarily

motivated by the main objective of the study, examining replay events in visual

cortex.

Event-related fields

To calculate the event-related fields (ERFs), MEG epochs were segmented

around trial onset for each trial and baseline-corrected using the mean activity in

the time window of [−0.3 s, 0] before trial onset. Only 48 planar gradiometers of

"Occipital" sensors were used to calculate the ERFs. The planar-combined ERF

activity was then averaged for each condition. To minimize potential baseline

confounds caused by the short interval between every two RDKs, we further

calculated the ERF peak amplitude using the mean activity in the time window of

[−0.3 s, 0] before the onset of the corresponding RDK as the baseline.

Subsequently, peak amplitude during each of the four RDK intervals was calculated

as the difference between the peak and its corresponding baseline.

Multivariate pattern analyses

Multivariate pattern analyses (MVPA) were performed to classify the neural

activity patterns elicited by the motion directions of the four RDKs in the main

phase. A one-versus-rest Lasso-regularized logistic regression model was trained

using the MEG signals from the 72 occipital sensors in the functional localizer

phase. Specifically, we trained a five-class classifier, including four classes from

trials in which the four RDKs were presented, and an additional class comprising an

equivalent amount of null data extracted from the 1–1.5 s ITI. Null data were

included to reduce the spatial correlation among the classes, thereby concurrently

lowering the decoding probabilities for all classes (Liu et al., 2021a; Nour et al.,

2021). Class weights were balanced by adjusting inversely proportional to class

frequencies (i.e., trial numbers) in the training procedure. To reduce noise, MEG

signals were averaged across every five trials within the same class before

decoding.

The trained classifier was then applied to MEG signals at each time point in

the main phase, and decoding probabilities were computed as the outputs of the

classifier. Different from the conventional decoding accuracy, where the classifier

predicts a single label at each time point, decoding probabilities provide a likelihood

estimate for each class. In our study, the classifier outputs a 5-column matrix for all

trials, where each row represents a single trial and each column represents the

probability for one class (i.e., one of the four RDKs and blank ITI). The probabilities

in each row sum to 1, reflecting the relative likelihoods across all classes for that

trial. The highest probability determines the decoded label when computing

decoding accuracy. Finally, averaging these probabilities across trials yields five

values that indicate the overall likelihood of the predicted stimulus belonging to a

given class.

Optimal time point of motion direction representation

The optimal time point of motion direction representation was considered as the

time point with the highest decoding accuracy in the functional localizer data. To index

the optimal time point of each motion direction for each participant, we conducted a

time-resolved motion direction decoding analysis on the functional localizer data using

Lasso-regularized logistic regression models. A leave-one-trial-out cross-validation

procedure was used to train the classifier to determine one of the four motion directions,

yielding a decoding accuracy at each time point for each participant. Finally, the time

point with the highest decoding accuracy was independently extracted for each

participant and each motion direction. These time points are referred to as optimal time

points and were used for training feature-specific decoding models.

Sequenceness measure

To identify sequenceness during the post-cue blank period in each trial, we

trained models to detect transient spontaneous neural reactivation of each motion

direction representation. Therefore, a one-versus-rest Lasso-regularized logistic

regression model was trained separately for each participant and motion direction using

the functional localizer data. MEG signals from the 72 occipital sensors obtained

throughout all localizer scanning sessions were used to train the decoding models. As

our aim was to quantify the evidence of feature-specific sequence, for each motion

direction, we trained a binomial classifier, using positive instances from trials in which

that feature (e.g., 0°) was presented and negative instances from trials in which all other

features (e.g., 90°, 180°, and 270°) were presented, together with an equivalent amount

of null data from the 1–1.5 s ITI. The sensor distributions of beta estimates and the

spatial correlation among the classifiers are shown in Figure S1.

The analysis pipeline of sequenceness is illustrated in Figure 3. We first applied

the trained models to MEG signals at each time point during the blank period, to

generate a [time x motion direction] reactivation probability (i.e., decoding probability)

matrix for each trial (Figure S2). The TDLM framework was then used to quantify

evidence for sequential reactivations consistent with the exposed motion sequence (Liu

et al., 2021b, 2021a; Nour et al., 2021).

TDLM is a multiple linear regression approach to quantify the extent to which a

lagged reactivation time course of one motion direction i, $(X(\Delta t)_i$, Δt indicates lag time)

can predict the reactivation time course of another motion direction j, (X_j). Two steps

were included in this pipeline. First, we performed multiple separate regressions using

the reactivation time course of each motion direction as a dependent variable ($j \in [1: 4]$)

and the historical (i.e., time-lagged) reactivation time courses of all motion directions ($i \in$

$[1: 4]$) as predictor variables.

$$X_j = \sum_{i=1}^{4} X(\Delta t)_i * \beta(\Delta t)_{i,j} + C \tag{1}$$

The predictor $X(\Delta t)_i$ was a Δt-lagged copy of the reactivation time course of X_i.

The regression coefficients $\beta(\Delta t)_{i,j}$ quantify the evidence for each empirical motion

direction $\rightarrow$ motion direction reactivation pattern at a given time lag, Δt. For example, if

X_j represents the reactivation time course of $0°$ during the blank period in the main

phase, and X_i represents the reactivation time course of $90°$ during the same period,

then $\beta(\Delta t)_{i,j}$ is the coefficient that captures the unique variance in the reactivation time

course of $0°$ explained by the Δt-lagged reactivation time course of $90°$ (Figure 3D, 1st

level GLM analysis). Finally, all such first-level coefficients were placed in a lag-specific

[4 x 4] empirical transition matrix B, representing evidence for all possible transitions at

a specific lag.

In the second step, we quantified evidence supporting that the empirical (lag-

specific) transition matrix, B, can be predicted by the sequences of interest, i.e., the

exposed motion sequence. All transitions of interest were specified in a model transition

matrix (Figure 3D, 2nd level GLM analysis; 1 for transitions of interest and 0 otherwise),

separately for forward (as in the motion sequence) (T_F), and backward (opposite to the

motion sequence) direction (T_B). Then, the strength of all sequences of interest was

measured by:

$$B = \sum_{r=1}^{4} Z_r * T_r \qquad (2)$$

where B is the [4 x 4] empirical (lag-specific) transition matrix that acquired from

the data, T_r is the [4 x 4] predictor transition matrix (for regressor r), and Z_r is the scalar

regression coefficient quantifying the evidence that the hypothesized transitions T_r

predict the empirical transitions, B. We included four predictor matrices: (1) T_F:

transitions as in the motion sequence in the forward direction (transitions corresponding

to [0° → 90° → 180° → 270°]), (2) T_B: transitions opposite to the motion sequence in the

backward direction ([270° → 180° → 90° → 0°], T_B is the transpose of T_F), (3) T_{auto}: self-

transitions to control for autocorrelation ([4 x 4] identity matrix), and (4) T_{const}: a constant

matrix to model away the average of all transitions, ensuring that any weight on T_F and

T_B was not due to general background neural dynamics.

Notably, the estimate of sequence strength, Z_r, is a relative quantity. An estimate

of zero for 0° to 90°, for example, does not mean lack of replay from 0° → 90° but no

stronger replay of 0° → 90° than that of other transitions. Repeating the regression in

Equation 2 at each time lag (Δt = 4, 8, 12, ..., 600 ms) results in both forward and

backward sequence strength as a function of time lag. Shorter lags indicate greater time

compression, corresponding to faster speeds.

In the current study, sequenceness was defined based on the contrast between

the evidence for the motion sequence replay in the forward ([0° → 90° → 180° → 270°])

versus backward ([270° → 180° → 90° → 0°]) direction (i.e., $Z_1 - Z_2$); thus, removing

between-participant variance in the sequential replay per se (Liu et al., 2021a; Nour et

al., 2021). Positive sequenceness values indicate replay in a predominantly forward

direction, whereas negative sequenceness values indicate replay in a predominantly

backward direction.

For statistical inference, we used nonparametric permutation tests involving all

possible permutations of the stimulus labels at the second-level regression, equivalent

to permuting the rows and columns together of the transition matrices used to calculate

sequenceness. For each permutation, we calculated the peak absolute mean sequence

strength over participants and across lags (controlling for multiple comparisons across

lags). Sequence strength in unpermuted data was considered significant if its absolute

magnitude was >95% of the within-permutation peak.

Identifying replay onsets and analyzing time-frequency

Having identified that the replay transition at the group level was a 32 ms-lag, we

next identified replay onsets. Replay onset was defined as the time point when a strong

reactivation of one motion direction (e.g., 0°) was followed by a strong reactivation of

the next motion direction (e.g., 90°) in the sequence, with a 32 ms-lag (Liu et al., 2019;

Nour et al., 2021). We first generated a matrix Orig as

$$Orig = X * T \tag{3}$$

where X is the [time x motion direction] reactivation matrix, and T is the backward

transition matrix. The transition matrix T defines the mapping between the motion

direction corresponding to column i in X, and column i in Orig. For example, column 1 in

X is the reactivation time course of the motion direction 0°, while column 1 in Orig is the

reactivation time course of the motion direction 90°. We then shifted each column of X

by Δt = 32 ms, to generate another matrix Proj,

$$Proj = X(\Delta t) \tag{4}$$

754   where row i in Proj corresponds to row i + 32 ms in X. Next, we multiplied Proj

755   and Orig elementwise, summing over the columns of the resulting matrix; therefore,

756   creating a [time x 1] vector, R, in which each element (i.e., row) indicates the strength of

757   replay with a 32 ms-lag at a given time.

$$R_t = \sum_{i=1}^{4} Orig_{ti} * Proj_{ti} \tag{5}$$

Finally, we identified putative replay event onsets by thresholding R at its 95th

percentile preceded by a 100 ms pre-onset baseline exhibiting a low probability of

replay at each time point.

We then epoched MEG data in the blank period surrounding each onset and

computed a frequency decomposition (wavelet transformation) in the time window of

−100–150 ms using all sensors. We aimed to find evidence for the power increase in

the high-frequency (120–160 Hz) region of interest at replay onset, compared with a

pre-onset baseline (−100–−50 ms from onset), similar to a previous study (Liu et al.,

2019). Finally, we generated two separate [time x frequency] matrices (i.e., using

forward and backward transition matrices separately) by averaging estimates over

sensors and events, capturing the typical spectrum-resolved power change at replay

onset.

MEG source reconstruction

We identified the neural sources associated with increased ripple power at

putative replay onsets. Forward models were generated based on a single shell using

the superposition of basic functions that approximately correspond to the plane

tangential to the MEG sensor array. Linearly constrained minimum variance

beamforming (Van Veen et al., 1997) was used to reconstruct the epoched MEG data to

a grid in MNI space (grid step, 5 mm). The sensor covariance matrix for beamforming

was estimated using broadband power data across all frequencies. The baseline activity

was the mean activity averaged over −100–−50 ms relative to replay onset. All

nonartifactual replay epochs were baseline-corrected at source level. We obtained

whole-brain results for voxels predicted by participant-specific ripple power at replay

onset. Nonparametric permutation tests were performed on the volume of interest to

compute the multiple comparison p values of clusters >10 voxels (whole-brain

corrected, cluster-defining threshold; t = 3.1, n = 5000 permutations).

Data and Code availability

Data used to generate the findings of this study will be available upon request (subject

to participant consent) to the Lead Contact. TDLM MATLAB Code available at:

https://github.com/YunzheLiu/TDLM. Custom computer code will be made available

upon request to the Lead Contact.

AUTHOR CONTRIBUTIONS

Conceptualization, T.H., Y.L., and F.F.; Investigation, T.H., Q.W., X.G., and X.Z.; Writing

– Original Draft, T.H.; Writing – Review & Editing, T.H., Q.W., X.Z., Y.L., and F.F.

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DECLARATION OF INTERESTS

The authors declare no competing interests.

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 SUPPLEMENTARY INFORMATION

Figure S1. Sensor maps and spatial correlation of trained Lasso logistic regression models (related to Figure 3)

(A) Left: Sensor map for each state decoding model in Experiment 1; MEG signals from

72 occipital sensors were selected as features while training the classifier. Right:

Correlation matrix among classifiers. No spatial correlation was found among trained

classifiers (highest correlation; $r < 0.12$). **(B and C)** Same as (A), except the sensor

maps and correlation matrix correspond to Experiments 2 (Panel B, highest correlation;

$r < 0.1$) and 3 (Panel C, highest correlation; $r < 0.1$), respectively.

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Figure S2. Decoded feature representations in start- and end-only conditions

(related to Figures 3)

Time series of reactivation probability output from the four regression models on an example trial. Time zero corresponds to trial onset. Both participants S08 and S12 were exposed with a fixed motion sequence of $0° \rightarrow 90° \rightarrow 180° \rightarrow 270°$.

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Figure S3. Sequenceness distribution across participants (related to Figures 4 and 5)

(A) In Experiment 1, of the 18 participants, 14 showed backward replay at 32 ms-lag for

the start-only condition (left), and 13 showed backward replay at 32 ms-lag for the end-

only condition (right). Each dot represents an individual participant. Backward and

forward sequenceness is denoted by blue and yellow dots, respectively. The inset

histogram shows the distribution of deviations from the unity line. **(B)** Same as

described in (A), but for the results of Experiment 2 in the 2nd-only (left) and 3rd-only

(right) conditions, respectively. **(C)** Same as described in (A) but for the results of

Experiment 3 in the start- (left) and end-only (right) conditions, respectively.

Figure S4. Sequenceness for each of the 24 possible orders (related to Figure 4)

(A) In the start-only condition of Experiment 1, only the backward sequence [4 → 3 → 2 → 1] with lags at 28–40 ms was significantly different (gray panels). The horizontal

1025   dashed lines represent significance thresholds derived from state label permutation at

1026   the 2nd-level GLM matrix of temporally delayed linear modeling. **(B)** Same as described

1027   in (A) but for the end-only condition in Experiment 1.

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Figure S5. Stimuli and experimental procedures in Experiments 2 and 3 (related to

Figure 5)

(A) We presented the second (2nd-only condition) or third RDK (3rd-only condition) as a

cue in Experiment 2. **(B)** The procedure of Experiment 3 was identical to that of

Experiment 1 except for removing the exposure phase.