Human statistical learning dynamically shapes

the hippocampal processing of temporal associations

Wei Tang[1, *], Morten H. Christiansen[2,3], Zhenghan Qi[4]

[1] Department of Computer Science and Department of Psychological and Brain Sciences, Indiana University-Bloomington, Bloomington, IN 47408

[2] Department of Psychology, Cornell University, Ithaca, NY 14853

[3] Haskins Laboratories, New Haven, CT 06511

[4] Department of Communication Sciences and Disorders and Department of Psychology, Northeastern University, Boston, MA 20115

* Corresponding author, wt1@iu.edu

Abstract

Recent research suggests that the human hippocampus becomes sensitive to temporal associative patterns during statistical learning (SL). And yet, how such sensitivity develops throughout the course of learning remains unclear. We investigated this question in human adults performing a visual SL task, in which the transition probability among stimuli varied systematically across learning sessions. Hippocampal activity acutely reflected this cross-session change and was predictive of behavioral performance. Using a hidden Markov model to decode stimulus patterns from hippocampal activity, we observed a "mere-exposure" effect, such that brief experience with the temporally structured stimuli was sufficient to alter the hippocampal processing. As learning unfolded, feedback from the repeated exposure shaped the hippocampal processing to better accommodate the environmental statistics. Our findings provide direct

evidence of how experience with associative patterns sculpts hippocampal activity in a nonlinear

fashion, bringing new insight to its active role in SL.

Introduction

Humans are able to learn patterned regularities in the environment based on the co-

occurrence of events in the spatial and temporal domains, an ability called statistical learning

(SL). In the temporal domain, participants in an SL task become sensitive to the associative

relationships between sequentially presented stimuli (Fiser & Aslin, 2002; Saffran et al., 1996;

Turk-Browne et al., 2005). Such temporal association can be characterized (at least in part) by

the transition probability (TP) between events, indicating how likely one stimulus is followed by

another. Behaviorally, high TP facilitates on-line sensory processing and later memory

measurements of the learned events. For example, the reaction time (RT) to probes during

learning is often shorter for stimuli with high TP than those with low TP (Batterink, 2017;

Schneider et al., 2020). Post-learning memory tests show that higher accuracy is associated with

items that were presented with a statistically patterned temporal structure during study, compared

with those in random order (Isbilen et al., 2020; Turk-Browne et al., 2005).

Recent neuroimaging studies provide preliminary evidence for a multi-level framework

across cognitive and sensory domains underlying the SL processes (Conway, 2020). At the

sensory level, statistical regularity facilitates modality-specific processing, inducing short-term

plasticity in early visual and auditory regions (Karuza et al., 2013; McNealy et al., 2006; Turk-

Browne et al., 2009, 2010). At the cognitive level, higher functions such as long-term memory,

working memory, and selective attention help develop sensitivity to statistical regularities

beyond sensory features. For example, both linguistic and nonlinguistic learning activates

associative prefrontal regions (Bischoff-Grethe et al., 2000; Furl et al., 2011; Karuza et al.,

2013). To achieve the learning goals, the brain network engages both domain-general and

modality-specific processing, allowing expectations to be generated from learned regularities to

facilitate perception (Conway, 2020; Reber, 2013; Thothathiri & Rattinger, 2015; Frost et al.,

2015).

The multi-level processing framework is yet to be extended to account for hippocampal

activity in SL. Hippocampal neurons code for association among objects/events that occur close

in time and space (Burgess et al., 2002; Ekstrom & Ranganath, 2018; O'Reilly & Norman, 2002;

Schiller et al., 2015). These conjunctive representations are insensitive to specific visual or

auditory features. Moreover, hippocampal activations reorganize in response to temporally

associated stimuli after only minutes of exposure (Schapiro et al., 2012). Thus, an intriguing

possibility is that hippocampal neurons may play an important role in the rapid development of

sensitivity to statistical regularities (Schapiro et al., 2017; Turk-Browne, 2019). The learned

sensitivity to statistical regularities can then further serve additional functions such as context-

based memory retrieval or prediction of future states/rewards (Sherman & Turk-Browne, 2020).

Nevertheless, the formation of a neural representation does not necessitate an active role

in learning *per se*. For example, conjunctive representations can be stored in the hippocampus as

a learning *outcome*, while the learning *process* is carried out elsewhere in the brain. Indeed, some

have argued that an effective long-term memory storage is critical for successful SL (Batterink &

Paller, 2017). In contrast, Schapiro and colleagues (Schapiro et al., 2017) used biologically

inspired computational models to show that circuits of hippocampal subfields can actively track

and update TP, whose activity accounts for the observed representational similarity. However,

despite the model predictions, experimental evidence of this rapid learning process has been

scarce, and the exact role of hippocampus in SL remains an open question (Covington et al.,

2018; Schapiro et al., 2014).

In the current study, we investigated whether the human hippocampus actively tracks and

changes its processing of temporal regularity in the course of visual SL. If the hippocampus

plays an active role in learning, we expect an acute change in activity in response to the change

of structure/randomness in the stimuli. Moreover, SL-related brain activity is expected to reflect

SL-related behavior. To assess the SL effect, we measured the similarity of hippocampal blood-

oxygenation-level-dependent (BOLD) activity during the presentations of neighboring stimuli

(i.e., lag-1 autocorrelation) and examined the change of this similarity over the course of

learning. Our approach was inspired by previous functional magnetic resonance imaging (fMRI)

work showing that multi-voxel BOLD activity exhibited similar spatial patterns in response to

stimuli that co-occurred in time (Schapiro et al., 2012, 2013). This multi-voxel similarity implies

lag-1 BOLD autocorrelation in individual voxels. Importantly, the autocorrelation measure

allowed us to build formal models to link hippocampal activity with stimulus transition patterns

(as we will describe in detail below). Specifically, we asked whether this autocorrelation pattern

would 1) be sensitive to the TP difference between structured and random stimuli, 2) reorganize

to reflect the switch between structure and randomness in the stimuli, and 3) correlate with SL-

related behavior. Although different factors can contribute to BOLD autocorrelation, including

the shape of the hemodynamic response function, only SL-related factors would drive the change

of autocorrelation to reflect the change of temporal patterns among stimuli. To further delineate

the hippocampal sensitivity to specific temporal statistics, we used a Hidden Markov Model

(HMM) to decode stimulus transition patterns from the autocorrelated hippocampal activity. In

the HMM, we assumed that the level of BOLD autocorrelation depended on the stimuli's

positions in a structured sequence rather than their sensory features. By examining the decoding

efficacy at different learning stages, we explored how hippocampal sensitivity to TP was shaped

by repeated exposure to structured and random sequences. We further examined whether the

BOLD autocorrelation and decoding patterns differed among hippocampal subfields. The results

suggest that the dynamic process of learning shapes the hippocampal representations of statistical

temporal patterns, bringing new insight into the active role of the hippocampus in visual SL.

Results

Learning effects in behavior and hippocampal activity. Subjects viewed sequentially

presented images while responding to target images embedded in each sequence (Fig. 1). The

sequence blocks varied along two dimensions: the stimuli in each block were (1) linguistic or

non-linguistic and (2) temporally arranged into triplets (S-block) or presented in random order

(R-block) (see Online Methods, *stimuli*). The temporal structure was set to systematically differ

between linguistic and nonlinguistic stimuli across two learning sessions (Fig. 1A). In Session 1,

linguistic S-blocks were pseudo-randomly intermixed with non-linguistic R-blocks. In Session 2,

the structure/randomness was switched, such that linguistic R-blocks were pseudo-randomly

intermixed with nonlinguistic S-blocks.

The target was set to be the third element of a randomly selected triplet from either the

linguistic or nonlinguistic sequences, and its location followed no systematic pattern in each

block either within- or across-participants (see Online Methods, *procedure*). Subjects were not

informed of the temporal structure and completed the target-detection task with an average hit

rate of $97.4 \pm 4.7\%$ (cross-subject median $\pm$ s.d.), false-alarm rate of $0.2 \pm 0.2\%$, and reaction

time (RT) of 446 ± 52 ms. Comparisons of RT between S- and R-blocks showed a facilitation

effect on target detection. For both linguistic and nonlinguistic stimuli, across subjects, the trial-average RT was significantly lower in S-blocks than in R-blocks (Fig. 1B; linguistic: Wilcoxon signed-rank $W = 23.0$, $p < 0.00028$; nonlinguistic: $W = 71.0$, $p < 0.024$). Correspondingly, for both types of stimuli and across hippocampal voxels (Fig. 1C & D), lag-1 autocorrelation of BOLD activity was significantly greater in S- than in R-blocks (Fig. 1D; linguistic: paired-sample $t_{11262} = 83.58$, $p < 1 \times 10^{-6}$; nonlinguistic: $t_{11262} = 77.12$, $p < 1 \times 10^{-6}$).

Temporal evolution of RT and BOLD autocorrelation during learning. We examined the change of RT and BOLD autocorrelation with respect to the change of temporal structure among stimuli. By design, within each session, TP differed between S- and R-blocks; across sessions, there was a switch between structure/randomness for each type of stimuli. Accordingly, the variation of RT across mini blocks depended on the session and block type (Fig. 2A, within-subject, block × session × block type interaction, $F_{1, 452.99} = 4.84$, $p < 0.028$). The cross-session RT change for each type of stimuli can account for this three-way interaction. Linguistic stimuli were presented as triplets in Session 1 (S-blocks, numbered 1-6 in Fig. 2A) and randomly in Session 2 (R-blocks, numbered 7-12). Correspondingly, within-subject, RT significantly increased from Session 1 to Session 2 (Wilcoxon test for median RT between-session, $W = 9.0$, $p < 0.00033$). In contrast, Nonlinguistic stimuli were presented randomly in Session 1 (R-blocks, numbered 1-6) and then in triplets in Session 2 (S-blocks, numbered 7-12). RT significantly decreased from Session 1 to Session 2 within-subject (Wilcoxon test for median RT between-session, $W = 30.0$, $p < 0.0051$). Indeed, the cross-subject median RT continuously decreased across concatenated mini blocks (slope $b = -4.27$, $p < 0.0059$).

The BOLD autocorrelation across mini blocks also varied depending on the session and block type (Fig. 2A, within-subject, block × session × block type interaction, $F_{1, 434.00} = 12.83$, p

< 0.00038). Although visual inspection suggested cross-session differences, the same post-hoc

comparisons as those in the RT analysis were not significant. This was not surprising because

autocorrelation could be affected by a variety of physiological factors that were unrelated to the

task. For example, intrinsic brain dynamics during the resting state could also lead to lag-1

autocorrelation in the BOLD activity (Huang et al., 2018). These dynamics would not change

according to environmental statistics, and thus, could contribute to the insignificance of cross-

session comparisons of the autocorrelation. Alternatively, the contrast between S- and R-blocks

would eliminate the common factors and leave those that were sensitive to the temporal

statistical difference among stimuli.

Thus, as an alternative post-hoc test to break down the three-way interactions found by

the ANOVA, we examined the S-R contrast of BOLD autocorrelation across mini blocks. In

each session, nonlinguistic blocks were subtracted from linguistic blocks, resulting in an S-R

difference for Session 1 and R-S difference for Session 2 (Fig. 2B). This subtraction order was

kept to facilitate the detection of cross-session differences: if the BOLD autocorrelation rapidly

adapted to the new temporal statistics, we would expect a sign-flip in Session 2. Indeed, the sign-

flip was evident in both the RT and BOLD autocorrelation measurements (Fig. 2B, within-

subject main effect of session, RT: $F_{1, 19} = 21.44$, $p < 0.00018$; BOLD autocorrelation: $F_{1, 82.30} =$

9.00, $p < 0.0035$). Across individual blocks, the hippocampal and behavioral measurements were

correlated, such that the S-R contrast of BOLD autocorrelation was linearly proportional to that

of RT (Fig. 2C, Spearman's $r = -0.62$, $p < 0.035$).

HMM decoding with simulated data. The BOLD autocorrelation measures the overall

similarity of the signal at neighboring points of a sequence. By design it does not resolve

temporal structures within a sequence. We were interested in the hippocampal sensitivity to

triplet boundaries in addition to its overall sensitivity to structure versus randomness. Therefore,

we devised an HMM approach to further test whether TP information can be decoded from the

hippocampal activity (Fig. 3A). The conditional-Gaussian HMM designed for this study

contained three hidden states: *start*, *mid* and *end*, corresponding to the three positions in a triplet.

Importantly, the model distinguishes stimuli by their positions but not the specific visual content.

If the BOLD autocorrelation differed with respect to stimulus position (e.g. within-triplet vs.

between-triplets), the HMM decoding would yield a triplet transition matrix (Fig. 3A). For the

random sequences, there was no inherent *start-mid-end* structure, and the BOLD autocorrelation

was assumed to be uniform among all stimuli. The corresponding decoding result was expected

to randomly deviate from the triplet transition matrix.

We first tested the model's decoding efficacy with simulated data (Fig. 3B-D), in which

the hidden states were strictly ordered into *start-mid-end* triplets. Theoretically, the decoding

accuracy is affected by two parameters: the sensitivity index d' and the autocorrelation ρ. The

sensitivity index indicates how well the observed BOLD data can be separated by their mean

values with respect to the hidden states. For example, given mean values of 0.4, 0.5 and 0.6

corresponding to *start, mid* and *end*, with an equal standard deviation of 0.1, the d' of this

pseudo-BOLD data equals 1.3 (see Eq. 11 in Online Methods) and indicates a good separation.

The pattern of BOLD autocorrelation ρ determines the extent to which the BOLD data is

indicative of the underlying temporal structure among states. For example, given ρ of 0.8 for

state pairs *start-mid* and *mid-end*, 0.2 for *end-start*, and 0 for any other pair, this autocorrelation

pattern represents high temporal association within a triplet, low temporal association across

triplets, and zero association for implausible state transitions. When the observed data are

generated with a high sensitivity index and triplet-representing ρ values, the HMM decoding will

yield a triplet transition matrix among hidden states. To demonstrate the model efficacy, we set

d' and ρ to the values described above and generated surrogate observations (Fig. 3B). We

compared the decoded transition matrix against the ground truth using normalized mean-squared

error (n-MSE, Eq. 10 in Online Methods), which grows proportionally with the deviation from

the ground truth. The HMM successfully decoded the transition matrix with 100% accuracy (n-

MSE = 0) in each of the 100 runs.

This simulated example represented an ideal case for decoding. However, the parameters

used here may differ from those of the veridical BOLD data. We next conducted numerical

experiments on different parameter settings to test how they affected decoding accuracy (Fig. 3C

& D). First, we varied the sensitivity index while keeping ρ fixed to the ideal values. When d'

dropped below 0.7, the decoding became imperfect (n-MSE > 0) (Fig. 3C, scatter plot), such that

nonzero probabilities were assigned to invalid transitions (i.e. false alarms). Nonetheless, even

with false alarms, the decoded transition matrix was qualitatively consistent with the ground

truth, such that the cells corresponding to the valid transitions had the highest probability (Fig.

3C, matrix plot). We then fixed the sensitivity index to the average value of the veridical BOLD

data (0.23 ± 0.11 across subjects, marked as shaded area in Fig. 3C). We tested the decoding

accuracy by varying the ρ values (Fig. 3D, scatter plot). The decoding accuracy decreased when

ρ decreased, but the decoded transition matrix remained qualitatively accurate at each ρ level

(Fig. 3D, matrix plot). The simulation results suggested that if data autocorrelation encoded

temporal association, and given the parameter ranges of veridical BOLD data (mean $\pm$ s.d.,

shaded areas in Fig. 3C & D), the HMM decoding would be imperfect but remain qualitatively

consistent with the true transition patterns. In the following text, for brevity, we refer to decoded

matrices that were qualitatively similar to the triplet-transition matrix as "triplet-like", and those

randomly deviating from it as "uninformative".

HMM decoding with BOLD data. We applied the HMM approach to decode transition

patterns in the veridical BOLD data, in each block throughout the task. As a cautionary step, we

validated the assumption that the BOLD autocorrelation reflected hippocampal sensitivity to TP

structure. We tested this assumption by comparing the conditional-Gaussian HMM, which uses

both the mean and autocorrelation values for decoding, with a standard univariate-Gaussian

HMM, which uses only the mean values for decoding (see Online Methods, *Hidden Markov*

modeling). Both analyses were performed on the voxel-averaged hippocampal BOLD time series

from the S-blocks in each subject. Across subjects, the conditional-Gaussian HMM showed

significantly greater accuracy than the univariate-Gaussian HMM (Fig. 4A, Wilcoxon $W = 0.0$, p

$< 8.8 \times 10^{-5}$), demonstrating the improvement of decoding using autocorrelation information.

Thus, the following analyses were focused on the conditional-Gaussian HMM.

Averaged across subjects and sessions, the decoding accuracy significantly varied across

S-blocks (one-way ANOVA block effect, $F_{5,114} = 2.98$, $p < 0.014$) but not R-blocks (one-way

ANOVA, main effect of block, $F_{5,114} = 0.92$, $p < 0.43$). Curiously, the highest accuracy occurred

in the early blocks (Fig. 4B). Decoded patterns of the transition matrix confirmed this

observation (Fig. 4C): triplet-like transition pattern was found in the second and last S-blocks, as

well as the first two R-blocks. Importantly, the decoded pattern was triplet-like for the last S-

block and noninformative for the last R-block. The triplet-like pattern in early S- and R-blocks

suggested that learning occurred rapidly and might be history-dependent. We further examined

this possibility. The rationale was that at the beginning of the task, the hippocampus might not be

sensitive to the S-R difference. The exposure to S-block(s) could have caused a mere-exposure

effect, such that the hippocampus responded to the early R-blocks as if there was a triplet

structure. If this was true, we should find triplet-like R-block decoding patterns exclusively in the

subset that was presented after S-blocks. We examined this possibility with data from Session 1.

Among the twenty subjects, ten (Group 1) started with an S-block and ten (Group 2) started with

an R-block (Fig. 4D). Separately for each group, we examined the decoded transition matrix of

the first block (outlined in black in Fig. 4D) as well as the following S- and R-blocks (outlined in

magenta and blue, respectively). The decoding outcome showed a presentation order effect (Fig.

4D), such that the decoded pattern was triplet-like if the first block was structured, and

uninformative if random. Moreover, the S- and R-blocks that followed the firstly presented S-

block both showed triplet-like decoded patterns. In contrast, the S- and R-blocks following the

firstly presented R-block showed uninformative decoded patterns.

BOLD autocorrelation and decoding accuracy across hippocampal subfields. We

compared the autocorrelation and decoding results among the hippocampal subfields (Fig. 5A).

Averaged across sessions and blocks, the greatest S-R difference of BOLD autocorrelation was

observed in subfields CA2/3 and CA4/DG (Fig. 5B, one-way ANOVA main effect of subfield,

$F_{4, 95} = 2.75$, $p < 0.032$). The decoding accuracy for S-blocks did not significantly differ across

subfields (one-way ANOVA on the n-MSE, main effect of subfield, $F_{4, 95} = 1.06$, $p < 0.38$).

Consistently, the decoded transition matrices showed similarly uninformative patterns for each

subfield (Fig. 5B). We further examined the autocorrelation and decoding results in the first and

last blocks, averaged across sessions (Fig. 5C & D). For the first block, the S-R difference of

BOLD autocorrelation did not significantly differ across subfields (one-way ANOVA main

effect of subfield, $F_{4, 95} = 0.58$, $p < 0.68$), while a triplet-like transition matrix was decoded in

CA1 from the S-block (Fig. 5C). For the last block, the greatest S-R difference of BOLD

autocorrelation was found in the parasubiculum (one-way ANOVA, near-significant main effect

of subfield, $F_{1, 48} = 3.59$, $p < 0.064$), in which a triplet-like transition matrix was decoded from

the S-block (Fig. 5D).

Discussion

This study investigates how processing in the human hippocampus changes to

accommodate temporal transitions of sequentially presented visual stimuli during an SL task. We

found that hippocampal activity is sensitive to the temporal structures in the visual inputs,

adaptive to the changes of the embedded temporal patterns, and rapidly alters processing as

reflected by a mere-exposure effect. Hippocampal sensitivity to temporal structure was reflected

in the autocorrelation of the hippocampal BOLD activity. We observed increasing

autocorrelation of hippocampal BOLD activity during the exposure to structured versus

randomly ordered stimuli. When the structure/randomness flipped across learning sessions, the

autocorrelation pattern swiftly reflected this change. Importantly, the autocorrelation pattern was

correlated with the RT pattern across mini blocks throughout the task, demonstrating a link

between the SL-related hippocampal activity and the SL-related behavior. Moreover, we found

that the triplet patterning can be decoded from the hippocampal activity, and that the decoding

outcome was history-dependent. At the beginning of the task, the decoding results depended on

the sequential order of S- and R-blocks being presented; at the end of the task, the decoding

results accurately separated the S- and R-blocks. Overall, these results suggest that the

hippocampus is not a passive storage for learning outcomes but plays an active role in the

learning process.

A long-standing theory is that the hippocampus codes for associations between

objects/events that occur close in time or space (Burgess et al., 2002; Cohen & Eichenbaum,

1995; Ekstrom & Ranganath, 2018; O'Keefe et al., 1978). In accordance with previous work, we

found that hippocampal activity became self-similar (measured by lag-1 autocorrelation) when

the presented items were highly associated in time (Fig. 1D). Moreover, our findings

demonstrate a direct link between hippocampal processing and the triplet patterning of the

stimuli. To learn a statistical pattern, in addition to detecting the co-occurring items, the brain

also needs to estimate how often such co-occurrence happens. In this study, this frequency

information is reflected by TP, which equals 1 within a triplet, 0.25 between triplets, and 1/12

among randomly ordered stimuli. The hippocampus is sensitive to the general difference of TP

between structured and random sequences, demonstrated by the S-R difference in the BOLD

autocorrelation.

However, the autocorrelation measurement cannot resolve TP differences within a

sequence, which critically determines the patterning of a triplet structure. Thus, we designed the

HMM analysis to test the hippocampal sensitivity to triplet patterning. The conditional-Gaussian

HMM assumes that the similarity of BOLD activity for neighboring stimuli can be differentiated

by their locations (*start*, *mid*, or *end* position of a triplet) but not necessarily the specific visual

content (linguistic or nonlinguistic). Thus, successful HMM decoding requires the BOLD

autocorrelation of items within a triplet to be different from that of items across triplets. The

simulation results validated this assumption by showing that when a hidden sequence generated

data with said autocorrelation structure (Fig. 3B), the transition matrix was correctly decoded

with 100% accuracy. Under noisy conditions that approximated true BOLD data, the decoded

patterns were robustly triplet-like (Fig. 3C & D). Thus, the successful decoding in veridical

BOLD time series was evidence that the hippocampal activity was sensitive to the triplet TP

pattern within-sequence (Fig. 4C & D).

Importantly, the HMM decoding results suggest a proactive learning process in which the

current sensitivity to structure depends on prior experience. The history-dependence is implied in

the decoding result of early R-blocks. At the beginning of the task, if one or more S-blocks

preceded the first R-block, the decoded R-block transition matrix contained triplet information

(Fig. 4D, subject group 1). In contrast, in R-blocks that did not follow any S-block, as well as the

first S-block that followed R-block exposure, no triplet transition pattern was decoded (Fig. 4D,

subject group 2). This result indicates that initially presented stimuli may shape the processing of

the subsequent state of the environment. This is in broad accordance with theories of context

dependency and predictive computations in the hippocampus, such as successor representations

in reinforcement learning (Dayan, 1993; Stachenfeld et al., 2017), temporal context models in

episodic memory (Howard & Kahana, 2002; Sederberg et al., 2008), and the Tolman-

Eichenbaum machine in unifying spatial and nonspatial tasks (Whittington et al., 2020). A

common conceptual element of these models is the associative maps stored in memory and

employed to predict the future. Our observations are direct evidence that associative-pattern

learning from experience results in changes to the processing of such regularities. SL as a

common mechanism underlying the multifaceted hippocampal activity in navigation, cognitive

map, memory and learning is worth exploring in future work.

Moreover, the decoding results across mini blocks (Figs. 4B & C) suggest that feedback

from the environment shapes processing biases to better accommodate the true associative

statistics. As learning unfolded, the decoded transition matrices deviated from the initial patterns

and fluctuated across both S- and R-blocks. At the end, a triplet-like pattern was decoded in the

last S-blocks and but not the R-blocks (Fig. 4C). This decoding pattern depicts a learning process

that does not progress in a linear fashion. A possible interpretation is that learners are particulary

sensitive to the temporal statistics during the initial encountering of the stimuli (Jungé et al.,

2017) and may "over-generalize" after this brief exposure (Kemp et al., 2007; Perfors et al.,

2010; Wonnacott, 2011). The triplet-like decoded pattern in early R-blocks could reflect this

strategy. The varying decoding efficacy in later blocks likely reflects the adjustment of learners'

representations by repeated exposure to the same stimuli. The mini block design in this task can

be a general experimental approach to further explore how learning evolves across stages.

Finally, our results suggest different roles of hippocampal subfields in SL and the

resulting experience-driven changes to TP processing. The CA2/3 and CA4/DG showed the

greatest task-average S-R difference in the BOLD autocorrelation, while the CA1 and

parasubiculum showed sensitivity to the triplet TP pattern at the beginning and end of the

learning session, respectively (Fig. 5C & D). These results intersect with the proposed functional

dichotomy between the monosynaptic and trisynaptic pathways (O'Reilly & Norman, 2002;

Turk-Browne, 2019). On one hand, the sensitivity of the CA2/3 and CA4/DG to the general

structural difference between sequences fits their role in the trisynaptic pathway for pattern

separation (Leutgeb et al., 2007). On the other hand, the CA1 is postulated to explicitly code for

associative patterns in the monosynaptic pathway through SL (O'Reilly & Norman, 2002;

Schapiro et al., 2017), which can account for its TP sensitivity in this study. Indeed, previous

work has observed CA1 place-cell coding for environmental context based on experience (Plitt

& Giocomo, 2021), supporting the history-dependent TP processing in our results. We postulate

that the learned changes to TP processing in the CA1 is conveyed to the parasubiculum for

output to downstream inferotemporal and visual regions. This output could alter the tuning for

objects in these regions through prediction-based processing adaptations to facilitate target

detection (Turk-Browne, 2019). The current study cannot test these hypotheses due to limited

temporal resolution of BOLD fMRI. Nonetheless, these results encourage further investigation

into the orchestrated dynamics among hippocampal subfields during SL.

Online Methods

Participants. Twenty-eight adults (mean age = 20.79 years, SD = 2.89 years, 7 males)

participated in this study. All participants were right-handed, native English speakers with no

history of neurological or psychiatric disorders. All participants gave written consent to perform

in accordance with the Institutional Review Board at the University of Delaware. All participants

were compensated for their participation. Due to incomplete MRI data resulting from acquisition

issues (N = 2), early termination of the study by the participants (N = 2), poor performance with

< 20% hit rate in at least one scanning session (N = 4), twenty participants in total (mean age =

20.16 years, SD = 0.81 years, 7 males) were included in the data analyses.

Stimuli. The stimuli contained two types of images: *Letters* (linguistic stimuli) were 12

images each containing an alien character holding up a sign with a capital letter (see

Supplemental Figure for all the stimuli). The alien character was held the same while the letter

differed among the 12 images. *Pictures* (nonlinguistic stimuli) were 12 different standalone alien

cartoons (Supplemental Figure). Both Letters and Pictures were presented in two types of mini

blocks. An *S-block* was composed of 48 stimuli, organized into 4 unique triplets each repeated 4

times. The presentation order was pseudo-randomized among triplets, so that the TP between

stimuli was 1 within-triplet and 1/4 between-triplet. No Letter triplet contained any words,

common acronyms, or initialisms. An *R-block* contained 12 unique stimuli each repeated 4 times.

In other words, an R-block was the randomization of all stimuli in an S-block of the same

stimulus type. The presentation order of individual stimuli was pseudo-randomized, and no

combination of three stimuli was presented more than once. Thus, the TP among stimuli was

uniformly 1/12. For both S- and R-blocks, each stimulus was presented at the center of the screen

for 0.8 s with an 0.2 s interval between stimuli. Each mini block lasted 48 s.

Procedure. Participants watched two sessions of stimuli presentation, organized into

mixed S- and R-blocks described above, while performing a target detection task. Session 1

contained 6 linguistic S-blocks, 6 nonlinguistic R-blocks and 6 12-second resting periods;

Session 2 contained 6 nonlinguistic S-blocks, 6 linguistic R-blocks and 6 12-second resting

periods. The order of S-blocks, R-blocks and resting periods was pseudo-randomized, and the

randomization was independent for each session and each participant. Adjustment was made to

ensure that no more than two blocks of the same type were adjacent to each other. Before each

block, a cue for the target was presented at the center of the screen for 2 s. For both linguistic and

nonlinguistic stimuli, the cue was a stimulus chosen from the last position of a triplet in an S-

block. The cue choice was randomized among participants. Throughout the task, in each

participant and for each type of stimuli, the same cue was used for all S- and R-blocks. Because

the order of stimuli and that of blocks were randomized independently for each participant, the

target location among stimuli followed no systematic pattern either within- or cross-participants.

Structural and functional MRI data acquisition. Data were acquired on a Siemens 3T

Magnetom Prisma scanner with a 64-channel phased array head coil at the Center for Brain and

Biomedical Imaging at The University of Delaware. Whole-head, high resolution structural

images, including a T1-weighted, magnetizations-prepared rapid gradient-echo (MPRAGE)

anatomical volume (TR=2500 ms, TE=2.9 ms, TI=1070 ms, flip angle=8.0°, voxel

resolution=1.0 mm isotropic, FOV=256 × 256, 176 sagittal slices) and a T2-weighted anatomical

volume (TR=3200 ms, TE=565 ms, flip angle=2.0°, voxel resolution=1.0 mm isotropic,

FOV=256 × 256, 32 sagittal slices) were collected. Functional images were acquired using

simultaneous multi-slice, T2*-weighted echo-planar imaging scans (TR=800 ms, TE=32 ms, flip

angle=61°, FOV=21 cm, matrix=64 × 64, acceleration factor=6). We acquired sixty adjacent

slices in an interleaved sequence with 2.5 mm slice thickness resulting in an in-plane resolution

of 2.5 × 2.5 × 2.5 mm^3.

Image preprocessing. MRI data were preprocessed using the analysis pipeline of

FreeSurfer (7.1.0) (Fischl, 2012) involving intensity normalization, registration to Talairach

space, skull stripping, segmentation of white matter, tessellation of the white matter boundary,

and automatic correction of topological defects (Fischl & Dale, 2000). BOLD preprocessing used

a combination of SPM12 (Penny et al., 2011), FSL (5.0.9) (Smith et al., 2001), AFNI (Cox &

Hyde, 1997) and FreeSurfer scripts implemented in Nipype 1.1.9 (Gorgolewski et al., 2011).

Preprocessing steps involved dropping the 4 initial frames, despiking (AFNI), slice time

correction (SPM SliceTiming), motion correction (SPM Realign), removal of linear and

quadratic trends (FSL TSNR), and artifact detection based on movement or deviation in intensity

(Nipype ArtifactDetect). Because of our subsequent autocorrelation and HMM analyses, no

smoothing was applied to avoid distortion in the BOLD time series. Lastly, each subject's T1

volume was co-registered to the functional space using boundary-based registration with 9

degrees of freedom (FreeSurfer bbregister).

Segmentation of the hippocampal subregions. Hippocampal subregions were segmented

from the T1-weighted images using an automated algorithm provided by FreeSurfer 7.1.0. This

algorithm utilizes a probabilistic atlas built upon ultra-high resolution (0.13mm isotropic, on

average), ex vivo MRI data from 15 cases (Iglesias et al., 2015). The segmentation yielded the

following subregions: parasubiculum, presubiculum, subiculum, CA1, CA3, CA4, granule cell

layer of dentate gyrus (DG), hippocampus-amygdala-transition-area (HATA), fimbria, molecular

layer, hippocampal fissure, and tail. The HATA and following subregions, as well as the

parasubiculum, are fine-grained structures that occupy only a few voxels. Thus, we grouped

parasubiculum and presubiculum into one subregion "parasubiculum", and CA4, DG, HATA and

following subregions into one "CA4/DG".

BOLD autocorrelation. BOLD images were coregistered to the FreeSurfer preprocessed

anatomical images in each subject's native space. BOLD time series were extracted from

hippocampal voxels using the segmentation masks. Each data point is the closest after stimulus

onset. For each voxel, lag-1 autocorrelation *r* was calculated following the formula:

423
$$r = \frac{[X(t) - \bar{X}] \cdot [X(t) - \bar{X}]}{\left\| X(t) - \bar{X} \right\| \cdot \left\| X(t-1) - \bar{X} \right\|}, \qquad (1)$$

where $X(t) = [x_1, x_2, x_3, \ldots, x_T]$ is a vector of BOLD activity at time points 1 to T, $X(t\text{-}1)$ denotes

the BOLD activity shifted by one time step, $\bar{X}$ is the temporal mean of $X(t)$, $\|\ \|$ denotes the vector

norm, and dot denotes inner product. Each mini block contained 48 stimuli, and thus, 48 time

points in the BOLD time series, and 47 time points after balancing for the lag. For cross-subject

comparison in Fig. 1, the BOLD autocorrelation was resampled to the MNI152 T1 1mm template

in FSL using FreeSurfer's mri_robust_register algorithm. For all other analyses the BOLD

autocorrelation was extracted from each subject's native space.

Linear mixed models and ANOVA. Within-subject treatment effects were analyzed with

linear mixed models (Bates et al., 2015). Repeated-measures ANOVA were applied to the model

output for assessing effect size and significance, with type III sums of squares and the

Satterthwaite approximation for degrees of freedom. Implementation of these analyses used the

lme4 package in R (4.0.5) with the RStudio interface (1.4.1106; Rstudio Inc.).

Hidden Markov modeling. The intuition behind this HMM came from the hypothesis that

hippocampal neurons should be sensitive to the TP between stimuli regardless of their sensory

features. Thus, the hidden states of this HMM contained 3 variables to represent the triplet

sequence:

$$Q = \{q_1, q_2, q_3\}, \quad (2)$$

where q_1, q_2, q_3 denote the first (*start*), second (*mid*) and third (*end*) positions in a triplet. A 48-

stimulus triplet sequence would correspond to 16 repetitions of the $q_1q_2q_3$ set. The probability of

transition between the hidden states was defined by the matrix

$$A = \begin{bmatrix} 0 & 1 & 0 \\ 0 & 0 & 1 \\ 1 & 0 & 0 \end{bmatrix}. \quad (3)$$

Each element on the *i*-th row and *j*-th column represents the probability of transition from q_i to

q_j. By construction, the transition for a triplet sequence was only possible from *start* to *mid*, *mid*

to *end*, and *end* to *start*. Thus, *A* had a probability of 1 in the cells corresponding to these

transitions and 0 elsewhere. Because a structured sequence always begins with *start*, the initial

probability of *Q* was set to $\pi = [1, 0, 0]$. Together, *Q* and *A* define the triplet sequence that drove

the SL-related BOLD activity. The goal of the HMM analysis was to test whether *A* can be

decoded from the observed BOLD activity.

In the HMM framework, observations are linked to hidden states via an emission

probability function. For the analysis in this study, the BOLD signal was averaged across

hippocampal voxels (within subfields for the subfield analysis) and treated as a univariate time

series, *O(t)*. The relationship between the hidden states and *O(t)* was determined by the emission

probability function $b(o_t|q_j)$, which represented the probability of observing activity o at time t

given state j. Critically, $b(o_t|q_j)$ was hypothesized to follow a conditional Gaussian distribution:

$$b(o_t|q_j) \sim N(m_j + \rho_{ij}(o_{t-1} - m_i), (1 - \rho_{ij}^2)\sigma^2), \quad (4)$$

where m_i and m_j denote the mean BOLD activity corresponding to hidden states i and j,

respectively, ρ_{ij} the correlation of the BOLD activity corresponding to hidden states i and j, and σ

the standard deviation that was assumed to be constant in the BOLD time series. The right-hand

side of Eq. 4 is the conditional distribution of o_t (corresponding to state j) given o_{t-1}

(corresponding to state i), assuming a bivariate Gaussian joint distribution. Note that o_t and o_{t-1}

denote measurements at adjacent time points, and therefore their correlation ρ_{ij} is the lag-1

autocorrelation of the BOLD activity. Eq. 4 states that given hidden state j, the probability of

observing o_t was a function of state-dependent variables m_i, m_j and ρ_{ij}. For triplet sequences,

because the transitions among states were deterministic, the i-j pairs were unique, i.e., there was

a unique m_i for each m_j. Thus, $b(o_t|q_j)$ was unique for each state j at time t.

The presence of autocorrelation ρ_{ij} in $b(o_t|q_j)$ set apart our conditional-Gaussian HMM

from previous univariate-Gaussian HMMs that had been applied to task fMRI studies

(Baldassano et al., 2017; Duan & Man, 2012). In a univariate-Gaussian framework, the setup for

hidden states Q, transition matrix A and initial states π would be the same as above, but the

emission probability would depend only on the mean and variance of the BOLD signal at time t:

$$b_{univ}(o_t|q_j) \sim N(m_j, \sigma^2). \quad (5)$$

In Eq. 5, the only state-dependent parameter was the mean BOLD activity m_j. The

univariate-Gaussian HMM assumes that different hidden states resulted in different mean levels

of BOLD activity. In contrast, the conditional-Gaussian HMM relaxes this assumption by adding

influences of autocorrelation ρ_{ij} to $b(o_t|q_j)$. In Eq. 4, the hidden states could be linked to o_t via

either the mean values or ρ_{ij} (or both). For example, if all three states corresponded to the same

mean BOLD activity, then ρ_{ij} would become the only state-dependent parameter to affect the

emission probability, and the hidden state information would be entirely encoded in the

autocorrelation. Alternatively, if SL did not affect the autocorrelation, i.e. $\rho_{ij} = 0$ for all state

pairs, then Eq. 4 would become Eq. 5 and the hidden state information would be entirely

encoded in the mean values. Thus, the conditional-Gaussian HMM allowed us to test whether the

BOLD autocorrelation contained any information of the hidden triplet sequence.

The test was achieved through a decoding procedure, in which the hidden-state transition

matrix A was treated as an unknown parameter to be estimated, or decoded, from the observed

data. By comparing the decoded matrix with the ground truth in Eq. 3, one could assess the

model efficacy in capturing the relationship between the hidden states and the BOLD data. If the

conditional-Gaussian HMM yielded more accurate results than the univariate-Gaussian HMM, it

would be evidence that the relationship specified by Eq. 4 was more plausible than that by Eq. 5,

i.e., the BOLD autocorrelation contributed to encoding the hidden state information. The

accuracy of the decoded matrix would reflect the accuracy of the hypothesized encoding scheme

in Eq. 4. For the decoding procedure, the definition of our conditional-Gaussian HMM fulfills

conventional HMM assumptions and its parameters can be estimated via the standard Baum-

Welch algorithm (Jurafsky & Martin, 2009). To initiate this algorithm, an uninformative prior of

A was set up such that each element in the matrix was 1/3. The prior of $b(\cdot)$ was set up using

parameters calculated from the empirical BOLD data. For each mini block, the observation $O(t)$

contained 48 BOLD data points corresponding to the 48 stimuli:

$$O_t = \{o_1, o_2, o_3, \ldots, o_{48}\}, \quad (6)$$

The initial values of parameters in Eq. 4 were calculated from O_t:

$$\hat{m}_i = \frac{1}{16} \sum_{\forall t: q_t = q_i} o_t$$

502   ,   (7)

$$\hat{\sigma} = \sqrt{\frac{1}{47} \sum_{t=1}^{48} (o_t - \hat{m})^2}$$

504   ,   (8)

505   where $\hat{m}_i$ denotes the mean of o_t across time points corresponding to state i, and $\hat{m}$ the mean

506   across all time points. The initial value of ρ_{ij} was calculated separately for each i-j pair:

$$\hat{\rho}_{ij} = \frac{\sum_t (o_{t-1,i} - \hat{m}_i)(o_{t,j} - \hat{m}_j)}{\sqrt{\sum_t (o_{t-1,i} - \hat{m}_i)^2 \sum_t (o_{t,j} - \hat{m}_j)^2}}$$

507   ,   (9)

508   in which the summation was over all time points associated with state pair i-j. With the initial

509   setup of A and $b(\cdot)$, the Baum-Welch algorithm will re-estimate A and $b(\cdot)$ recursively through an

510   expectation-maximization procedure until the results converge. To compare the BOLD-estimated

511   transition matrix with the ground-truth represented by Eq. 3, a difference score was computed in

512   the form of a normalized mean squared error (n-MSE):

$$\text{n-MSE} = \sum_{i,j} (a_{ij} - \hat{a}_{ij})^2 / \sum_{i,j} a_{ij}^2$$

513   ,   (10)

where a_{ij} denotes elements in the ground-truth matrix (Eq. 3), and $\hat{a}_{ij}$ elements in the estimated

transition matrix.

Simulation. Stochastic processes were generated from the HMMs for testing decoding

efficacy. HMMs are generative models, i.e. given π, A, and $b(\cdot)$, samples of o_t can be drawn from

the conditional-Gaussian emission probability distribution up to an arbitrary time point T. To

match the task setup, we let $T = 48$ for all simulated time series. To test the decoding efficacy

under different parameter settings, two quantities were systematically varied: the sensitivity

index d' and autocorrelation ρ_{ij}. Sensitivity index is a function of m_i, m_j, and σ:

522
$$d' = (|m_1 - m_2| + |m_2 - m_3| + |m_3 - m_1|)/3\sigma. \qquad (11)$$

It measures the joint effect of m_i and m_j, i.e., how well the observations corresponding to the

three states can be separated by their mean values. For each parameter set, 100 time series were

generated and a Baum-Welch decoding algorithm was applied to each sample. The accuracy was

measured by n-MSE averaged across the 100 samples.

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Figure 1. Repeated exposure to structured stimuli corresponded with reduced RT for target
detection and increased lag-1 autocorrelation in the hippocampal activity. (A) Schematic of
the task design. Subjects viewed 2 sessions of stimuli presentation, each composed of 12 randomly
intermixed mini blocks (middle row). Session 1 contained 6 S-blocks (blue) of linguistic stimuli
and 6 R-blocks (orange) of nonlinguistic stimuli, and in Session 2 the S/R block type was switched
between linguistic and nonlinguistic stimuli. Each mini block started with a cue for the target,
followed by 48 stimuli presented sequentially. Each stimulus was presented at the center of the
screen for 0.8 s, followed by a 0.2 s inter-stimulus interval. Subjects responded to the target with
a button press. Top row: an example S-block, in which linguistic stimuli were organized into
triplets. Bottom row: an example R-block, in which nonlinguistic stimuli were randomly ordered.
(B) RT of the structured and random sequences for each type of stimuli, shown as median ± s.e.
across subjects. Black dots on the swarm plot represent the cross-trial median of individual subjects.
Asterisks mark the significance level of the Wilcoxon test, ***: $p < 0.001$, *: $p < 0.05$. (C) The
hippocampus mask on the MNI152 T1 brain template from FSL. (D) Histograms of BOLD
autocorrelation across hippocampal voxels in the MNI152 template. The BOLD autocorrelation
from each subject's native MRI space was transformed to the MNI152 template. The
autocorrelation value of each voxel in the template was averaged across subjects.
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Figure 2. Behavioral and hippocampal SL effects were correlated across mini blocks. (A) RT
(median ± s.e. across subjects) and BOLD autocorrelation (mean ± s.e. across subjects) varied
across mini blocks. The blocks were numbered by the order of presentation throughout the entire
task. Background color separates the two sessions, with gray indicating the session in which the
stimuli were structured. (B) The S-R difference of RT (median ± s.e. across subjects) and BOLD
autocorrelation (mean ± s.e. across subjects) showed a sign-flip across sessions. The mini blocks
were numbered in the same manner as in A. (C) The S-R difference of RT and that of BOLD
autocorrelation were linearly correlated across mini blocks.

Figure 3. Conditional-Gaussian HMM decoding with simulated data demonstrated the
model efficiency in capturing the triplet transition pattern. (A) Schematic of the conditional-
Gaussian HMM structure. Three hidden states *start, mid, end* are defined according to the position
of each stimulus in a triplet, regardless of their specific visual features. The TP among states is
defined by the transition matrix (abbreviations: s. for start, m. for mid, e. for end). Each hidden
state is linked to the observed BOLD signal by the emission probability, which follows a
conditional-Gaussian distribution. (B) A simulated time series from the HMM, with parameters
$m_1 = 0.4$, $m_2 = 0.5$, $m_3 = 0.6$, $\sigma = 0.1$, $\rho_{12} = \rho_{23} = 0.8$, $\rho_{31} = 0.2$. (C) Decoded transition matrices
(top) and decoding accuracy (bottom) with varying *d'* and $\rho_{12} = \rho_{23} = 0.8$, $\rho_{31} = 0.2$, averaged
across 100 runs. Error bars represent s.e. across runs. Shaded area marks the mean and s.d. of *d'*
in the veridical BOLD data. (D) Decoded transition matrices (top) and decoding accuracy (bottom)
with varying $\boldsymbol{\rho}$ and *d'* = 0.23, while keeping the relationship $\rho_{12} = \rho_{23} = 4\rho_{31}$. The results were
averaged across runs, with error bars representing s.e. across runs. Shaded area marks the mean
and s.d. of $\boldsymbol{\rho}$ in the veridical BOLD data.

Figure 4. HMM decoding with veridical BOLD data showed varying accuracy across the mini blocks. (A) Decoding accuracy was significantly higher from the conditional-Gaussian than the univariate-Gaussian HMM. The symmetric contours represent the subject-wise histograms of n-MSE. Dashed lines mark the first, second and third quantiles of the distribution. (B) Decoding accuracy (mean ± s.e. across subjects) significantly varied across mini blocks. The n-MSE scores were averaged across sessions. (C) Decoded transition matrices for each mini block, averaged across sessions. Effective decoding was present in the early (first and/or second) S- and R-blocks, as well as the last S-block. (D) Decoding outcomes showed a mere-exposure effect. The letter arrays list the structure/randomness of the first 4 mini blocks (columns) in each subject (rows). The subjects were divided into two groups according to the stimulus structure in the first mini block (black boxes). For Group 1, decoded matrices are shown for the first S-block, and the first S- (magenta boxes) and R-blocks (blue boxes) that followed the first S-block, averaged across subjects. For Group 2, decoded matrices are shown for the first R-block, and the S- and R-blocks that immediately followed the first R-block, averaged across subjects. The selection strategy for Group 2 was based on the concern that any S-block between the first and second R-blocks might affect the hippocampal response to the second R-block, according to the mere-exposure assumption. Therefore, only the S- and R-blocks that immediately followed the first R-block were chosen for decoding comparison. Group 1 showed effective decoding in all selected blocks while Group 2 showed none.

Figure 5. SL effects differed across hippocampal subfields. (A) Segmentation of hippocampal subfields shown in a subject's native space. Abbreviations: PrS=parasubiculum, S=subiculum. (B-D) S-R difference of BOLD autocorrelation (z-normalized, i.e., mean-subtracted and divided by s.d. across subfields) and decoded transition matrix for each subfield, using data averaged across all mini blocks (B) and from the first (C) and last (D) mini blocks. Colors in the bar charts correspond to the color scheme in A. Magenta boxes highlight effective decoding results.

Supplemental Figure. All the linguistic (left) and nonlinguistic (right) stimuli used in the task.
For each type of stimuli, the three items on each row represent a triplet used in S-blocks.