Cell assemblies in the cortico-hippocampal-reuniens network during slow oscillations

Abbreviated title: Sequences in Nucleus Reuniens

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Author Contributions:

M.F. and P.P.Q. designed the experiments. M.F. and P.P.Q. performed the experiments. A.G. assisted in performing experiments. D.A.G., M.F. and P.P.Q. interpreted the results and performed the analysis. All the authors reviewed and edited the manuscript.

Keywords:

Cell assemblies; Hippocampus; Prefrontal cortex; Nucleus reuniens; Slow oscillations; Sequences.

Abstract The nucleus reuniens (NR) is an important anatomical and functional relay between the medial prefrontal cortex (mPFC) and the hippocampus (HPC). Whether the NR controls neuronal assemblies, a hallmark of information exchange between the HPC and mPFC for memory transfer/consolidation, is not known. Using simultaneous LFP and unit recordings in NR, HPC and mPFC in rats during slow oscillations under anesthesia, we identified a reliable sequential activation of NR neurons at the beginning of UP states, which preceded mPFC ones. NR sequences were spatially organized, from dorsal to ventral NR. Chemical inactivation of the NR disrupted mPFC sequences at the onset of UP states as well as HPC sequences present during sharp-wave ripples. We conclude that the NR contributes to the coordination and stabilization of mPFC and HPC neuronal sequences during slow oscillations, possibly via the early activation of its own sequences.

Significance Statement (117 words)

Neuronal assemblies are believed to be instrumental to code/encode/store information. They can be recorded in different brain regions, suggesting that widely distributed networks of networks are involved in such information processing. The prefrontal cortex, the hippocampus and the thalamic nucleus reuniens constitute a typical example of a complex network involved in memory consolidation. In this study, we show that spatially organized cells assemblies are recruited in the nucleus reuniens at the UP state onset during slow oscillations. Nucleus reuniens activity appears to be necessary to the stability of prefrontal cortex and hippocampal cell assembly formation during slow oscillations. This result further highlights the role of the Nucleus Reuniens as a functional hub for exchanging and processing memories.

Introduction

Information exchange between the hippocampus (HPC) and the medial prefrontal cortex (mPFC) is essential for different memory processes, including consolidation (Siapas and Wilson, 1998; Frankland and Bontempi, 2005; Preston and Eichenbaum, 2013; Maingret et al., 2016; Eichenbaum, 2017; Kitamura et al., 2017; Latchoumane et al., 2017; Ferraris et al., 2018). In both regions, the representation of information is supported by the recruitment of cell assemblies of neurons, which fire in a fine time-resolved manner (Skaggs and McNaughton, 1996; Nadasdy et al., 1999; Lee and Wilson, 2002; Euston et al., 2007; Luczak et al., 2007; Pastalkova et al., 2008; Peyrache et al., 2009; Battaglia et al., 2011). These neuronal assemblies display a precise temporal sequential activation, which reflects the encoded information (Nádasdy, 2000; Foster and Wilson, 2006; Ji and Wilson, 2007; Davidson et al., 2009; Marre et al., 2009; Pfeiffer, 2017). Such activity occurs mainly during sleep (Foster, 2017), particularly during non-REM sleep, and is organized in space and time by a set of oscillations, such as hippocampal sharp-wave ripples, cortical slow oscillations and spindles (Sirota et al., 2003; Sirota and Buzsáki, 2005; Staresina et al., 2015; Maingret et al., 2016). How these cells assemblies are finely coordinated between the HPC and the mPFC to support memory consolidation is not well understood. The thalamic nucleus reuniens (NR), which bi-directionally connects the HPC and the mPFC (Herkenham, 1978; Van der Werf et al., 2002; Vertes, 2006; Varela et al., 2014), plays a key role in memory consolidation (Loureiro et al., 2012; Cassel et al., 2013; Pereira de Vasconcelos and Cassel, 2015; Dolleman-van der Weel et al., 2019). In particular, the NR synchronizes gamma bursts, which can open temporal windows for information exchange (Buzsáki and Draguhn, 2004), between the HPC and the mPFC during non-REM sleep (Ferraris et al., 2018). The NR is

therefore ideally poised to orchestrate the dynamics of cell assemblies in both regions.

We have recorded and manipulated the NR–mPFC–HPC circuit during slow oscillations under anesthesia, which resembles sleep patterns and provide long periods for analysis. We find that cell assemblies are recruited in the NR at the onset of the UP state of slow oscillations and that NR activity is needed for the stability of HPC and mPFC cell assemblies.

Materials and Methods

Contact for Reagent and Resource Sharing

Further information and requests for resources may be directed to and will be fulfilled by the Lead Contact, Dr. Pascale P. Quilichini (pascale.quilichini@univ-amu.fr).

Experimental Model and Subject Details

All experiments were performed in accordance with experimental guidelines approved by Aix-Marseille University Animal Care and Use Committee. All rats were group housed to avoid social isolation-induced stress (Manouze et al., 2019). A total of 18 rats were used in this study. Part of these data (14 Wistar Han rat data) was used in a previously published study (Ferraris et al., 2018), and 4 Wistar Han rats are original data. They include local field potentials (LFPs) and single-unit recordings made in the mPFC, HPC and NR of anesthetized rats.

Animal surgery

Wistar Han IGS male rats (250-400g; Charles River; RRID:RGD_2308816) were anesthetized with urethane (1.5 g/kg, i.p., Cat#U2500, Sigma-Aldrich) and ketamine/

xylazine (20 and 2 mg/kg, i.m., Renaudin Cip: 3400957854195 and CENTRAVET Cat#ROM001, respectively), additional doses of ketamine/xylazine (2 and 0.2 mg/kg) being supplemented during the electrophysiological recordings. The heart rate, breathing rate, pulse distension and the arterial oxygen saturation were also monitored with an oximeter (MouseOx, Starr Life Science) during the entire duration of the experiment to ensure the stability of the anesthesia and monitor the vital constants. The head was secured in a stereotaxic frame (Kopf #962, Phymep) and the skull was exposed and cleaned. Two miniature stainless-steel screws, driven into the skull, served as ground and reference electrodes. Up to three craniotomies were performed to target, from bregma: the prelimbic area of the medial prefrontal cortex (mPFC) at +3 mm AP and +0.8 mm ML; the CA1 field of the intermediate hippocampus (HPC) at -5.6 mm AP and +4.3 mm ML; and the nucleus reuniens (NR) at -1.8 mm AP and -2 mm ML. Silicon probes (NeuroNexus) were used to record from these structures: a A1x32-Edge-5mm-20-177-H32-15 probe placed at [-2.5 -3.1] mm from brain surface to reach mPFC layer 5; a A1x32-Edge-10mm-20-177-H32-50 32-site probes placed at -7.2 mm from brain surface to reach the NR; a HPC A1x32-6mm-50-177-H32-15 probe placed at [-2.8 -3.0] mm perpendicularly to the CA1 field from *stratum oriens* to *stratum lacunosum moleculare* in the HPC. All the probes were lowered inside the brain with a motorized manipulator (IVM single, Scientifica).

For NR inactivation experiments (n = 5 rats), a local injection of a fluorophore-conjugated muscimol (BODIPY-MSCl TMR-X Conjugate, Cat#M23400, Invitrogen) was performed in the NR and data from the mPFC and the HPC (CA1) were simultaneously acquired. The injection needle (33 gauge, Cat#87930, Cat#7803-05, Hamilton) was inserted in the NR (using the same depth coordinates as the probes and mounted on the same micromanipulator) and 0.70 nmol of muscimol in 0.3 µl of PBS (Ferraris et al., 2018) was delivered over 60s through a micropump

(UltraMicroPump UMP3-1, WPI). The needle was left in place for 3 additional minutes to allow for adequate diffusion of the drug, then carefully removed.

At the end of the recording, the animals were injected with a lethal dose of Pentobarbital Na (150mk/kg, i.p.) and perfused intracardially with 4% paraformaldehyde solution in phosphate buffer (0.12M). The position of the electrodes (DiIC18(3), Cat#46804A, InterChim) was applied on the back of the probe before insertion) was confirmed histologically on Nissl-stained 60 μm sections (NeuroTrace 500/5225 Green Fluorescent Nissl Stain, Cat#N21480, Invitrogen). Only experiments with appropriate position of the probe were used for analysis (Fig. 2A).

Data collection and initial analysis

Extracellular signal recorded from the silicon probes was amplified (1000x), bandpass filtered (1 Hz to 5 kHz) and acquired continuously at 32 kHz (64-channel DigitalLynx; NeuraLynx) at 16-bit resolution. Raw data was preprocessed using a custom-developed suite of programs (Csicsvari et al., 1999). After recording, the signals were downsampled to 1250 Hz for the LFP. Spike sorting was performed automatically, using KLUSTAKWIK ((Harris et al., 2000); http://klustakwik.sourceforge.net/; RRID:SCR_008020; RRID:SCR_014480), followed by manual adjustment of the clusters, with the help of autocorrelogram, cross-correlogram and spike waveform similarity matrix (KLUSTERS software, RRID:SCR_008020; (Hazan et al., 2006). After spike sorting, the spike features of units were plotted as a function of time, and the units with signs of significant drift over the period of recording were discarded. Moreover, only units with clear refractory periods and well-defined cluster were included in the analyses. Recording sessions were divided into brain states of theta and slow oscillation periods. The epochs of stable slow oscillations (SO) periods were visually selected respectively

from the ratios of the whitened power in the slow oscillations band (0.5-2 Hz) and the power of the neighboring band (20-30 Hz) of mPFC or NR LFP and assisted by visual inspection of the raw traces (Quilichini et al., 2010).

Neurons were assigned as "NR neurons" by determining the approximate location of their somata relative to the recording sites, the known distances between the recording sites, the histological reconstruction of the recording electrode tracks and subsequent estimation of the recording sites. All the neurons recorded from sites located near the close contour of the NR were discarded. Neurons located at a minimal distance of 200μm of NR border and located within contours of the ventro-median, submedian or antero-median thalamic nuclei were classified as "other thalamic neurons" and used in the analysis (Ferraris et al., 2018).

Data post-processing

All the analysis was performed using custom-written MATLAB (RRID:SCR_001622; RRID: SCR_008020) scripts. From the spike times, the instantaneous firing rates of each cell was calculated by counting the number of spikes inside a window of 50 ms, in overlapping intervals of 10 ms. The population firing rate was estimated averaging the single cell firing rates at each 10ms interval. Both the population rate and the single cell firing rate were smoothed with a Gaussian kernel of 5ms width. Peaks of population activity (AP) were identified as the points where the population rate amplitude was larger than the average rate plus 1 standard deviation. A separation between two consecutive peaks of at least 600 ms was imposed on the peak detection algorithm to avoid multiple peaks of activity within one up-state. For visualization purposes, all firing rate's heat maps were normalized with the peak firing rate for each cell to guarantee a variation between [0:1].

Phase analysis and sequence identification

LFP signals (from mPFC or NR) during slow oscillation phase were band-pass filtered between 0.5 Hz and 2 Hz with a second order Butterworth filter to extract only the UP/DOWN transitions. The time evolution of the phase during the UP/DOWN cycle was extracted performing the Hilbert transform of the filtered LFP (from the mPFC or the NR, where the [0.5 2] Hz oscillation is most likely volume-conducted from the cortex). We used Rayleigh circular statistics (Berens, 2009; Ferraris et al., 2018) to compute the mean phase at which each neuron fires ("preferred phase") and to build their firing-phase histograms (Fig. 1Ba). For visualization purposes, the resulting histograms depicted in the heat maps were normalized with the peak value of the distribution. For each cell, we then calculated a resultant vector characterized by an angle describing the average preferred phase and a magnitude with values between [0 1], quantifying the coherence of the phases. The template order was obtained by organizing the average preferred phase in increasing order (0 to 2π).

Sequence reliability, participation index and sequence delay

We extracted the UP state duration as described in Ferraris et al. (2018). At each UP state, the local activation order was calculated by measuring the time lag between the first activation of each neuron relative to the AP in a +/- 200ms window. Ordering the latencies in increasing order resulted in the local activation order. The reliability of a given sequence within a population peak was quantified as the Spearman rank correlation of the template order and the local activation order in that particular UP state. Sequences were considered reliable above 99% significance level. For each reliable sequence, a robust fit between the local activation order and the activation latency was computed. To assess whether a neuron participated in a given sequence, the outliers in the robust fit were identified as those whose residual value were larger than twice the standard deviation of the residuals in the robust fit. The

participation index was then calculated as the mean fraction of neurons that participated in the sequences of a recording session. A second linear fit between the local activation latency and the site of the linear probe closest to the neuron gave a measure of the directionality of the sequence activation. A slope of the fit different than 0 implies an activation towards a given direction. Since this slope has units of time/electrode site, we multiplied the resulting value by 32 (the number of sites in the electrode) to account for the time required for a sequence (sequence delay) to travel along the electrode.

Activity peak triggered histogram (APTH)

Once the APs have been identified according to the algorithm described in Data post-processing section, we store the values of activation time lag relative to the AP at each UP state, allowing us to compute the so-called activity peak triggered histogram for each cell. With the APTH it is possible to assess the statistics of the firing latencies in the neighborhood of the UP state (+/- 200 ms). Once the APTH was obtained, one can calculate the variability of the activation lags as a measure of the firing extent around the peak reported in Fig. 3C. Variability of the APTH can be calculated in analogy to the variance of a probability distribution function where τ is the mean value of the APTH (See Fig. 3C insert).

SPW-Rs detection and analysis

The procedure of SPW-Rs detection in the HPC stratum pyramidale LFP was based on those described previously (Isomura et al., 2006; Ferraris et al., 2018). Briefly, the LFP was digitally bandpass filtered [80 250] Hz, and the power (root-mean-square) of the filtered signal was calculated. The mean and SD of the power signal were calculated to determine the detection threshold. Oscillatory epochs with a power of 5 or more SD above the mean were detected. The beginning and the end of oscillatory

epochs were marked at points where the power fell 0.5 SD. Once the SPW-Rs were detected, the SPW-R half time was calculated as the average between the start and the end of it. Then, the APTH for each cell was computed, taking each SPW-R half time as the activity peak. To test whether a cell robustly participates in the SPW-Rs, we compared the APTH against a uniform distribution with identical mean and standard deviation. Cells whose APTH were different from the flat distribution above a 95% level were considered as robustly participating in the SPW-Rs. Cell participation ratio was obtained dividing the number of robustly participating neurons by the total number of recorded neurons for that session. Organizing the average time lag for robustly participating neurons in increasing order defined the template order for the ripple. For each SPW-R, the reliability of activation respect to the template was calculated via the Spearman rank correlation between the activation order of that SPW-R and the template order. To calculate this correlation, only the neurons that belong to the template order were considered. A correlation above 95% confidence interval was considered reliable.

Statistics

All results reported are based on a significance threshold α=0.05, otherwise stated, and all groups included enough samples to enable rejection of the null hypothesis at that level. We used two sample Kolmogorov-Smirnov test to assess differences between distributions, and t-Student's test to evaluate differences in the mean of distributions. In parallel, results from small samples data (n < 10) were evaluated using estimation statistics providing the effect size and the 95 % confidence interval (CI) of the median difference between two groups. Raw data was processed online in https://www.estimationstats.com website directly giving both results and graphs (Calin-Jageman and Cumming, 2019). The median difference between two groups is shown with a Gardner-Altman estimation plot. Raw data is plotted on the left axis.

The median difference is plotted on a floating axis on the right as a bootstrap sampling distribution (Fig. 1CD 2D, 3CE, 4B). Five thousand bootstrap samples were taken; the confidence interval is bias-corrected and accelerated. The mean difference is depicted as a dot; the 95 % confidence interval is indicated by the ends of the vertical error bar. If 95% CI includes 0, differences are considered as non-significant. The p value(s) reported are the likelihood(s) of observing the effect size(s), if the null hypothesis of zero difference is true. Correlation tests involving ranked variables (neuron indices and electrode sites) were performed via a Spearman rank correlation. We tested significant differences between percentages with a two proportion Z-test.

Results

Cell assemblies are recruited in NR at UP state onset

We first assessed the behavior of NR (n=166 cells, n=5 rats), mPFC (n=496, n=7) and HPC (n=163, n=4) neurons during slow oscillations (SO) in anesthetized rats. Slow oscillations during anesthesia share similar features as non-REM sleep (Hutt, 2011; Ferraris et al., 2018) and offer the advantage of long duration stable recordings necessary to identify statistically significant sequences. Neighboring thalamic neurons (antero-median and ventro-median nuclei) were also recorded as a control group ("TH", 89 neurons, n=4).

During SO, most of the neurons fired at the onset of the UP states, mainly in mPFC and NR (Fig. 1A) (Ferraris et al., 2018). We found a robust recruitment of neuronal activity at the UP state onset in NR (median percentage of recruited neurons per UP state 70.4 %, min/max: 61.1 / 85.9 %), and mPFC (median: 62.9 %, min/max: 42.1 / 70 %). In comparison, HPC neurons showed less participation during UP state (median: 33.8 %, min/max: 29.3 / 46.4 %. To identify any ordering in this firing

activity, we ranked each neuron according to its mean preferred SO phase (Fig. 1Ba). In contrast, HPC neurons displayed better sequential activation during sharp-wave ripples (SPW-Rs), which tend to occur during the UP state (Sirota et al., 2003; Battaglia et al., 2004; Isomura et al., 2006; Buzsáki, 2015; Maingret et al., 2016; Khodagholy et al., 2017) (median: 45 %, min/max: 35 / 64 %). Neurons were organized in increasing order of average preferred phase, defining a *template order* (Fig. 1Bb). Sequences in NR always preceded mPFC ones (Fig. 1B). As a negative control, we analyzed the activity of neighboring thalamic nuclei (TH) neurons. Although TH neurons displayed a strong entrainment by SO (Ferraris et al., 2018), there was poor sequential activity (Fig. 1B). The sequential activity at UP state onset therefore appears to be specific of NR and mPFC neurons in the areas investigated here. As a further control, we analyzed the activity of NR neurons during epochs dominated by theta (4-6 Hz) oscillations. We did not find stable sequential neuronal assemblies during theta (data not shown). Together, these results show that, at the beginning of UP states, cell assembly formation occurs first in the NR, then in the mPFC and marginally in the HPC and neighboring thalamic nuclei.

We then quantified the reliability of these sequences. To do so, we computed at each UP state the activation latency of each neuron relative to the population peak activity (see Methods). The local activation order of the cell assembly was determined from the ascending sorting of the activation latencies. We then calculated the Spearman rank correlation between the local activation order and the template order. A rank correlation r = 1 indicates that the sequential activation follows exactly the template order determined previously by the average phase preferences. For NR neurons, the template order was significantly expressed in 32% of the total number of UP states (min/max = 15 / 37 %, Spearman test, p < 0.01), while mPFC reliable sequences were found in 25 % of the UP states (min/max = 5 / 31 %, Fig. 1C). In contrast, reliable HPC sequences were only detected in 2 % of the

UP states (min/max = 1 / 6 %) and in 2 % for sequences in TH neurons (min/max = 1 / 14 %). Although the proportion of reliable sequences in the NR and mPFC were not significantly different (KS-test, p = 0.15), the average Spearman correlation values of the reliable sequences were significantly larger in the NR than in the mPFC (KS-test, p < 0.01, Fig. 1D), indicating that the NR cell assembly sequential organization is the most consistent. To estimate the participation of neurons in sequence generation, we calculated a participation index, defined as the probability for a given neuron to be involved in a given sequence (see Methods). The participation index was consistently larger across experiments in the NR (median: 0.63, min/max: 0.55 / 0.81) than in the mPFC (median: 0.52, min/max: 0.36 / 0.60; KS-test, p < 0.05). The sequential firing of neurons could just be the result of an increased excitability of early firing neurons as compared to neurons firing later after the onset of the UP state (Buzsáki, 1989). We thus tested the correlation between the firing order and the firing rate for each NR and mPFC neuron. We found that 2/5 NR recordings were significantly correlated while only 1/7 mPFC recordings showed significant correlations (Spearman test, p<0.05).

Together, these results show that mPFC and NR neurons display highly stable patterns of sequential neuronal activations at the onset of the UP state, with NR sequences being more reproducible. The sequential activation cannot be solely explained by increased excitability of early firing cells.

NR cell assemblies are spatially distributed

We also investigated the spatial distribution of the NR and mPFC cell assemblies. To do this, we correlated the template order of each neuron to their anatomical dorso-ventral localization, since the silicon probes was sampling NR from the dorsal to ventral pole (Fig. 2A). The slope of a linear fit of such correlation provides the information on whether the activity is propagating in a given direction (Luczak et al.,

2007) (see Methods, Fig. 2B). Furthermore, we quantified the degree of spatial distribution of the activity for each UP state, calculating a linear fit between the activation latency of each neuron relative to the peak activity and the dorso-vental localization. The slopes of the fit for each UP state, properly multiplied by the number of sites of the probe resulted in the sequence delay (see methods). Positive or negative delays are fingerprints of propagating activity in certain direction. Even though both NR and mPFC can show some degree of propagation in a preferential direction (Fig. 2C), NR average delays were consistently larger than mPFC ones (KS-test, $p < 0.05$, Fig. 2D), reflecting a dorsal-to-ventral direction of the sequential activation. To assess the consistency of the preferential direction of propagation, we calculated the Spearman rank correlation between the rank of the template order and the anatomical location of the corresponding recording site. We found a highly significant correlation in NR recordings in most cases (n = 4 out of 5, median | r | value across all n: 0.65, $p < 0.001$), whereas it was virtually absent in mPFC (n = 1 out of 7, |r| = 0.35, $p < 0.01$) . There are thus more spatially organized sequences in NR as compared to mPFC (z-test, $p < 0.05$, Fig. 2E).

Altogether, these results demonstrate that NR displays at the UP state onset robust sequential activations, which are spatially organized in a dorso-ventral pattern. This is particularly surprising as it suggests a topical spatial organization in the NR.

NR activity is necessary to the mPFC and HPC sequences stability

The fact that NR cell assemblies occur earlier than mPFC ones raises the possibility that the NR may be involved in the sequential activation of mPFC neurons. To test this hypothesis, we used Muscimol to inactivate chemically the NR (see Methods) and recorded the activity of mPFC neurons (n=190 neurons, n=5 experiments). Following NR inactivation, mPFC neurons showed a reduced firing activity during the UP state as compared to non-inactivation, control recordings (Fig. 3A) (Ferraris et

al., 2018). During SO, the distribution of interspike intervals is characterized by the presence of two peaks: one around 1 Hz corresponding to the frequency of the SO, and one around 100 ms, which indicates that cells fire bursts of action potentials (Fig. 3B). The distribution of interspike intervals was modified in inactivation conditions with a smaller peak around 100 ms, reflecting reduced bursting activity during UP state (Fig. 3B). We then calculated the distribution of the pooled variability of the activity peak triggered histogram (APTH, see methods and Fig. 3C insert) in control and NR inactivation conditions. The median APTH variability across experiments was not significantly changed when NR was inactivated (control median: 9.62, min/max: 8.98 / 10.60, NR inactivation median: 9.02, min/max: 8.67 / 13.73, KS-test, p =0.4428), as shown in Fig. 3C. This reveals the fact that, even though bursting is reduced, the time extent to which neurons in mPFC fire is maintained. However, the UP state duration was slightly shorter (control: 0.55 s, NR inactivation: 0.48 s, Mann-Whitney test, p < 0.001), an effect that can be attributed to the extra peak in UP state duration distribution around 0.25 s, which did not exist in control data (Fig. 3D). We then evaluated the outcome of the mPFC sequences in the NR inactivation condition. Inhibition of NR activity reduced the capacity of mPFC to generate reliable sequences as compared to the control condition, since only 5.6% of them (min/max: 1.4 / 8.4%) were reliable (as compared to control median: 25%, min/max: 5 / 30 %, KS-test, p=0.0228 Fig. 3E). Moreover, the fraction of participating neurons in reliable sequences decreased in the inactivation condition (control: 53%; inactivation: 37%; KS-test, p < 1e-5; Fig. 3F). These results support the proposal that NR activity is important for the stability and reliability of mPFC sequences.

The UP state of the SO poorly modulates the activity of HPC neurons and the formation of timely organized cell assemblies. However, sequential activation of HPC

neurons consistently occurs during SPW-Rs (Buzsáki, 2015). We therefore investigated the consequences of NR inactivation on HPC sequences during SPW-Rs. NR inactivation did not affect HPC neurons firing rate (control median: 70 Hz, min/max: 41 / 85 Hz, NR inactivation median: 52.59 Hz, min/max: 33 / 71 Hz, KS-test, p=0.97, Fig. 4A). In addition, NR inactivation did not alter the frequency of SPW-Rs occurrence (median control: 0.087 Hz, min/max: 0.043 / 0.11 Hz, NR inactivation: 0.049 Hz, min/max: 0.022 / 0.111 Hz, KS-test, p=0.8254, Fig. 4B). Similarly, neither their power (mean normalized power control: 157 ± 2, NR inactivation: 144 ± 5, T-test, p=0.295; Fig. 4C bottom panel) nor their inner frequency (mean frequency control: 193 ± 1 Hz, NR inactivation: 213 ± 3 Hz, T-test, p=0.084, Fig. 4C top panel) were modified. This result shows that NR inactivation does not change either SPW-R properties or global HPC cell firing. In contrast, the number of reliable sequences found within SWP-Rs was drastically reduced when the NR was inactivated (control median: 15%, NR inactivation median: 3%, z-test, p<0.001; Fig. 4D). In this condition, the remaining cell assemblies recruited a significantly lower number of neurons (control median: 37.5 %, min/max: 12.5% / 80%, NR inactivation median: 27.8 %, min/max: 0% / 74.4%, KS-test, p<0.001, Fig. 4E). These findings suggest that the NR also controls the sequential organization of neuronal firing in the HPC during SWP-Rs.

Discussion

In this study, we show that two third of NR neurons fire within robustly spatially and temporally organized cell assemblies at UP state onset; and that NR activity controls cell assemblies' stability in mPFC at UP state onset and in HPC during SPW-Rs. These results further support the concept that the NR is a key functional hub in memory networks involving the medial prefrontal cortex and the hippocampus.

The sequential activation of neuronal assemblies is believed to constitute a core feature of information processing in the brain (Tonegawa et al., 2018). Cell assemblies are found in the archicortex (Lee and Wilson, 2002; Harris et al., 2003; Dragoi and Buzsáki, 2006; Pastalkova et al., 2008; Villette et al., 2015; Malvache et al., 2016), cortical areas (Kenet et al., 2003; MacLean et al., 2005; Ferezou et al., 2006; Euston et al., 2007; Luczak et al., 2007; Luczak and Maclean, 2012), as well as in striatum (Lansink et al., 2009). They constitute a way to code/encode/store information (Maass, 2016; Kitamura et al., 2017). During non-REM sleep, cortical activity is dominated by the sequential activation of cortical neurons at the onset of the UP state, while, in the hippocampus, the sequential activation mostly occurs during SPW-Rs (Sirota et al., 2003; Battaglia et al., 2004; Peyrache et al., 2009; Maingret et al., 2016; Khodagholy et al., 2017). Similar sequential firing occurs during the slow oscillations measured during anesthesia (Luczak et al., 2007), supporting the view that such brain state shares many features with non-REM sleep (Tung and Mendelson, 2004; Isomura et al., 2006; Clement et al., 2008; Quilichini et al., 2010; Hutt, 2011; Ferraris et al., 2018). Whether sequences represent internally generated representations or preconfigured cell assemblies (Pastalkova et al., 2008; Dragoi and Tonegawa, 2012; Liu et al., 2019) or a functional template of offline replay in the framework of memory consolidation (Lee and Wilson, 2002; Buzsáki, 2015; Pfeiffer, 2017) still remains to be elucidated. Our work demonstrates that sequences can also be recorded in a thalamic nucleus. This property appears to be specific to the NR, on the basis of recordings of other thalamic nuclei recorded in the vicinity. A systematic study of all thalamic nuclei should now be performed, to determine whether this property is specific to the NR. In the present work, we used anesthesia conditions, as the identification of sequences requires long lasting stable recordings (Bermudez Contreras et al., 2013). Our results should also be validated during long sleep sessions .

The way mPFC and HPC sequences are generated remains poorly understood. Our results demonstrate that NR activity is a key regulator of mPFC and HPC sequence stability. Interestingly previous (Ferraris et al., 2018) and present results demonstrate that the NR does not regulate oscillatory patterns, as NR inactivation does not change the frequency of SO, gamma oscillations and SPW-Rs. However, the NR seems to organize the synchronization of gamma oscillations between the HPC and the mPFC (Ferraris et al., 2018) as well as the sequential activation of cell assemblies in the mPFC during SO or in the HPC during SPW-Rs. The NR is ideally located for such fine-tuning of cell activity, as it is bi-directionally connected to the mPFC and HPC (Vertes, 2006; Cassel et al., 2013; Varela et al., 2014). The fact that NR sequences always precede mPFC ones at UP state onset suggests that NR cells may directly drive mPFC cells. In keeping with this proposal, NR neurons have an excitatory action on HPC and mPFC (Dolleman-Van der Weel et al., 1997; Dolleman-Van der Weel and Witter, 2000; Di Prisco and Vertes, 2006; Dolleman-van der Weel et al., 2019) by modulating/activating both interneurons and principal cells. However, the control of sequences during hippocampal SPW-Rs is more difficult to explain as SPW-Rs occur at variable times after UP state onset (Sirota et al., 2003; Battaglia et al., 2004). This rather suggests that the NR is an important hub in a wider network of networks, from which oscillations can emerge.

The presence of sequences specifically in the NR is quite remarkable. Since NR neurons are involved in reference memory consolidation (Loureiro et al., 2012) and in spatial memory (Jankowski et al., 2014; Ito et al., 2015; Jankowski et al., 2015; Ali et al., 2017; Cholvin et al., 2018), NR sequences may constitute an activity template used to organize information at the beginning of the UP state. Such a mechanism may enable the transmission of information to target areas in a packet-based manner (Luczak et al., 2015); a default activity in a default mode (Luczak and Maclean, 2012; Sanchez-Vives and Mattia, 2014).

Another remarkable feature of NR sequences is that their dorso-ventral organization, suggesting a precise topological organization in terms of afferences and efferences, despite the fact that NR does not have a layered organization (Jones, 1985; Bokor et al., 2002; Van der Werf et al., 2002). The NR probes was inserted in the brain with an angle to properly target the NR (Fig.. 2Aa). As a result, the bottommost sites were located more medially than the topmost ones. Although the medio-lateral distance was less that 250 µm between the top and bottom sites, we cannot fully rule out that the spatial organization we describe is medio-lateral rather than dorso-ventral (or both). The origin of the dorso-ventral organization of these sequences remains to be elucidated. While NR outputs to the subiculum are topographically organized along the dorso-ventral axis, no study has reported such precise topological organization of NR inputs (Dolleman-Van Der Weel and Witter, 1996). Only differences involving the rostro-caudal axis have been so far reported (Cassel et al., 2013), as well as a preferential targeting of NR fibers to prelimbic and infralimbic deep layers (5 and 6) and superficial layer 1 in mPFC (Berendse and Groenewegen, 1991; Vertes, 2004; Hoover and Vertes, 2012). The fact that we could not find any dorso-ventral pattern in mPFC sequences does not mean that there is none to be found as, given the spatial organization of the mPFC, we could only target one layer. There is also no available information on the local connectivity among NR neurons, except a caudal to rostral pathway (Dolleman-Van der Weel et al., 1997). The NR includes difference cell types, but the lack of specific molecular markers prevents, so far, a proper optogenetic investigation (Bokor et al., 2002; Walsh et al., 2017).

In conclusion, our results further support the concept that the NR plays a key role as an anatomical and functional hub between the mPFC and the HPC. The control it exerts on mPFC and HPC information packets suggests that it strongly participates in the organization of information in both regions but also in the transfer of

information from the HPC to the mPFC. Its internal organization allows the genesis of information packet sequences, which may represent similar features as those coded in the mPFC and the HPC.

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FIGURE LEGENDS

Figure 1. Sequential dynamics of cell assemblies.

A) The upper trace depicts the [0.5-2] Hz filtered LFP in **(a)** mPFC and **(b)** NR (which is volume-conducted from the cortex), where the UP states correspond to troughs in the LFP signal (gray boxes), from two typical recordings. The corresponding template heat maps (bottom) of normalized NR unit activities with **(a)** mPFC or **(b)** HPC normalized unit activities show the repetition of neuronal assemblies particularly in the NR and the mPFC. The identification number of the recorded neurons are indicated on the left axis of the heat-maps.

B) (a) Phase distribution of the NR, mPFC, HPC and TH population firing (grouped data, all recordings) with reference to the SO phase (depicted by the dashed curve). The dashed line indicates the most preferred phase of NR neurons, showing that their firing precedes other cells located in the mPFC, HPC and TH at the onset of the UP state. **(b)** Heat maps of normalized distribution of neurons preferred SO phase for sample representative firing patterns of NR (same neurons as in Aa), mPFC (same as in Aa), HPC (same as in Ab) and TH (representative example). The activation order for each neuron is calculated as the average preferred phase. Heat maps are ordered according to the increasing value of average preferred phase (colored circles), defining a template order. The sequential activation is clearly apparent in NR and mPFC neurons, but neither in the HPC nor in the TH (the phase distribution extends over at least p/2 for each neuron).

C) Top panel: The largest percentage of reliable sequences is found in the NR and mPFC as compared to the HPC and the TH. Bottom panel: the median differences for 6 comparisons are shown in the above Cumming estimation plot. Each median difference is plotted on the lower axes as a bootstrap sampling distribution. Median

differences are depicted as dots; 95% confidence intervals are indicated by the ends of the vertical error bars.

D) The average Spearman correlation shows that there are more reliable sequences in the NR than in the mPFC. The estimation plot on the right side shows the median difference (dots) and the 95% confidence interval.

Figure 1. Sequential dynamics of cell assemblies.

Figure 2. Spatial organization of NR and mPFC cell assemblies.

A) Representation of the position of a linear silicon probe with 32 recording sites in **(a)** the NR and **(b)** mPFC. NR and mPFC contours (and layers) are delimited by the white dashed line over the green fluorescent Nissl staining (fmi: forceps minor of the *corpus callosum*; 3rd V: third ventricle; PrL: prelimbic area; IL: infralimbic area). The red-orange staining corresponds to the DiI that was deposited at the back of the silicon probe before insertion (D: dorsal; L: lateral).

B) Relationship between anatomical dorso-ventral location of NR (left panel) and mPFC (right panel) neurons (location defined by the site of the probe recording the maximum amplitude of the action potentials) and the template rank showing a linear correlation for NR neurons but not for mPFC ones in a template experiment.

C) Distribution of spatial delays (from the beginning to end of a sequence) across the probe from the topmost site to the bottom-most one for mPFC (red) and NR (blue) neurons.

D) Mean sequence delays (with median difference and 95% confidence interval) across experiments for NR and mPFC.

E) The percentage of experiments showing a dorso-ventral (DV) orientation of sequences was larger in the NR than in the mPFC.

Figure 2. Spatial organization of NR and mPFC cell assemblies.

A

a

NR

3rd
V

250 µm

b mPFC

PrL

I II III V VI

fmi

IL

250 µm

Nissl
DiI

D

L

B

Recording site (NR, left plot):
Slope=-1.06
R²=0.77

NR Template Index

dorsal

ventral

Recording site (mPFC, right plot):
Slope=-0.45
R²=-0.06

mPFC Template Index

620 µm:
32 sites @ 20µm spacing

C

x10⁻³

P(time)

Sequence delay (ms)

D

Mean sequence delays (ms)

Median difference

NR   mPFC   diff.

*

E

% of DV orientation

NR   mPFC

*

Figure 3. Inactivation of NR impairs reliable sequential activation of mPFC neurons at the beginning of the UP state.

A) Two template heat maps of normalized mPFC unit activities in control and during NR inactivation condition showing less stable cell assemblies. The upper traces depict the [0.5-2] Hz filtered LFP in mPFC, where the UP states correspond to troughs in the LFP signal.

B) Inter-spike interval distribution of mPFC neurons grouped data for control (red) and NR inactivation conditions (purple), showing that a decrease in burst firing when the NR is inactivated.

C) Grouped data with median difference and 95% confidence interval of the variability of the APTH for control (CTR) and NR inactivation conditions (MSCI). Insert: APTH: the activity peak triggered histogram quantifies the lag distribution with respect to the population activity peak for each neuron.

D) Distribution of the duration of the UP state in control (red) and NR inactivation (purple) condition, where a peak at short values indicates the emergence of shorter UP states.

E) The number of reliable sequences found in mPFC was decreased when the NR was inactivated (MSCI) as compared to the control (CTR) condition (grouped data, n=5 experiments).

F) The fraction of neurons participating to reliable sequences was decreased when the NR was inactivated.

Figure 3. Inactivation of NR impairs reliable sequential activation of mPFC neurons at the beginning of the UP state.

Figure 4: Inactivation of NR impairs reliable sequential activation of HPC neurons during SPW-Rs.

A) Two template heat maps of normalized HPC unit activities in control (CTR) and NR inactivation (MSCI) condition(bottom of each panel). The upper traces depict the [0.5-2] Hz filtered LFP in mPFC, where the UP states correspond to troughs in the LFP signal, and the black arrow mark the presence of a SPW-R event, which are visible in the [80 200] Hz filtered LFP (bottom trace).

B) SPW-Rs occurrence did not changed when NR was inactivated (grouped data, n=5 experiments, n=913 SPW-R events in CTR, n=149 SPW-R events in MSCI).

C) NR inactivation did not change the normalized power and inner frequency of SPW-Rs.

D) The proportion of reliable cell assemblies during SPW-Rs was considerably reduced when the NR was inactivated.

E) The fraction of reliably participating HPC neurons to sequences during SPW-R events was decreased in NR inactivation condition.

Figure 4: Inactivation of NR impairs reliable sequential activation of HPC neurons during SPW-Rs.