1	COMPUTING HUBS IN THE HIPPOCAMPUS AND CORTEX
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12	Abstract
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13	Nouvel computation which valies on the active stars go and showing of information
14	Neural computation, which relies on the active storage and sharing of information,

15 occurs within large neuron networks in the highly dynamic context of varying 16 brain states. Whether such functions are performed by specific subsets of neurons 17 and whether they occur in specific dynamical regimes remains poorly understood. 18 Using high density recordings in the hippocampus, medial entorhinal and medial 19 prefrontal cortex of the rat, we identify computing substates, or discrete epochs, 20 in which specific computing hub neurons perform well defined storage and 21 sharing operations in a brain state-dependent manner. We retrieve a multiplicity 22 of distinct computing substates within each global brain state, such as REM and 23 nonREM sleep. Half of recorded neurons act as computing hubs in at least one 24 substate, suggesting that functional roles are not firmly hardwired but

dynamically reassigned at the second timescale. We identify sequences of substates
whose temporal organization is dynamic and stands between order and disorder.
We propose that global brain states constrain the language of neuronal
computations by regulating the syntactic complexity of these substate sequences.

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30 Information processing in the brain can be approached on three different levels: 31 biophysical, algorithmic and behavioral (1). The algorithmic level, which remains the 32 least understood, describes the way in which emergent functional computations can be 33 decomposed into simpler processing steps, with architectures mixing serial and 34 massively parallel aspects (2). At the lowest level of individual system components -35 here, in single neurons, such building blocks of distributed information processing can 36 be modeled as primitive operations of storing, transferring, or non-linearly integrating 37 information streams (3).

38 In resting state conditions, both BOLD and EEG signals are characterized by discrete 39 epochs of functional connectivity or topographical stability, defined as resting state 40 networks and microstates, respectively (4-5). The transitions between these large-scale 41 epochs are neither periodic nor random but occur through a not yet understood syntax, 42 which is fractal and complex (5). Does such organization at the macroscopic scale 43 (whole brain and networks of networks for resting state networks and microstates, 44 respectively) also exist at the microscopic scale? Said differently, is neuronal activity 45 at the microcircuit level organized in discrete epochs associated to different "styles" of 46 information processing? Our first goal is to determine whether information processing 47 at the local neuronal circuit level is structured into discrete sequences of substates, and 48 whether such sequences have an observable syntax, whose complexity could be a 49 hallmark of computation. Here we focus on low-level computing operations, performed 50 by individual neurons such as basic information storage and sharing (6-7). To reduce

51 external perturbations, such as sensory inputs, and to establish if primitive processing 52 operations and their temporal sequences are brain state-dependent, we study two 53 conditions: anesthesia and natural sleep, which are characterized by alternating stable 54 brains states, theta (THE)/slow oscillations (SO) and rapid eye movement 55 (REM)/nonREM sleep, respectively. We consider the CA1 region of the hippocampus, 56 the medial entorhinal cortex (mEC) and the medial prefrontal cortex (mPFC) to 57 determine whether algorithmic properties are shared between regions with different 58 cytoarchitectures.

59 The second goal is to determine whether primitive processing operations are 60 localized, or on the contrary, distributed within the microcircuit as proposed for 61 attractor neural networks (8) and liquid state machines (9). This raises two key 62 questions: Are certain operations driven by a few key neurons, similar to hub cells in a 63 rich club architecture (10)? and Do neurons have pre-determined computing roles, such 64 as 'sharer' or 'storer' of information', as well as rigidly prescribed partners in their 65 functional interactions? Said differently - is information routed through a hardwired 66 'neuronal switchboard system' like in early days of telephony? Or dynamically via 67 different addressable nodes like in decentralized peer-to-peer services?

68 Here we demonstrate the existence of a multiplicity of distinct computing substates 69 at the microcircuit level within each of the probed global brain states in both anesthesia 70 and natural sleep. The low-level algorithmic roles played by individual neurons change 71 from one substate to the other and appear largely independent from the underlying 72 cytoarchitecture, with roughly half of the recorded neurons acting as transient 73 computing hubs. Furthermore, we reveal complexity not only at the level of information 74 processing within each substate but also at the level of how substates are organized into 75 temporal sequences, which are neither regularly predictable nor fully random. Substate sequences display an elaborate syntax in all the probed anatomical regions, whosecomplexity is systematically modulated by changes in global brain states.

Taken together, our findings suggest a more distributed and less hierarchical style of
information processing in neuronal microcircuits, more akin to emergent liquid state
computation than to pre-programmed processing pipelines.

81

82 **RESULTS**

83 Analysis design

84 Neurons were recorded simultaneously from the CA1 region of the hippocampus 85 and the medial entorhinal cortex (mEC) under anesthesia (18 recordings from 16 rats), 86 and from the CA1 region and the medial prefrontal cortex (mPFC) during natural sleep 87 (6 recordings from 3 rats, see Figures 1A, S1 and S2 for more details on recordings). 88 We focus on two elementary processing functions: *information storage*, i.e. how much 89 information a neuron buffers over time that it has previously conveyed, as measured by 90 the active information storage (3); and *information sharing*, i.e. how much a neuron's 91 activity information content is made available to other neurons, as measured by mutual 92 information (see e.g. in 7). We use the term *feature* to discuss the metrics we use; i.e. 93 firing, information storage or sharing (Figure 1B-C). We use the same analysis design 94 for all features. The FeatureVector (t_a) contains the values for the descriptive 95 features as measured in window t_a (Figure 1B). For example, for firing features, if 20 96 cells are recorded, FeatureVector (t_a) contains 20 values, representing the firing 97 density of each neuron during window t_a . We first correlate feature vectors for a given 98 window pair (t_a, t_b) . Here, a high correlation value means that the two feature vectors 99 are very similar to one another, i.e. that the features measured at t_a are also found at t_b . 100 After, we build a *feature similarity matrix*, a collection of correlation values between 101 feature vectors for all window pairs, organized in time (Figure 1D). A block along the 102 diagonal indicates a stable state for a given feature, e.g., a period over which units fire, 103 store or share information in a consistently preserved pattern. The axes of the similarity 104 matrix represent time, and repetitions of a block structure along a horizontal or vertical 105 line mean that a stable state for a given feature is reoccurring over time. We then use a 106 simple clustering technique to extract different stable states, which we call *substates*, 107 and display their switching behavior during the recording session (Figure 1D). Finally, 108 we define *computing hubs* as neurons that more heavily participate to the buffering 109 (storage hubs) or the funneling (sharing hubs) of information streams (Figure 1D, see 110 Material and Methods). This notion of computing hub generalizes previously 111 introduced notions of "hubness" (25, 26) beyond the ability to synchronize firing 112 toward more general types of influence on information processing.

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114 Identification of brain global states

115 Unsupervised cluster analysis of the spectral features of the fields recorded in the 116 various brain regions allowed a clear identification of typical global oscillatory patterns 117 (Figure S3), which we call global brain states. In the following, all brain states are 118 identified by the clustering analysis of field recordings performed in the CA1 region 119 (stratum oriens to stratum lacunosum moleculare). Unsupervised clustering identified 120 two states for anesthesia corresponding to epochs dominated by slow (SO state) and 121 theta (THE state) oscillations; and two states during sleep corresponding to REM vs 122 nonREM episodes.

123

124 Brain state-dependent firing substates

As subsets of cells tend to fire spontaneously together in stereotypical patterns (*11*-*126 12*), we first analyzed neuronal firing assemblies. Figure S4 shows that the firing rate, the burst index and entrainment by the phase of the ongoing oscillations were brain 128 region- and brain state-dependent as previously reported (13-14). A simple visual 129 inspection of firing behavior revealed the probable existence of different firing sets, as 130 some neurons tended to fire together during certain epochs; with these epochs repeating 131 themselves over time (Figure S5). To quantify this observation, we constructed the 132 feature vectors $Firing(t_a)$, whose entries are given by the average firing rate of each 133 neuron within the window of analysis t_a . The complex block structure of the similarity 134 matrix revealed a repertoire of state transitions much richer than the one associated to 135 global brain states (Figure 2). In this example, unsupervised clustering revealed a total 136 of six firing substates in mEC (Figure 2A) and five in mPFC (Figure 2D) during THE 137 and REM, respectively, for the two animals. Figure 2B demonstrates that a given brain 138 state was characterized by the switching between different firing substates. Figure 2E 139 shows that a subset of firing substates was shared between brain states, and importantly 140 that the switch from one firing substate state to another did not necessarily coincide 141 with a change in the brain global state (and vice versa). Quantification over all 142 recordings revealed that firing substates occurred 87% of the time during either one of 143 the possible global brain states (Figures 2C and 2F). Substates were found in the mEC, 144 CA1 and mPFC and we found an average of ~5 substates for all brain regions and brain 145 states (Table 1). These results reveal that, although field recordings show stereotyped 146 oscillatory behavior during a given brain state, the firing behavior of neurons display a 147 richer dynamic repertoire. Their activity is compartmentalized in a small number of 148 firing substates, with discrete switching events from one substate to another. The firing 149 substates are brain state and brain region specific, and they are not strictly entrained by 150 the global oscillatory state.

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154 Storage of information is dynamic within a brain state

155 At any given time, neuronal activity conveys an amount of information that can be 156 measured by Shannon entropy. We first focused on active information storage, which 157 measures the fraction of information carried by a neuron *i* at a time *t* that was present 158 in the past activity history of *i* itself (Figure S6A). For storage features, we extract 159 several substates (6 for the mEC in the animal shown in Figure 3A, and 7 for CA1 in 160 the animal shown in Figure 3D), with an average of ~4 states across all animals (Table 161 1). As before, there was no strict alignment between brain state transitions and storage 162 substate transitions (Figures 3A, B and C, E). Yet, brain state specificity of storage 163 states was 80% for all regions (Figures 3C and F and Table 1).

164 Under anesthesia, the absolute storage values were stronger in mEC than in CA1, 165 particularly in layers 3 and 5 of mEC (Figure S7). During natural sleep, however, 166 storage values for CA1 were two orders of magnitude larger than during anesthesia and 167 were as strong as in mPFC (Figure S7). Storage tended to be weaker for all probed 168 regions and layers in THE with respect to SO during anesthesia, but not during natural 169 sleep (Figure S7). Therefore, information storage is dynamically distributed in discrete 170 substates and is brain state- and region-dependent. In particular, the involvement in 171 storage of a neuron could vary substantially along time without being necessarily 172 paralleled by a comparable change in firing rate (Figures 3B and 3E).

173

174 Information sharing is dynamic within a brain state

A primitive processing operation complementary to information storage is information sharing, providing a pseudo-directed metric of functional connectivity between any two circuit units (7). For each neuron *i* we quantified both "shared-in" (*i* acts as a sharing target, with information shared from *j* neurons' past activity, Figure S6B) and "shared-out" information (*i* acts as a sharing source and information is shared

180 to *j* neurons' future activity). We first constructed the feature vector $Sharing(t_a)$ 181 containing the total amount of information funneled through each given neuron 182 (integrated in- and out-sharing strengths, represented by big arrows in Figure 4A), 183 irrespective of whom the information was being shared with. Since in- and out-sharing 184 strengths were strongly correlated (average Pearson correlation >0.9), we ignored the 185 distinction between in- and out-sharing and speak generically of sharing substates. 186 Representative sharing similarity matrices and state sequences are shown in Figure 4B 187 (top) for mEC during anesthesia and mPFC during sleep and in Figure S8 for CA1. 188 Here, we studied only information sharing within regions, because the number of pairs 189 of simultaneous units in different regions that showed significant sharing was too small 190 to reach robust conclusions. We found ~4 sharing substates on average across animals. 191 Sharing states displayed an 86% specificity for a given brain state (Figure 4D, Table 192 1).

193 During anesthesia, we measured a stronger absolute sharing values in CA1 than in 194 mEC, a pattern reversed with respect to storage values, particularly in stratum radiatum 195 (SR) and stratum pyramidale (SP) of CA1, even though mEC layer 5 had a sharing 196 strength comparable to CA1s SR and SP (Figure S9). During natural sleep, the 197 participation to information sharing of SO in CA1 increased by an order of magnitude 198 and was as large as the one of mPFC, notably layer 4 (Figure S9). As for storage, the 199 involvement of a neuron in sharing could vary along time even without corresponding 200 variations of its firing rate (Figures S8B and S8E).

201

202 Sharing assemblies are "liquid"

The previous analysis is focused on sharing strengths at the single cell level. We then determined with which neurons sharing cells were exchanging information, i.e. the detailed network neighborhood of sharing, or *sharing assembly* (cartoon networks in

206 Figure 4A). Two striking features were apparent. First, both the block structure of the 207 sharing assemblies and the state transition sequences are nearly matching the sharing 208 strength ones (Figure 4B), as evidenced by a relative mutual information value of 98% 209 on average. Second, in contrast to sharing strengths, the blocks in the sharing assembly 210 similarity matrix were of a light blue color, indicating a strong variability of sharing 211 assemblies within a given substate. This phenomenon was quantified by *liquidity* 212 analysis, with liquidity being a measure bounded between 0 and 1, where a value of 0 213 represents an absence of internal variability within a substate and a value of 1 214 representing completely random variability (see *Materials and Methods*). The liquidity 215 values of sharing assemblies for all sharing substates throughout all recordings lied 216 below the diagonal (Figure 4C). This result can be better understood considering the 217 toy examples of Figure 4A. The cartoons represent snapshots at three different, non-218 sequential times of a given hub neuron in its sharing network environment. The three 219 considered time frames all fall within the same substate, therefore the overall in- and 220 out-sharing strengths, represented by the orange and grey arrows respectively, are 221 constant (meaning stability). However, the sources and targets of the funneled 222 information can widely vary in time (meaning instability). Although the sum of in-223 going and out-going information remained overall constant within each sharing 224 substate, information was shared over different cell assemblies from one time period to 225 the next. All three brain regions displayed remarkable liquidity in sharing assemblies 226 through all brain states, and liquidity was brain region- and brain state-specific (Figure 227 4C). As reported in Table 2, the largest liquidity was observed for mPFC sharing 228 assemblies during natural sleep (~94%). CA1 displayed a substantial reduction in the 229 liquidity of sharing assemblies when moving from anesthesia to sleep (dropping from 230 \sim 86% in anesthesia to \sim 57% in sleep). Finally, as for the other features, information 231 sharing substates were brain state specific (Figure 4D).

232

233 Loose coordination of substate transitions between brain regions

234 Single units were recorded simultaneously in two regions (CA1 and mEC; CA1 and 235 mPFC). We thus assessed whether substate transition events in one region matched the 236 transition in the other region. We computed the relative mutual information between 237 substate sequences of a given type (e.g. firing, storage or sharing) observed in one 238 region and the other. We did not find significant differences for these measures across 239 the three features (firing, storage, sharing) and therefore pooled them together. The 240 median relative mutual information between substate transitions in the probed cortical 241 and hippocampal regions was 18% during anesthesia (between mEC and CA1) and 42% 242 during natural sleep (between mPFC and CA1). These levels of coordination between 243 substate sequences denoted a lack of perfect parallelism between transitions in the 244 different regions, but they were still well above chance level (Figure S10). Thus, 245 substate dynamics display some coordination between CA1 and mPFC during sleep 246 (Table 3), which is in keeping with the fact that information exchange occurs between 247 the two regions during sleep (15). The weak coordination under anesthesia suggests that 248 circuits may operate more independently from one another in this condition (but still 249 not completely).

250

251 A large fraction of cells can act as computing hubs

Functional, effective, and anatomical hub neurons (mostly GABAergic) have been identified in the brain (*16*). We complement the concept, introducing *storage* and *sharing hubs*, i.e. neurons displaying an elevated storage or sharing values, respectively (see Methods). In contrast to the sparsity of functional, effective, and anatomical hubs, a large fraction of cells acted as a computing hub in at least one substate, as illustrated in Figure 5A. Computing hubs could be recruited across all probed regions and layer 258 locations (Figure 5B and C). As summarized in Figure 5B, the probability of serving as 259 computing hub – storage or sharing confounded – was of 40% or more on average for 260 almost all layers, apart from the, possibly under-sampled, stratum lacunosum 261 moleculare and stratum radiatum in CA1. We observed a general tendency for 262 inhibitory interneurons to have a larger probability to serve as computing hubs than for 263 excitatory cells. This tendency was particularly strong for cortical regions and was 264 notably significant in layer 5 of mEC (during anesthesia) and layer 3 of mPFC (during 265 sleep), for which the probabilities of inhibitory interneurons serving as computing hub 266 in at least one substate approached 70%. The probability of serving as a computing hub 267 at least once was relatively similar when evaluated separately for storage or sharing. In 268 particular, 43% of the neurons serving as a storage hub in a substate could serve as a 269 sharing hub in another substate, but in general, not simultaneously as only 12% of the 270 neurons were "multi-function" hubs.

Despite this large flexibility in the dynamic assignment of hub roles, the notion of hub continued to make sense within each individual substate. Within a substate, on average only ~9% of cells acted as hub (storage or sharing pooled), so still a strict "elite" (although not a permanent one but appointed just within the associated state). The set of recruited hubs constituted thus at each time a characteristic fingerprint of the active substates (with only 4% of the substates being "hubless").

We also studied the probability that a computing hub emerged in a given layer (Figure 5C). During anesthesia, all probed layers of CA1 and mEC showed a ~20% uniform probability for a storage and sharing computing hub to emerge. Natural sleep was associated to an enhanced recruitment of computing hubs. The probabilities of hub emergence exceeded ~40% for storage hubs in layer 5 of mPFC and in SP of CA1. The analysis of deep or superficial CA1 SP principal neurons, which are involved in different microcircuits (*17-18*), did not reveal an intra-layer distribution of computing

hubs (not shown). These results suggest that the probability that a neuron serves ascomputing hub is not correlated to its anatomical region or layer location.

286 Finally, we tested whether computing hubs were characterized by high firing rates. 287 Using the same procedure utilized to extract computing hubs, we found that 62% of the 288 cells were high-firing at least in one firing substate with 70% being putative 289 interneurons. Remarkably, there was a poor overlap between computing hubs and high 290 firing rate cells. Table 3 already shows that storage and sharing substate sequences are 291 only loosely coordinated with firing substate sequences (cf. as well firing rate 292 information in Figures 3B-E and S8B-E). Furthermore, being a high firing rate cell does 293 not guarantee that this cell will also be a computational hub (or the other way around). 294 This is also shown in Figure 5C, where the yellow levels over the histogram bars 295 indicate the fraction of storage and sharing hubs which also happen to be high firing 296 cells. We conclude that a storage hub can have a normal or even smaller than average 297 firing rate.

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300 The syntax of substate sequences is complex and brain state-dependent

301 Collectively, our results demonstrate the existence of substate sequences in three 302 different brain regions during anesthesia and natural sleep. Using a linguistics analogy 303 (Figure S11A), we assign a *letter* to each identified substate (represented by a color in 304 the figures). The temporal sequence of substates thus translates into a stream of *letters*. 305 However, if we consider the three features simultaneously, we obtain a stream of 3 letter 306 words (Figure S11B). All combinations of possible letters from our 3 features define 307 the dictionary of words that can be expressed. We represent a stream of words as a 308 switching table (Figure 6A). This allows us to explore two aspects of the "neuronal 309 language": the statistics of the words and the statistics of the transitions between the 310 words (the syntax). We found that the words were mostly (85%) brain state-specific, as 311 expected since the substates *letters* are already brain state-specific (cf. Figures 3C and 312 F, 4D). Although the syntactic rules structuring the production of *words* are unknown, 313 we can quantify their complexity. Algorithmic information theory (19), the minimum 314 description length framework (20) and the Lempel-Ziv method (21) link complexity to 315 the notion of compressibility. As illustrated in Figure 6B, an ordered, regular switching 316 table requires a short description, as a small list of instructions can be written to 317 reproduce the table (e.g. word D 100 times, followed by word B 88 times, etc.). At the 318 opposite extreme, a completely random switching table would need a lengthy 319 exhaustive description -as many instructions as the length of the table itself. A complex 320 switching table stands between regularity and randomness and requires a description 321 that is compressed, longer than a regular table but shorter than a random table.

Figure 6C shows that the syntax was complex (between 0 and 1) for all brain regions and brain states and that THE/REM states were more complex than SO/nonREM states. We added two recordings from mPFC under anesthesia for comparison. Figure 6D shows that the measured complexity was significantly larger than the upper threshold for regularity and significantly smaller than the lower threshold for randomness (p<0.05, Bonferroni Corrected, direct c.i. comparison).

Finally, we assessed whether switching from SO to THE or from nonREM to REM increased the complexity. As shown in Figure 6E, the tendency was toward an increase of complexity in all cases, from +30% for mEC during anesthesia and mPFC during anesthesia or sleep to roughly +10% for CA1 during anesthesia or sleep. This relative increase was always significant (p<0.05, Bonferroni Corrected, c.i. comparison) apart from CA1, for which two recordings displayed increased complexity during nonREM sleep. We conclude that the syntax is complex and brain state-dependent.

335

336 What determines complexity?

We then investigated which factors contribute to complexity. Different durations of *words* may account for variations in complexity. Although *word* dwell times were different by one order of magnitude between anesthesia and sleep with median ~18 min (~10 min 1st quartile, ~28 min 3rd quartile) during anesthesia and ~1.4 min (~1 min 1st quartile, ~2.1 min 3rd quartile) during sleep, complexity values for anesthesia and natural sleep were similar.

We also evaluated the burstiness coefficient, *B* (22), of the stream of *words*. This coefficient ranged between $-1 \le B \le 1$, with B = -1 corresponding to a perfectly periodic stream of *words*, B = 0 to a Poisson train and B = 1 to a maximally bursting stream. We found a positive correlation between burstiness and complexity (Figure S11A, *p*<0.01, Bootstrap c.i). Burstiness was greater during THE/REM (0.15) than during SO/nonREM (0.09 *p* = 0.03, Kruskal-Wallis test), which may contribute to the increased complexity found during THE/REM.

The richness of the dictionary also affects complexity (21). We therefore evaluated the Used Dictionary Fraction, i.e. the ratio between the number of observed *words* and the maximum theoretical number of *words*, i.e. the *dictionary*. We find a significant positive correlation between the Used Dictionary Fraction and complexity (Figure S12B, p<0.05, Bootstrap c.i). The richness of the dictionary was greater during THE/REM (21%) than during SO/nonREM (14%, p = 0.032, Kruskal-Wallis test),

356 which may also contribute to the increased complexity found during THE/REM.

357 A bivariate linear regression of complexity over burstiness and Used Dictionary 358 Fraction revealed a correlation of 0.62 (p<0.05, Bootstrap c.i) between predicted and 359 observed complexity, demonstrating that complexity is largely explained by burstiness 360 and the Used Dictionary Fraction.

Finally, we verified that our results did not depend on the measure of complexity. Redoing analyses using Lempel-Ziv complexity (21), which was previously used to analyze neural activity (23-24), lead to qualitatively equivalent results. Lempel-Ziv complexity also strongly correlated with our measure of complexity (Figure S11C Pearson correlation 0.84, p<0.001, bootstrap c.i.).

366

367 **Discussion**

368 Here we demonstrate two levels of organization of brain activity. At the single cell 369 level, we find that a large proportion of recorded neurons act as computing hubs during 370 discrete time epochs (substates) within a given stable brain state (e.g. REM and 371 nonREM). At the microcircuit level, we find a rich repertoire of computational substates 372 characterized by temporally structured sequences, whose complexity was modulated by 373 the global brain oscillatory state. Such type of organization was shared between three 374 anatomical different brain regions: the hippocampus, the medial entorhinal cortex and 375 the medial prefrontal cortex.

376

The "hubness" of a neuron may be determined by fixed features, e.g. an exceptional 377 378 extension of axonal arborizations (25); a suitable location in the circuit wiring diagram 379 facilitating the control of synchronization (26); or yet, some developmental "droit 380 d'aînesse" (16). During natural sleep and anesthesia, however, we find that >40% of 381 the recorded neurons act as a computational hub during at least one substate, meaning 382 that computing hubs form a rather open and not so elitist club. The computational 383 hubness is dynamic - a neuron acting as a hub in a given substate may not be a hub in 384 a different substate or may swap its nature (e.g. converting from a storage to a sharing 385 hub). The stronger tendency for putative inhibitory cells to serve as hubs (>70%) is in 386 keeping with the known role of GABAergic cells in orchestrating network activity (16,

387 25-26). Furthermore, because our analysis was limited to few brain states, the
388 proportion of putative principal and GABA neurons acting as computational hubs may
389 even be an underestimate. Perhaps all neurons act as computational hubs during specific
390 brain states (including exploration and quiet awakening).

391

392 That hubs share information with ever changing source and target neurons is in 393 apparent contradiction with the existence of sequential firing of cell assemblies in 394 cortical and hippocampal circuits, including during nonREM sleep (11,16, 27-37). Our 395 information-theoretical analyses require the use of at least 5 s long sliding windows, 396 which is not sufficient to detect fast sequences of activation, as replay events occur 397 within 500 ms (38). Interestingly, replay sequences are not strictly stable as they 398 demonstrate inter-cycle variability (39), which may reflect liquidity. The liquid nature 399 of information sharing suggests that neuronal activity is not frozen at fixed-point 400 attractors as in classic artificial neural networks (40) but may be sampling the broad 401 basin of attraction of shallow attractors (8) or higher-dimensional manifolds "at the 402 edge of chaos", as found in reservoir computing schemes (41-43). In this case, 403 information is shared across extremely volatile assemblies within a given substate. The 404 assembly dynamics are thus "liquid" – i.e. neither frozen into crystallized patterns, nor 405 fully random as in a gas – and are only mildly constrained to robustly maintain the 406 computational role of sharing hubs while preserving entropy of firing patterns, and 407 therefore bandwidth for local computations (44). This preservation of hub function in a 408 heterogeneous and reconfiguring circuit can be seen as a form of homeostasis of the 409 computing role, generalizing the concept of homeostasis of circuit behavior evidenced 410 in invertebrate systems (45-46). While this latter homeostasis preserves the functional 411 level, in our case homeostasis would extend down to the algorithmic level, referring to 412 the three-level hierarchy proposed by Marr & Poggio (1).

413

414 During a "stable" behavior such as resting state, analysis of BOLD and EEG signals 415 consistently revealed the presence of temporal sequences of resting state networks and 416 topographical microstates, respectively (4-5). Here, we find that an analogous switching 417 between discrete states occurs at a completely different scale of microcircuits. During 418 a "stable" oscillatory regime (e.g. theta rhythm), neuronal computation is indeed 419 organized in temporal sequences of computational substates. Interestingly, while field 420 oscillations constrain neuronal firing and neuronal firing produces field oscillations 421 (47), we find only a loose match between the switch from one oscillatory mode to the 422 other and the switch from one substate to the other. Transitions between global states – 423 related to the scale of mesoscale collective dynamics- sometimes anticipate and 424 sometimes follow transitions between firing, storage or sharing substates --related to the 425 scale of microscopic firing dynamics-, as if dynamic changes occurring at either one of 426 the scales had destabilizing effects on the dynamics at the other scale (in both directions, 427 meso- to micro-scale and micro-to meso-scale). The behavior of CA1 cells may reflect 428 specific internal dynamics, not tightly controlled by the CA1 local field which mostly 429 reflects synaptic inputs originating from outside the CA1 region. Importantly, the 430 repertoire of computing substates is brain state specific. Beyond proposals that 431 oscillations are central for the routing of information between regions (47-48), we thus 432 suggest here that global oscillatory states could also organize information processing 433 within local regions by enforcing the use of their own state-specific "languages" 434 (expressed in terms of combinations of alternative intrinsic substates).

435

436 Signatures of computation can be identified even if the function and meaning of the
437 computation are unknown and even when system states are sampled partially, as it is
438 the case for the present study. This allowed us to extract a symbolic representation of

substates (*letters*) for a given feature, which make *words* when considering several features. The syntax of the substate word sequences is complex, standing between order and randomness (as it was already the case for the sharing dynamics within each substate). The capacity to generate complex sequences of patterns is a hallmark of selforganizing systems and has been associated to their emergent potential to perform universal computations (70). Moreover, dynamics at the "edge of chaos" confer advantages for information processing (41-43).

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447 Importantly, we find that the syntactic complexity of substate sequences is brain 448 state-dependent as it was the case for the substate dictionaries, and more complex 449 during theta oscillations/REM sleep than during slow oscillations/nonREM sleep, 450 suggesting an increased load of computation in the former brain state. Remarkably, the 451 temporal complexity of activation sequences was also shown to be modulated by brain 452 states at the macro-scale level of whole-brain dynamics (49). In keeping with the view 453 that slow/theta oscillations measured during anesthesia share general properties with 454 nonREM/REM sleep (50-52), we found similar rules of organization in terms of 455 substate sequences and their complexity, despite the fact that the word dwell times in 456 anesthesia are one order of magnitude greater than during natural sleep. We speculate 457 that the nature of the undergoing oscillation (slow vs theta) constrains the repertoire of 458 words used and their syntax, modulating the type of computation performed by the 459 recruitment of varying computing hubs. Sleep, oscillatory patterns, and neuronal firing 460 are altered in numerous neurological disorders, including epilepsy (18, 53-56) and 461 therefore it will be important to assess whether the repertoire of substates and the syntax 462 are likewise affected.

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464 In conclusion, our results reveal a rich algorithmic-level organization of brain 465 computations during natural sleep and anesthesia, which combines a complex 466 combinatorics of discrete states with the flexibility provided by liquidly reconfiguring 467 assemblies. While we cannot yet prove that this substate dynamics is functionally 468 relevant, it has the potential to serve as a substrate for previously undisclosed 469 computations. The next aim will be to perform the similar analysis during specific 470 behavioral tasks, such as goal-driven maze navigation. Words and/or their sequence 471 may sign specific cognitive processes. The fact that the algorithmic instructions and 472 primitive processing operations are similar in three brain regions with different 473 architectural organizations suggests the existence of a basic architecture for low-level 474 computations shared by diverse neuronal circuits.

475

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672

- 673 **Tables**
- 674 *Table 1 Number of states and their oscillatory mode specificity*
- 675

Oscillatory mode specificity	Median # states
1.0	2
0.88 (0.78, 0.95)	5
0.80 (0.69, 0.88)	4
0.86 (0.78, 0.95)	4
	1.0 0.88 (0.78, 0.95) 0.80 (0.69, 0.88)

676 *Median* (1st, 3rd quartile), all regions and conditions

677

678 Table 2 – Sharing assembly liquidity across regions and conditions

679

Liquidity sharing strengths	Liquidity sharing assemblies
0.31(0.22, 0.45)	0.84 (0.72, 0.89)
0.39 (0.36, 0.47)	0.86 (0.77, 0.89)
0.04 (0.03, 0.05)	0.57 (0.50, 0.68)
0.18 (0.11, 0.26)	0.94 (0.89, 0.96)
	0.31 (0.22, 0.45) 0.39 (0.36, 0.47) 0.04 (0.03, 0.05)

680 *Median* (1st, 3rd quartile), SO/THE and non-REM/REM states confounded

682 Table 3 – Matching between substate sequences of different types across

683 conditions and regions

684

nfounded) Storage substates		
Storage substates		
C	0.38 (0.32, 0.53)	
Sharing substates	0.47 (0.36, 0.61)	
Sharing substates	0.38 (0.31, 0.49)	
mEC substates with HPC substates (all types)		
confounded)		
Storage substates	0.50 (0.46, 0.59)	
Sharing substates	0.45 (0.38, 0.66)	
Storage substates	0.42 (0.35, 0.57)	
HPC substates with mPFC substates (all types)		
(Sharing substates substates (all types) confounded) Storage substates Sharing substates Storage substates Storage substates	

685 *Median (1st, 3rd quartile), SO/THE and non-REM/REM states confounded*

687 Materials and Methods

688 Data information. We use in this work a portion of the data (13 out of 18 689 experiments) initially published by Quilichini et al. (2010), which includes local field 690 potentials (LFPs) and single-unit recordings obtained from the dorsomedial entorhinal 691 cortex (mEC) of anesthetized rats. We also use a portion of the data (2 out of 16 692 experiments) initially published by Ferraris et al. (2018), which includes LFPs and 693 single-units recorded in the medial prefrontal cortex (mPFC) under anesthesia. Seven 694 recordings are original data in both mEC and dorsal hippocampus (HPC) under 695 anesthesia, and 10 recordings in 4 animals during natural sleep in HPC and mPFC. See 696 Figures S1 and S2 for details on recordings, number of cells, and layers recorded.

697 Animal surgery. We performed all experiments in accordance with experimental 698 guidelines approved by the Rutgers University and Aix-Marseille University Animal 699 Care and Use Committee. We performed experiments on 13 male Sprague Dawley rats 700 (250–400 g; Hilltop Laboratory Animals), 8 male Wistar Han IGS rats (250-400g; 701 Charles Rivers) and 3 male Long Evans rats (350-400g; Charles River). We performed 702 acute (anesthesia) experiments on the Sprague Dawley and 7 of the Wistar rats, which 703 were anesthetized with urethane (1.5 g/kg, i.p.) and ketamine/xylazine (20 and 2 mg/kg, 704 i.m.), additional doses of ketamine/xylazine (2 and 0.2 mg/kg) being supplemented 705 during the electrophysiological recordings. We performed chronic (natural sleep) 706 experiments on one Wistar and the Long Evans rats, which were anesthetized using 707 isoflurane 2% in 11/min of O₂ for the surgery procedure. In both cases, the body 708 temperature was monitored and kept constant with a heating pad. The head was secured 709 in a stereotaxic frame (Kopf) and the skull was exposed and cleaned. Two miniature 710 stainless-steel screws, driven into the skull, served as ground and reference electrodes. 711 To reach the mEC, we performed one craniotomy from bregma: -7.0 mm AP and +4.0 712 mm ML; to reach the CA1 area of HPC, we performed one craniotomy from bregma: -

713 3.0 mm AP and +2.5 mm ML in the case of HPC coupled to mEC recordings, and from 714 bregma: -5.6 mm AP and +4.3 mm ML-3.0 mm in the case of HPC coupled to mPFC 715 recordings; to reach the mPFC, we performed one craniotomy from bregma: +3 mm AP 716 and +0.8 mm ML. We chose these coordinates to respect known anatomical and 717 functional connectivity in the cortico-hippocampal circuitry (51, 57-59). We used 718 different types of silicon probes to record the extracellular signals. In acute experiments, 719 the probes were mounted on a stereotaxic arm. We recorded the dorso-medial portion 720 of the mEC activity using a NeuroNexus CM32-4x8-5mm-Buzsaki32-200-177 probe 721 (in 8 experiments), a 10-mm long Acreo single-shank silicon probe with 32 sites (50 722 µm spacing) arranged linearly (in 5 experiments), or a NeuroNexus H32-10mm-50-177 723 probe (in 5 experiments), which was lowered in of the EC at 5.0-5.2 mm from the brain 724 surface with a 20° angle. We recorded HPC CA1 activity using a H32-4x8-5mm-50-725 200-177 probe (NeuroNexus Technologies) lowered at 2.5 mm from the brain surface 726 with a 20° angle (in 4 experiments), a NeuroNexus H16-6mm-50-177 probe lowered at 727 2.5 mm from the brain surface with a 20° angle (in 2 experiments) and a E32-1shank-728 40µm-177 probe (Cambridge Neurotech) lowered at 2.5 mm from the brain surface 729 with a 20° angle (in 1 experiment). We recorded mPFC activity using NeuroNexus 730 H32-6mm-50-177 lowered in the layer 5 at 3 mm perpendicularly from the brain surface 731 (in 2 experiments). In chronic experiments, the probes were mounted on a movable 732 micro-drive (Cambridge Neurotech) fixed on the skull and secured in a copper-mesh 733 hat. We recorded HPC CA1 activity (probes lowered perpendicularly at 2.5 mm from 734 the brain surface) using a Neuronexus H32-Poly2-5mm-50-177 probe (in 2 735 experiments), a Cambridge Neurotech E32-2shanks-40µm-177 probe (in 1 experiment) 736 and a NeuroNexus H32-4x8-5mm-50-200-177 probe (in 1 experiment). We recorded 737 mPFC activity (probes lowered perpendicularly at 3.0 mm from the brain surface) using 738 a NeuroNexus H32-4x8-5mm-50-200-177 probe (in 2 experiments), and a Neuronexus

H32-Poly2-5mm-50-177 probe (in 1 experiment). The on-line positioning of the probes
was assisted by: the presence of unit activity in cell body layers and the reversal of theta
([3 6] Hz in anesthesia, [6 11] Hz in natural sleep) oscillations when passing from L2
to L1 for the mEC probe, and the presence in stratum pyramidale either of unit activity
and ripples (80-150 Hz) for the HPC probe, and the DV depth value and the presence
of intense unit activity for the mPFC.

745 At the end of the recording, the animals were injected with a lethal dose of 746 Pentobarbital Na (150mk/kg, i.p.) and perfused intracardially with 4% 747 paraformaldehyde solution. We confirmed the position of the electrodes (DiI was 748 applied on the back of the probe before insertion) histologically on Nissl-stained 40 µm 749 section as reported previously in detail (60). We used only experiments with appropriate 750 position of the probe for analysis. The numbers of recorded single units in different 751 anatomical locations for the different retained recordings are summarized in Figure S2.

752

753 Data collection and spike sorting. Extracellular signal recorded from the silicon 754 probes was amplified (1000x), bandpass filtered (1 Hz to 5 kHz) and acquired 755 continuously at 20 kHz with a 64-channel DataMax System; RC Electronics or a 258-756 channel Amplipex, or at 32 kHz with a 64-channel DigitalLynx; NeuraLynx at 16-bit 757 resolution. We preprocessed raw data using a custom-developed suite of programs (61). 758 After recording, the signals were downsampled to 1250 Hz for the local field potential 759 (LFP) analysis. Spike sorting was performed automatically, using KLUSTAKWIK 760 (http://klustakwik.sourceforge.net (62)), followed by manual adjustment of the clusters, 761 with the help of auto-correlogram, cross-correlogram and spike waveform similarity 762 matrix (KLUSTERS software package, http://klusters.source-forge.net (63)). After 763 spike sorting, we plotted the spike features of units as a function of time, and we 764 discarded the units with signs of significant drift over the period of recording.

765 Moreover, we included in the analyses only units with clear refractory periods and well-766 defined clusters. Recording sessions were divided into brain states of theta and slow 767 oscillation periods. The epochs of stable theta (THE in anesthesia experiments, REM 768 in natural sleep experiments or slow oscillations (SO in anesthesia experiments, non-769 REM in natural sleep experiments) periods were visually selected from the ratios of the 770 whitened power in the theta band ([3 6] Hz in anesthesia, [6 11] Hz in natural sleep) 771 and the power of the neighboring bands ([1 3] Hz and [7 14] Hz in anesthesia, [12 20] 772 Hz in natural sleep) of EC layer 3 LFP, which was a layer present in all the 18 anesthesia 773 recordings, or layer 5 mPFC recordings in natural sleep recordings, and assisted by 774 visual inspection of the raw traces (60) (Figure S3). We then used band-averaged 775 powers over the same frequency ranges of interest as features for the automated 776 extraction of spectral states via unsupervised clustering, which confirmed our manual 777 classification.

We determined the layer assignment of the neurons from the approximate location of their somata relative to the recording sites (with the largest- amplitude unit corresponding to the putative location of the soma), the known distances between the recording sites, and the histological reconstruction of the recording electrode tracks.

782

783 Characterizations of single unit activity. We calculated pairwise cross-784 correlograms (CCGs) between spike trains of these cells during each brain state 785 separately (60, 64-65). We determined the statistical significance of putative inhibition 786 or excitation (trough or peak in the [+2 5] ms interval, respectively) using the 787 nonparametric test and criterion used for identifying monosynaptic excitations or 788 inhibitions (60, 64-65), in which each spike of each neuron was jittered randomly and 789 independently on a uniform interval of [-5 5] ms a 1000 times to form 1000 surrogate 790 data sets and from which the global maximum and minimum bands at 99% acceptance

P1 levels were constructed. Inspection of CCGs thus allowed to identify single units as
putatively excitatory or inhibitory, an information which we used to perform the
computing hub characterizations in Figure 5B.

To perform the analyses of Figure S4 we then computed the burst index and the phase modulation of units. Burst index denotes the propensity of neurons to discharge in bursts. We estimated the amplitude of the burst spike auto-correlogram (1 ms bin size) by subtracting the mean value between 40 and 50 ms (baseline) from the peak measured between 0 and 10 ms. Positive burst amplitudes were normalized to the peak and negative amplitudes were normalized to the baseline to obtain indexes ranging from -1 to 1. Neurons displaying a value of 0.6 were considered bursting.

801 To establish the phase modulation of units, we concatenated different epochs of slow 802 or theta oscillations, and estimated the instantaneous phase of the ongoing oscillation 803 by Hilbert transform of the [0.5 2] Hz or [3 6] Hz in anesthesia and [6 11] Hz in natural 804 sleep filtered signal, for slow or theta oscillations respectively. Using linear 805 interpolation, we assigned a value of phase to each action potential. We determined the 806 modulation of unit firing by Rayleigh circular statistics; p < 0.05 was considered 807 significant. We first assessed circular uniformity of the data with a test for symmetry 808 around the median (66) and we performed group comparison tests of circular variables 809 using circular ANOVA for uniformly distributed data and using a nonparametric multisample test for equal medians "CM-test", similar to a Kruskal-Wallis test, for non-810 811 uniformly distributed data (Berens, 2009; 812 https://philippberens.wordpress.com/code/circstats), and p < 0.05 was considered 813 significant.

814

815 Feature-based state extraction. We performed a sliding-window analysis of the
816 recorded LFP time-series and single unit spike trains, extracting in a time-resolved

817 manner a variety of different descriptive features. For all the considered features (see 818 specific descriptions in later sub-sections), we use similar window sizes and overlap for 819 the sake of a better comparison. For anesthesia recordings, we adopted a long window 820 duration of 10 s – demanded by the estimation needs for the most "data-hungry" 821 information-theoretical features – with an overlap of 9 s. For natural sleep recordings, 822 we adopted a window duration of 10 s with an overlap of 9 s.

We computed each set of descriptive features and compiled them into multi-entry
vectors FeatureVector(t) for every time-window centered on different times t.

825 We then compute a similarity matrix M_{sim} , to visualize the variability over time of 826 the probed feature set. The entries $M_{sim}(t_a, t_b)$ are given by the Pearson correlation 827 coefficient between the entries in the vectors $FeatureVector(t_a)$ and 828 FeatureVector(t_b), treated as ordered sequences, and are thus bounded between -1 829 and +1. Blocks of internally elevated correlation along the similarity matrix diagonal 830 denote epochs of stable feature configurations. Similar configurations are detected by 831 the presence of off-diagonal highly-internally correlated blocks and the existence of 832 multiple possible configurations by the poor correlation between distinct blocks.

833 We then extracted feature-based states using a standard iterative K-means algorithm 834 (67) to cluster the different vectors FeatureVector(t), based on the correlation 835 distance matrix defined by 1- M_{sim} . We defined the substates of different types as the 836 different clusters obtained for different feature types. We chose the number of clusters 837 K by clustering using $K = 2, 3, \dots 20$ and first guessing K using a maximal silhouette 838 criterion (68) across all Ks. We also inspected dendrograms from single-linkage 839 clustering as a cross-criterion. Using both pieces of information the K was manually 840 adjusted case-by-case (up to ± 2 clusters with respect to the unsupervised silhouette 841 criterion) to best match the visually apparent block structure of the similarity matrix 842 M_{sim} , which results in an optimized K selection for each recording.

843

844 Feature Robustness. To compute the robustness of the feature computation, the 845 original spiking times were randomly shuffled 1000 times and the features recomputed 846 for each instance for 2 files, one in anesthesia and one in natural sleep. To compare it 847 to the original features computed, the k for each recording and each feature was kept 848 the same. The information retained after shuffling was computed by dividing the mutual 849 information between the shuffled features and the original by the entropy of the new 850 feature set. We found a significant difference for both anesthesia and in natural sleep 851 across all features, and the results have been quantified in Table S1.

852

853 Global oscillatory states. We defined eight different unequally-sized frequency 854 ranges, which were manually adjusted recording-by-recording to be better centered on 855 the recording-specific positions of the slow-wave and theta peaks and of their 856 harmonics (e.g., 0-1.5 Hz, 1.5-2 Hz, 2-3 Hz, 3-5 Hz, 5-7 Hz, 7-10 Hz, 10-23 Hz and 857 23–50 Hz for the anesthesia spectrogram and the similarity matrix of Figure S3A). We 858 averaged the spectrograms over all channels within each of the layers in the 859 simultaneously recorded regions (e.g. EC and CA1 for anesthesia) and then we coarse-860 grained the frequencies by further averaging over the eight above ranges. We compiled 861 finally all these layer-averaged and band-averaged power values into time-dependent 862 vectors Spectra(t), with a number of entries given by eight (number of frequency 863 bands) times the number of layers probed in the considered recording, i.e. up to eight 864 (CA1 stratum oriens (SOr), stratum pyramidale (SP), stratum radiatum (SR) and 865 stratum lacunosum moleculare (SLM); EC layers 2, 3 and 5; and PFC layers 1,2, 3 and 866 5), yielding at most 64 entries. We then processed these spectral features as described 867 in the previous section to extract global oscillatory states –as any other substate type– 868 via unsupervised clustering.

869

870 Firing sets and firing hubs. Not all neurons are equally active in all temporal 871 windows. To determine typical patterns of single neuron activation we binned the 872 spiking data for each unit in 50 ms windows – if a neuron fired within that window the result was a '1', if it did not fire the result was a '0'. We enforced a strictly binary 873 874 encoding, i.e. we attributed to a bin a '1' symbol even when more than one spike was 875 fired within this bin. Our bin size choice however was such to maintain the loss of 876 information when ignoring multiple firing events within a bin was less that 5%. Note 877 furthermore that for the majority of spike trains multiple firing events were extremely 878 rare, i.e. apart from a few cases the information loss was way smaller than 5%. We then 879 averaged over time this binned spike density, separately for each single unit and within 880 each time window and compiled these averages into time-dependent vectors 881 Firing(t), with N entries, where N is the overall number of single units probed within 882 the considered recording. We constructed separate feature vectors for each of the 883 simultaneously recorded regions. Firing substate prototypes were given by the centroids of the clusters extracted from the similarity matrix M_{sim} resulting from the stream of 884 885 Firing(*t*) feature vectors.

We then defined a neuron to be a *high-firing cell* in a given state if its firing rate in the state prototype vector was higher than the 95% percentile of all concatenated state prototype vector entries.

889

Active Information Storage. Within each time-window we computed for each single unit an approximation to the Active Information Storage (AIS). AIS is meant to quantify how much the activity of a unit is maintaining over time information that it was conveying already in the past (*3*, *6*). This information-theoretical notion of storage is distinct from the neurobiological notion of storage in synaptic weights. It is indeed

an activity-based metric (hence the adjective "active"), able to detect when temporal
patterns in the activity of a single unit can serve the functional role of "memory buffer".
AIS is strictly defined as:

898

899 $AIS_i = MI[i(t), i(\rightarrow t)]$

900

901 i.e. as shared information between the present activity i(t) of a single unit *i* and its past 902 history of activity $i(\rightarrow t)$ (cf. Figure S6A). Prior to computing mutual information, we 903 binned all spike trains with method as for determining the Firing(*t*) descriptive 904 feature vector. The limited amount of available data within each temporal window 905 makes necessary to introduce approximations. Therefore, we replaced the full past 906 history of activity $i(\rightarrow t)$ with activity at a time in the past $i(t-\tau)$ and then summed over 907 all the possible lags:

- 908
- 909 $\widehat{AIS}_i = \Sigma_{\tau} MI[i(t), i(t-\tau)]$
- 910

911 where the lag τ was varying in the range $0 \le \tau \le 0.5T_{\theta}$, where T_{θ} is the phase of the theta 912 cycle. Note that MI values were generally vanishing for longer latencies (cf. Figure 913 S13A). We evaluated MI terms using a 'plug-in' function estimator on binarized spike-914 trains, which takes the binned spike trains of two neurons for a defined time window 915 and computes the mutual information and entropy values of the two variables (6). Concretely speaking, we estimated the probability p that a bin includes a spike and the 916 917 complementary probability 1 - p that a bin is silent for each unit, by direct counting of 918 the frequency of occurrence of "1"s and "0"s in the binned spike trains of each unit. 919 These counts yielded the probability distributions P(i) and P(j) that two neurons i and j 920 fire or not. Analogously, we sampled directly from data the histogram P(i,j) of joint

921 spike counts for any pair of two units *i* and *j*. These histograms were then directly922 "plugged in" (hence the name of the used estimator) into the definition of MI itself:

923
$$MI(i,j) = \sum_{i} \sum_{j} P(i,j) \log_2 \frac{P(i,j)}{P(i)P(j)}$$

924 We then subtracted from each MI value a significance threshold (95-th percentile of 925 MI estimated on shuffled binarized trains, 1000 replicas), putting to zero non-926 significant terms (and thus negative after bias subtraction). Although such corrected 927 plug-in estimator is very rough, it is sufficient in our application in which we are not 928 interested in quantitatively exact values of MI but just in relative comparisons of values, 929 finalized to state clustering over a large amount of observations. We compiled the N930 resulting \hat{AIS}_i values into time-dependent vectors Storage(t), constructing separate 931 vectors for each of the simultaneously recorded regions. We then constructed storage 932 substates through unsupervised clustering based on the M_{sim} matrices, as previously 933 described. We defined a neuron to be a *storage hub* in a given state if its \widehat{AIS}_i value in 934 the state prototype vector was higher than the 95% percentile over all entries of 935 concatenated cluster prototype vectors. Such conservative threshold guarantees that 936 only neurons with exceptionally high AIS values are labeled as hubs. While we may 937 have some false negatives -i.e. neurons with values in the right tail of the AIS 938 distribution not labeled as hubs-, we are thus protected against false positives.

AIS absolute values varied widely between the different recordings. To compare AIS measures and their relative changes between global oscillatory states across recordings, we first averaged AIS for all the units within a specific anatomic layer. We then normalized these average AIS values by dividing them by the average AIS value in the SO state (in anesthesia) or the nonREM state (in natural sleep) for the specifically considered recording and layer. The results of this analysis are shown in Figure S7, where different lines correspond to different recordings.

946

947 Information sharing networks and strengths. Within each time-window we 948 computed time-lagged Mutual Information $MI[i(t), i(t-\tau)]$ between all pairs of spike 949 density time-series for different single units *i* and *j* (evaluated via the same binning 950 method for determining the Firing(t) descriptive feature vector). Although MI is not 951 a directed measure, a pseudo-direction of sharing is introduced by the positive time-lag, 952 supposing that information cannot be causally shared from the future. Thus, for every 953 directed pair of single units i and j (including auto-interactions, with i = j), we defined 954 pseudo-directed information sharing as:

- 955
- 956

 $I_{shared} (j \rightarrow i) = \Sigma_{\tau} \operatorname{MI}[i(t), j(t-\tau)]$

957

958 where the lag τ was varying in the range $0 \le \tau \le 0.5T_{\theta}$, where T_{θ} is the phase of the 959 theta cycle. Once again, we estimated MI terms via direct plug-in estimators on 960 binarized spike trains, as with storage, subtracting a significance threshold (95-th 961 percentile of MI estimated on shuffled binarized trains, 400 replicas) and zeroing not significant terms. All these I_{shared} $(j \rightarrow i)$ entries were interpreted as weights in the 962 963 adjacency matrix of an information sharing directed functional network, and we defined 964 as *sharing assembly* formed by a neuron *i* the star-subgraph of the information sharing 965 network composed of *i* and all its immediate neighbors. We compiled all the overall N^2 different values of I_{shared} $(j \rightarrow i)$ into time-dependent feature vectors Sharing A(t), 966 967 describing thus all the possible sharing assemblies at a given time. We then also 968 computed *information sharing strengths* by integrating the total amounts of information 969 that each single unit was sharing with the past activity of other units in the network 970 ("sharing-in"):

972
$$I_{shared} (\rightarrow i) = \Sigma_j I_{shared} (j \rightarrow i)$$

973

974 or with the future activity of other units in the network ("sharing-out"):

975

976
$$I_{shared} (i \rightarrow) = \Sigma_j I_{shared} (i \rightarrow j)$$

977

978 In other words, the integrated amount of shared information was given by the in-979 strength and the out-strength of a node in the information sharing network with 980 individual link weights I_{shared} $(j \rightarrow i)$. We compiled the N incoming I_{shared} $(\rightarrow i)$ and N 981 outgoing I_{shared} ($i \rightarrow$) values into time-dependent vectors Sharing S(t). We computed 982 separate Sharing A(t) and Sharing S(t) for each of the simultaneously recorded 983 regions. We then performed as before unsupervised clustering based on the associated 984 M_{sim} matrices to extract sharing substates. Since the block structure displayed by thee 985 M_{sim} matrices for sharing assemblies and strengths are nearly perfectly overlapping we 986 conducted all substate analyses based on Sharing S(t) vectors only. We defined a 987 neuron to be a *sharing hub* in a given state if its I_{shared} (* \rightarrow i) and/or I_{shared} (i \rightarrow *) values 988 in the state prototype vector were higher than the 95% percentile of all concatenated 989 cluster prototypes entries (again protecting against false positive detection).

The relative comparisons of information sharing between SO and THE (REM and nonREM) epochs for different recordings shown in Figure S9, are based, as in the case of AIS in Figure S7, on averaged and scaled values. We first averaged the total I_{shared} (i.e. sharing in plus sharing out) over all the units within a specific anatomic layer. We then normalized these average total I_{shared} values by dividing them by the average total I_{shared} value in the SO state (in anesthesia) or the nonREM state (in natural sleep) for the specifically considered recording and layer.

997 Liquidity of sharing. The M_{rec} matrices for Sharing Assemblies display light blue 998 (low internal correlation) blocks while the M_{rec} matrices for Sharing Strengths have 999 similar blocks but red-hued (higher internal correlation). We quantify this visual 1000 impression by evaluating liquidity of sharing strength and sharing assembly substates. 1001 For a given recording and a given associated M_{rec} matrix (e.g. the one for the 1002 Sharing A or the Sharing S features), we define T_{α} as the set of times t for which 1003 the system is in a given substate α relative to the considered feature of interest. We then 1004 evaluate the liquidity $\Lambda(\alpha)$ of this substate α as:

1005
$$\Lambda(\alpha) = \sum_{t}^{t \in T_{\alpha}} \sum_{t' < t}^{t' \in T_{\alpha}} (1 - |M_{rec}(t, t')|) / {\binom{\#T_{\alpha}}{2}}$$

1006 where $|\cdot|$ denotes the absolute value operator and $\#T_{\alpha}$ is the number of elements of the 1007 set T_{α} . Liquidity values are thus bounded in the interval $0 \le \Lambda(\alpha) \le 1$, with 1 indicating 1008 maximum liquidity (i.e. maximum internal variability) of a substate.

1009

1010 Oscillatory mode specificity and hub distributions. For each substate (firing, 1011 storage, sharing) we computed the fraction of times that the substate was observed 1012 during a SO or a THE state (in anesthesia) or a nonREM or REM state (in natural sleep). 1013 We defined the largest among these fractions as the oscillatory specificity of this 1014 substate. Oscillatory specificities close to 1 indicate that a substate occurs mostly within 1015 one of the two possible global states observed in each recording, while specificities 1016 close to 0.5 indicate that the substate do not occur preferentially in one of the global 1017 states.

To evaluate the probability that a hub emerges in a given anatomical layer, we computed for every recording the fraction of cells recorded in each layer that were labeled as hubs at least in one computing substate (storage or sharing). We computed separately these fractions layer-by-layer, for excitatory and inhibitory cells and for

1022 anesthesia or sleep. These fractions were equal to unit when all the excitatory (or 1023 inhibitory) cells in a layer happened to be hubs at least once. We then evaluated the 1024 general probability that a hub emerges in a layer, which is different from the previous 1025 one, because it takes in account as well the fact that some cells may be labeled as hubs 1026 more often than others. We then considered the list of all hubs of a given type (storage 1027 or sharing) across all substates, including repetitions (if a neuron was hub in more than 1028 one substate then it appeared multiple times in the list) and evaluated the fraction of 1029 times in which a hub in this list was belonging to a given layer. We computed separately 1030 these fractions layer-by-layer, for storage or sharing hubs and for anesthesia or sleep. 1031 95% confidence intervals for the mean fractions above were evaluated as 2.996 times 1032 their sample standard deviation over the different recordings for which they could be 1033 computed. We considered two mean fractions to be different when their 95% 1034 confidence intervals were fully disjoint.

1035

1036 Coordination between substate transitions. To compare sequences of substates 1037 of different types or in different regions we introduced a symbolic description of substate switching. Each substate was assigned a *letter* symbol, i.e. a label $s^{(p)}$ where p 1038 1039 can stand for firing, information storage or sharing and $s^{(p)}$ is an arbitrary integer label 1040 different for every substate. We could thus describe the temporal sequences of the 1041 visited substates of each different type as an ordered list of integers $s^{(p)}(t)$. We quantified 1042 the degree of coordination between the sequences of substates of different types (e.g. p 1043 = 'storage' vs q = 'sharing') or in different regions (e.g. p = 'storage in EC' vs q = 1044 'storage in CA1') by evaluating the relative Mutual Information term:

1045

1046
$$MI[s^{(p)}(t), s^{(q)}(t)] / max[H(s^{(p)}(t)), H(s^{(p)}(t))]$$

1048 normalized between 0 –full statistical independence between the two substate 1049 sequences– and 1 –complete overlap between the two substate sequences–, by dividing 1050 it by the entropy H of the most entropic among the two symbolic streams. We evaluated 1051 these MI and H terms using direct plug-in estimators on the joint histograms of substate 1052 labels. We estimated chance expectations for the level of coordination by repeating the 1053 same procedure for substate sequences with shuffled substate labels and then finding 1054 the 99th percentile over 1000 permutation replicas of the computed MI/H.

1055

1056 Mutual Information Measure's Dependence on Bin Size. The original decision 1057 for the bin size was chosen such that when discretized, the information content lost by 1058 counting 2 or 3 spikes on the same neuron within a given bin as a '1' was less than 5%. 1059 On average, the information content lost was less than 1% across all recordings. To 1060 analyze the dependence on bin sizes, one example recording was chosen in the PFC 1061 during natural sleep in different bin sizes, 25 ms, 33 ms and 66 ms and computed 1062 substates using the same methods described above. To make the comparison focused 1063 on bin size, the same number of clusters per feature was chosen to reflect the original 1064 number. We then computed the amount of information about the substate sequences 1065 computed with the original binsize were retained by corresponding substate sequences 1066 derived for each different bin size. To do so, we used the same procedure described in 1067 the previous section to quantify coordination between sequences for different types of 1068 states or between different regions. Notably we computed mutual information between 1069 the substate sequences for different bin-sizes (normalized by the entropy of the original 1070 sequence) and compared this relative mutual information with chance expectation 1071 (obtained via shuffling substate sequences, as above). We found that the mutual 1072 information between corresponding sequences for different bin sizes was two order of 1073 magnitudes above chance level (Figure S13B), denoting high robustness of our

procedure for extracting substates. Correspondingly, we also found that, for matched
substates between sequences extracted for different bin-sizes, the identification, number
and anatomical localization of hubs were only marginally altered.

1077 **Complexity of substate sequences.** After converting sequences of substates into 1078 symbolic streams of letters, we defined substate *words* as the triplets of letters 1079 corresponding to the firing, the information storage and the information sharing 1080 substates simultaneously observed at each time t, i.e.:

1081

1082
$$S(t) = s^{(firing)}(t) \ s^{(storage)}(t) \ \dots \ s^{(sharing)}(t)$$

1083

1084 We then constructed a *switching table* T in which the temporally ordered columns 1085 provide the sequence of substate words S(t) along time. We compiled separate 1086 switching tables for each recording and for each of the simultaneously recorded regions. 1087 The total set of substate words effectively found in a specific switching table constitutes 1088 its associated *dictionary* of substate combinations. We defined then the *used dictionary* 1089 fraction, as the ratio between the number of observed words and the maximum 1090 theoretically possible number of words that could have been composed given the 1091 available substate letters (depending on how many firing, storage or sharing substates 1092 have been effectively extracted).

We then evaluated the complexity of substate word sequences using a procedure inspired from the notion of Kolmogorov-Chaitin complexity (*19*) and minimum description length approaches (MDL; *20*). The basic concept is that, for a regular symbolic sequence (as our streams of substate words), it will be possible to design a tailored "compression language" such the sequence will admit a much shorter description when reformulated into this language with respect to the original length in terms of number of words. On the contrary, a random symbolic sequence will be poorly

1100 compressible, i.e. its descriptions in terms of a generative language will be nearly as 1101 long as the original list of symbols appearing in the sequence. A complex symbolic 1102 sequence will stand between these two extremes - still admitting a compressed 1103 generative description but not as short as for regular sequences. Departing from 1104 universal compression approaches, as the original MDL formulation (20) or the 1105 Lempel-Ziv approach (21), we introduce here a "toy language" for generative 1106 description, specialized to compress state transition tables as the ones of Figure 6A. Our 1107 choice is conceptually compliant with the MDL approach but - for the sake of 1108 pedagogy- avoids technical steps as the use of binary prefix coding.

1109 Let $\Omega = \{S_1, S_2, ..., S_{\omega}\}$ be the dictionary of substate words appearing in the switching 1110 table T which we want to describe. We first define the *exhaustive list description* (D_{list}) 1111 of T as a string of the following form:

1112

1113
$$D_{\text{list}} \coloneqq \mathsf{S}_1 t_{1,1} t_{1,2} \dots t_{1,k1} \mathsf{S}_2 t_{2,1} t_{2,2} \dots t_{2,k2} \dots \mathsf{S}_{\omega} t_{\omega,1} t_{\omega,2} \dots t_{\omega,k\omega}$$

1114

In such a description the symbol of each substate word S_q (counting as one description unit) is followed by an exhaustive list of all the k_q times $t_{q,1}, t_{q,2}, ..., t_{q,kq}$ (each time index counting as an extra description unit) at which the recorded system produced the matching substate word. If the number of analyzed time windows is $K = k_1 + k_2 + ... +$ k_{ω} , then the length of the exhaustive list description will be $|D_{\text{list}}| = K + \omega$ description units (*K* time stamps, plus ω substate word symbols).

1121 Let then define the *block-length description* (D_{block}) of the stream of substate 1122 codewords, as a description of the following form:

1124
$$D_{\text{block}} \coloneqq S_1 w_{1,1} l_{1,1} w_{1,2} l_{1,2} \dots w_{1,m1} l_{1,m1} S_2 w_{2,1} l_{2,1} w_{2,2} l_{2,2} \dots w_{2,m1} l_{2,m1} \dots$$

1125
$$S_{\omega} w_{\omega,1} l_{\omega,1} w_{\omega,2} l_{\omega,2} \dots w_{\omega,m\omega} l_{\omega,m\omega}$$

1126

1127 In such a description the symbol of each word S_q (always counting as one description 1128 unit) is followed by a list of stepping instructions for a hypothetical "writing head" 1129 moving along different discrete positions on an idealized tape, similarly to computing 1130 automata as the Turing Machine (69). At the beginning the machine is initialized with 1131 the head on the first position on the tape. The integers $w_{q,n}$ –at odd positions (1st, 3rd, 1132 etc.) after the word symbol- indicate for how many steps the machine head must shift 1133 on the tape toward the right *without* writing, but just skipping positions. The integers $l_{q,n}$ –at even positions (2nd, 4th, etc.) after the word symbol– indicate instead for how 1134 1135 many steps the machine must also write on the tape the symbolic string S_q before then 1136 shifting to the next position on the right. Every time that a new symbol S_q is met when 1137 parsing the step lengths description, the position of the writing head is reset to the 1138 leftmost starting position on the tape. Such parsing grammar is obviously more complex 1139 than the one for a simpler "parrot machine", designed to parse exhaustive list 1140 descriptions as the ones described above. The length in symbols of this block-length 1141 description is variable and depends on how regular the word sequence is to compress and regenerate. The block-length description segment $S_q w_{q,1} l_{q,1} \dots w_{q,mq} l_{q,mq}$ will be 1142 shorter than the matching exhaustive list description segment $S_q t_{q,1} \dots t_{q,kq}$ whenever 1143 $2m_p < k_p$, which can happen if transitions for the different types of substate letters are 1144 1145 regularly aligned, in such a way that the resulting switching table have long alternating 1146 blocks with repeated substate words.

1147 The syntactic complexity of a sequence of substate words can then be evaluated by 1148 quantifying how much the program to generate the switching table T via a "smart" 1149 compressing machine interpreting block-length descriptions is shorter than the program

1150	to generate the same table T via a "dumb" parrot machine interpreting exhaustive length
1151	descriptions. We define the <i>description length complexity</i> of a switching table T as:
1152	
1153	$DLC = D_{block}(T) / D_{list}(T) $
1154	
1155	To give a toy example, let's consider the sequence T = "AAAAAAA BBBB AAAAA
1156	CCCCC DDD BBBBBB", built out of four possible collection of substate words $S_1 =$
1157	A, $S_2 = B$, $S_3 = C$ and $S_4 = D$. The exhaustive list description for this sequence will be:
1158	
1159	$D_{\text{list}}(T) = A \ 1 \ 2 \ 3 \ 4 \ 5 \ 6 \ 7 \ 12 \ 13 \ 14 \ 15 \ 16 \ B \ 8 \ 9 \ 10 \ 11 \ 25 \ 26 \ 27 \ 28 \ 29 \ 30$
1160	C 17 18 19 20 21 D 22 23 24
1161	
1162	with length $ D_{\text{list}}(T) = 34$ descriptive units. Its step lengths description will be:
1163	
1164	D_{block} (T) = A 0 7 4 5 B 7 4 13 6 C 16 5 D 21 3
1165	
1166	with length $ D_{block}(T) = 16$ descriptive units, i.e. $ D_{block}(T) < D_{list}(T) $.
1167	Given the noisiness of data, we dropped from both the exhaustive list description
1168	and the step lengths description the segments corresponding to exceedingly rare words
1169	$S_{q.}$ In particular, ranking the code words from the least to the rarest, we dropped all the
1170	words S_r with $r \ge R$, such that removing all of their occurrences in the word stream
1171	reduced the stream's overall length of no more than 10% (lossy compression).
1172	We computed confidence intervals for DLC values via a Jacknife construction in
1173	which we drop one word at random position from the temporal stream $S(t)$ made of K
1174	symbols, generating up to K Jackknife surrogate streams, each with K -1 symbols. The

1175 confidence interval was then given by the 5-th and the 95-th percentile over the 1176 complexities evaluated from these Jackknife surrogates.

1177 Appropriate reference criteria were then required to discriminate complex vs ordered 1178 or random switching tables. We need to compare the empirically observed DLC values 1179 against two thresholds. Complex switching tables should have indeed a DLC below a 1180 threshold for randomness testing and above a threshold for regularity testing. Given a 1181 switching table T, we constructed a randomized version T_{rand} by randomly permuting 1182 independently the entries of each of its rows. For each recording, we constructed 1000 1183 instances of T_{rand} and evaluated DLC for all of them, identifying as upper threshold for 1184 complexity the 5-th percentile $DLC_{rand} = q_{5\%}[DLC(T_{rand})]$. We then constructed an 1185 enhanced-regularity version T_{regular} of each T by lexicographically sorting entries row-1186 by-row (to get blocks of homogeneous code-words as long-lasting as possible based on 1187 exactly the same building bricks). We then arbitrarily defined a lower threshold for 1188 complexity $DLC_{regular} = 2 DLC(T_{regular})$. The thresholds $DLC_{regular}$ and DLC_{rand} varied for every switching table. However, the criterion DLC_{regular} < DLC < DLC_{rand} was fulfilled 1189 1190 for all the considered recordings, whose state transitions sequences could then be 1191 certified to be complex (in our arbitrary but quantitative and operational sense).

1192 Importantly, we could restrict the evaluation of complexity to sub-table restricted to 1193 words occurring during selected different global oscillatory states only. In this way we 1194 could compare the complexity of sequences occurring within different global states, 1195 e.g. REM vs nonREM. We plot in Figure 6E relative complexity variations between 1196 two global states α and β :

1197

1198
$$\Delta (DLC) = (DLC_{\alpha} - DLC_{\beta}) / (DLC_{\alpha} + DLC_{\beta})$$

We evaluated once again confidence intervals for relative complexity variations viaone-leave-out Jacknife on global state-restricted switching table columns.

1202

1203 Burstiness of state sequences. We also characterize switching tables in terms of 1204 their "style" of transitions, looking at two different temporal statistics. First, we 1205 computed all inter-transition times from a table T, i.e. the number of time-steps 1206 occurring between one block (continuous time-interval with a same substate word 1207 maintained in time) to the next. Note that these inter-transition times are precisely the 1208 $l_{p,n}$ integers appearing in the block-length description D_{block} (T) of the table T. After 1209 computing the mean μ_l and the standard deviation σ_l of these inter-transition times, we 1210 then evaluated the burstiness coefficient (22):

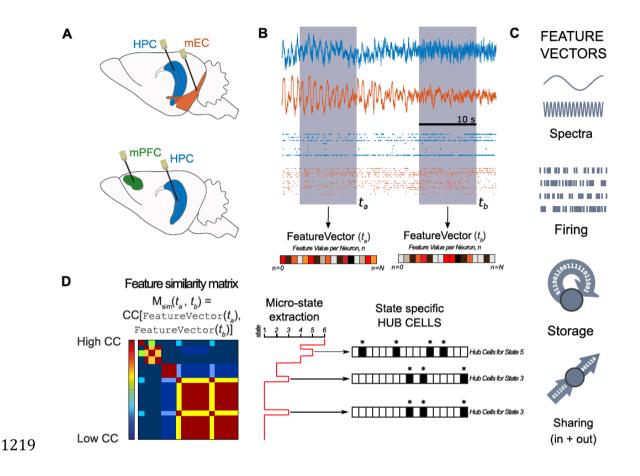
1211

1212
$$B = \frac{\frac{\sigma_l}{\mu_l} - 1}{\frac{\sigma_l}{\mu_l} + 1}$$

1213

1214 Such a coefficient is bound between -1 < B < 1 and is equal to 0 when transitions 1215 between substate words follow a Poisson statistic, negative when the train of transitions 1216 is more periodic and positive when more bursty than for a Poisson train.

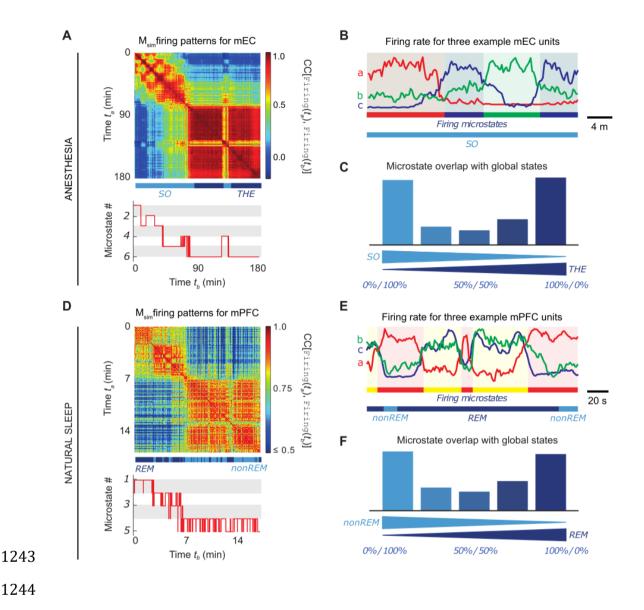
1218 Figures



1220

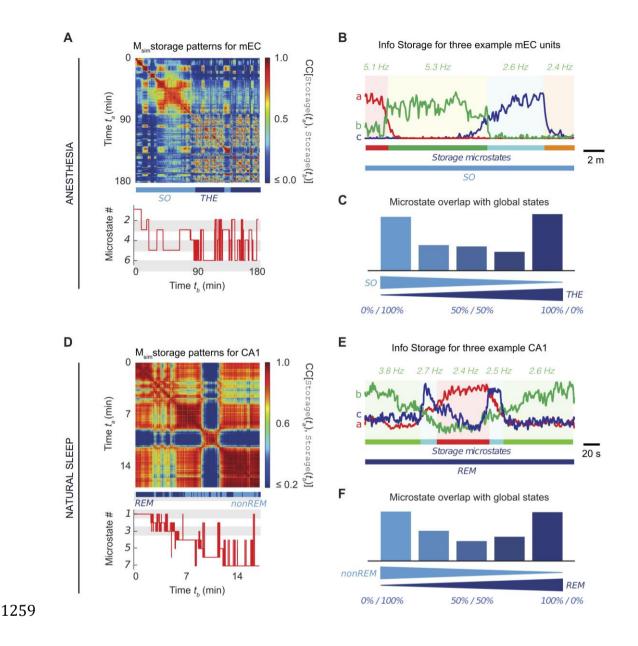
1221 Figure 1. Unsupervised extraction of states and hubs. A. Cartoon representing the 1222 approximate recording locations (mEC and CA1, mPFC and CA1) during 2 experiment 1223 types in anesthesia and sleep. **B**. Example LFP trace taken from the 32 channels in 1224 CA1 (blue) and 32 channels in mEC (orange). Below are examples of isolated unit 1225 activity taken from the same recording. For each time window (t), we extract different 1226 features represented by the FeatureVector(*t*), which has a feature value for each 1227 channel or single unit recorded. C. We consider four features: spectral band averaged 1228 powers (from LFP channels); single unit firing rates; information storage and 1229 information sharing. **D.** Left panel: A cartoon representation of M_{sim} . To extract substates and their temporal dynamics, we construct a *feature similarity* matrix M_{sim} in 1230 1231 which the entry $M_{sim}(t_a, t_b)$ measures Pearson correlation between the vectors

1232 FeatureVector(t_a) and FeatureVector(t_b). Time flows from the top-left corner 1233 horizontally to the top-right corner and vertically to the bottom-left corner. A block 1234 (square) along the diagonal in the resulting image identifies a period of feature stability. 1235 i.e. a substate. A block appearing several times horizontally or vertically indicates that 1236 a feature is repeated several times. Middle panel: Unsupervised clustering identifies the 1237 different substates (indicated by a number) and their temporal dynamics (the vertical 1238 axis corresponds to that of the similarity matrix). Right panel: We identify computing 1239 hub cells, i.e. neurons that display exceptionally high values for a given feature, 1240 associated with given substates. Note that reoccurring states have the same hub cells 1241 (state 3 in this example).



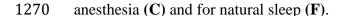
1245 Figure 2. Firing substates. Examples of similarity matrices M_{sim} obtained from 1246 Firing(t) at different times in mEC during anesthesia (A) and in mPFC during natural 1247 sleep (**D**), measured in two animals. The bar below M_{sim} indicates the transitions 1248 occurring between THE/REM (dark blue) and SO/nonREM (light blue). Although there 1249 were only two global brain states, six (A) and five (D) firing substates were identified. 1250 Panels (B) and (E) show examples of the firing density of three neurons (a, b and c) 1251 recorded in mEC and mPFC, respectively, with amplitude normalized for visualization. 1252 Neurons tended to fire in specific substates, indicated here with a color code. These 1253 examples also illustrate the switching between different firing substates inside a given

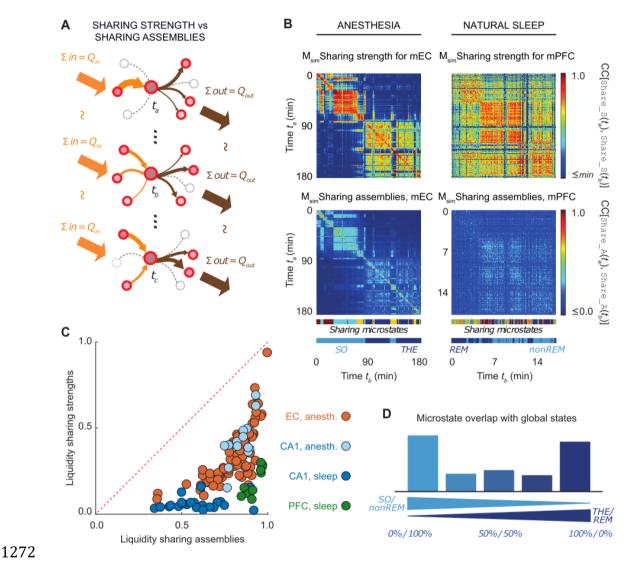
- 1254 global oscillatory state, and their overlap across different global oscillatory states. The
- 1255 analysis of all recordings revealed that a majority of firing substates tended to occur
- 1256 during a preferred global oscillatory state, as indicated by the bimodal histograms
- 1257 during anesthesia (C) and natural sleep (F), respectively.



1260 Figure 3. Information storage substates. Examples of similarity matrices M_{sim} 1261 obtained from Storage(t) at different times in mEC during anesthesia (A) and CA1 1262 during natural sleep (**D**). As for firing substates, we identified more storage substates (6 and 7, respectively, in the shown examples) than global oscillatory states. We show 1263 1264 in panels (B) and (E) that the participation of three individual neurons to information 1265 storage (indicated in arbitrary units for visualization) was substate-dependent. The 1266 values reported above the plots correspond to the average firing rate of the neuron b 1267 (green color) during the corresponding epochs within consistent storage substates. The 1268 analysis of all recordings showed that storage substates tended to occur during a

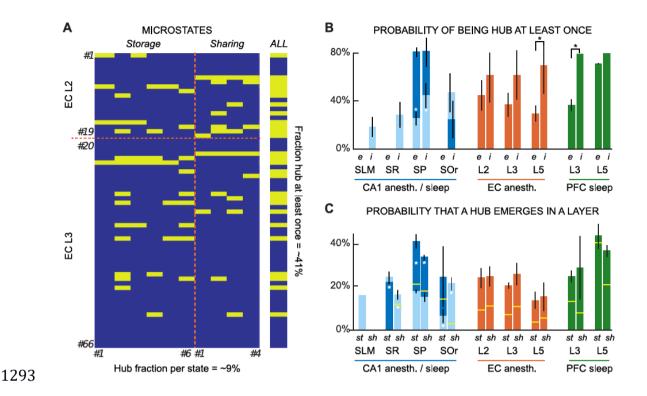
1269 preferred global oscillatory substate, as indicated by the bimodal histograms for





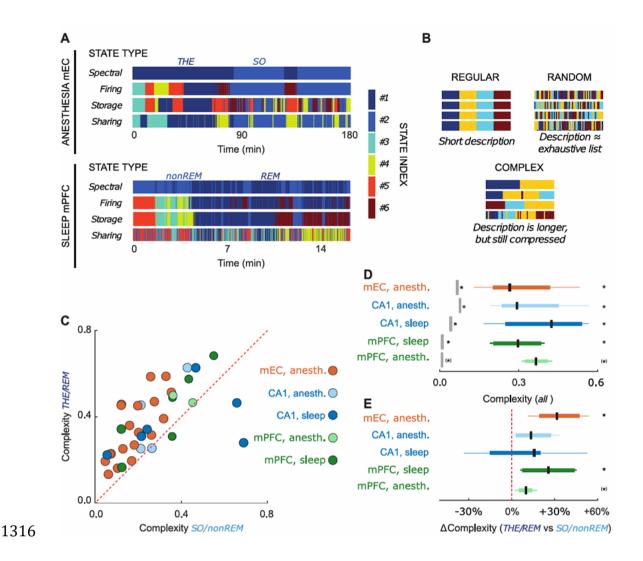
1273Figure 4. Information sharing substates. The cartoon in panel A shows an example1274of sharing assembly for a given sharing hub neuron across 3, non-sequential1275occurrences of the same substate. The total strength of in- and out-going sharing is equal1276(large, external arrows) during t_a , t_b , and t_c while the assembly changes (smaller, internal1277arrows). The changing size of internal arrows represent the sharing strength of that1278particular functional connection between the sharing hub and its source and target

1279 neurons. (B) Similarity matrices M_{sim} for sharing strengths Sharing S(t) (top) and 1280 sharing assemblies Sharing A(t) (bottom), in mEC during anesthesia (left) and 1281 mPFC during natural sleep (right). We identified a multiplicity of substates within each 1282 global oscillatory state as shown by the colored bars below the feature similarity 1283 matrices. The similarity matrices for sharing strengths and assemblies have a matching 1284 block structure. However, sharing strengths were very stable within a substate (red-1285 hued blocks), while sharing assemblies were highly volatile (light blue-hued blocks). 1286 This is quantified for each sharing assembly substate by a *liquidity* coefficient (\mathbf{C}). As 1287 shown in (C), for all observed sharing substates across all regions and global oscillatory 1288 states in all animals, the liquidity of sharing assemblies was much larger than the one 1289 of sharing strengths. Finally, (D) demonstrates that most sharing substates occurred preferentially during a preferred global oscillatory state for both anesthesia and natural 1290 1291 sleep combined (see Figure S7 for separated histograms for the two conditions).



1294 Figure 5. A democracy of computing hubs. (A) Within every computing substate 1295 some neurons exhibited significantly strong values of information storage or sharing 1296 (computing hubs). However, these computing hubs did generally change from one 1297 substate to the other, as shown in this example. Different rows correspond to different 1298 single units recorded in mEC during anesthesia and different columns correspond to 1299 different computing substates (left, storage substates 1 to 6; right, sharing substates 1 1300 to 4). An entry is colored in yellow when the neuron is a computing hub within the 1301 corresponding substate. In the shown example, while ~9% of neurons on average were 1302 simultaneously acting as computing hub, over 40% of the recorded units were recruited 1303 as hubs for at least one substate, when considering all the computing substates together 1304 (vertical bar on the right). (B-C) The probability that a neuron acted as hub depended 1305 only loosely on its anatomical localization. Panel **B** shows that for all regions and layers 1306 the probability that a neuron act as computing hub at least once was always larger than 1307 30%. Inhibitory (i) neurons tended to be recruited as hubs more frequently than 1308 excitatory (e) neurons. Analogously, panel C shows that none of the layers display a

specialization in either one of the two processing operations of information storage or sharing. Stars denote statistically significant comparisons (lack of overlap between 95% confidence intervals for the probability, reported as vertical ranges on top of the histogram bar). In panel **C** a yellow horizontal line indicates the fraction of computing hub cells which also happen to simultaneously be high-firing rate cells. Many computing hubs have thus average or low firing rate. In panel **B-C** in CA1 light blue represents anesthesia and dark blue represents natural sleep.

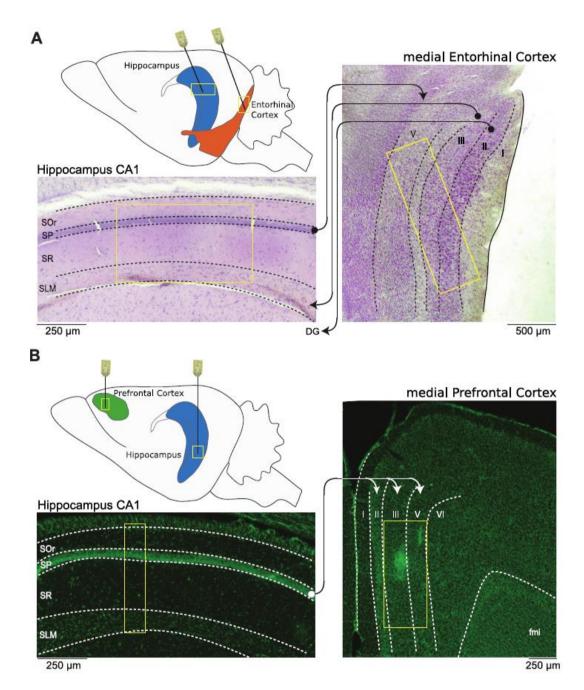


1317 Figure 6. Complexity of substate sequences. State switching found for each feature 1318 (firing, storage, sharing) did not align in time. This can be visualized by state switching 1319 tables, whose different rows graphically represent transitions between global brain 1320 oscillatory states and firing, storage, and sharing substates. Examples of switching 1321 tables are shown in (A) for mEC during anesthesia (top) and for mPFC during natural 1322 sleep (bottom, note the different time scales). Switching tables were neither perfectly 1323 regular (**B**, top left), nor random (**B**, top right), but they were "complex", displaying 1324 organized patterns escaping simple description (**B**, bottom). (**C**) The complexity of the switching tables was larger for THE/REM than for SO/nonREM for most recordings. 1325 1326 We included two recordings from mPFC under anesthesia for comparison. (D) 1327 Switching tables were complex in all cases. Complexity values were significantly above

1328	the upper threshold	for regularity	and below the	lower threshold	for randomness.	(E)

- 1329 The increase of complexity was significant for mEC when transitioning from SO to
- 1330 THE and for mPFC from nonREM to REM sleep. This trend in CA1 was not
- 1331 statistically significant (significance assessed in terms of lack of intersection between
- 1332 95%-confidence intervals and threshold values for both panels **D** and **E**; a (*) symbol
- 1333 indicates that the number of recordings in this category was not enough to assess
- 1334 significance but that the median value lied below or above the considered threshold).

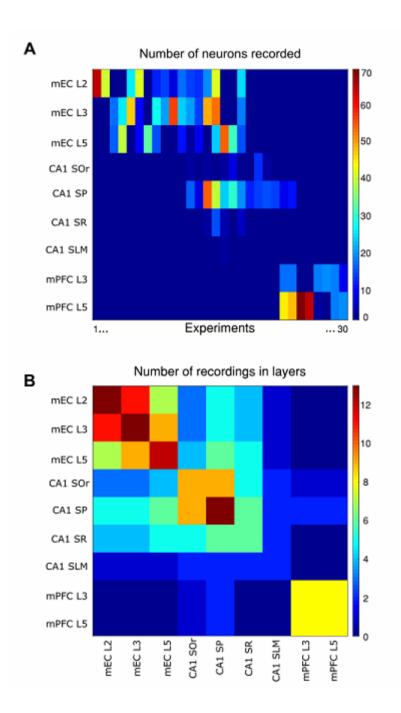
1335 Supplementary figures



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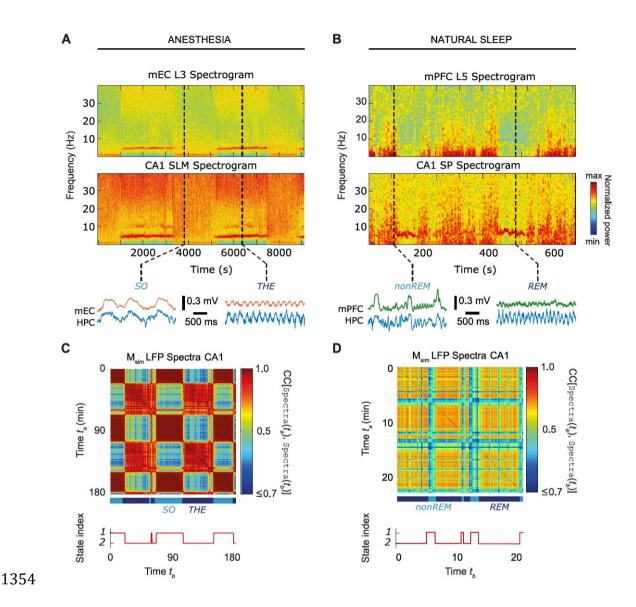
Figure S1. Recording paradigm. Schematic representation of the (**A**) simultaneous mEC/HPC recording setup and (**B**) simultaneous mPFC/HPC during anesthesia and natural sleep. The Nissl stained sections display the anatomical regions recorded by the different silicon probes used (yellow boxes). Arrows represent the anatomical connectivity (\bullet : source layer, \rightarrow : target layer) between the dorsal hippocampus CA1

- 1342 region (SOr: stratum oriens; SP: stratum pyramidale; SR: stratum radiatum; SLM:
- 1343 stratum lacunosum moleculare) with the dorso-medial entorhinal cortex (mEC, layers I
- to VI) and medial prefrontal cortex (mPFC, layers I to VI).



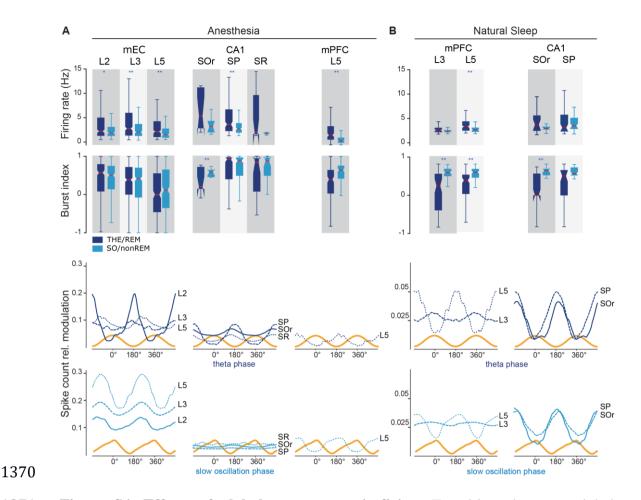
1346

Figure S2. Information about recordings. We analyzed data from 30 different recordings performed in 24 different rats. In each recording we identify single units in different anatomical locations. (**A**) Number of recorded single units (color coded on the right scalebar) per anatomical layer (rows), for each of the 30 recordings (columns). (**B**) Number of recordings (color coded on the right scalebar) simultaneously targeting pairs of two different anatomical layers.

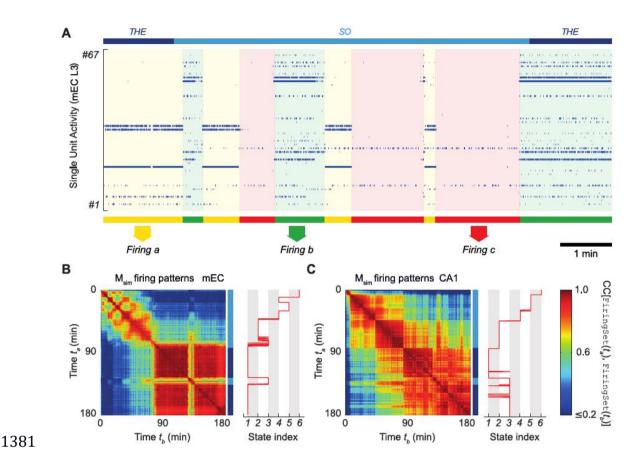


1355 Figure S3. Global brain oscillatory states. We performed a time-frequency spectral 1356 analysis of the LFP signals from all channels. Time-frequency spectrograms of LFPs 1357 are shown in (A) from mEC III and CA1 SLM layers during anesthesia and in (B) for 1358 mPFC layer V and CA1 SP. A characteristic alternation is visible between epochs 1359 dominated by SO/THE rhythms and REM/nonREM. Example LFP traces at time points 1360 corresponding to the dashed vertical lines are magnified and shown below the 1361 spectrograms. To characterize global oscillatory states in an unsupervised manner 1362 within each time-window of analysis (Figure 1), we averaged the power across different 1363 frequency bands and compiled all LFP channels into the feature vector Spectra(t). 1364 The similarity matrices corresponding to (A) and (B) are shown in (C) and (D),

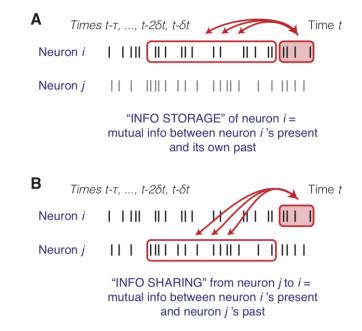
- 1365 respectively. The alternation between SO and THE epochs in (C) and between nonREM
- 1366 and REM epochs in (**D**) is well visible in the marked block structure of the feature
- 1367 similarity matrices. Unsupervised clustering identified 2 states under anesthesia or
- 1368 natural sleep.



1371 Figure S4. Effects of global states on unit firing. Transitions between global 1372 oscillatory states in both anesthesia and sleep significantly modulated the median firing 1373 rate in mEC, mPFC and CA1 layers (A, top). Burstiness (B, bottom) was significantly 1374 modulated during natural sleep only. Both firing rate and bursting indices were 1375 heterogeneous across neurons, as emphasized by long box-plot whiskers. Single units 1376 in mEC, mPFC and CA1 layers fired preferentially at well-defined phases of the 1377 ongoing theta (**B**, top) and slow oscillation (**B**, bottom) rhythms as visualized by phase-1378 binned histograms of spike count relative modulation, compared with reference average 1379 LFP waveform cycles. Note that phase modulations were an order of magnitude 1380 stronger during anesthesia than natural sleep.

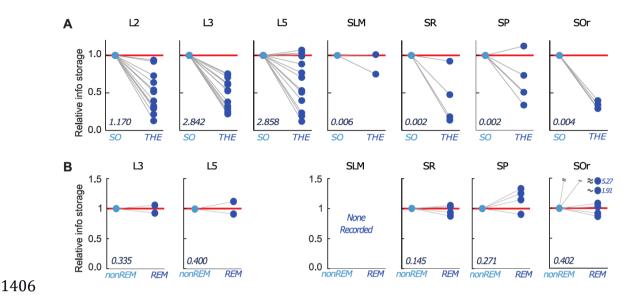


1382 Figure S5. Firing substates. (A) displays the firing, represented by blue dots, of 67 1383 neurons recorded in layer 3 in mEC. The solid lines above indicate the global brain 1384 states (THE and SO) identified using unsupervised clustering of the spectral features of 1385 the field potential (as described in Figure S3). Unsupervised clustering identified three 1386 sets of co-firing neurons (indicated by yellow, green and red solid lines), which are 1387 clearly visible, during the recording time shown here. Note the alternation of the firing 1388 substates, and the fact that global oscillatory state transitions (THE \rightarrow SO \rightarrow THE) do not 1389 correspond to transitions between firing substates. Below, (B) and (C) represent the 1390 dynamics of firing sets in mEC and CA1 during the whole recording session, where 1391 squares across the diagonal represent different firing substates, as in (A).

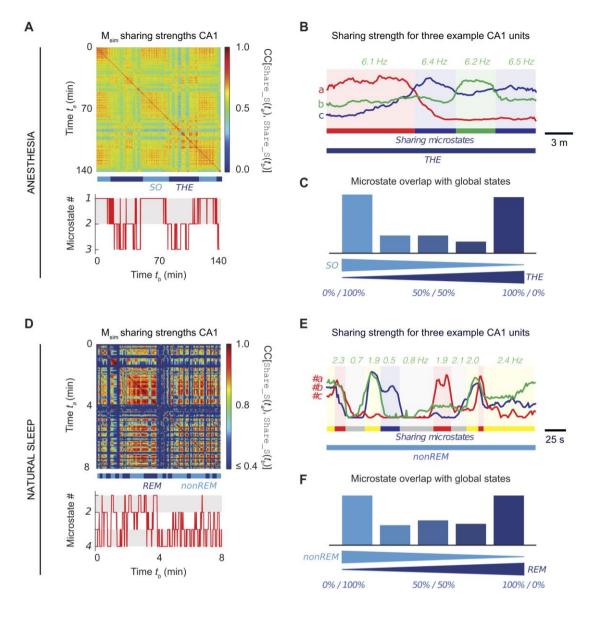


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1393 Figure S6. Primitive operations in information processing: explanatory cartoons. 1394 The information conveyed by the activity of a unit at a given time may have different 1395 sources. A fraction of the total information conveyed at time t by a neuron i may have 1396 already been present in i's past activity (A). Therefore, we say that this fraction of 1397 information is being *actively stored* into neuron *i* by its activity, implementing a 1398 "memory buffer". The involvement of a unit into this primitive information processing 1399 operation is quantified by its Active Information Storage score (see Figure 3). A 1400 complementary fraction of the total information conveyed at time t by a neuron i may 1401 have been present already into the past activity of a different neuron i (**B**). We say in 1402 this case that this fraction of information is *shared* from *j* toward *i*, with time-lagged 1403 mutual information providing a pseudo-directed measure of functional connectivity 1404 (see Figure 4).



1407 Figure S7. Information storage is brain state-dependent. (A) Relative variation of 1408 active information storage values in the different mEC and CA1 lavers between SO and 1409 THE states during anesthesia. Different lines correspond to different rats. All values are 1410 normalized to SO values for better visualization of the size and direction of effects. The 1411 absolute values of active information storage during SO are indicated in the lower left 1412 corner of each subpanel. During anesthesia, mEC layers have larger active information 1413 storage values than CA1 layers. Generally, switching from SO to THE state tends to 1414 reduce active information storage values. (B) Same as A but for mPFC and CA1 layers 1415 during natural sleep. The active information storage has the same order of magnitude 1416 for mPFC and CA1 but is lower than mEC during anesthesia. Furthermore, during 1417 natural sleep there is no major general difference between REM and nonREM (as 1418 opposed to SO/THE in A).

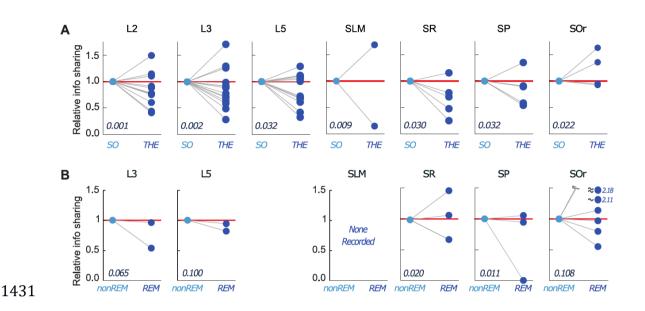




1420 Figure S8. Substates of information sharing: additional information. We show here 1421 additional typical feature similarity matrices M_{sim} obtained from sharing strengths 1422 Sharing S(t) in CA1 during anesthesia (A) and natural sleep (D) (see also Figure 4). 1423 As in firing and storage substates, (C) and (E) show that the participation of different 1424 neurons to the information sharing was varying along time in a switching fashion 1425 (arbitrary normalized units, smoothed time-series). The values reported above the plots 1426 correspond to the average firing rate of the neuron b (green color) during the 1427 corresponding epochs within consistent sharing substates. Sharing substates tended to

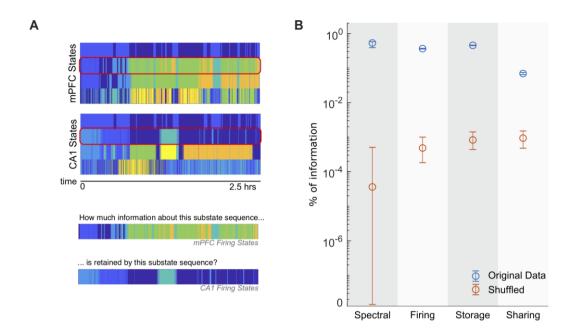
- 1428 occur during a preferred global oscillatory substate, as indicated by the bimodal
- 1429 histograms in (C) for anesthesia and (F) for natural sleep (see also Figure 4D).

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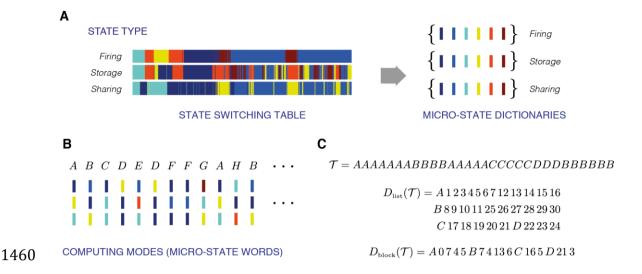
1432 Figure S9. Variations of information sharing as a function of the global brain 1433 oscillatory states. (A) Relative percent variation of information sharing total strength 1434 values between SO/THE states during anesthesia, averaged over different layers in 1435 mEC and CA1. Different lines correspond to different rats and average values of sharing 1436 strength in SO state are normalized to allow a simpler comparison of the size and 1437 direction of effects for different layers, but absolute values of sharing strength in the 1438 SO state, averaged over the different rats, are indicated in the lower left corner of each 1439 subpanel. (B) Same as above but for different mPFC and CA1 layers during natural 1440 sleep. We did not observe any systematic direction of change for sharing strengths when 1441 switching from SO to THE states during anesthesia or from nonREM to REM during 1442 natural sleep. mEC layers II and III during anesthesia were associated to the weaker 1443 absolute values of sharing strengths and mPFC layer IV and CA1 SO during natural 1444 sleep to the stronger values.

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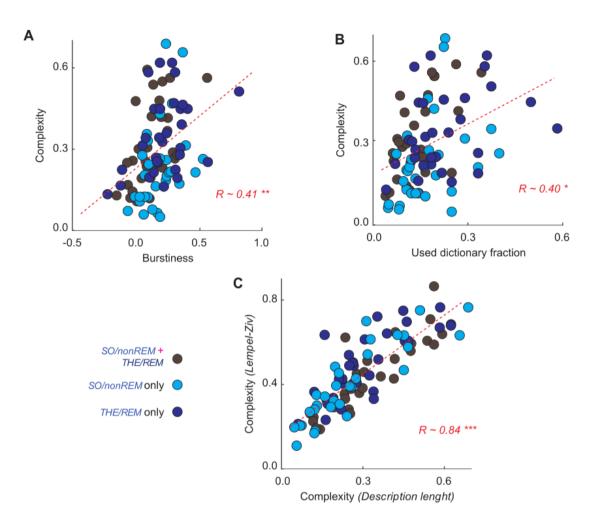




1447 Figure S10. Coordination of substate transitions between brain regions. Sequences 1448 of firing, storage and sharing substates do not show a perfect match between 1449 simultaneously recorder regions, as visible from the state transition tables of a 1450 representative simultaneous CA1/mPFC recording during natural sleep (A, top). To 1451 quantify the level of interregional coordination between substate transitions we 1452 evaluated the mutual information between matching substate sequences in the two 1453 simultaneously probed regions (A, bottom). We also estimated the corresponding 1454 chance levels of coordination, by repeating the same procedure on shuffled state 1455 transitions sequences. For all features, we find that the sequences are loosely coupled 1456 between regions, but still far above chance level (B). Vertical bars denote the 99% 1457 confidence interval (bootstrap with replacement for original data, permutation-based 1458 for shuffled data, 1000 replicas in both cases). This specific graph is built using the 1459 example in (A).



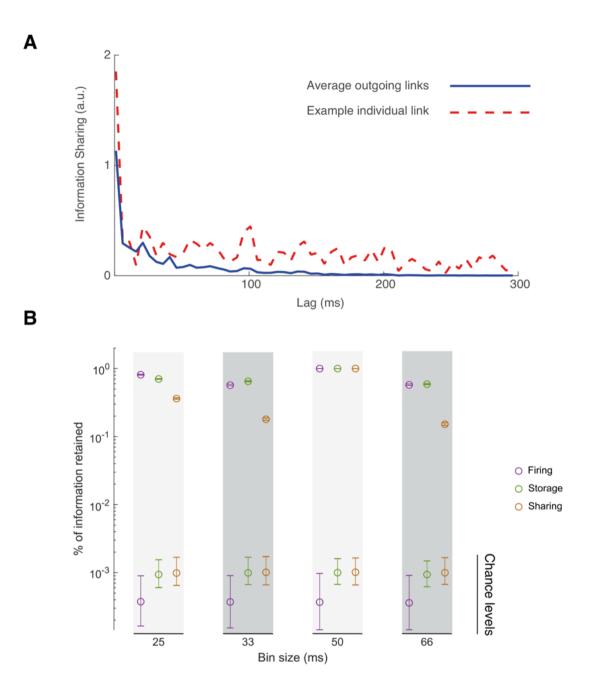
1461 Figure S11. Calculation of complexity of the state switching table. (A) A given 1462 recording can be represented by the superposition of 3 sequences of substates for each 1463 feature (firing, storage, sharing). (B) At any point in time, the triplet of substates can be 1464 represented as a word made of three substate letters. For simplicity, the letters are color-1465 coded, and a word is represented by a letter from the alphabet (A, B, C etc.). (C) A state switching table \mathcal{T} is the temporal sequence of *words* with one word per analysis window 1466 1467 t_A as defined in Figure 1). There are two ways to represent the sequence of words. $D_{\text{list}}(\mathcal{T})$ lists the positions at which the *words* appear (e.g. at window 1, 2, 3, 4, 5, 6, 7, 1468 12, 13, 14, 15, 16...). This representation is exhaustive, but not compact. $D_{block}(\mathcal{T})$ lists 1469 1470 how many positions one should skip from the start of the sequence before writing and 1471 in how many consecutive positions the considered word should be printed. In the 1472 example shown here, the *word* "A" occurs at the beginning of the string – zero positions skipped – and is then printed seven times. After, 4 positions are skipped, and it is written 1473 1474 again 5 times, etc. This representation is more compact. Complexity is given by the ratio between the lengths of the descriptions $D_{\text{block}}(\mathcal{T})$ and $D_{\text{list}}(\mathcal{T})$. 1475



1477

1478 Figure S12. Burstiness and Used Dictionary Fraction explain complexity. 1479 Complexity linearly increased with burstiness (A) and Used Dictionary Fraction (B). 1480 (C) Lempel-Ziv (LZ) complexity as a function of our measure of complexity. The two 1481 complexities were highly linearly correlated, and results of complexity analyses were 1482 thus qualitatively the same using either one of the two measures. We plot together 1483 results for complexity analyses restricted to SO/nonREM states (light blue dots), to 1484 THE/REM states (dark blue dots) or over all states combined (grey dots), as no 1485 significant differences were observed between the three groups with respect to the 1486 plotted linear trends.

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1490 Figure S13. Additional robustness analyses. (A) Lagged mutual information terms 1491 between spike trains, the building block terms of both storage and sharing features, 1492 quickly decay as a function of the considered lag τ , justifying our choice to integrate 1493 MI only for latencies up to an average theta cycle period ($\sim 125 - 250$ ms depending on 1494 recordings). The red dashed line refers to a representative individual link, associated to 1495 a specific pair of units, and show some additional peak. However, these secondary 1496 peaks are much smaller than the main peak for very short latency and are not aligned 1497 for different pairs of units so that they are averaged out away when averaging over

1498	multiple outgoing links originating from a same unit (solid blue line). (B) Our procedure
1499	for substate extraction is robust against changes of the bin size. We considered four
1500	different choices of bin size different from the original choice of 50 ms, extracted
1501	substate sequences for each of these new bin choices and computed mutual information
1502	between the newly obtained and the original corresponding substate sequences. Shown
1503	here are the relative fraction of retained information for different bin sizes and substate
1504	types (evaluated on a representative recording, mPFC, natural sleep, the same as for
1505	figure S10). Across all features, the fraction of information retained about the substate
1506	sequences for the main reference bin size of 50 ms is two orders of magnitude above
1507	chance levels, with sharing being most effected. Vertical bars denote 99% confidence
1508	interval (bootstrap with replacement for original data, permutation-based for shuffled
1509	data, 1000 replicas in both cases).
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- 1512 Supplementary tables
- 1513

1514 Table S1 – Percent of information about substate sequences in empirical

1515 recordings retained after shuffling

Substate	Anesthesia	Natural Sleep
Spectral	2.55×10 ⁻⁹ , 1.58×10 ⁻⁴	1.47×10 ⁻⁸ , 5.43×10 ⁻⁴
Firing	6.13×10 ⁻⁵ , 5.71×10 ⁻⁴	9.78×10 ⁻⁵ , 5.71×10 ⁻⁴
Sharing	4.68×10 ⁻⁴ , 1.13×10 ⁻³	2.66×10 ⁻⁴ , 1.42×10 ⁻³
Storage	9.49×10 ⁻⁴ , 1.15×10 ⁻³	4.15×10 ⁻⁴ , 1.61×10 ⁻³

1516 0.1% and 99.9% percentile of MI/H for state sequences. The very low values of these

1517 amounts of relative information indicate that the observed state sequences strongly

1518 deviate from a null hypothesis of lack of temporal structure