## 1 Title: Weak Coupling Between Spontaneous Local Cortical Activity State Switches Under

### 2 Anesthesia Leads to Strongly Correlated Global Cortical States.

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### 14 Abstract:

15 Under anesthesia, neural dynamics deviate dramatically from those seen during wakefulness. 16 During recovery from this perturbation, thalamocortical activity abruptly switches among a small 17 set of metastable intermediate states. These metastable states and structured transitions among 18 them form a scaffold that guides the brain back to the waking state. Here, we investigate the 19 mechanisms that constrain cortical activity to discrete states and give rise to abrupt transitions 20 among them. If state transitions were imposed onto the thalamocortical system by changes in the 21 subcortical modulation, different cortical sites should exhibit near-synchronous state transitions. 22 To test this hypothesis, we quantified state synchrony at different cortical sites in anesthetized 23 rats. States were defined by compressing spectra of layer-specific local field potentials (LFPs) in 24 visual and motor cortices. Transition synchrony, mutual information, and canonical correlations 25 all demonstrate that most state transitions in the cortex are local and that coupling between sites is weak. Fluctuations in the LFP in the thalamic input layer 4 were particularly dissimilar from 26 27 those in supra- and infra-granular layers. Thus, our results suggest that the discrete global cortical 28 states are not imposed by the ascending modulatory pathways but emerge from the multitude of 29 weak pairwise interactions within the cortex.

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#### 31 Introduction:

32 Brain activity arises as a result of interactions amongst billions of neurons and synapses. Each 33 component in this vast network exhibits complex nonlinear dynamics (Hodgkin and Huxley, 1952; 34 Pan and Zucker, 2009). Generically, such complex nonlinear dynamical systems can dramatically 35 change their collective behavior after small changes in parameters or perturbations to their 36 ongoing activity (Canavier et al., 1993; Destexhe et al., 1994; Ermentrout, 1998; Izhikevich, 2007; 37 Strogatz, 2015). Furthermore, because nonlinear systems generally have multiple steady state behaviors, there is no guarantee that after a dramatic perturbation, the system will recover to its 38 39 previous state once the perturbation subsides.

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41 These considerations suggest that brain activity ought to be quite fragile and unable to withstand 42 dramatic perturbations. Contrary to this intuition, there is ample evidence that the brain is 43 remarkably robust to perturbations. Seizures, for instance, are a paradigmatic example of 44 aberrant brain activity, being characterized by extreme synchronization in neuronal firing and 45 subthreshold voltage fluctuations (Timofeev et al., 2004). While seizures can be followed by a 46 transient postictal period characterized by abnormal brain activity and function (Fisher and Engel, 47 2010), normal brain function is eventually restored. Another classic example of the brain's ability 48 to recover from an extreme perturbation is general anesthesia (Brown et al., 2010). Every year, 49 millions of patients undergo general anesthesia. While some patients experience aberrant brain 50 activity, which manifests as delirium upon emergence (Saczynski et al., 2012), most eventually 51 recover normal brain activity and cognitive function. During general anesthesia, the brain may 52 exhibit dramatically abnormal activity patterns, such as burst suppression, which is caused by the 53 hyperpolarization and silencing of more than 90% of cortical neurons (Amzica, 2009; Civillico and 54 2012; Contreras and Steriade, 1997). Occasionally, complete isoelectric Contreras. 55 electroencephalogram (EEG) is observed in surgeries requiring circulatory arrest (Stecker et al., 56 2001). Nevertheless, once anesthetic delivery is stopped, the brain regains normal function. Given 57 this and the fact that anesthetic delivery can be precisely controlled, general anesthesia is a good 58 model system to address the general question of how the brain is able to restore normal activity 59 patterns after a dramatic perturbation.

60

Several converging lines of evidence strongly argue that recovery from anesthesia cannot be explained by anesthetic washout alone. The first is that recovery of consciousness after anesthesia occurs at a lower anesthetic concentration than induction of anesthesia across taxa, from *Drosophila* (Joiner et al., 2013) to mice (Friedman et al., 2010) and humans (Warnaby et al., 2017). Furthermore, this neural inertia can be modulated by factors altogether unrelated to the

concentration of anesthetic, such as single gene mutations (Friedman et al., 2010) and
manipulations of specific neuronal populations (Kelz et al., 2008; Reitz et al., 2021; Zhou et al.,
2018). Together, these results strongly argue that recovery from anesthesia is not simply the
byproduct of anesthetic washout. They do not, however, directly shed light on the mechanisms
that allow the brain to recover after general anesthesia.

71

72 In order to recover from anesthesia, the brain must follow a path through the state space that 73 begins in the deeply anesthetized state and eventually leads back to the pre-anesthetic 74 conditions. The neurophysiological processes that allow the brain to navigate this path efficiently have been addressed by Hudson et al. (2014). Specifically, they show that en route to recovery 75 76 of consciousness, brain activity is constrained to a low-dimensional space. In this space, most 77 activity is confined to a small number of discrete activity patterns, and the transitions between 78 these patterns are highly structured. In sum, these mechanisms greatly constrain the number of 79 possible paths through the activity space that can lead to wakefulness and allow the brain to 80 recover consciousness on a physiological time scale. Abrupt transitions between discrete activity 81 states have been observed in rodents (Hudson et al., 2014), non-human primates (Ballesteros et 82 al., 2020; Ishizawa et al., 2016; Patel et al., 2020) and human patients (Chander et al., 2014) after 83 exposure to a variety of anesthetics with distinct mechanisms of action. Abrupt transitions 84 between different activity patterns at a fixed anesthetic concentration are observed not only at the 85 level of the local field potentials (e.g., Hudson et al., 2014), but also in the activity of individual 86 cortical neurons (Lee et al., 2020). These discrete activity patterns and structured transitions 87 between them serve as a scaffold that guides the brain back towards normal patterns of activity 88 after it has been profoundly disrupted by anesthetics.

89

90 Given that state transitions are critical for reinstating consciousness, it is of fundamental 91 importance to determine the neuronal mechanisms that give rise to transitions between discrete

92 activity states during recovery from a dramatic perturbation. Previous work on anesthesia 93 (Chander et al., 2014; Hudson et al., 2014; Ishizawa et al., 2016) and sleep (Gervasoni et al., 94 2004) defined different activity patterns on the basis of oscillatory activity observed in the local 95 field potentials (LFPs) of firing of individual neurons (Lee et al., 2020). Much of this oscillatory 96 activity is coordinated via thalamo-cortical loops (Contreras and Steriade, 1997; Liu et al., 2015; 97 Schiff, 2008; Steriade et al., 1993b). An extensive body of work shows that the thalamocortical 98 circuitry is modulated by the arousal pathways ascending from the brainstem and basal forebrain 99 to produce oscillations at different characteristic frequencies (Destexhe et al., 1994; Jones, 2003; 100 Steriade et al., 1993a). Indeed, during constant anesthetic concentration, fluctuations in the firing 101 rates of individual neurons within these arousal nuclei co-vary with fluctuations in the spectra of 102 cortical LFPs (Gao et al., 2019). Direct manipulations of neuronal activity within the reticular 103 activating system can elicit profound changes in the oscillations observed in the cortical LFP (Gao 104 et al., 2019; Moruzzi and Magoun, 1949; Steriade et al., 1993a; Vazey and Aston-Jones, 2014). 105 Thus, one distinct possibility is that the discrete oscillatory patterns of activity observed under 106 fixed anesthetic concentration are imposed onto the thalamocortical networks by fluctuating 107 modulatory tone. If this is the case, because modulatory systems project broadly across the 108 thalamus and cortex (Jones, 2003), we expect to find that abrupt transitions between distinct 109 oscillations occur in close temporal proximity across the different cortical layers and regions. 110 Alternatively, it is possible that the oscillatory activity in different cortical regions is largely 111 coordinated through short-range thalamo-cortical and cortico-cortical interactions. In this case, 112 we expect to find that transitions between different oscillatory patterns are largely local.

113

Here, we provide direct experimental evidence for this latter possibility by simultaneously recording abrupt transitions between different states across cortical layers and across distant cortical areas at a constant anesthetic concentration. Using a complementary combination of analytic techniques, we show that state transitions across different cortical sites are only weakly

118 coupled. Furthermore, we demonstrate that state transitions in layer 4 (L4)—the layer that directly receives input from the thalamus—are particularly decoupled from state transitions observed in 119 120 other layers. This suggests that cortico-cortical interactions rather than fluctuations in the broad 121 modulatory tone play a crucial role in controlling state transitions under anesthesia. Remarkably, 122 we also show that the multitude of weak pairwise interactions between local state transitions is 123 sufficient to constrain the overall brain activity to just a few states embedded in a low-dimensional 124 space. Thus, our results suggest that the highly coordinated, low-dimensional macroscopic brain 125 dynamics that allow the brain to recover from a dramatic perturbation emerge as a consequence 126 of a multitude of weak pairwise interactions between different cortical sites.

### 127 Materials and Methods:

#### 128 Animals

129 All experiments were performed using ten male Sprague-Dawley rats, each two to three months 130 of age (250–350 g) (Charles River Laboratories, Wilmington, MA). Two animals were excluded 131 from further analyses because of excessive burst suppression or noise, respectively. One 132 additional animal was excluded after current source density analysis revealed that the V1 probe 133 was inserted too deeply to clearly identify cortical L4 and the supragranular layers. Rats were 134 housed under a conventional 12:12 h, light:dark cycle and given food and water ad libitum. All 135 experiments were performed in accordance with the Institutional Animal Care and Use Committee at the University of Pennsylvania and conducted in accordance with the National Institute of 136 137 Health Guidelines.

138

### 139 Surgery

140 All surgeries were performed under aseptic conditions. Each animal was weighed immediately 141 prior to surgery. Animals were induced with 2.5% isoflurane in oxygen and secured in a 142 stereotaxic frame (Kopf Instruments, Los Angeles, CA) in the prone position. Core body temperature was maintained at 37 (± 0.5) °C using a temperature controller (TC-1000 143 144 Temperature Controller, CWE, Incorporated, Ardmore, PA). Prior to surgery, isoflurane 145 concentration was reduced to 1.5% (flow rate 1 L/min), and dexamethasone (0.25 mg/kg) was 146 delivered subcutaneously. Bupivacaine (5 mg/mL) was injected under the scalp to provide local 147 anesthesia. Throughout the surgery, the lack of response to a toe pinch was used to assess 148 proper anesthetic depth.

149

The scalp was retracted and two 2 x 2 mm craniotomies were performed using a dental drill: one centered over -5.52 mm AP, 4 mm ML of bregma and another centered over -1.26 mm AP and 1.55 mm ML of bregma for V1 and M1 sites respectively. Dura was removed and Gelfoam (Pfizer, 153 New York, NY) was placed on the exposed cortical tissue to prevent the tissue from desiccating. 154 Prior to insertion, both linear probes (Cambridge NeuroTech, Cambridge, UK; H3 acute 64-155 channel linear probe) were dipped in Dil to allow for subsequent track tracing and lowered to 1.2 156 mm into the brain. Prior to electrode insertion, Dura Gel (Cambridge NeuroTech) was applied to 157 each craniotomy and isoflurane concentration was lowered again to 1% (flow rate 1 L/min) for 158 recordings. Immediately following electrophysiological recordings, animals were perfused trans-159 cardically with 4% paraformaldehyde under 4% isoflurane. Brain was harvested and processed 160 for electrode track tracing.

161

### 162 Histological confirmation of recording sites

Brains were sectioned at 80µm on a vibratome (Leica Microsystems, Wetzlar, Germany). Sections
were mounted with medium containing a DAPI counterstain (Vector Laboratories, Burlingame,
CA). Electrode tracks were manually identified and localized using epifluorescence microscopy
(Olympus, Tokyo, Japan; BX41) at 4x magnification.

167

### 168 Electrophysiology and Preprocessing

169 All recordings were performed at 1% isoflurane, after allowing the anesthetic concentration to 170 equilibrate for at least 30 minutes. Signals were amplified and digitized on an RHD2132 171 headstage (Intan, Los Angeles, CA) and streamed to a PC using an Omniplex acquisition system 172 (Plexon, Dallas, TX) at a rate of 40,000 samples per second per channel. All recordings were 173 performed using a ground skull screw as reference. Local field potentials (LFP) were extracted 174 from raw signals online using the bandpass filter with a passband of 0.1-300 Hz. Offline, LFP 175 were decimated to 1 kHz and filtered using a custom acausal FIR 0.1-200 Hz bandpass filter. 176 Noisy channels were removed by visual inspection of the signals. Before subsequent analyses, 177 data were re-referenced to the mean computed over all clean channels on the laminar probe. All 178 data analysis was completed using custom built MATLAB (MathWorks, Natick, MA) code unless otherwise stated. In total, 29.88 hours of recordings were used to generate all data in thismanuscript.

181

### 182 Current Source Density and Channel Selection

183 In order to facilitate cortical layer localization, a series of 10 ms light flash stimuli was presented 184 from a green LED positioned about one inch from the eye contralateral to the craniotomy over V1. 185 Interstimulus intervals were drawn from a uniform distribution between 3 and 5 seconds to prevent 186 stimulus entrainment. Current source density (CSD) analysis was then applied to the post-187 stimulus LFP to identify layers in V1. The CSD C<sub>t</sub> at time *t* was calculated by computing a 188 smoothed second spatial derivative (a representative example is shown **Figure 5**):

$$egin{split} C_t(z) &= V_t(z) \, st \, K(z) \ K(z) &= rac{z^2 - \sigma^2}{\sigma^5 \sqrt{2\pi}} \mathrm{exp}\left(rac{-z^2}{2\sigma^2}
ight) \end{split}$$

189

Here, *z* is the channel depth,  $\sigma = 280 \ \mu m$  is the distance along the electrode from *z* at which the kernel changes sign, *V<sub>t</sub>* is the mean voltage over all light flash trials at time *t* relative to flash onset, and \* indicates convolution. The electrode closest to the center of L4 was identified manually from the CSD as the earliest current sink. Once L4 was identified, supra- and infragranular channels were selected for analysis at 140 µm intervals above and below L4.

195

### 196 Time-Frequency Analysis

Spectrograms of selected channels were calculated from LFP signals using the multitaper method with 17 Slepian tapers and time-bandwidth product (NW) = 9. A 6-second sliding window with a step size of 100 ms was used. Windows containing signal artifacts were identified and removed using a combination of automatic burst suppression detection based on the root-mean-square of LFP in a moving exponential window and manual inspection of multitaper spectrograms. Each window was zero-padded to 65.536 s to increase the frequency resolution and input a power-of203 2 number of samples to the Fourier transform. In order to sample frequencies of greater interest more densely, 279 frequencies were selected from 0.14 to 300 Hz, spaced on a log scale from 204 205 0.14 to 10 Hz and on a linear scale above 10 Hz. The multitaper spectrograms were then 206 smoothed over frequencies with a median filter spanning 10 frequency steps (up to 17.5 Hz) and 207 over time with an exponential (Poisson) window spanning 2 minutes. In order to remove baseline 208 differences in power across frequencies (such as power-law scaling) and emphasize temporal 209 fluctuations, each spectrogram was rank-order normalized along the time axis. At each frequency 210 bin, the time window with the highest power was given the value of one. Each other window was 211 given the value of (r-1)/(N-1), where r is that window's sorted index among the N windows. Thus, 212 the smallest power value at each frequency was represented as zero, and the largest as one.

213

### 214 Dimensionality Reduction

215 Dimensionality reduction was performed on each channel's spectrogram individually, in order to 216 obtain high reconstruction accuracy and ensure that any characteristic differences in activity 217 patterns between sampled regions and cortical depths were preserved. Non-negative matrix 218 factorization (NMF) (Lee and Seung, 1999; Mankad and Michailidis, 2013) was used to compress 219 the rank-ordered spectrograms. The NMF output represents the signal at each time as a short 220 vector of K non-negative coefficients (scores) that weight a sum of corresponding frequency 221 components (loadings) to reproduce the original spectrum. Given a spectrogram A of size 279 x 222 N, NMF produces a loading matrix U of size 279 x K and a score matrix V of size N x K. The 223 product  $UV^{\mathsf{T}}$  reconstructs A with some error E, quantified relative to the norm of A as:

$$E=rac{\|A-UV^{ op}\|_F}{\|A\|_F}$$

224

Where  $\|\cdot\|_F$  is the Frobenius norm. To select an appropriate number of components (*K*) for each channel, a cross-validation approach was employed (Owen and Perry, 2009). First, spectrograms 227 were downsampled across time by a factor of 20, for computational efficiency. Then, a random 228 subset of 20% of the rows and columns were selected to be withheld. Starting with K = 1 and 229 increasing to 15, NMF was applied to the down-sampled matrix after the random subset of rows 230 and columns had been removed. This iteration provides both a loading and score matrix. Next, 231 NMF was run again on the data with only the pre-selected rows withheld. In this iteration, the 232 loading matrix from the first round was fixed and only a new score matrix was calculated. In the 233 third and final run of NMF, NMF was run on the data with only the pre-selected columns removed, 234 fixing the score matrix from the first round and calculating only a new loading matrix. Finally, the 235 loading and score matrices produced in the second and third run of NMF, respectively, were 236 multiplied to generate an estimate of the original dataset and calculate error as a function of K. 237 This procedure was repeated for five replicates for each value of K, and the optimal K was chosen 238 such that increasing K by one would reduce mean reconstruction error by less than 1%. In our 239 dataset, the optimal value for K ranged from five to nine for different channels. After the cross-240 validation procedure, each channel's full, normalized spectrogram was subjected to NMF using 241 the channel's optimal K, resulting in a mean reconstruction error of 14.8% across all channels 242 (~85% of the variance captured by NMF for each spectrogram). Note that NMF does not constrain 243 the relative scales of the loading vectors: for any invertible diagonal K x K matrix D,  $UV^{T} = UD(VD^{-1})^{T}$ . To remove these degrees of freedom, U and V were rescaled by a matrix D 244 245 such that the rescaled loadings had unit L<sub>2</sub> norm.

246

### 247 Transition and Discrete State Identification

The rescaled score matrix  $VD^{-1}$  is the basis for defining each channel's state over time. For each channel, at each time point, the component with the highest score was taken as the state of the brain near that channel's recording site, and samples where the state changed were marked as local transition times. In order to prevent an arbitrarily high number of transitions during periods when two or more components had similar scores, transitions that were likely to reflect transient

253 fluctuations were ignored and the state assignments between them were updated accordingly. 254 Specifically, suppose one time segment between two transitions was assigned state "A" and either 255 the previous or next segment was assigned state "B." If the first segment was less than 100 256 seconds long and, within the first segment, the mean score for NMF component A was less than 257 1.1 times the mean score for component B (*i.e.*, if the state assignment was sufficiently 258 ambiguous), the transition between the two segments was ignored and the combined segment 259 was assigned state B. If a segment could be merged with either the previous or next segment, 260 the tie was broken by ignoring the transition with a smaller magnitude of change in the full NMF 261 score vector from the 3 seconds before the transition to the 3 seconds following it. A matrix of 262 state transition frequencies was computed by tabulating how often each discrete state followed 263 each other state over the duration of the recording using the table of discrete state transitions for 264 each channel.

265

### 266 Markov-based Shuffled Null Model

267 When testing whether pairs of channels are synchronized in the sense that they preferentially 268 occupy certain combinations of discrete states, apparent synchrony could arise due to the 269 channels' individual NMF score distributions, independent of the relative timing of transitions. To 270 control for this possibility, a discrete-time Markov chain (the "null model") was fit to the transition 271 frequencies of each channel independently. The channel's null model was then used to simulate 272 1000 new discrete state sequences of the same length as the original data. For each pair of 273 channels, these "null" state sequences were then used to fit distributions of transition synchrony 274 and normalized mutual information (see corresponding sections below). This distribution reflects 275 the probability of observing a given state synchrony and mutual information under the assumption 276 of complete independence between different recording sites. To obtain a null distribution of 277 canonical correlation-based synchrony (see below), full score matrices were generated from each 278 channel's null state sequences as follows: for each of the K states k, at each sample with null

discrete state assignment *k*, the corresponding row of the null score matrix was randomly drawn
from the set of rows of the original data score matrix where the original discrete state was equal
to *k*. These random sequences for all pairs of channels were then subjected to canonical
correlation analysis.

After fitting normal distributions for each of the three channel pair interaction measures (transition synchrony, normalized mutual information, and canonical correlations) to the shuffled surrogates, the values obtained for the real data were tested against these distributions to estimate whether they would be expected by chance, given the statistics of the data (see "Statistical Tests" below).

287

#### 288 Transition Synchrony

289 To quantify how frequently channels transitioned together we employed the SPIKE-290 synchronization score ("synchrony score"), a method for quantifying synchrony between two 291 simultaneously recorded sequences of events (Kreuz et al., 2015). At its core, this method is a 292 coincidence detector in which the coincidence window is derived from the dataset. The adaptive 293 definition of the coincidence window means that this method for quantifying synchrony is equally 294 well-suited for state transitions as it is to spike trains. Each transition r is assigned a local window 295 length  $\tau(r)$ , which is defined as half the smaller of the inter-transition intervals directly before and 296 after r. For a pair of channels i and j, if transition  $r_i$  in j was the closest transition to transition  $r_i$  in j 297 and vice versa, and the time between  $r_i$  and  $r_i$  is less than min( $\tau(r_i), \tau(r_i)$ ), both transitions have a 298 synchrony score of 1. All other transitions have a score of 0. This measure is extended to the 299 multi-channel case by assigning each transition a synchrony score equal to its mean pairwise 300 synchrony score with the nearest transitions in all other channels. Both pairwise and all-channel 301 synchrony scores were computed for all discrete state transitions in each recording, and then 302 averaged over all transitions to obtain pairwise and global mean synchrony measures.

- 303
- 304

### 305 Normalized Mutual Information

306 Mutual information of discrete states was used to quantify the synchrony of states themselves 307 rather than just the timing of their transitions. Specifically, this measure was implemented to 308 quantify how well one could predict the state in one channel, given the state of another channel 309 at the same time point. Since NMF was performed separately on each channel, states labeled 310 with the same index in different channels are not necessarily the same with respect to the 311 frequency characteristics of the signal. Regardless, mutual information is able to reveal temporal 312 relationships between channel pairs because it does not assume any particular relationship 313 between the state assignments of the different channels and is, therefore, agnostic to the 314 assignments themselves.

315

316 Mutual information I(X; Y) between two channels X and Y with N observations and sets of classes

317  $K_X$  and  $K_Y$  was computed pointwise as follows:

$$I(X;Y)=I(Y;X)=H(Y)-H(Y\,|\,X)$$

$$H(Y) = -\sum_{k\in K_Y} P(Y=k)\log_2 P(Y=k)$$

$$H(Y \,|\, X) = -\sum_{j \in K_X} P(X=j) \sum_{k \in K_Y} P(Y=k \,|\, X=j) \log_2 P(Y=k \,|\, X=j)$$

$$P(Y=k) = egin{array}{l} rac{|\{t\,|\,Y[t]=k\,\}|}{N} \ P(Y=k\,|\,X=j) \ = egin{array}{l} rac{|\{t\,|\,X[t]=j,\,Y[t]=k\}}{|\{t\,|\,X[t]=j\}|} \end{array}$$

318

Mutual information is not a pure measure of the predictability of one variable given the other; it also increases with the entropy of each variable. For example, if channels X and Y each occupy a wider distribution of states and, as a result, have higher entropy than both channels W and Z, then I(X; Y) > I(W; Z). This is true even if the state of X is perfectly predictable given Y, Y given X, W given Z, and Z given W. In order to control for this, mutual information was normalized by

324 the sum of the entropies of the two channels, giving the normalized mutual information, or 325 symmetric uncertainty (Witten et al., 2011):

$$U(X,\,Y)\,=\,2rac{I(X;\,Y)}{H(X)+H(Y)}$$

326

329

327 Using another definition for mutual information in terms of the individual and joint entropies of *X*328 and Y, we can write:

$$U(X,Y) \,= 2 rac{H(X) \,+\, H(Y) - H(X,Y)}{H(X) + H(Y)}$$

Thus, normalized mutual information can be understood as twice the fraction of the sum of individual entropies, H(X) + H(Y), that exceeds (is redundant to) the joint entropy H(X, Y) due to mutual information between X and Y. For example, if X and Y are identical, U(X, Y) = 1 and 50% of H(X) + H(Y) is redundant, as only one of the variables carries unique information.

334

### 335 Canonical Correlation

336 Both the transition synchrony and normalized mutual information measures assume that LFP 337 signals at each channel form discrete states and that the sequence of NMF components with the 338 largest magnitude at each time point is informative about this state. However, there may be cases where multiple components must be considered. For instance, consider a situation in which NMF 339 340 component A in channel i is characterized by strong activity in two frequency bands, and 341 components B and C in channel *j* are characterized by strong activity in one of those frequency 342 bands each. If only the "top" component determines the discrete state, there could be artificially 343 low synchrony and mutual information between channels *i* and *j*. This is because, during a bout 344 of state A in channel *i*, there could be frequent switching between states B and C in channel *i*. 345 even though the overall signal characteristics in channel *j* remain largely static. To address this 346 kind of ambiguity and compute a state synchrony measure that softens the artificially sharp boundaries between "discrete states," canonical correlation analysis (CCA) was applied to the 347

348 NMF score matrices of pairs of channels. Intuitively, CCA allows each score matrix to be linearly 349 transformed to optimally match components between channels. CCA maximizes the correlations 350 between the matched, transformed components. These correlations are used to derive a measure 351 of state similarity.

352

The procedure for computing CCA-based synchrony is as follows: let  $V \in \mathbb{R}^{N\times L}$  and  $W \in \mathbb{R}^{N\times M}$  be the NMF score matrices two channels, and let  $K = \min(L, M)$ . At each step *i* from 1 to K, CCA finds coefficient vectors  $a_i$  and  $b_i$  to maximize the correlation  $\rho_i = \operatorname{corr}(Va_i, Wb_i)$ , with the constraints that  $a_i$  is uncorrelated with all previous vectors  $a_1, ..., a_{i-1}$ , and likewise for  $b_i$ . The MATLAB function *canoncorr* was used to perform this algorithm and the canonical correlation coefficients  $\rho_1, ..., \rho_K$  were averaged to obtain a state similarity measure.

359

#### 360 Statistical Tests

361 This section describes the procedure used to establish the statistical significance of interactions 362 between recordings sites as measured by the synchrony score, normalized mutual information, 363 and canonical correlation analysis. For each channel pair under consideration and each of these 364 three interaction measures, the measure was computed both on the experimental dataset and on 365 a set of 1000 null-model datasets generated from discrete Markov models of each channel's transition statistics, as described above. The values of each measure were approximately 366 normally distributed across null-model datasets. To test statistical significance, the deviation of 367 368 each measure obtained in the experimental dataset from those generated from null-model 369 datasets was expressed as a z-score. The one-tailed p-value was then directly computed from 370 the z-score. The significance threshold was set at  $\alpha$ =0.05. Bonferroni correction was applied to 371 account for multiple comparisons over all channel pairs in each animal. The percentage of pairs 372 for which each interaction measure was different from chance after Bonferroni correction is reported in the manuscript, and non-significant pairs are graved out in Figures 6-8. 373

374

375 To compare interaction measures between different sets of channel pairs, special consideration 376 must be paid to the statistical dependence between observations. In a recording with *n* channels, 377 for any channel k, one would not in general expect the values of a distance-like measure on the 378 pairs (k, 1), ..., (k, k-1), (k, k+1), ..., (k, n) to be independent. For example, if channel k were an 379 outlier, all n-1 pairs would take extreme values due to what is statistically only one extreme 380 observation. If pairwise statistics were compared naively, e.g., using a two-sample *t*-test, these 381 dependencies would result in an overestimation of effective sample size and thus significance. 382 Instead, a Monte Carlo permutation procedure was used to establish null distributions for 383 comparisons of pairwise measures between groups of channel pairs. This procedure randomly 384 shuffled group assignments while preserving the dependency structure inherent in the matrix of 385 pairwise measures by only shuffling rows and columns. For each such comparison, 10<sup>7</sup> 386 permutations of only the channels of each recording that were included in that comparison were 387 conducted, and the difference of group means was computed after each permutation. The 388 frequency with which these null differences exceeded the difference of means of the unpermuted 389 groups was taken as the *p*-value of the comparison.

390

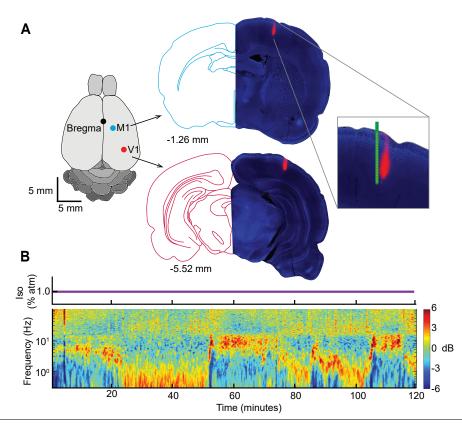
391 Finally, when comparing the interaction measures for between-region channel pairs in M1/V1 392 recordings to those in bilateral V1 recordings, the method of permuting channel labels cannot be 393 used because there are no data for pairs of channels that mix different recordings. Instead, the 394 distribution for the difference of means of the measure over pairs between the two sets of 395 recordings was estimated by bootstrapping over channels. Specifically, each group in such a 396 comparison consists of a set of rectangular matrices, containing values of the measure for each 397 pair of one channel along the rows and one channel along the columns. By resampling both rows 398 and columns with replacement in each such matrix, the dependencies along rows and columns 399 were preserved, but the variance in the mean could be estimated thanks to the principles of

- 400 bootstrapping. A total of 10<sup>6</sup> bootstrapped estimates of the group mean difference were computed
- 401 in this manner for each interaction measure and used to obtain a *p*-value for the one-tailed
- 402 hypothesis that the measure is greater on average between hemispheres of V1 than between M1
- 403 and V1.

### 404 Results:

### 405 State transitions under constant anesthetic can be local.

406 We sought to determine whether state transitions under a fixed concentration of isoflurane (1% 407 atm.) occur simultaneously across different cortical regions and across layers within the same 408 cortical region. This concentration was chosen based on previous work (Hudson et al., 2014) 409 showing that burst suppression is not likely to occur at this concentration, but that state transitions 410 in the spectral characteristics of the LFP are frequently observed. Here we focused on the local 411 field potentials (LFPs) recorded using two laminar probes that sampled signals across all cortical 412 layers. In half of the experiments, both electrodes were inserted into the right hemisphere: one in 413 the primary visual area (V1) and the other in the motor cortex (M1) (n = 3) (Figure 1A). In the



**Figure 1: Experimental setup. A.** Verification of Electrode placement into V1 and M1. DAPIstained histological section showing tracks of the Dil-dipped electrode (right) juxtaposed with the corresponding section from the rat brain atlas (left). The zoomed cutout includes an image to show electrode channel layout. **B.** Time-resolved spectrogram recorded from V1 under 1% isoflurane general anesthesia (concentration shown above spectrogram). Spectrogram is plotted as deviations from temporal mean.

other half of experiments, bilateral V1 recordings were performed (n = 4). Postmortem localization
of electrodes (Methods) in a representative experiment is shown in Figure 1B. Consistent with
previous findings (Hudson et al., 2014), at 1% isoflurane, the power spectrum of the LFP
fluctuated between several discrete states (Figure 1C).

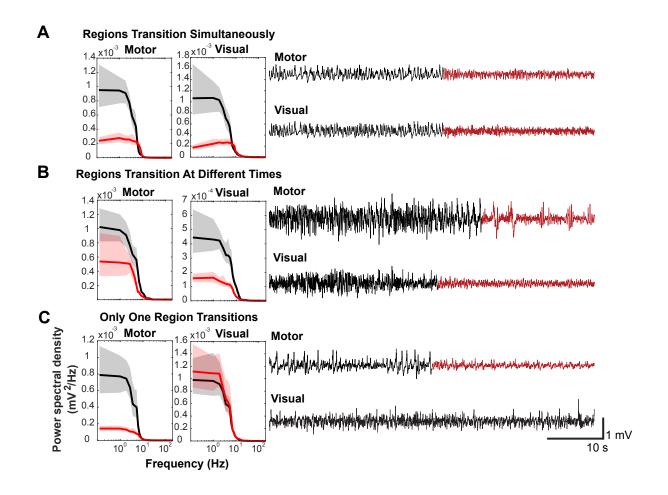
419

420 State transitions can be readily identified in the raw LFP (Figure 2). The top and bottom LFP 421 traces show one minute of recordings from a single M1 and V1 electrode, respectively. The 422 accompanying spectra were calculated using a multitaper spectral estimate. These spectra were 423 averaged across two second windows of LFP with a one second step size, sampled either from eight to two seconds prior to transition (black, pre-transition) or from two to eight seconds after 424 425 the transition (red, post-transition). Spectral estimates are shown as mean ± 95% confidence 426 interval computed from 1000 bootstraps. In some instances, state transitions occur approximately 427 simultaneously in the motor and visual cortices (Figure 2A). However, this was not always the 428 case. For instance, Figure 2B shows an example of a state transition that occurs first in the visual 429 cortex and, only after a delay of approximately 10 seconds, is seen in the motor cortex. Thus, 430 abrupt changes in the LFP characteristics need not occur simultaneously in different brain 431 regions. Figure 2C shows a more extreme example of this phenomenon. A state transition is 432 clearly seen in the motor cortex, but in the visual cortex, the LFP characteristics remain 433 unchanged. These observations suggest that, while some state transitions may indeed be global, 434 there is a previously unappreciated degree of independence between state fluctuations observed 435 in the cortex during fixed anesthetic administration.

436

437 Multitaper analysis and non-negative matrix factorization extract states and their
 438 transitions across cortical layers and regions.

439 To quantify the degree of coupling between state transitions at different recording sites, we 440 developed a methodology to automatically detect state transitions at the level of individual



**Figure 2: Examples of state transitions. A-C** Right: LFP traces (1 minute) recorded simultaneously from right M1 and V1. Visually apparent abrupt transitions in the character of the LFP are indicated by shifts of color from black to red. Left: spectra computed from the red and black time periods respectively to indicate that the abrupt switches in the features of the signals are associated with changes in the spectra. A. An example where both M1 and V1 LFPs appear to change state simultaneously. **B.** An example where both M1 and V1 signals change states but with an appreciable time delay (~10 s). **C.** An example where a state transition is observed in M1 but not in V1. In this case for the purposes of computing the spectrum (left, red) in V1, the time segment highlighted in red for the M1 electrode was used.

441

442 channels (Methods). We then deployed this methodology to determine the degree to which

transitions in different cortical sites are coupled. **Figure 3** is a flowchart of the initial analysis steps.

444 The first step in the analysis is to compress the LFP recording into a low-sample-rate, low-

dimensional matrix that accurately captures fluctuations in oscillatory activity. The right side of the

figure presents an example five-minute window of data from one recording site to demonstrate

the outcome of each step. Briefly, wideband data were filtered between 0.1 and 300 Hz to extract

the LFP signal. (Figure 3A) LFP signals were converted to frequency domain using multitaper spectral analysis, (Figure 3B). Raw power spectra were then normalized such that the power contained in each frequency band was mapped onto a value between 0 (smallest observed power) and 1 (largest observed power) (Figure 3C). Non-negative matrix factorization (NMF) was used to further decompose the signal into a set of loadings and associated scores across time (Figure 3D-E).

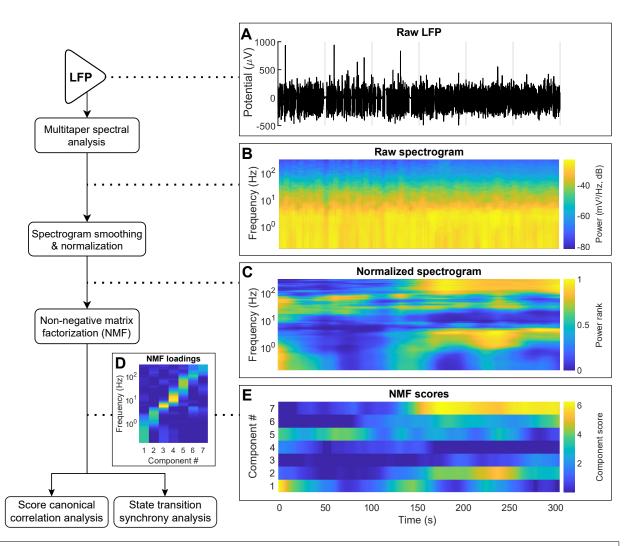
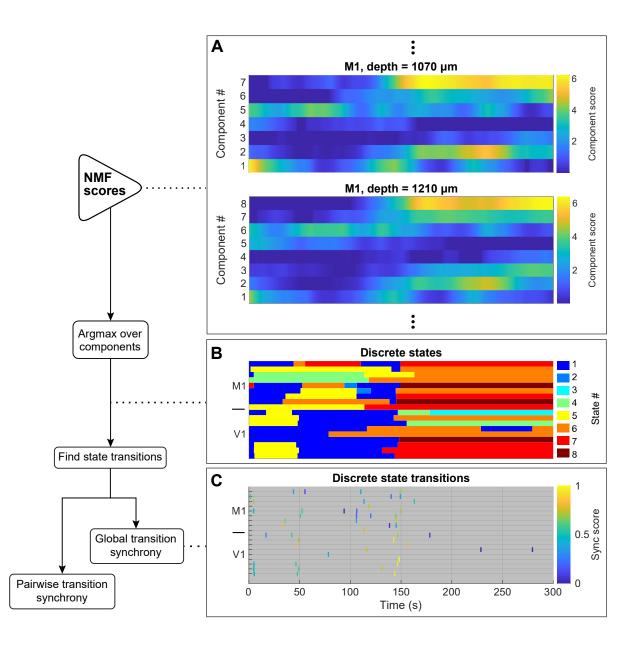


Figure 3. Schematic of LFP analysis, through NMF calculation. Left: Flowchart of analysis steps. Right: A. Five minutes of raw LFP signal centered around a state transition.
B. Power spectrogram of LFP, computed using the multitaper method. C. The spectrogram from panel B after smoothing and rank-order normalization across time (Methods). D-E The loading (D) and score (E) matrices generated using NMF showing the spectral characteristics of each component and its relative contribution to the signal across time, respectively. The number of NMF components was optimized individually for each channel (Methods).

455 NMF can be thought of as a "soft" clustering algorithm. Previous work on state transitions under 456 anesthesia (Hudson et al., 2014) and sleep (Gervasoni et al., 2004) used k-means clustering of 457 the spectrograms to assign the state of the brain. Our first approach to state assignment used a 458 similar strategy-the index of the NMF component with the highest score in each time window 459 was defined as the state of the LFP at each recording site (Methods). This assumption was 460 relaxed in subsequent stages of the analysis (see below). Figure 4A shows the score matrices 461 for two different channels recorded simultaneously from two contacts along the same electrode 462 in the motor cortex. The upper matrix is the same as **Figure 3E**, and the lower matrix was 463 generated from data collected by a contact 140 um deeper inside the cortex. Notice that these 464 matrices resemble one another but are not identical. Figure 4B shows state classifications for 18 465 channels of simultaneously recorded data: nine from an electrode in V1 and nine from an 466 electrode in M1. Note again that some state transitions are observed around the same time in 467 most of the electrodes. There are, however, many instances where state transition is observed in 468 just a subset of the recording sites.

469

470 One way to characterize the coupling between state transitions is to quantify the propensity of 471 state transitions to occur simultaneously across different recording sites. Brain state transitions 472 were defined as time points at which consecutive windows from the same channel have different 473 brain state assignments (Methods). Figure 4C shows an example of this analysis. There are 474 many transitions that appear in only one or very few channels, while others appear to be more 475 global. Figure 4C is a raster plot of transitions. The color of each line shows the synchrony score 476 of that transition with all other channels (Methods). Consistent with the observations in Figure 2 477 and 4C, the synchrony score reflects the fact that most state transitions are localized to a small 478 subset of electrodes.

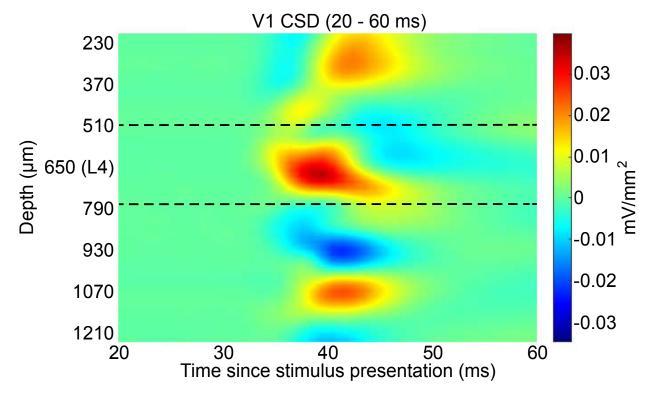


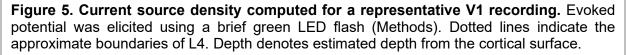
**Figure 4. Schematic of NMF score analysis to define state transitions and synchrony.** Left: Flowchart of analysis steps. Right: **A.** The NMF score matrix presented in Figure 3E (upper) and another NMF score matrix from simultaneously collected LFP from a neighboring channel (lower). Note that while nearby channels share similar characteristics across time, they are not identical. Also, the two channels have different optimal number of components, since NMF was performed and optimized (Methods) independently for each channel. **B.** State assignments across example time window from 18 simultaneously recorded signals: 9 signals from an M1 (top rows) electrode and 9 from V1 (bottom rows). State # indicates the NMF component with the highest score in each time window, after removing state segments that were both short and ambiguous due to small score fluctuations (Methods). **C.** Raster plot of all transition times from the channels presented in panel **B**. Transitions are colored according to their synchrony (sync score) with transitions in all other channels (Methods).

480 As we show below, coupling between state transitions depends on the cortical layer. Layer 481 assignment in V1 was performed using current source density (CSD) analysis computed 482 immediately following brief light stimulus (Methods). Figure 5 shows a representative example of 483 CSD in V1 showing the stereotypical pattern of response to visual stimuli. The first current sink 484 occurs approximately 33 ms following stimulus presentation in L4. A short time after, additional 485 sinks and sources appear above and below, revealing interlaminar communication. The channel 486 where the initial sink occurred was defined as the center of L4. The dashed black lines in this 487 figure mark the approximate boundaries of L4 based on the average thickness of this layer in rats 488 and the spacing between channels (Einevoll et al., 2013; Quairiaux et al., 2011; Self et al., 2013). 489 In the motor cortex, we did not estimate the location of cortical layers directly. Instead, we 490 estimated the depth of each recording electrode relative to the cortical surface.

491







### 494 State transitions in different cortical sites exhibit weak synchrony.

495 We used three different analytical techniques to quantify the tendency of oscillatory states and 496 the transitions between them to be coordinated across recording sites. Each technique relies on 497 a different set of assumptions and was performed on a different feature of the data. First, we 498 quantified the synchrony of transitions, as demonstrated in Figure 4 (Methods). Figure 6A-B 499 shows the cumulative distribution of synchrony scores (red curves) computed over all channel 500 pairings and across all animals (M1/V1: 3 animals, 16-18 electrodes/animal, median of 99 501 transitions/electrodes/animal; bilateral V1: 4 animals, 15–19 electrodes/animal, median of 175.5 502 transitions/electrode/animal).

503

504 In order to compare the synchrony scores (Figure 6A-B) to those expected by chance, we 505 generated shuffled datasets constrained to have the same state transition statistics. This was 506 accomplished by simulating a Markov process defined by the state transition probability matrix 507 derived from state assignments for each recording (Methods). This control preserves the statistics 508 of each recording site, while destroying any coordination between them. The cumulative 509 distributions of the synchrony scores obtained in these shuffled controls are shown in Figure 6A-510 **B** (blue curves; shading shows 95% confidence intervals computed over 1000 shuffled datasets). 511 Both in the experiments involving M1 and V1 (Figure 6A) and in those involving bilateral V1s, we 512 find that the synchrony score is consistently higher than expected by chance (p < 0.001, z-test 513 based on means of shuffled datasets). Despite this large deviation from the null hypothesis, state 514 transitions do not typically occur at the same time in different cortical sites (mean synchrony score 515  $\approx$  0.35 for both M1/V1 and bilateral V1 recordings). This implies that while state transitions 516 observed across different cortical sites are not completely independent, coupling between 517 channels is weak.

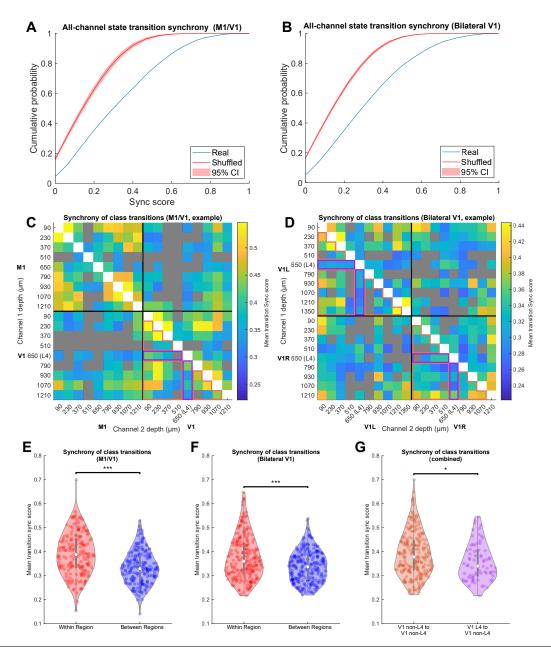


Figure 6. Transition synchrony between channels in the same anatomical region is higher than between channels in different regions. A-B The cumulative distribution of SPIKE-synchronization (synchrony) scores across all channels, in real recordings (blue) and the median  $\pm$  95% CI of 1000 shuffled recordings (red), for M1/V1 experiments (**A**) and bilateral V1 experiments (**B**). C-D Mean synchrony score across transitions for all channel pairs from a representative M1/V1 (**C**) and bilateral V1 (**D**) recording. Channel pairs whose synchrony scores were not significantly different from shuffled controls after Bonferroni correction are colored gray. **E-F** Channel pairs in which both channels are in the same region (red) have higher synchrony scores than those in which the channels are in different regions (blue) for M1/V1 (**E**, p = 1e-7, permutation test) and bilateral V1 (**F**, p = 2e-7, permutation test) recordings. **G.** Channel pairs in which one channel was within L4 and the other was not had lower synchrony scores than pairs in which neither channel was in L4 (p = 0.015, permutation test). Data included in these comparisons for the representative experiments are outlined in orange and purple, respectively, to highlight that only data from V1 electrodes were used.

519

520 Data in Figure 6A-B aggregate the transition synchrony scores calculated between all channel 521 pairs—both pairs of channels in the same cortical region and those located in different cortical 522 sites. We hypothesized that, because most cortical connectivity is local, nearby electrodes would 523 tend to have a higher propensity to change state at the same time. **Figure 6C-F** shows that state 524 transitions are indeed more synchronous between electrodes within a cortical region than 525 between regions. Figure 6C-D shows synchrony scores between all channel pairs in a 526 representative pair of experiments: an M1/V1 experiment (Figure 6C) and a bilateral V1 527 experiment (Figure 6D). Pairs with scores that did not reach significance compared to the shuffled 528 datasets, after Bonferroni correction for multiple comparisons, are shown in gray. Across all 529 experiments, 57.0% of channel pairs from M1/V1 experiments and 80.2% of pairs from bilateral 530 V1 experiments had significantly synchronous transitions at the corrected p < 0.05 level. The 531 synchronization scores for all channel pairs from all experiments are quantified in Figure 6E-F. 532 for M1/V1 and bilateral V1 experiments respectively. Both panels show the synchrony scores for 533 within-region channel pairs (red) and between-region channel pairs (blue). In both types of 534 recordings, within-region pairs had significantly larger synchrony scores than between-region 535 pairs (p = 1e-7 for M1/V1 and p = 2e-7 for bilateral V1, compared to  $10^7$  random permutations of 536 the relevant channels (Methods)).

537

L4 is the thalamic input layer and has fewer horizontal connections than the supragranular or infragranular layers, which are rich in horizontal connections (Zilles and Palomero-Gallagher, 2017). To test whether layer organization affects transition synchrony, from each V1 recording (in which L4 was identified using CSD), we separated channel pairs in which one channel was in L4 from pairs in which neither channel was in L4. **Figure 6G** presents synchrony scores from all channel pairs from all experiments in which one channel was in L4 and the other was not (purple) and all channel pairs from all experiments in which neither channel was in L4 (orange). In **Figure** 

6E and F. the specific channel pairs that were included in the "L4" and "non-L4" groups are 545 546 outlined in purple and orange, respectively. We found that synchrony between channel pairs with 547 one channel in L4 tended to be lower than between pairs in which neither channel was in L4 (p =548 0.015, compared to  $10^7$  random permutations of the relevant channels (Methods)). Therefore, 549 transition times in channels from L4 tend to be relatively uncoupled from the specific timing of 550 transitions in channels from other layers. This observation suggests that it is unlikely that 551 thalamocortical input is the principal driver of state transitions in the cortex. If it were, one would 552 expect that the thalamic input layer (L4) would transition in synchrony with the rest of the cortex. 553 Therefore, these results imply different mechanisms, such as cortico-cortical interactions, are 554 likely responsible for the timing of these spatially localized transitions.

555

556 Our final analysis using synchrony scores was performed to build upon these L4 results and 557 determine whether the type of subcortical input to a cortical region has an influence on transition 558 synchrony. It is typically assumed that switches of the oscillatory activity in the cortical LFP 559 critically involve interactions with the thalamus (Contreras and Steriade, 1997; Herrera et al., 560 2016; Liu et al., 2015; Schiff, 2008; Steriade et al., 1993a, 1994). In light of this, one may expect 561 two regions receiving similar thalamic input to exhibit greater synchrony of state transitions than 562 two regions that interact with the thalamus in different ways. Therefore, we tested whether 563 between-region comparisons for the bilateral V1 experiments had higher synchrony scores than 564 the between-region comparisons for the M1/V1 experiments. Contrary to our hypothesis, we were 565 not able to detect any increase in synchrony scores calculated between the bilateral V1s relative 566 to M1/V1 experiments (p = 0.35, percentile bootstrap over channels (Methods)).

567

### 568 **Discrete states in different cortical sites have weak correspondence.**

569 Until this point, our analysis was based on transition synchrony, a measure that is sensitive to the 570 timing of transitions but not the identities of the states. In what follows, we shift our focus away

571 from the timing of state transitions and quantify the consistency of LFP-defined states at different sites. We accomplish this using normalized mutual information (MI), a measure of the amount of 572 573 information obtained about one random variable by observing another random variable 574 (Methods). In our case, these random variables are the time series of discrete states of two 575 channels. High MI between these time series represents a large reduction in uncertainty about 576 the state in channel *j* given the state in channel *i*. Two channels do not need to be in the same 577 brain state to have high mutual information; indeed, since states are defined for each channel 578 independently, there is no definition of different channels being in the "same" state. Rather, there 579 must only be a consistent mapping from the states in one channel to those in the other. For 580 example, if channel *i* is always in state A whenever channel *j* is in state D, one can predict the 581 state of channel *i* from the state of channel *i*, and the MI between these channels would be high. 582 As noted in the Methods, we normalized MI by the total entropy of the state distributions in the 583 two channels over time in order to obtain a measure that was comparable across channels with 584 different state distributions.

585

586 Figure 7A-B shows the normalized MI between all channel pairs in the same representative 587 M1/V1 and bilateral V1 experiments as those in Figure 6C-D. 81.9% of channel pairs from M1/V1 588 experiments and 96.9% of pairs from bilateral V1 experiments had normalized MI that was 589 significantly higher than for shuffled data, after Bonferroni correction for multiple comparisons (z-590 test based on distribution of shuffled data). The summary of normalized MI across all animals is 591 shown in Figure 7C-D, for M1/V1 and bilateral V1 experiments respectively. In both types of 592 recordings, within-region channel pairs had significantly higher normalized MI than between-593 region pairs (p = 1e-7 for M1/V1 and p = 1e-7 for bilateral V1, compared to  $10^7$  random 594 permutations of the relevant channels (Methods)). Note that, while for most channel pairs MI was 595 higher than for a shuffled dataset, the amount of information about the state of one channel 596 contained in the state of another was small. Normalized mutual information varies between 0 and

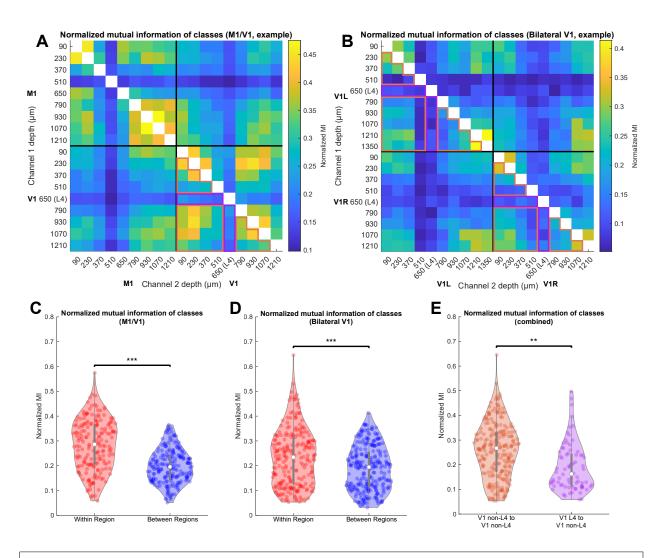


Figure 7. Normalized mutual information (MI) between channels in the same anatomical region is higher than between channels in different regions. A-B Normalized MI between state assignment vectors for all channel pairs from a representative M1/V1 (A) and bilateral V1 (B) recording. All normalized MI values are significantly different from shuffled controls after Bonferroni correction. C-D Channel pairs in which both channels are in the same region (red) have higher normalized MI than those in which the channels are in different regions (blue) for M1/V1 (C, p = 1e-7, permutation test) and bilateral V1 (D, p = 1e-7, permutation test) recordings. E. Channel pairs in which one channel was within L4 and the other was not had lower normalized MI than pairs in which neither channel was in L4 (p = 0.002, permutation test). Data included in these comparisons for the representative experiments are outlined in orange and purple, respectively, to highlight that only data from V1 electrodes were used.

597

598 1, where 1 denotes that the two channels carry identical information. Yet, even in a pair of

- channels within a single cortical region, the mean MI is about 0.3. One way to interpret this statistic
- 600 (Methods) is that no more than 15% of the combined information carried by the states of any two

601 channels is redundant. Thus, most of the information about the state of one channel cannot be602 extracted from observing the state of a nearby channel in the cortex.

603

604 As with transition synchrony, we did not detect a higher mean normalized MI in left/right V1 605 channel pairs compared to M1/V1 channel pairs (p = 0.70, percentile bootstrap over channels 606 (Methods)). Additionally, as with the transition synchrony analysis, pairs including a channel in L4 607 did have lower normalized MI than pairs where neither channel was in L4 (p = 0.002, compared 608 to 10<sup>7</sup> random permutations of the relevant channels (Methods)). These results show not only that 609 channels from the same brain region are more likely to undergo transitions at the same time, but 610 also that the broader structure of these state assignments across the entire recording is more 611 similar in channels from the same region. Furthermore, the conclusions regarding the differences 612 between L4 and other cortical layers are consistent between synchrony and mutual information 613 analyses.

614

Full compressed spectrograms of different sites have moderate correspondence,
 depending on distance and cortical layer.

617 In the previous analyses, to generate a single-value description of activity across time, we defined 618 brain state as the NMF loading with the highest score in each time window. This method was 619 convenient for comparing synchrony of transitions and mutual information of state sequences. 620 Parcellation of the LFP signals into discrete states is also supported by previous work (Hudson et 621 al., 2014) However, reducing the LFP to a single value eliminates much of the information in the 622 original signal. In order to incorporate more of this information, rather than collapsing the LFP 623 signal to a single value, we used the vector of NMF scores for the LFP in each temporal window 624 directly. Each score vector, once multiplied through by the appropriate loading matrix (Methods 625 and **Figure 3**), yields a good approximation of the actual spectrum of the LFP in that time window.

626

627 To test for correlated fluctuations in the spectral features of LFPs at different cortical sites, we 628 applied canonical correlation analysis (CCA) to the pair of score matrices derived from each pair 629 of channels. High canonical correlation indicates a close linear relationship between two sets of 630 variables. The mean of the vector  $\rho$  of canonical correlations between all pairs of canonical 631 variables was calculated to give a measure of overall state similarity that is invariant to invertible 632 linear transformations of each channel's state space. This method of taking the average across p 633 is explained further in Alpert and Peterson (1972). Figure 8A-B shows the CCA similarity measure 634 for all channel pairs from the same representative M1/V1 and bilateral V1 experiments that have 635 been shown previously. All channel pairs from both M1/V1 and bilateral V1 experiments had significantly higher CCA similarities than for shuffled data, after Bonferroni correction for multiple 636 637 comparisons (z-test based on distribution of shuffled data). The summary of CCA similarity across 638 all animals is shown in Figure 8C-D. These results are very similar to those for transition 639 synchrony and normalized MI and show that in both types of recordings, within-region channel 640 pairs had significantly higher CCA similarities than between-region pairs (p = 1e-7 for M1/V1 and p = 1e-7 for bilateral V1, compared to  $10^7$  random permutations of the relevant channels 641 642 (Methods)). Furthermore, as with the previous measures, channel pairs including a channel in L4 643 had lower CCA similarities than pairs in which neither channel was in L4 (p = 0.001, compared to 644 10<sup>7</sup> random permutations of the relevant channels (Methods)). We did not detect a higher mean 645 CCA similarity in left/right V1 channel pairs compared to M1/V1 channel pairs (p = 0.12, percentile 646 bootstrap over channels (Methods)).

647

### 648 Global brain state is low-dimensional, despite weak pairwise interactions.

All results shown up until this point were calculated on pairs of channels for which state assignments were computed independently. What we have shown is that channels within the same cortical region tend to be more similar in their activity patterns and state transition times than channels from different cortical regions. However, close inspection of the results shows that,

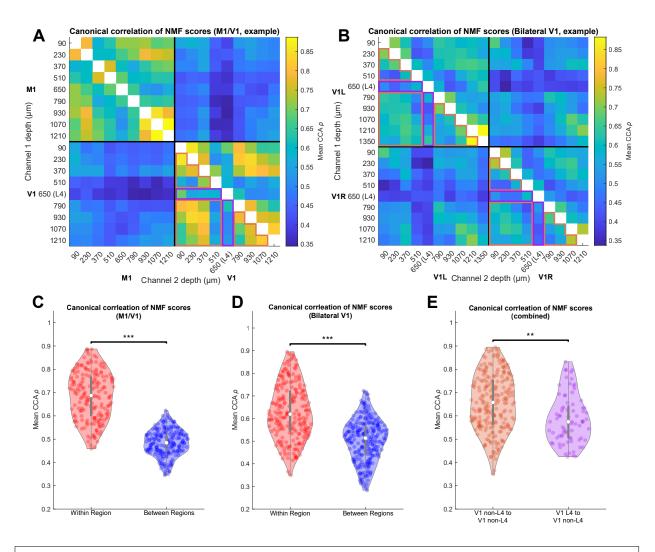


Figure 8. Canonical correlation analysis (CCA) reveals higher correspondence of overall activity between channels in the same anatomical region than between channels in different regions. A-B CCA measure on NMF scores for all channel pairs from representative M1/V1 (A) and bilateral V1 (B) recordings. C-D Channel pairs in which both channels are in the same region (red) have higher NMF score correspondence than those in which the channels are in different regions (blue) for M1/V1 (C, p = 1e-7, permutation test) and bilateral V1 (D, p = 1e-7, permutation test) recordings. E. Channel pairs in which one channel was within L4 and the other was not had lower NMF score correspondence than pairs in which neither channel was in L4 (p = 0.001, permutation test). Data included in these comparisons for the representative experiments are outlined in orange and purple, respectively, to highlight that only data from V1 electrodes were used.

653

even for the channel pairs within the same cortical region, only about one third of the information

- 655 contained within the discrete state sequences is shared between channels (Figure 7C). For
- 656 channel pairs from different cortical regions, the amount of mutual information in state sequences

657 is even lower. This weak coupling between channels could imply that spatially restricted regions 658 of the brain act independently of one another and there is no discernable global state of the brain 659 at any given time. Alternatively, it is possible that this weak coupling between channels, en masse, 660 gives rise to a complex, global state of activity that is differently expressed in the oscillation 661 patterns of spatially restricted regions of cortex. In this final analysis, we sought to directly 662 distinguish these possibilities by characterizing the global brain state. In a key distinction from the 663 previous work, rather than defining the global state on the basis of the concatenated spectra from 664 all recordings, we attempted to identify global macroscopic dynamics from the simplified dynamics 665 observed at each recording site. This was accomplished by first concatenating the NMF score 666 vectors from all simultaneously recorded channels at each timepoint into a single vector that 667 encodes the joint state of all channels. The resulting full matrix of joint states over time was then 668 subjected to principal component analysis (PCA).

669

670 We found that all but one recording required 10 or fewer components to account for 80% of the 671 variance in the concatenated NMF score matrices, which ranged in dimensionality from 91 to 136. 672 The recording that required greater than 10 required 15 components to reach the same threshold. 673 This is far outside the 95% confidence interval of expected cumulative explained variance, 674 computed on Markov-shuffled controls which ignore weak pairwise correlations between 675 fluctuations in different channels (Figure 9A, D). These results demonstrate that widespread 676 weak coupling is sufficient to give rise to a highly correlated global state. **Figures 9B** and **E** show 677 the loadings onto channels and frequencies (mapped back from corresponding NMF loadings) for 678 the top two principal components of a representative M1/V1 and bilateral V1 recording, 679 respectively. These data offer qualitative evidence that the global state is differentially reflected 680 in different regions and layers of the cortex. For example, the loadings of the second principal 681 component (PC2) of the M1/V1 recording in Figure 9B show that, while there is high power in the 682 higher frequencies for the V1 channels, the same is not true in the M1 channels. In contrast,

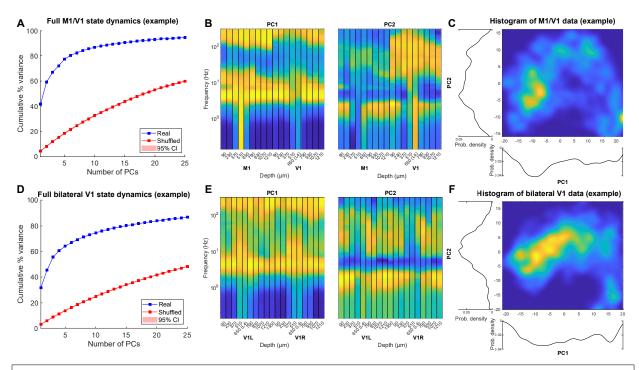


Figure 9. Weakly correlated fluctuations in different cortical sites give rise to highly correlated cortical states. NMF scores from all recorded channels were concatenated into a single state vector (median dimension across recordings = 106) and subjected to PCA. Fraction of total variance as a function of number of PCs is shown in **A** and **D** for M1/V1 and bilateral V1 example recordings respectively (blue). Shuffled surrogates (Methods) were subjected to the same analysis (red). **B** and **E** show loadings of the top 2 principal components, mapped back from each channel's NMF components to frequencies, for the two representative recordings. This projection reveals consistent differences between M1 and V1 (**B**) but is relatively consistent across bilateral V1s (**E**). In both instances, Layer 4 is distinct from supra-and infragranular layers. **C** and **F** show histograms of the data projected onto the top two PCs for the representative M1/V1 (**C**) and bilateral V1 (**F**) recordings. In both instances, the distribution of data is multimodal, suggesting the presence of discrete global cortical states.

683

684 Figure 9E shows that the loadings of PC1 of the bilateral V1 recording onto all channels of both 685 electrodes are fairly uniform, except for in channels near L4 where there is higher power in the 686 lowest frequency bands. Figures 9C and F show histograms of all samples from these representative recordings projected onto the first two principal components. Although more than 687 688 two dimensions would be necessary to fully visualize the landscape of the global dynamics, even 689 in this limited projection, a clustered pattern is visible, similar to previous results (Hudson et al., 690 2014). These data suggest that global brain states comprise regionally distinct oscillation patterns 691 that are weakly coupled with one another. Remarkably, these results show that discrete

- transitions between global cortical states (Ballesteros et al., 2020; Hudson et al., 2014; Patel et
- al., 2020) under a fixed anesthetic concentration arise from the multitude of weakly coupled local
- 694 fluctuations.

#### 696 **Discussion**:

697 Here we set out to determine how abrupt transitions between global thalamocortical states arise 698 at a fixed anesthetic concentration. Using several complementary analysis methods, we 699 demonstrate that correlated fluctuations in the oscillatory behavior observed at different cortical 700 sites are widespread, but that each pairwise interaction is weak. Thus, for instance, the ability to 701 infer the current state of one channel by observing the state of a nearby channel in the cortex is 702 limited. Remarkably, we provide evidence that abrupt transitions between discrete macroscopic 703 cortical activity patterns (Ballesteros et al., 2020; Chander et al., 2014; Hudson et al., 2014; 704 Ishizawa et al., 2016; Lee et al., 2020; Patel et al., 2020) emerge naturally from the multitude of 705 these quasi-independent local fluctuations. We also demonstrate that the strength of the 706 interactions between recording sites depends on the inter-electrode distance and on the cortical 707 layer. Specifically, we find that fluctuations in L4, the thalamic input layer, tend to be less 708 congruent with those in other layers. Altogether, these results argue that abrupt global state 709 transitions are not imposed on the thalamocortical networks by changes in the activity of broadly 710 projecting modulatory arousal systems, but rather are strongly influenced by the local cortico-711 cortical interactions.

712

713 It has been conjectured that structured transitions between discrete states constrain the space of 714 possible brain activity patterns and thereby allow the brain to efficiently recover its normal waking 715 state after a dramatic perturbation (Hudson et al., 2014). The idea that, in order to recover from a 716 perturbation, the space of possible activity states must be constrained by stabilization of a few 717 discrete activity patterns is not specific to recovery from anesthesia per se. For instance, 718 pharmacologically provoked recovery of consciousness in the setting of brain injury is also 719 characterized by abrupt transitions between quasi-stable activity patterns (Victor et al., 2011). 720 Sleep is also well known to consist of discrete activity patterns (e.g., Gervasoni et al., 2004). Thus,

it appears that abrupt state transitions among discrete activity states accompany recovery ofnormal consciousness in a variety of settings.

723

724 It is thus of great interest to determine how such state transitions arise and how they are 725 coordinated across thalamocortical networks. Here, in keeping with previous work (e.g., 726 Gervasoni, 2004; Hudson et al., 2014), we defined the state of each local recording site on the 727 basis of the power spectrum of the LFP. Since we focused on state transitions observed in the 728 anesthetized brain, most fluctuations occurred in the slow oscillations (< 1 Hz) (Steriade et al., 729 1993b), delta oscillations (1-4 Hz), and the spindle range of 8-14 Hz (Purpura, 1968). Multiple 730 distinct neurophysiological mechanisms contribute to the generation and coordination of the 731 various brain oscillations observed in the anesthetized brain. Slow oscillations, for instance are 732 thought to be primarily generated through local synaptic mechanisms in the cortex (Sanchez-733 Vives and McCormick, 2000; Steriade et al., 1993b). Thalamocortical and thalamic reticular 734 neurons reflect these slow oscillations and are phase locked to them (Steriade et al., 1993b). 735 However, the fact that slow oscillations are abolished in the thalamus of decorticated animals 736 (Timofeev and Steriade, 1996) but are observed in the cortex of athalamic animals (Steriade et 737 al., 1993b) strongly argues for the cortical origin of slow oscillations. Cortico-cortical interactions 738 are thought to underlie not just the generation of slow waves, but also the synchronization of these 739 waves across the cortex. Pharmacologic and surgical lesions of intra-cortical connections disrupt 740 the synchrony of slow waves (Amzica and Steriade, 1995).

741

The observation that slow oscillations are coordinated primarily through cortico-cortical interactions is consistent with our results. Many of the state transitions under isoflurane involve fluctuations in the power of slow oscillations. Using three distinct analysis methods, we consistently find that state fluctuations in L4 are relatively dissimilar to those observed in the infraand supragranular layers. L4 neurons are most directly affected by spatially localized inputs from

the thalamus, whereas supra- and infra-granular neurons are primarily driven by cortico-cortical connections and matrix projections from the thalamus (Jones, 2001). While anesthetics suppress both core and matrix thalamocortical inputs, their dominant effect is specifically suppressing cortico-cortical connectivity (Raz et al., 2014). It is thus likely that the local nature of state transitions in the slow oscillation range is a consequence of both weakened thalamocortical and cortico-cortical interactions in the anesthetized brain.

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754 Transitions between slow (< 4Hz) and faster EEG oscillations, occasionally observed even in the 755 anesthetized brain (e.g., Figure 2) are thought to arise as a result of the interaction of the thalamo-756 cortical networks with neuromodulatory projections from cholinergic neurons in the brainstem and 757 basal forebrain (Steriade, 2004). Noradrenergic neurons (Vazev and Aston-Jones, 2014) and 758 other brain stem and basal forebrain nuclei also contribute to the modulation of the oscillations 759 exhibited by the thalamocortical networks (Jones, 2003). Activity within the various arousal 760 promoting nuclei is coordinated by a group of medullary neurons, activation of which can trigger 761 prompt awakening from deep states of anesthesia (Gao et al., 2019). In the anesthetized brain, 762 fluctuations in the firing rate of these medullary neurons co-varies with the fluctuations in the 763 spectral characteristics of the cortical LFP (Gao et al., 2019). Thus, it is possible that the 764 spontaneous fluctuations of the LFP characteristics between the slower and faster oscillations are 765 in part mediated by fluctuations in the activity of the nuclei that modulate the thalamocortical 766 networks. However, most arousal nuclei have broad projections to the thalamus and the cortex (Jones, 2003). Thus, if the fluctuations in the state of the LFP were entirely driven by the 767 768 fluctuations in the activity of the modulatory projections, one would expect that the state of the 769 LFP would fluctuate coherently across the cortex. Instead, we observe that fluctuations in state 770 of the LFP are only weakly coupled between different cortical sites. This implies that the influence of the modulatory nuclei on the power of specific cortical oscillations, within the physiological 771 772 range, is not absolute. Rather, activation of the modulatory systems likely biases the cortex

towards a particular oscillatory state. The overall pattern of activity at each cortical site, however,
is strongly influenced by interactions within the thalamocortical networks.

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776 The experiments performed here cannot directly address the cellular and synaptic mechanisms 777 that give rise to local state transitions and their coordination across the cortex. They do, however, 778 offer clear insights into network mechanisms of global state transitions. Here, rather than 779 attempting to simplify the dynamics of the global signals directly (Hudson et al., 2014), we 780 embedded the dynamics of the local signals into a low-dimensional space. This analysis revealed 781 only weak interactions between local signals. Remarkably, assembling just the low-dimensional 782 projections of the local signals into a state vector recapitulated the low-dimensional dynamics and 783 discrete global cortical states. Thus, we show that the global states and abrupt transitions 784 between them arise because of weak coupling between local state fluctuations.

785

786 We are not the first to note that weak coupling among local fluctuations can give rise to coherent 787 macroscopic states. In the retina, weak correlations in spike timing co-exist with a conspicuously 788 high probability of certain large ensembles of neurons firing in synchrony (Schneidman et al., 789 2006). It may seem that a network with weakly correlated nodes can be well approximated by a 790 collection of completely independent nodes, but this is not the case. Weakly coupled elements 791 can yield highly correlated macroscopic states if the weak interactions are prevalent enough 792 throughout the network. Indeed, we find that while the correlations between different cortical sites 793 were weak, they were present and statistically significant for most electrode pairs.

794

The emergence of highly correlated global states from weak pairwise interactions has been investigated extensively in statistical mechanics using Ising models. It has been shown that an Ising model is mathematically equivalent to a maximum entropy models of the statistics of neural firing that are constrained only by the experimentally observed firing probabilities of individual

799 neurons and their pairwise correlations (Schneidman et al., 2006; Tkačik et al., 2006). The 800 maximum entropy approach has proved successful in diverse systems (Ohiorhenuan et al., 2010; 801 Tang et al., 2008; Tkačik et al., 2014; Yu et al., 2008). Although Ising models have traditionally 802 been applied to binary state spaces, such as the presence or absence of an action potential within 803 a small time window, the maximum entropy approach can be generalized to continuous variables 804 (Bialek et al., 2012), such as local fields. In this work, we did not explicitly attempt to construct a 805 maximum entropy model of local field fluctuations, as we are recording only a tiny fraction of all 806 cortical signals. Future work may sample of local field fluctuations more densely to determine 807 whether an Ising-type model suffices to explain the fluctuations of the global state of the brain 808 under anesthesia, or whether other mechanisms in addition to pairwise interactions are needed 809 (Ohiorhenuan et al., 2010; Tang et al., 2008). Regardless of the specific details of such a model, 810 however, we directly demonstrate that widespread weak correlations in local field fluctuations give 811 rise to coherent global cortical states. This conclusion is strongly supported by the observations 812 that locally defined cortical states yield highly correlated global behavior despite weak pairwise 813 interactions, whereas the shuffled controls do not.

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815 There are multiple parallels between our characterization of state transitions in the anesthetized 816 brain and those observed during slow wave sleep (NREM). While sleep and anesthesia are clearly 817 distinct phenomena, the neurophysiological mechanisms that give rise to oscillations in the 818 thalamocortical circuitry under anesthesia and during natural sleep share some essential 819 similarities (Steriade et al., 1993b; Steriade and Amzica, 1998). Many diverse anesthetics 820 promote activity in the sleep active subcortical nuclei and suppress activity in the wake active 821 ones (Jiang-Xie et al., 2019; Moore et al., 2012; Nelson et al., 2002; Zhang et al., 2015). 822 Furthermore, both sleep and anesthesia consist of several discrete states, each characterized by 823 a distinct pattern of oscillations in the cortex and thalamus (Saper et al., 2010). Based on the 824 original recordings at the microscopic level of single isolated neurons or, alternatively, on the 825 macroscopic level using EEG, it has long been hypothesized that sleep stages are brain-wide 826 phenomena and that the neurophysiological mechanisms that give rise to sleep stage switching 827 specifically prevent multiple sleep stages or sleep and wakefulness from coexisting at the same 828 time in different brain regions (Lu et al., 2006; Saper et al., 2010). Interestingly, at the mesoscopic 829 level of neuronal populations and local fields, sleep state transitions, much like in this work, can 830 be local (Nir et al., 2011; Poulet and Petersen, 2008; Vyazovskiy et al., 2011). Furthermore, it 831 has been suggested that antecedent neuronal activity driven by a specific task can increase the 832 propensity of a population of cortical neurons to exhibit local sleep-like slow oscillations (Huber et 833 al., 2004), implying that transitions between different oscillatory modes are strongly influenced by 834 local synaptic interactions. The degree of synchrony between cortical locations across naturally 835 observed state transitions, such as those between different sleep stages or between sleep and 836 wake, has not been directly quantified in a systematic fashion. Because sleep is strongly 837 influenced by both homeostatic and circadian influences, it will be challenging to disentangle 838 these global influences from the local interactions between different sites in the cortex. However, 839 analysis of cortical state transitions in the brain anesthetized with a fixed anesthetic concentration 840 is free from these complications. This analysis shows that the apparently global coordinated shifts 841 in cortical activity arise naturally out of weakly interacting local state switches.

# 843 Data and Software Availability

- 844 Datasets are available upon reasonable request. Code for time-frequency analysis of LFP and
- 845 interaction measure calculation is publicly available at <u>https://github.com/ProektLab/spec-state-</u>
- 846 trans and other public repositories linked from the README.
- 847

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- 854

### 855 Competing Interests

- 856 The authors declare that they have no competing interests.
- 857
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