

Supplementary Materials for

Conserved brain-wide emergence of emotional response from sensory experience in humans and mice

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Supplemental Note 1 Figs. S1 to S38 Tables S1 to S3 References

Other Supplementary Material for this manuscript includes the following:

MDAR Reproducibility Checklist

Supplemental Note 1

In the following supplemental note, we discuss correspondences between human and mouse brain regions that are described in the study. Notably, the study's primary claims regarding the nature of neural dynamics underlying emotional responses do not rest upon direct anatomical mappings between species. For example, the computational technique of extracting different coding dimensions is agnostic to the exact anatomical distribution of relevant neural dynamics. This neural activity dimension approach allows for conclusions to be made about neural dynamics which are present at the population-level across recorded brain regions in both species, without necessarily relying on the presence of a particular recorded brain region.

Cross-species correspondence of eyepuff assay

The core features of the eyepuff task are conserved between mouse and human. However, we also note the many ways in which the experience of human participants in the task may be quite different from the experience of mice. For example, humans certainly have conscious awareness of task context, and subjective experience of the task itself, whereas these aspects are less clear in mice. These potential distinctions underscore the value of developing a cross-species translational approach to allow direct assessment of the human experience while also leveraging the powerful tools available with mice. We do not intend to claim that the mouse experience is identical to the human experience, but rather that it has many similar features, especially on the fast timescales surrounding individual puffs, which suggest that some (if not all) aspects are conserved. Using mice allows us to investigate additional details that we cannot access in humans, but the combination with measurements from humans is complementary and allows us to further investigate the neural basis of the human experience.

Cross-species correspondence of specific brain regions

We focus below on aspects of cross-species conservation among specific brain regions, to elaborate upon certain of the regions where eyepuff-driven emotional responses may emerge (see regions with significant changes in Fig. 3C). We note that there are of course substantial differences between mouse and human brain anatomy and function, including the lissencephalic nature of the mouse brain, the differing scale of brain sizes, and the absence of circuits dedicated to language use. Especially in non-sensorimotor cortical regions (including but not limited to orbital frontal cortex, cingulate cortex), the mappings between rodent and human brain regions are unlikely to be one-to-one.

• Insula: This region has been shown to be involved in interoception across humans and mice, and in the assignment of valence to internal and external, current and future stimuli (Gogolla 2017, Craig et al. 2011). Across species, insular cortex is comprised cytoarchitecturally of granular, dysgranular and agranular subdivisions, and receives functional input from sensory, interoceptive, and limbic regions (Gehrlach et al. 2020). Relevant to our study's focus on the emergence of affect from sensation, humans and rodents with insula lesions have been shown to sometimes have "pain asymbolia," in which participants are aware of pain, but do not seem to attribute negative emotional valence to the experience (Berthier et al. 1988). Denoted "INS" in human Fig 3; "AIv" in mouse Fig. 4.

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• Orbitofrontal cortex: Studies have found the relevance of this region to reward valuation, emotional processing, and decision making in both humans and mice (Wallis 2011, Rudebeck et al. 2022). For example, causal experiments in mice and humans have shown the importance of orbital frontal cortex to reward devaluation tasks, in which participants must make choices based on the perceived value of choice outcomes (Lichtenberg et al. 2021, Howard et al. 2020). In humans, alterations in orbitofrontal cortex activation and connectivity have been linked to altered emotional processing, depression and other psychiatric conditions (Rolls et al. 2020). Notably, orbitofrontal cortex is among the supramodal (non-sensorimotor) regions likely present in early mammalian

ancestors of both mice and humans (Kaas 2011, Beauchamp et al. 2022). Denoted "ORB" in human Fig 3; "ORBm", "ORBl", "ORBvl" in mouse Fig 4.

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- Cingulate cortex: In the latest consensus definitions, human cingulate cortex can be roughly subdivided into anterior cingulate cortex (also in literature as ventral ACC), mid-cingulate cortex (also in literature as dorsal ACC), and posterior cingulate cortex (van Heukelum 2020). Across mice and humans, anterior cingulate cortex seems to be chiefly involved in processing affective and autonomic signals, exhibiting stronger structural connectivity to autonomic brainstem regions, amygdala, and frontal cortex (Etkin 2011, van Hout 2024). In contrast, the more caudal mid-cingulate cortex seems to be involved in aspects of cognitive control, and shows stronger structural connectivity to regions like the parietal and motor cortex (van Heukelum 2020). Posterior cingulate cortex and retrosplenial cortex have been shown to be involved in self-referential thought (Alexander 2023). Here, we adopt the updated ACC and MCC rostral-caudal sub-division nomenclature in both mouse and human, which has been shown to allow for more effective functional and anatomical homologies compared to the mouse subdivisions of Cg1, Cg2, infralimbic, and prelimbic cortices. Notably, cingulate cortex is among the supramodal (non-sensorimotor) likely present in early mammalian ancestors of both mice and humans (Kaas 2011, Beauchamp 2022). Anterior cingulate: "ACC" in human Fig. 3; anterior subdivisions of "ACAd", "ACAv" [area A24], "ILA" (infralimbic), "PL" (prelimbic) in mouse Fig 4. Midcingulate: "MCC" in human Fig. 3; posterior subdivisions of "ACAd", "ACAv" [area A24']. Posterior cingulate: "PMC" in human Fig. 3; "RSPv" in mouse Fig. 4.

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- **Supramarginal gyrus**: In humans, this sub-region of the posterior parietal cortex has been shown to be involved in functions as diverse as complex tool use, verbal working memory, and language processing (Oberhuber et al. 2016, Deschamps et al. 2014, Wandelt et al. 2022). In mice, the posterior parietal cortex has been shown to be involved in navigation, decision-making, and sensory processing (Lyamzin et al. 2019). Denoted "SMG" in human Fig. 3; not recorded from in mouse. *References:*

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- Motor cortex: Primary motor cortex demonstrates functional conservation across species, and is essential for the precise control of movement including for the orbicularis oculi muscles essential for motor control of the eyelid in humans and mice (Paradiso et al. 2005, Pronichev et al. 1998). Notably, mice have large primary motor cortices, dominating much of their frontal cortex, whereas the human motor cortex occupies a smaller fraction of relative area, located anterior to the central sulcus (Ebbesen et al. 2017). Relative to all other mouse and human cortical regions, sensorimotor cortex exhibits the highest overall correlation in the expression profile of homologous genes between species (Beauchamp et al. 2022). This is reflected in the high levels of conservation of primary motor cortex cell type hierarchies across humans and mice with notable differences, including the fact that the ratio of glutamatergic to GABAergic neurons in humans and mice are 2:1 and 5:1, respectively (Bakken et al. 2021). Denoted "MOT" in human Fig. 3; "MOp" and "MOs" in mouse Fig. 4.

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- **Hippocampus**: This region plays a conserved role in memory function across species, exhibiting structural and functional homologies (Manns et al. 2006). Pertaining to alterations in hippocampal eyepuff responses, hippocampal activity has been shown to be modulated by the emotional valence of stimuli in mice and humans (Meyer et al. 2019, Qasim et al. 2023). Concordant with the expansion of higher order association cortices in primates, the human hippocampus appears more functionally coupled (via fMRI correlations) to association cortices, whereas the rodent hippocampus appears more directly coupled to sensory cortices (Bergmann et al. 2016). Denoted "HIPP" in human Fig. 3; "DG-mo", "CA", "CA3" in mouse Fig. 4. *References*:
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- **Basal ganglia**: These are a cluster of subcortical nuclei involved in motor control and reward learning, conserved in many respects across humans and mice (Cox et al. 2019, Hardman et al. 2002). Based on transcriptomic homologies, mouse and human striatal regions show strong conservation, with mappings between mouse caudoputamen and human caudate and putamen (Beauchamp et al. 2022). Based on connectomic homologies from functional MRI, mice seem to lack certain subdivisions of caudate most strongly connected to frontal cortex (Balsters et al. 2020). Denoted "BG" in human Fig 3; "CP" (caudoputamen), "ACB" (nucleus accumbens), "LSr" (lateral septal nucleus) in mouse Fig 4. *References*:
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- Thalamus: Both humans and mice exhibit multi-nucleated thalamic structures, functioning in part to relay information from sensory to cortex, cortex to cortex, and from subcortex to cortex (Schmitt et al. 2017). Studies have demonstrated homologies between thalamo-cortical connectivity in mice and humans of primary sensory nuclei, including medial geniculate nucleus (Keifer et al. 2015). Notably, the ventral posteromedial nucleus is among the principal sensory nuclei in humans and mice (Morel et al. 1997). Given the size of the human intracranial electrodes and volume of tissue recorded relative to the size of thalamic nuclei, it is challenging to assign particular recordings to different human thalamic nuclei. Denoted "THAL ANT" and "THAL POS" in human Fig. 3; "MD" (mediodorsal nucleus), "VPM" (ventral posteromedial nucleus), "PO" (posterior complex), "LP" (lateral posterior), "TH" (other thalamic nuclei), "PoT" (posterior triangular nucleus) in mouse Fig. 4. References:
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- Notably, there are several regions we were able to record from in one species but not the other. In mice, we were uniquely able to record from brainstem and midbrain regions. In human participants, this category includes supramarginal gyrus and amygdala.

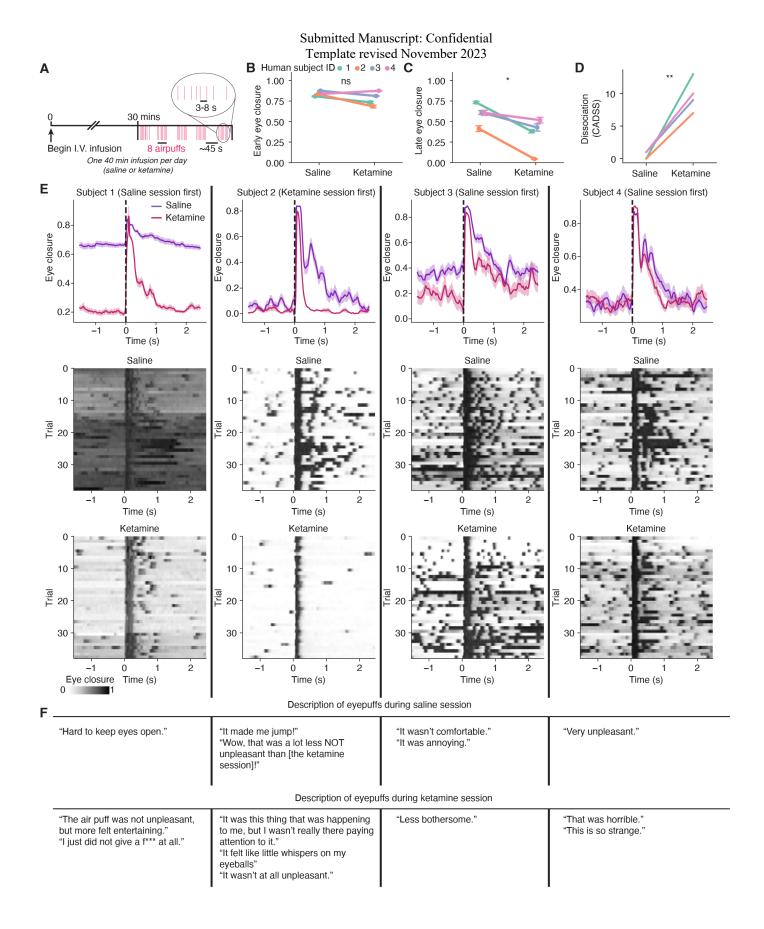


Fig. S1. Additional details about eyepuff assay in humans. (A) Eyepuff protocol during infusion of ketamine or saline. During each session, a total of 40 air puffs are directed at the subject's left cornea, with an inter-puff interval that varies

between 3 to 8 s to prevent subjects from predicting the timing of each air puff. To reduce stress for the subject, there is a 35 s interval after every eighth puff. To allow for direct comparison, the same inter-puff interval timing sequence was used in saline and infusion sessions and across subjects. For analysis, we do not include the first two trials in order to exclude an initial response that can differ from subsequent trials. (B) Early eye closure (during 0.1 to 0.2 s after puff onset) does not change between saline and ketamine sessions (n=38 trials per subject, median \pm 95% confidence interval). (C) Late eye closure (during 0.3 to 0.8 s after puff onset) decreases with ketamine relative to saline (n=38 trials per subject, median \pm 95% confidence interval). (D) Dissociation increases during ketamine, as measured using the Clinician-Administered Dissociative States Scale (CADSS). (E) Puff-triggered eye closure for each subject. top: summary, mean \pm s.e.m across trials. middle: saline individual puffs, bottom: ketamine individual puffs. (F) Self-reported descriptions of subjective experience of eyepuffs during saline or ketamine. ns, P-value \geq 0.05. *, P-value < 0.05. **, P-value < 0.01. ****, P-value < 0.001. See Supplementary table 3 for information on statistical analyses and sample sizes.

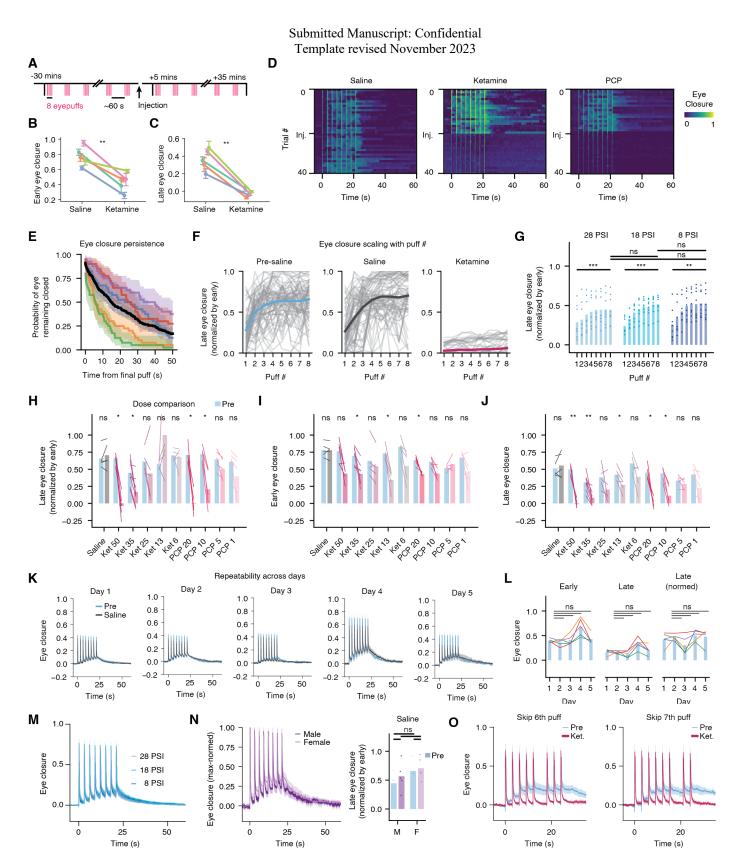


Fig. S2. Additional characterization of eyepuff assay in mice. (A) Eyepuff protocol with bolus drug infusion. Each trial consists of 8 air puffs with an inter-puff interval of 3 s. The inter-trial interval ranges from 50 to 70 s. Twenty trials are presented before infusion, and twenty trials after infusion. (B) Early eye closure (during 0.1 to 0.2 s after puff onset) decreases with ketamine but remains present (n=18 trials per subject, median \pm 95% confidence interval). (C) Late eye closure (during 0.3 to 0.8 s after puff onset) is abolished by ketamine (n=18 trials per subject, median \pm 95% confidence interval). (D) Eye closure on individual trials (each consisting of eight puffs) with saline, ketamine, or PCP infusion at trial

20. Saline elicits no change in eye closure, which consists of a fast eye blink in response to each puff and an increasing extended eye closure across puffs in a trial. In contrast, ketamine and PCP maintain the presence of the fast eye blink, but the extended eye closure is abolished. (E) Persistence of eye closure after final puff of each trial (Kaplan Meier survival time =15.16 s, n=100 trials from N=5 mice, each colored line is an individual mouse, median ± 95% confidence interval). (F) Scaling of late eye closure with number of puffs in a sequence, colored lines are mean across sequences, gray lines are individual sequences. The increase in late closure across the sequence of puffs has some variability on single trials, and on average scales upwards to a saturating level of closure. (G) Late (normed by early) eye closure scales with puff number but not puff intensity, comparing air puffs of different pressure (PSI, pounds per square inch). (H) Dose comparison of ketamine and PCP, specified in mg/kg for late normalized by early, (I) early, and (J) late eye closure. (K) Eye closure preceding and after saline infusion for the same mice across five days. While there can be some variability in the overall magnitude of eye closure, the early and late eye closures scale together (n=5 mice, mean \pm s.e.m.). (L) Early, late, and late normalized by early eye closure on each day does not significantly differ from on the first day. (M) Eye closure during puff sequence with different puff intensities (n=8 mice, mean ± s.e.m.). (N) Male and female mice exhibit similar eye closure, preceding and after saline infusion (n=5 mice, mean \pm s.e.m.). (O) Mice do not predict occurrence of a puff, and do not reflexively close their eye if a puff is skipped (n=5 mice, mean \pm s.e.m.), while late eye closure remains persistently elevated during skipped puff. ns, P-value ≥ 0.05. *, P-value < 0.05. **, P-value < 0.01. ***, P-value < 0.001. ****, P-value < 0.0001. See Supplementary table 3 for information on statistical analyses and sample sizes.

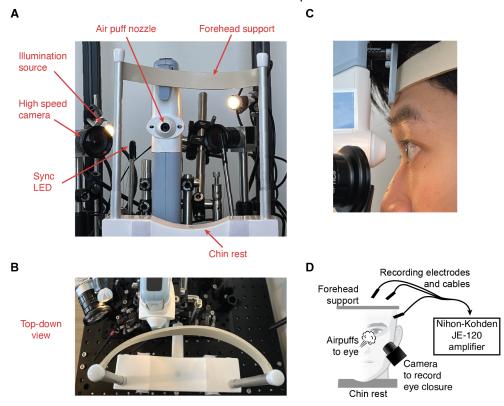


Fig. S3. Photos of the human eyepuff system. (A) Photo of eyepuff device, from the perspective of the subject before placement of head against the chin rest and forehead support. (B) Top-down view of the eyepuff device. (C) Example alignment of air puff nozzle towards subject's left eye, with the camera visible in the foreground of the image. (D) Schematic of iEEG recording during eyepuff assay.

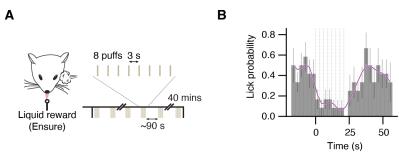


Fig. S4. Generalization of eyepuff-evoked aversive behavioral state to reduced consumption of liquid reward. (A) Schematic of protocol for measuring reward consumption during the eyepuff assay. Throughout the entire session, mice were free to collect a small amount of liquid reward (Ensure, a sweet caloric drink) from a spout within reach of their tongue, while the standard eyepuff assay was administered. A small amount of liquid was dispensed automatically every ~4.75 s (with exponentially distributed random time of [4.5, 5], mean 4.75). An additional small amount of water was dispensed if the spout was licked, with refractory period of 0.5 s. (B) Binned (3s) lick probability during eyepuff sequence. Purple line, smoothed probability estimate for ease of visualization. Lick probability decreased persistently during the air puff sequences and slowly recovered after the final air puff of the sequence, demonstrating that persistent behavioral responses to eyepuffs generalizes beyond eye closure, and providing additional evidence that eyepuffs induce a negative affective state.

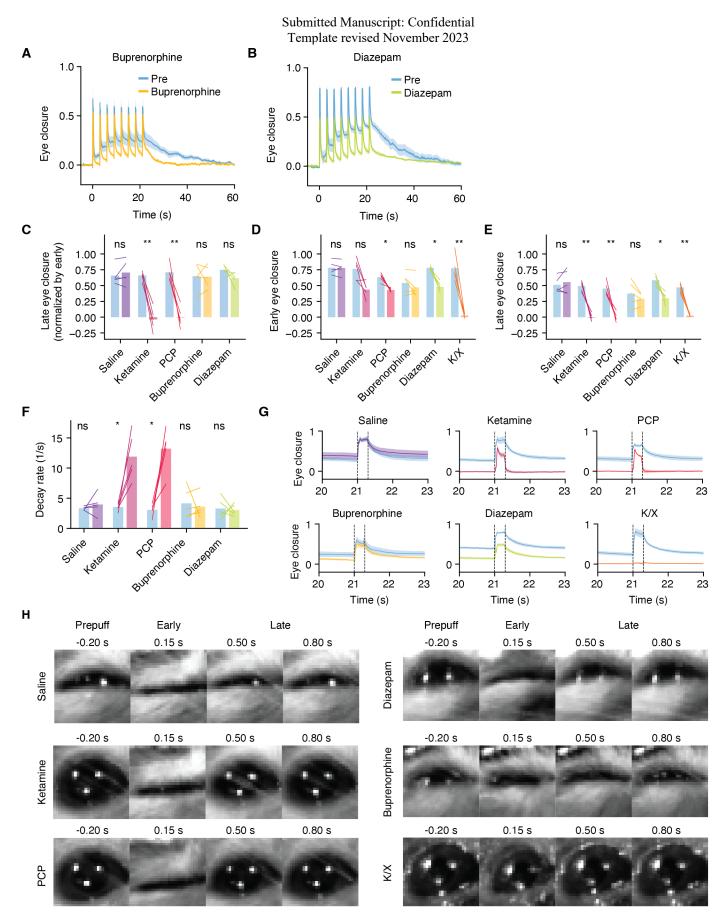


Fig. S5. Comparison of different drugs on eyepuff assay in mice. (A) Eye closure before and after injection of buprenorphine (an opioid analgesic, 2 mg/kg), and (B) diazepam (a benzodiazapene, 2 mg/kg). (A-B, n=5 mice, mean ± s.e.m). (C) Only ketamine (50 mg/kg) and PCP (20 mg/kg) decrease late normalized by early eye closure. (D) Change in

early eye closure and (E) late eye closure. Diazepam and K/X both yield significant decreases in early and late eye closure, as a consequence of overall eye closure decreasing but without late eye closure specifically exhibiting a greater reduction. (F) Decay rate of an exponential fit to eye closure during 0.2 to 2 s after the onset of the final (eighth) puff in each trial. (G) Eye closure during the final puff in each trial (which occurs 21 s after the onset of the first puff in a trial). Unlike with Saline, Buprenorphine, and Diazepam, with Ketamine and PCP there is no accumulation of extended eye closure across the sequence of puffs. (n=5 mice, mean \pm s.e.m). (H) Example images of eye preceding, during, and after the final puff of a trial, after infusion of indicated drug. With all drugs except for K/X, there is an early eye closure, but with Ketamine and PCP the eye quickly opens wide. ns, P-value \geq 0.05. *, P-value < 0.05. **, P-value < 0.001. ****, P-value < 0.001. ****, P-value < 0.001. See Supplementary table 3 for information on statistical analyses and sample sizes.

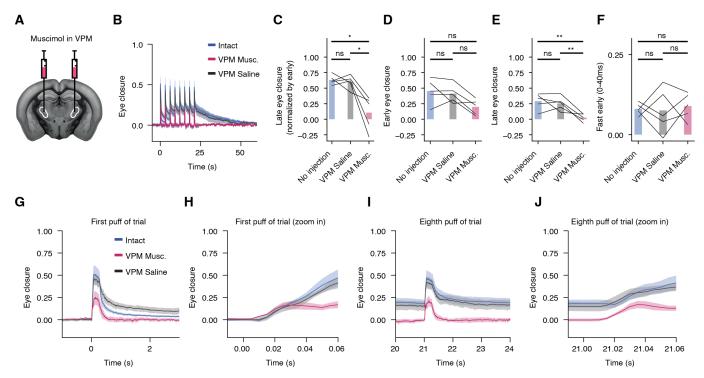


Fig. S6. Inhibition of mouse VPM thalamus with muscimol reduces affective eyepuff response. (A) Location of bilateral muscimol injection (coordinates: \pm 1.6 ML, -1.9 AP, -3.6 DV). (B) Eye closure during puff sequence in each condition. With muscimol, there is early eye closure after each puff, but no accumulating late eye closure. (C) Late (normalized by early) eye closure. (D) Early eye closure. (E) Late eye closure. (F) Fast early eye closure (within first 40 ms after the first puff of each trial). (G) Eye closure during first puff and (H) zoomed in. While the early eye closure does not reach the overall same magnitude, the fast component of the early eye closure is nearly identical to the intact and saline settings. (I) Eye closure during final puff (which occurs at 21 s after the onset of the first puff), and (J) zoomed in. All traces are mean \pm s.e.m., n=5 mice. ns, P-value \geq 0.05. *, P-value \leq 0.05. **, P-value \leq 0.01. ***, P-value \leq 0.001. ****, P-value \leq 0.0001. See Supplementary table 3 for information on statistical analyses and sample sizes.

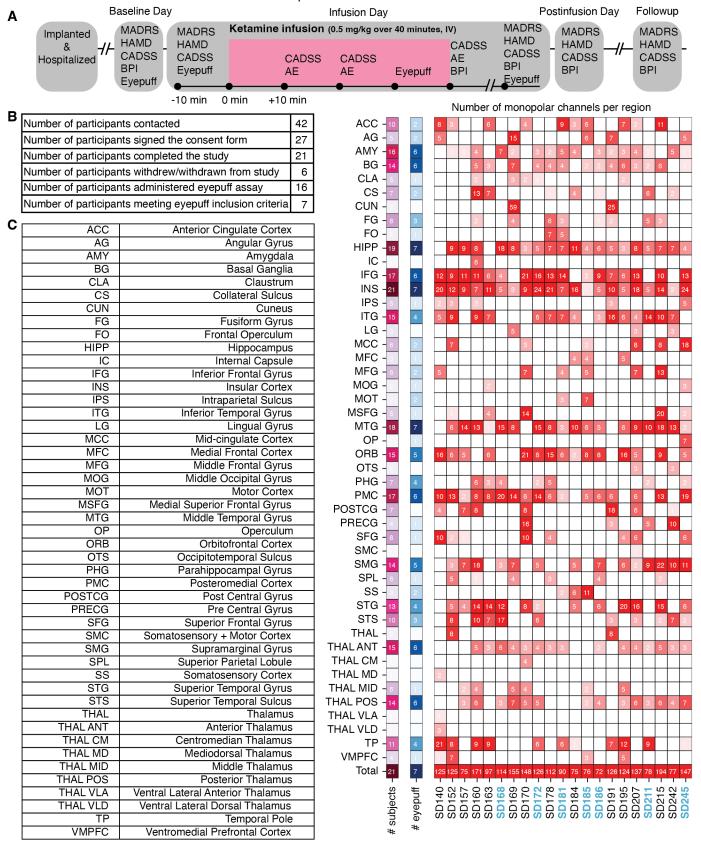


Fig. S7. Human iEEG study details. (A) Full study protocol, including psychiatric questionnaires, ketamine infusion, and eyepuff assay. (B) Participant study completion. See Methods for explanation of inclusion criteria. Note that we excluded from eyepuff analysis the 2 participants who received bolus infusions, because dosing and timing were not comparable to the other participants, and we thus considered 14 participants for the eyepuff assay in the main text. (C) Region acronym

legend. (D) Anatomical distribution for included subjects. From left: number of subjects with at least one channel in a given region; number of subjects meeting inclusion criteria for eyepuff analysis that have at least one channel in a given region; and number of monopolar channels per region for each subject. Eyepuff subjects meeting inclusion criteria are specified in blue.

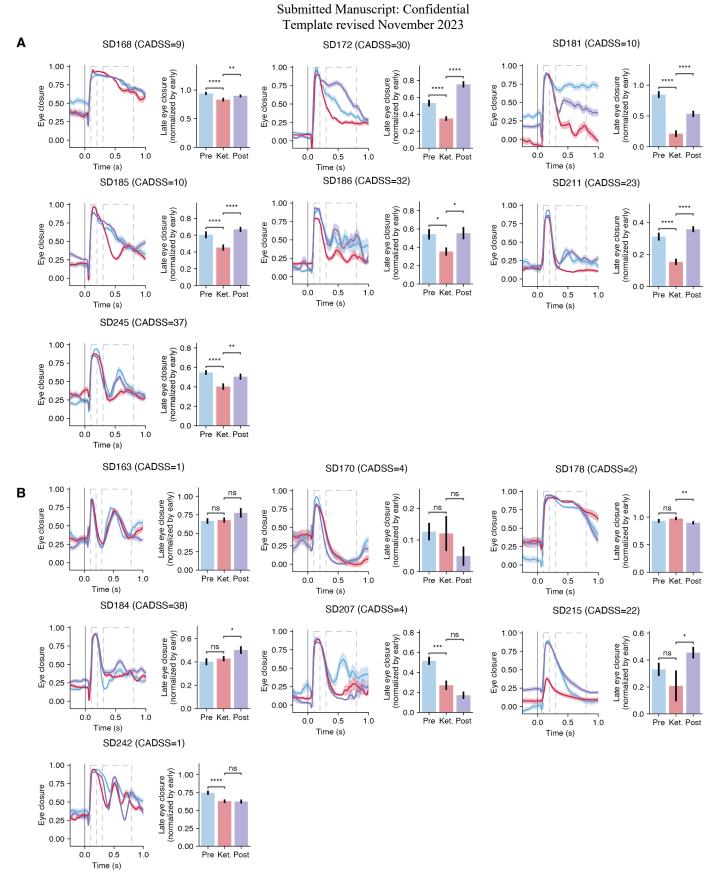


Fig. S8. Additional details for human eyepuff behavior. (A) Eye closure for each condition for subjects that meet the behavioral inclusion criteria for neural analysis of eyepuff-triggered dynamics. Left: eye closure timecourse (dashed boxes: left, window for quantifying early eye closure, right, window for quantifying late eye closure). Right: summary of late eye closure normalized by early eye closure. Dissociation symptoms during infusion (CADSS score) indicated for each subject.

(B) Eye closure of subjects not meeting the inclusion criteria. Inclusion criteria was defined as a subject exhibiting significant change in late eye closure normalized by early eye closure: preinfusion vs. ketamine, and ketamine vs. postinfusion (Methods). Timecourse plots, mean \pm s.e.m., n=12 to 39 trials depending on subject. Vertical dashed lines represent, starting from left: onset of early window, offset of early window, onset of late window, offset of late window. ns, P-value ≥ 0.05 . *, P-value < 0.05. **, P-value < 0.01. ***, P-value < 0.001. ***, P-value < 0.0001. See Supplementary table 3 for information on statistical analyses and sample sizes, and Methods for additional description of inclusion criteria.

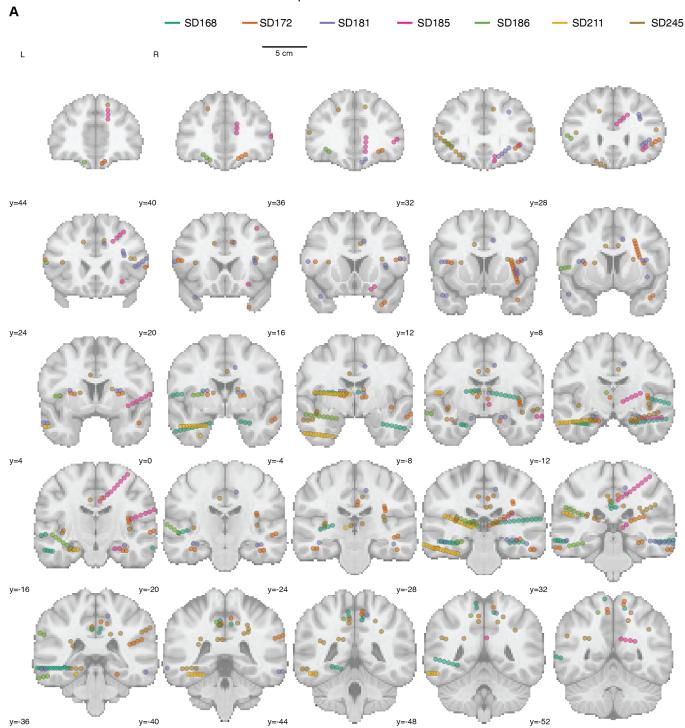


Fig. S9. Human recording sites for subjects that meet eyepuff assay behavioral inclusion criteria. (A) Recording sites transformed into MNI coordinates and plotted on coronal slices of MNI152 template.

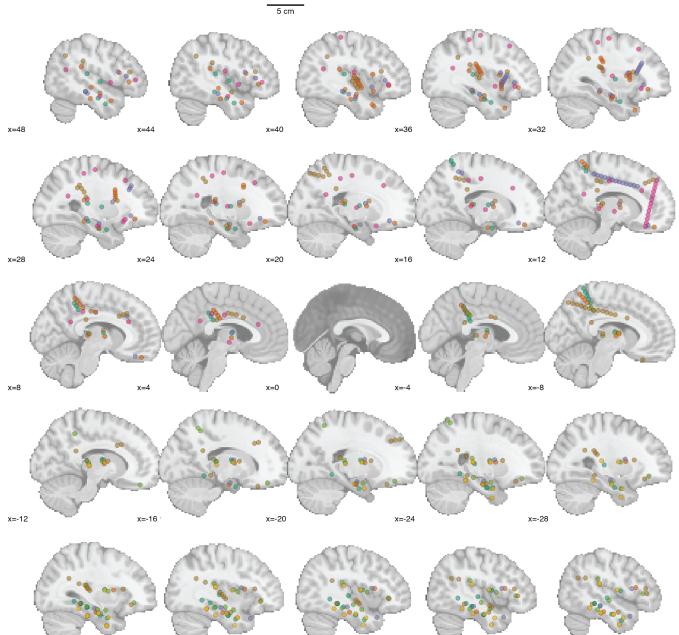


Fig. S10. Human recording sites for subjects that meet eyepuff assay behavioral inclusion criteria. (A) Recording sites transformed into MNI coordinates and plotted on sagittal slices of MNI152 template.

x=-44

x=-40

x=-32

x=-36

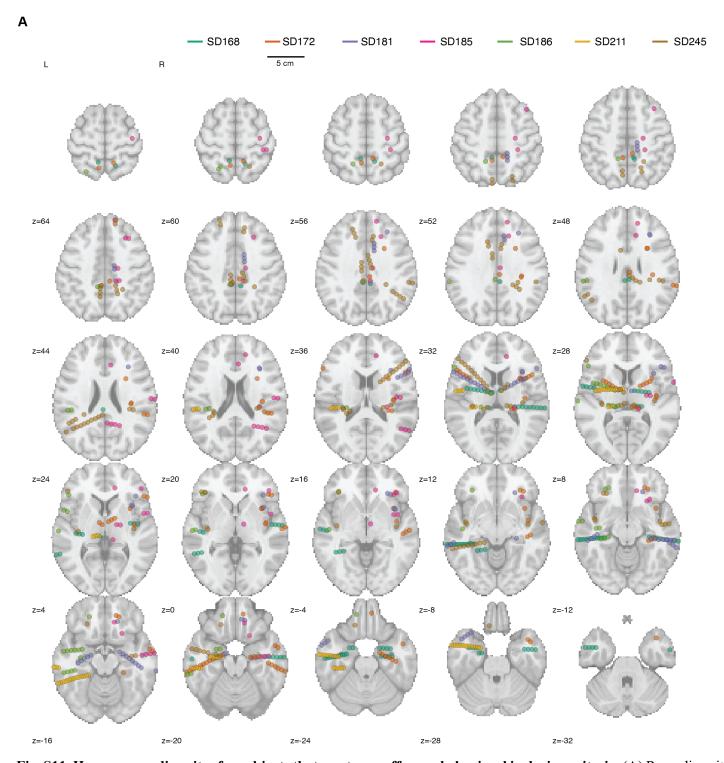


Fig. S11. Human recording sites for subjects that meet eyepuff assay behavioral inclusion criteria. (A) Recording sites transformed into MNI coordinates and plotted on horizontal slices of MNI152 template.

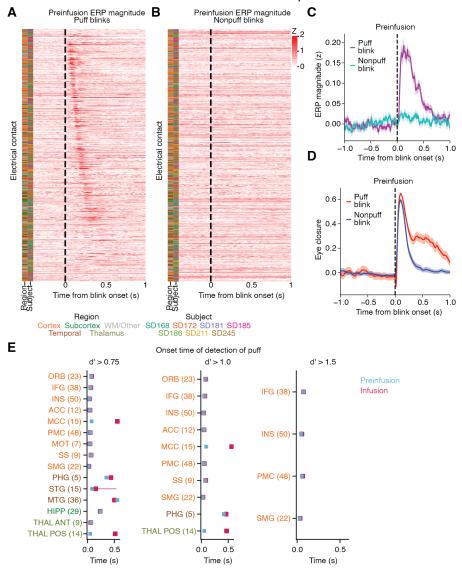


Fig. S12. Human puff-triggered electrical activity is different than activity during non-puff-triggered eyeblinks. (A) Preinfusion puff-triggered event related potential (ERP) for all channels (n=773 channels from N=7 subjects), sorted using peak time of a separate validation set of trials. (B) Same ordering, ERP of non-puff-triggered eyeblinks. (C) Magnitude of ERP is much larger during puff-triggered eyeblinks (n=773 total channels from 7 subjects, mean \pm s.e.m., ERP for each channel computed from 6 to 9 blink events). (D) Puff-triggered and non-puff-triggered eyeblinks reach the same magnitude of eye closure (n=52 trials from 7 subjects, mean \pm s.e.m.). (E) Onset time of detection (d' across trials) of puff by the average puff-triggered LFP trace across all channels in a region, for different d' thresholds. Regions across the brain, and especially cortex, exhibit fast detectability of the puff during both preinfusion and infusion. Error bars are bootstrap 95% confidence intervals. See Supplementary table 3 for information on statistical analyses and sample sizes.

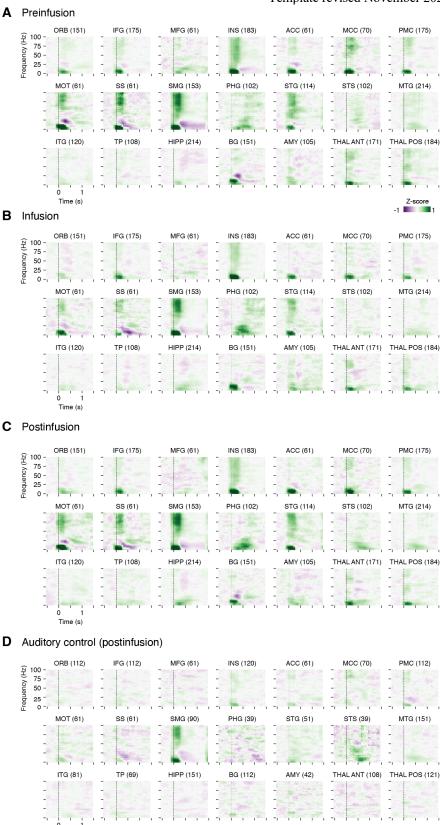


Fig. S13. Human puff-triggered spectrograms by brain region. (A) Preinfusion, (B) Infusion, (C) Postinfusion, (D) Auditory control (following Postinfusion assay, the eyepuff assay protocol is run but with the air puff nozzle pointed away from the subject). For each subject, a regional trace was computed as the average across channels in that region; all trials for a region were then stacked across subjects. Total number of trials indicated next to each region; different regions were

sampled in different subsets of subjects, so the number of trials differed across regions. Similar patterns of puff-triggered dynamics are exhibited by regions across the brain, suggesting there may be a low dimensional representation of the puff-triggered response. Dashed vertical line, air puff onset.

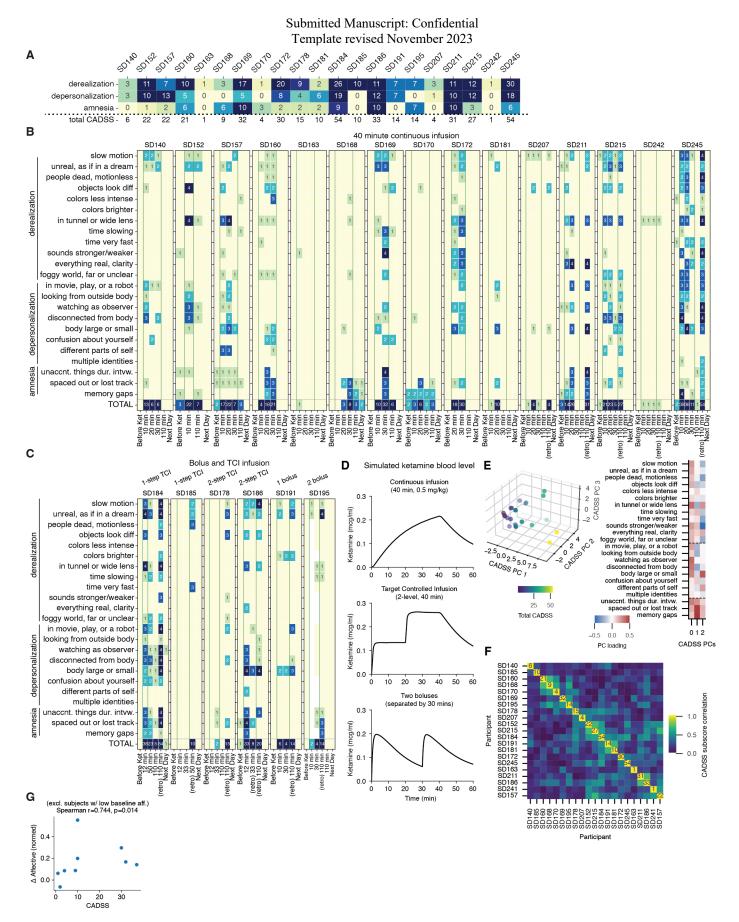


Fig. S14. Diverse Clinician-Administered Dissociative States (CADSS) symptom profiles across human participants receiving ketamine. (A) Summary of peak CADSS scores across study subjects. (B) CADSS scores across time points for

subjects receiving continuous 40-minute infusions (0.5 mg/kg). Retro denotes retrospective CADSS, in which a patient recounted after the infusion their peak experience during the infusion. (C) CADSS scores across time points for subjects receiving bolus or target-controlled infusions (TCI). (D) Simulated ketamine blood levels for different infusion types in (B) and (C), using StanpumpR PK/PD simulation program. (E) Left: Distribution of patients in dimensionality-reduced CADSS space (each point is a patient). Right: Loading of CADSS subquestions across first three PC axes. (F) Correlation of each patients' CADSS subscores. Patients are sorted using hierarchical clustering. Patient total CADSS scores are annotated along diagonal. There appear to be soft clusters of CADSS response-types, however this study was not powered to investigate those differences. (G) Spearman correlation between ketamine change in affective normalized by reflexive eye closure (Δ defined here as: infusion – mean(preinfusion, postinfusion)) vs. CADSS score. Only subjects with preinfusion normalized affective eye closure > 0.5 were included in this analysis, to ensure that there was an adequate baseline level from which to detect changes. See Supplementary table 3 for information on statistical analyses and sample sizes.

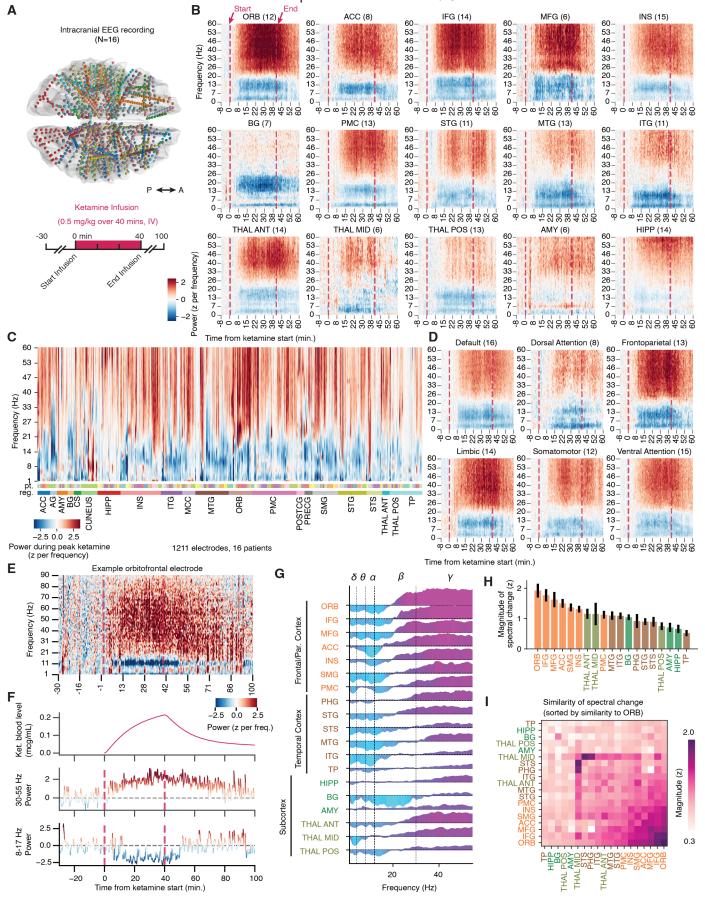


Fig. S15. Human intracranial spectral power changes during ketamine infusion. (A) Schematic of intracranial neural measurements conducted during 40-minute ketamine infusions. (B) Average spectrograms across regions of ketamine-induced power changes. Values are z-scored at each frequency to a 25-minute pre-ketamine baseline window. Numbers in parentheses denote patient count. (C) Spectral power changes, averaged during 30-40 minute peak ketamine window. Each column is an individual channel. (D) Average spectral power changes for Yeo7-defined functional networks. Numbers in parentheses denote patient count. (E) Example spectrogram from a single orbitofrontal bipolar channel. (F) Ketamine pharmacokinetics (top), compared to spectral power change over time shown in (E) for high (middle) and low (bottom) power bands. (G) Change in power (ketamine – preinfusion), z-scored at each frequency, median across subjects. Sorted anatomically, dark green: subcortical, lime green: thalamus, orange: frontal/parietal cortical, brown: temporal cortical. (H) Magnitude of spectral change, by region (mean ± s.e.m.). Frontal cortex exhibited the largest magnitudes of change, led by orbitofrontal cortex. (I) Similarity of spectral changes, sorted by similarity to orbitofrontal cortex. Cortical regions exhibited similar changes, especially frontal cortex, that were also reflected by anterior thalamus and to a lesser extent mid-thalamus. Other subcortical regions, such as basal ganglia, exhibited distinct changes.

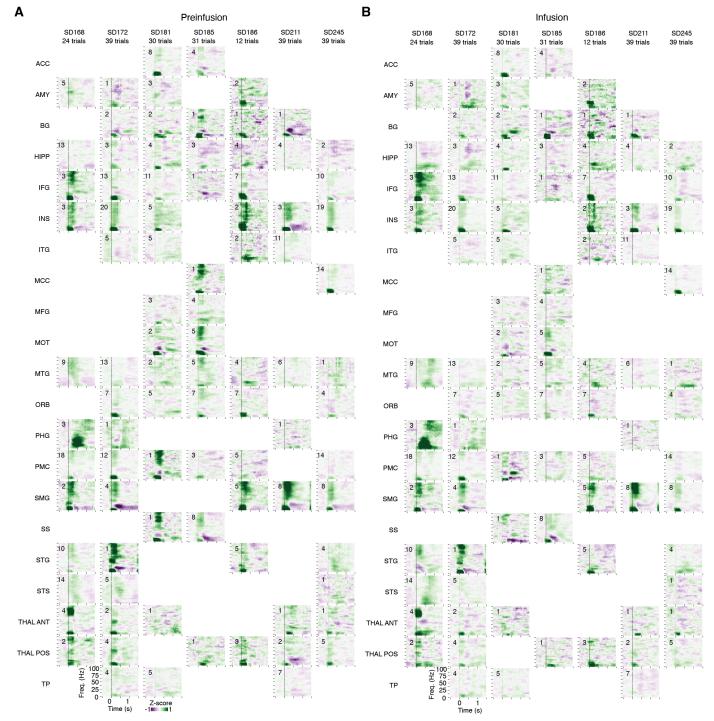


Fig. S16. Puff-triggered spectrogram, by human subject and region, averaged across trials. (A) Preinfusion and (B) Infusion. Number of channels indicated in top left corner of each spectrogram.

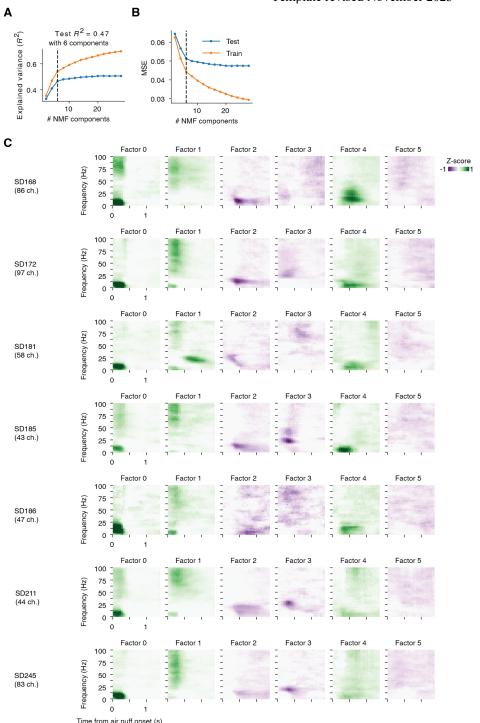


Fig. S17. Additional detail on human puff-triggered spectrogram factorization. (A) Explained variance on a train and test set (with the same channels, but using different sets of trials to compute the trial-averaged spectrogram, split ratio = 0.5) as a function of the number of NMF components. As indicated by performance on the test set, the optimal number of components is determined by the location of the 'elbow' to be six, past which there are diminishing returns for increasing the number of components. (B) Mean squared error (MSE) on a test set of trials as a function of number of NMF components. (C) Factors computed on each subject individually, showing that most factors are present in some form in all subjects. See Supplementary table 3 for information on statistical analyses and sample sizes.

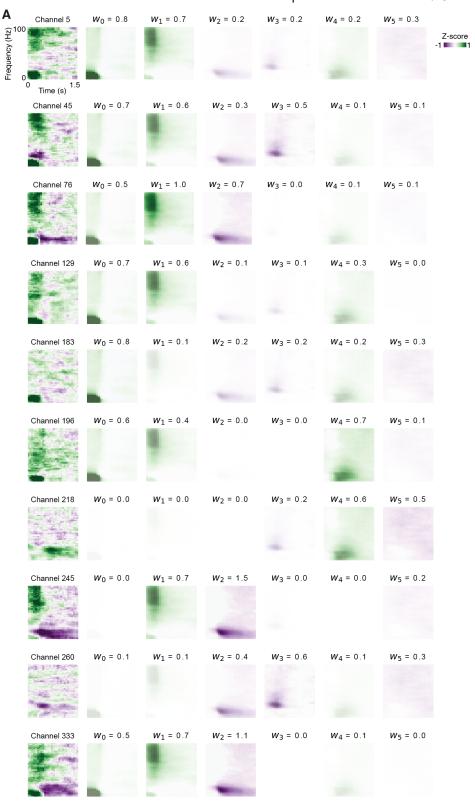


Fig. S18. Additional detail on human puff-triggered spectrogram factorization by channel. (A) Visualization of factor loadings for a set of example channels. Left column, the spectrogram of the channel (averaged across trials). Right columns, the factors and their loadings for that channel (denoted w_{factor}), with alpha transparency of the factor set according to loading.

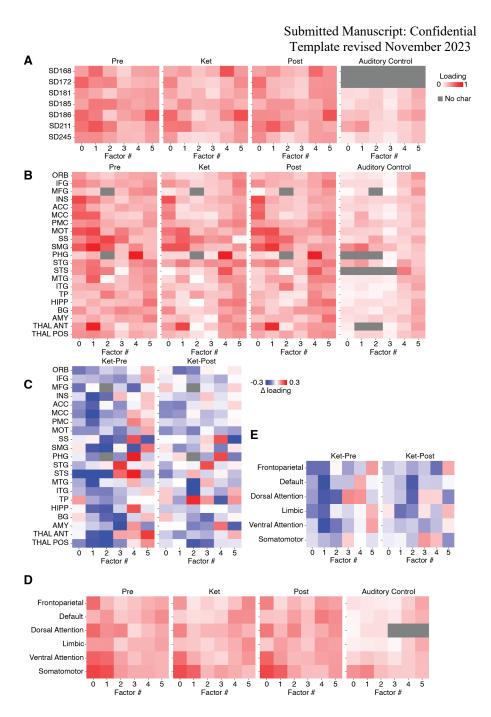


Fig. S19. Anatomical loading of each factor by condition. (A) Loading of each factor, averaged across channels within each subject, during preinfusion and infusion, indicating that most factors have substantial loading on multiple subjects. (B) Loading of each factor, averaged across channels within each region, during preinfusion, ketamine infusion, postinfusion, and auditory control conditions. The auditory control exhibits substantially lower loading overall. (C) Change in loading by region during infusion, compared with preinfusion and postinfusion. (D) Loading of each factor, averaged across channels within each Yeo7 resting state networks during each condition. (E) Change in loading by Yeo7 network during infusion relative to preinfusion and postinfusion.

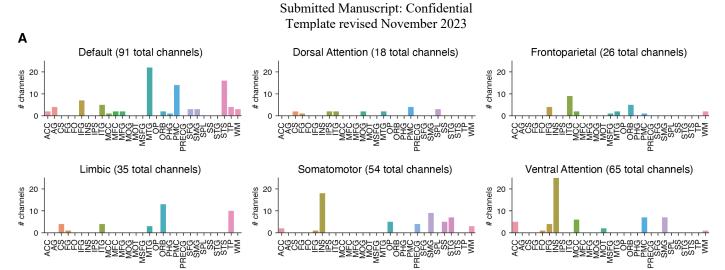


Fig. S20. Yeo7 human anatomical mapping. (A) Number of channels by region assigned to each Yeo7 network. Note that contribution to the somatomotor network is primarily composed of channels from insula.



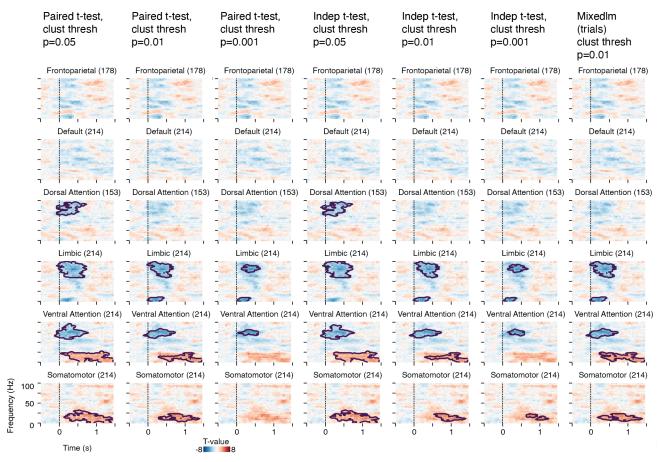


Fig. S21. More detail on permutation cluster test by human Yeo7 networks. (A) Significant changes (infusion – preinfusion) by Yeo7 resting state network, comparing different permutation cluster test parameters: cluster threshold p-value, and statistical test for comparing conditions. These results indicate that the significant clusters are robust to different permutation cluster hyperparameters. Vertical line, air puff onset.

Submitted Manuscript: Confidential Template revised November 2023 Α Paired t-test, Paired t-test, Paired t-test, Ind t-test, Ind t-test, Ind t-test, IndHat t-test, Paired t-test, Indep t-test, MixedIm clust thresh clust thresh clust thresh clust thresh clust thresh clust thresh p=0.05 p=0.001 tfce [0, 0.1] tfce [0, 0.1] clust thresh p=0.05p=0.01p=0.001p=0.01 p=0.001 p=0.01 ORB (151) 3 D S £.3 IFG (175) MFG (61) INS (183) SN ACC (61) MCC (70) 3 3 53 PMC (175) 5 望 0 8 2.3 PMC MOT (61) SS (61) 8 SMG (153) PHG (102) STG (114) E-SE-STS (102) STS (102) MTG (214) TG (120) ITG (120) TG (120) TG (120) ITG (120) ITG (120) TG (120) TG (120) TG (120) ITG (120) TP (108) HIPP (214) BG (151) BG (151) BG (151) **F** BG (151) **F** BG (151) BG (151) BG (151) BG (151) BG (151) BG (151)

BG

THAL ANT

100 50 AMY (105)

THAL ANT (171)

THAL POS (184)

AMY (105)

THAL POS (184)

AMY (105)

THAL POS (184)

THAL ANT (171) THAL ANT (171)

Fig. S22. More detail on permutation cluster test by human brain region. (A) Significant changes (infusion – preinfusion) by region, comparing different permutation cluster test parameters: cluster threshold p-value (0.05, 0.01, 0.001, and threshold-free cluster enhancement), and statistical test for comparing conditions (paired t-test, independent t-test, independent t-test with hat variance correction, and mixed-effects linear model across trials grouped by subject with paired t-test). Regions that exhibit significant changes robust to all hyperparameter settings are indicated at left. Vertical line, air puff onset.

AMY (105)

THAL ANT (171)

THAL POS (184)

AMY (105)

THAL ANT (171)

THAL POS (184)

AMY (105)

THAL ANT (171)

THAL POS (184)

AMY (105)

THAL POS (184)

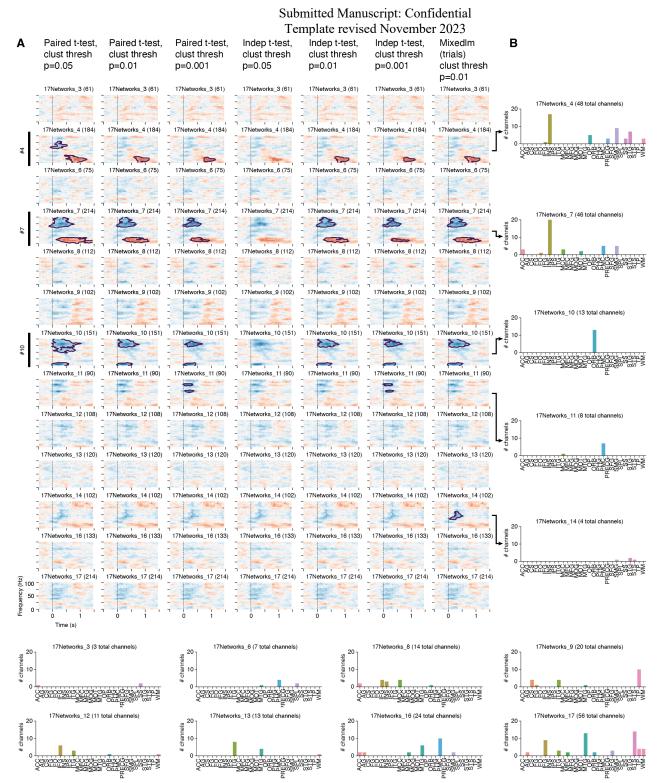


Fig. S23. Permutation cluster test by human Yeo17 network. A finer-grained 17-network resting-state atlas. (A) Significant changes (infusion – preinfusion) for each Yeo17 resting state network, comparing different permutation cluster test parameters: cluster threshold p-value, and statistical test for comparing conditions. Vertical lines, air puff onset. (B) Number of channels by region assigned to each Yeo17 network. Networks 4 & 7 (primarily insular), and 10 (primarily orbitofrontal) exhibited significant changes that were robust to hyperparameter settings. This analysis yielded similar results as the coarser-grained 7-network resting-state atlas, and further highlighted significant changes in orbitofrontal cortex (ORB) and insular cortex (INS), as well as supramarginal gyrus (SMG).

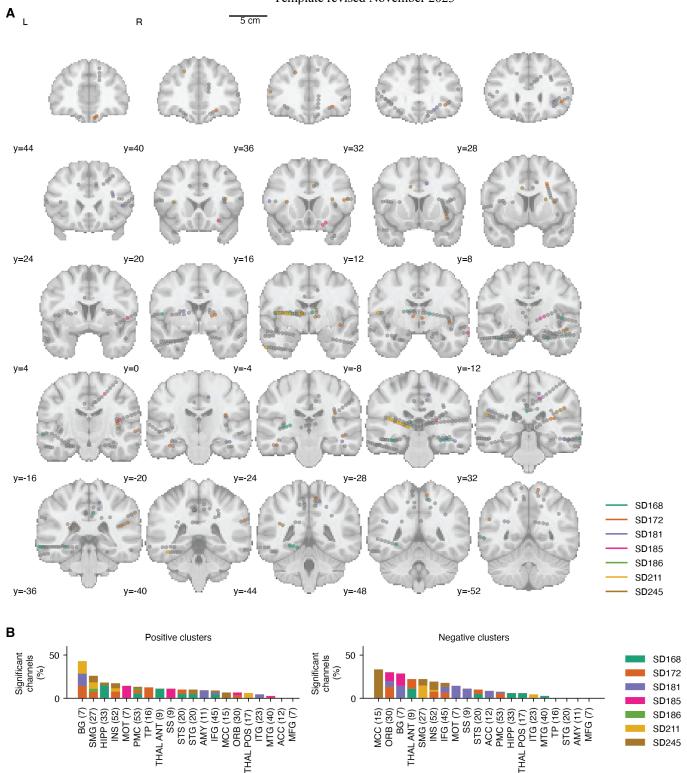


Fig. S24. Permutation cluster test, by individual human channel. (A) Location of channels with a significantly changing spectrotemporal cluster (infusion – preinfusion). (B) Percent of channels, by region and subject, that have a cluster of significantly increasing (positive) activation or significantly decreasing (negative) activation. Total number of channels per region in parentheses. Regions with a large fraction of significant channels are generally those that exhibited robust region-level significant spectrotemporal changes as shown in Fig. 3C.

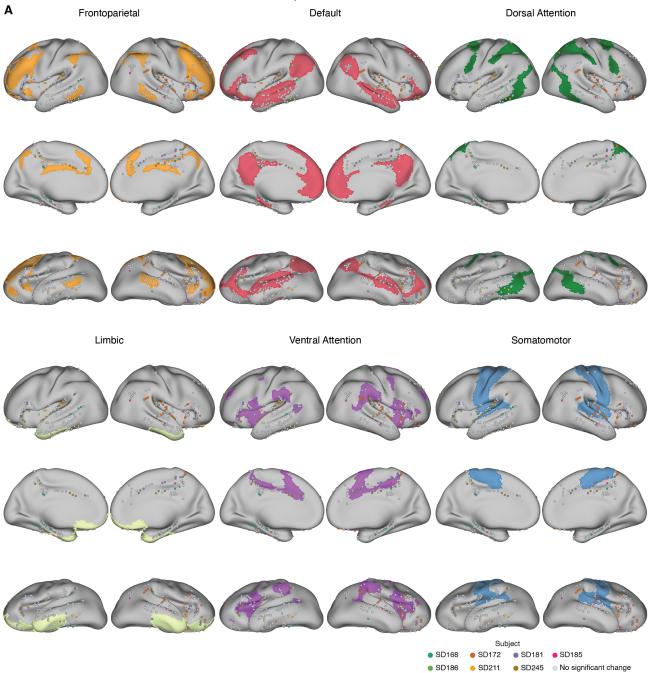


Fig. S25. Permutation cluster test by individual human channel, on Yeo7 parcellation. (A) Location of channels with a significantly changing spectrotemporal cluster (infusion – preinfusion) overlaid on Yeo7 resting-state network anatomical parcellation.

Streamline density of thalamocortical structural connectivity

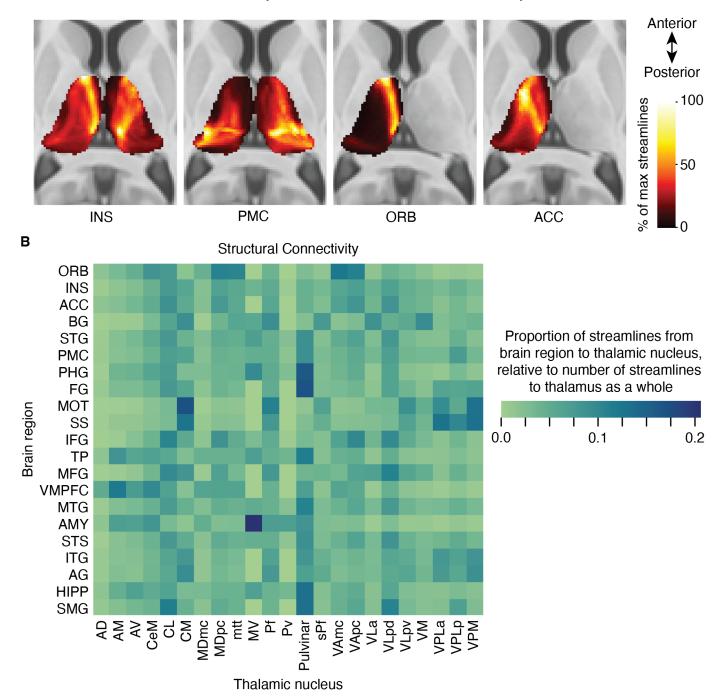


Fig. S26. Tractography-based structural connectivity to thalamus. The presence of shared stimulus-evoked spectrotemporal neural activity patterns suggests the possibility that brain regions with a ketamine-mediated change in eyepuff response directly coordinate their activity through anatomical connections. For example, certain specific regions, including several that are widely-separated across the brain, showed strikingly similar patterns of modulation—such as PMC and ORB, which exhibited same-direction modulation in distinct high and low frequency bands (Fig. 3C), resembling the ketamine-modulation response of the limbic network (Fig. 3B) associated with factor 2 (Fig. 2F, J). Unlike PMC and ORB, insula (INS) instead revealed the opposite-direction modulation in distinct high and low frequency bands (Fig. 3C) closely resembling ketamine-induced changes seen in the ventral attention network (Fig. 3B) associated with factors 2 and 3 (Fig. 2F, J). One possible mechanism for coordination of cortical region activity could be through shared thalamic connectivity. To investigate this, we used diffusion tractography to assess structural connectivity between thalamus and seed locations corresponding to patient contact locations. (A) Density of streamlines connecting thalamus to seeds placed on patient contact

locations in insular, posteromedial, orbitofrontal, and anterior cingulate cortices (Max Intensity Projection). INS and ORB exhibit stronger connectivity to anterior thalamus, whereas PMC exhibits stronger connectivity to posterior thalamus. ORB also exhibits some connectivity to posterior thalamus. This suggests that the strikingly similar effect of ketamine on PMC (exhibiting more robust connectivity to posterolateral thalamic structures) and ORB (with distinct anterior thalamic connectivity preference), both in terms of resting spectrotemporal profile (fig. S15 G-I) and eyepuff-evoked dynamics (Fig. 3C), may arise through mechanisms beyond shared thalamic connectivity. Alternative possibilities for coordination among these regions include direct ORB-PMC connectivity, intra-thalamic connectivity, finer-scale connectivity with thalamus not resolvable by diffusion tractography, and indirect coordination via other interconnected regions. (B) Summary of structural connectivity from non-thalamic brain regions to each thalamic nucleus.

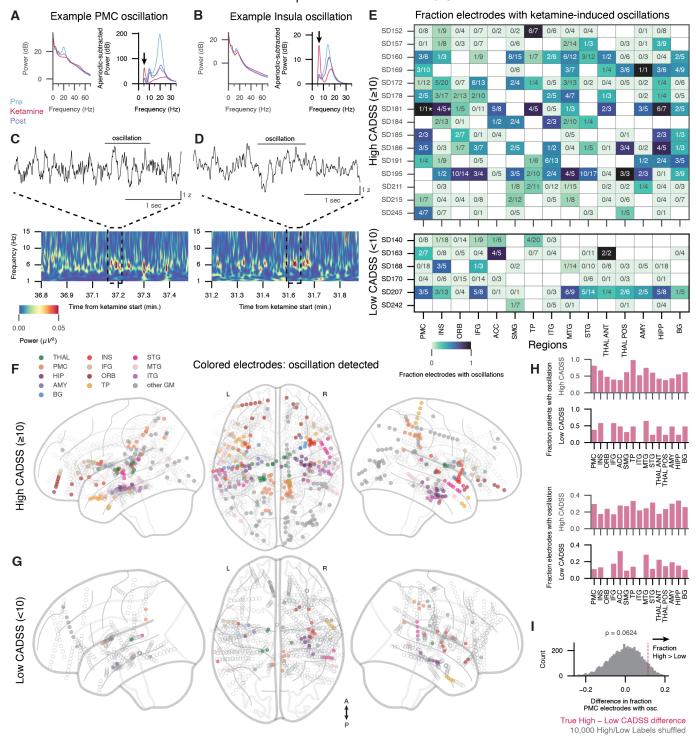


Fig. S27. Ketamine induces slow oscillations broadly across human brain regions. We previously reported that ketamine induces slow oscillations in mice and that seizure aura-associated dissociation is associated with slow oscillations in human PMC (28); subsequent iEEG investigation in humans also found a low frequency spectral power increase in PMC during rapidly-administered ketamine (0.5 mg/kg over 14 minutes, a ~3x faster infusion rate than our continuous infusion) (38). Here we investigate ketamine-induced band-limited, low-frequency oscillations (81) in our human participants. All oscillations in this figure refer to sub-10 Hz slow oscillations. (A) Left: Power spectral density (PSD) demonstrating a ketamine-induced slow oscillation for an example posteromedial cortex (PMC) bipolar contact, from SD181. Right: PSD with 1/f aperiodic spectrum subtracted, highlighting band-limited oscillatory peaks. Black arrow indicates a sub-10 Hz slow oscillation. (B) Same as (A), but for an example insula bipolar contact, from SD181. (C) Top: Bipolar voltage trace of PMC

contact in (A), showing example slow oscillation (black line). Bottom: spectrogram of PMC contact. (D) Same as (C), but for example insula bipolar contact in (B). (E) Distribution of ketamine-induced slow oscillations across regions and patients, subdivided by high and low CADSS score. Ketamine-induced oscillations are identified when their amplitude and/or frequency are different compared to pre- and post-infusion. (F) Spatial arrangement of ketamine-induced slow oscillations for high CADSS subjects (CADSS >= 10). Uncolored channels did not have detected oscillations. (G) Same as (F), for low CADSS subjects. (H) Comparison across high vs. low CADSS subjects of channel fraction with slow oscillations per region (top), and of patient fraction with > 0 slow oscillations (bottom). (I) Comparison of measured difference between high and low CADSS subject's total channel fraction with slow oscillations versus a null distribution in which high and low subject labels are randomly permuted 10,000 times. Despite heterogeneity across subjects (as expected from diversity in patient-specific responses to ketamine, as well as anatomical sampling differences dictated by clinically-guided sEEG electrode placement) (fig. S27 E, F), all subjects with strong dissociation (CADSS score >10) exhibited low-frequency (<10 Hz) oscillations induced by ketamine (most notably in PMC and TP; we also observed that subjects with high CADSS scores tended to exhibit a higher proportion of channels recording these low-frequency oscillations compared to subjects with low CADSS scores: fig. S27 E–I).

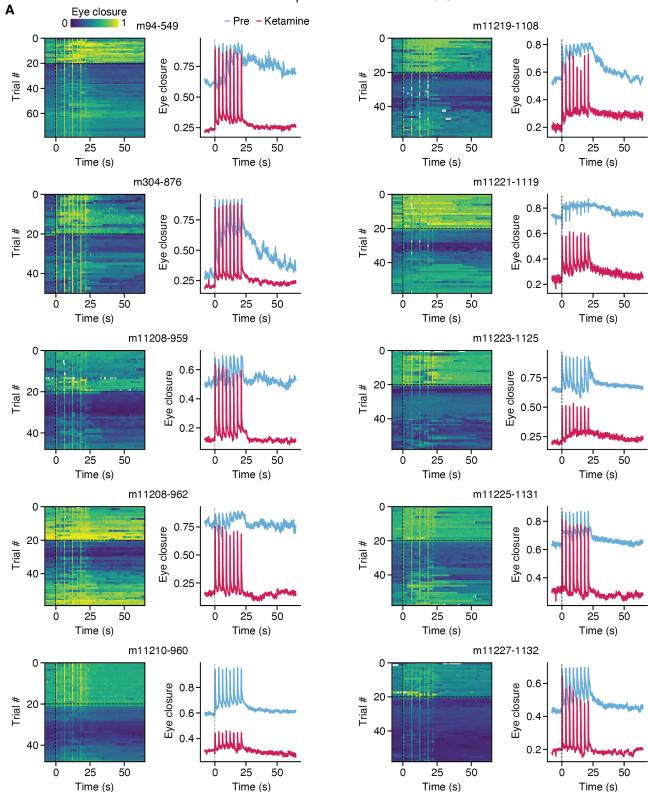


Fig. S28. Mouse eye closure during Neuropixels recording sessions. (A) Left, eye closure on individual trials (each consisting of eight puffs); Right, mean across trials during preinfusion and ketamine conditions. Bolus ketamine infusion occurs at trial 20.

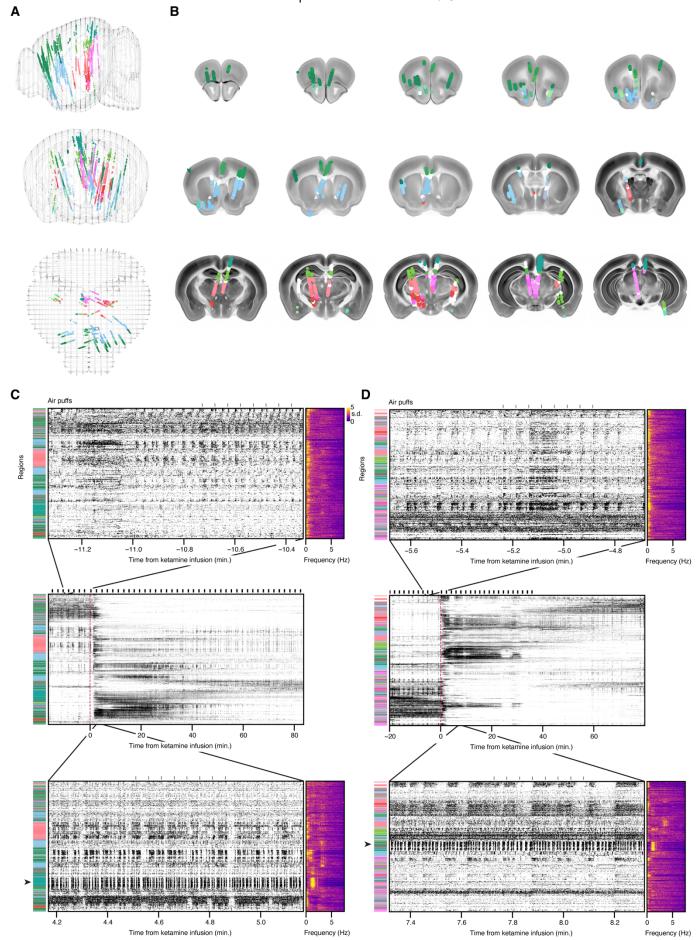


Fig. S29. Additional details on mouse Neuropixels recording during ketamine infusion. (A, B) Anatomical location of all recorded units. (C, D) Example recording. Red dashed line indicates time of bolus ketamine infusion. Units ordered by rastermap clustering algorithm. Top, preinfusion. Bottom, after infusion. Left, region labels, colored as in (B). Black arrow indicates retrosplenial cortex. Right, power spectral density of each channel. Retrosplenial cortex exhibits particularly strong rhythmic activity.

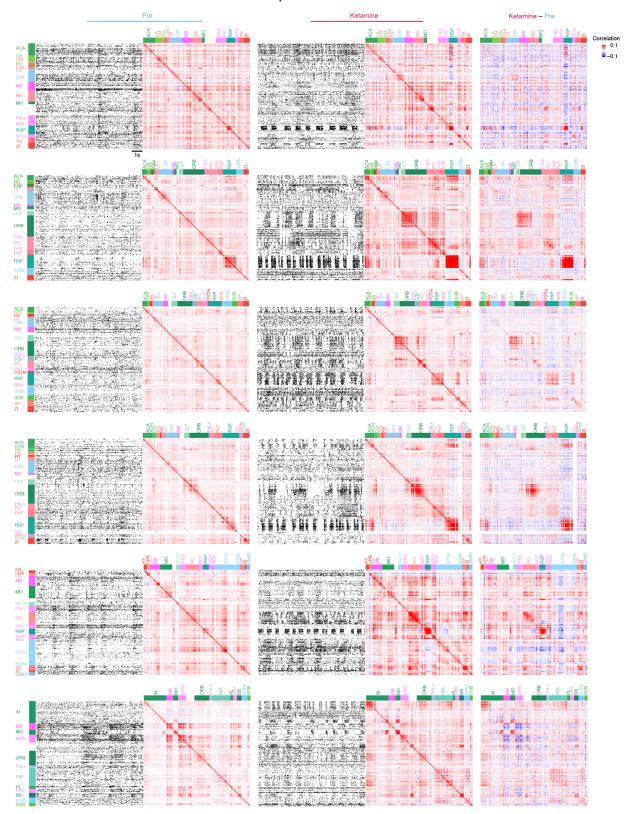


Fig. S30. Change of mouse neural activity correlation structure with ketamine. (A) Left: preinfusion spiking activity of simultaneously recorded units during example ten second window, and single unit correlation matrix across 10 min window. Middle: same, but following bolus ketamine infusion. Right: Change of correlation matrix (ketamine – preinfusion).

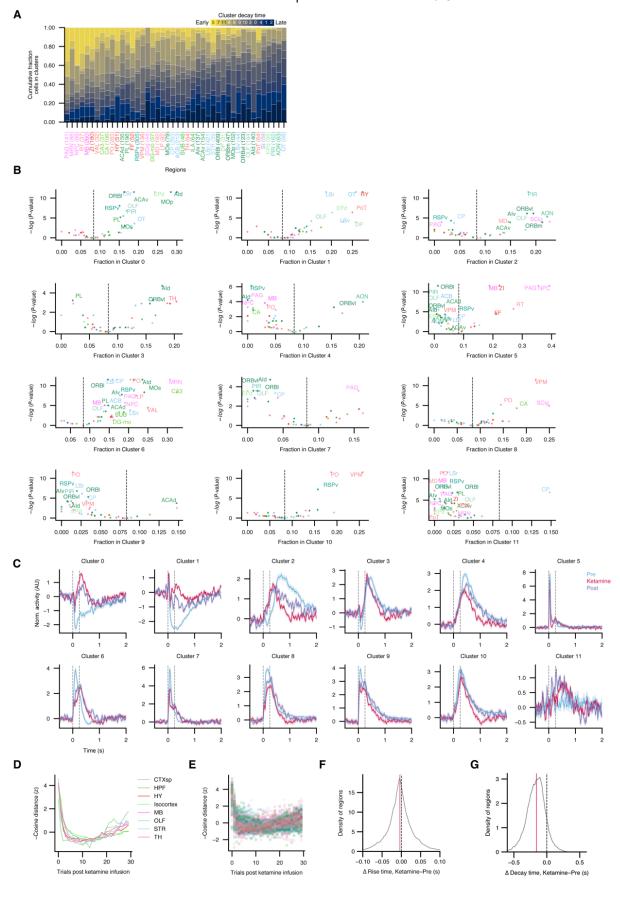


Fig. S31. Additional detail on mouse single unit activity and peri-puff changes on ketamine. (A) Cumulative fraction of cells in clusters, by region. Clusters are colored by the rank order time for average activity across all recorded cells in a cluster to decay to 0.25x the peak value post-puff. Clusters are numbered according to their order from top to bottom in Fig. 4D. Regions sorted in ascending order by a weighted average of cluster decay times, with weights equal to the fraction of cells in a cluster in a given region. Region acronyms colored by the Allen Brain Atlas colormap. (B) Representation of regions in each cluster. (C) Mean cluster activity (normalized within each condition) for all clusters. Mean ± 95% CI. (D, E) Similarity (negative z-scored cosine distance) between preinfusion cluster activity and activity on each trial following bolus ketamine infusion. Similarity averaged across mice within coarse anatomical grouping (D) and finer anatomical grouping (E, each region shown as individual dot for each time bin). Bootstrapped distribution of changes in peri-puff rise times (F) and decay times (G) across regions. Red line, median. See Supplementary table 3 for information on statistical analyses and sample sizes.

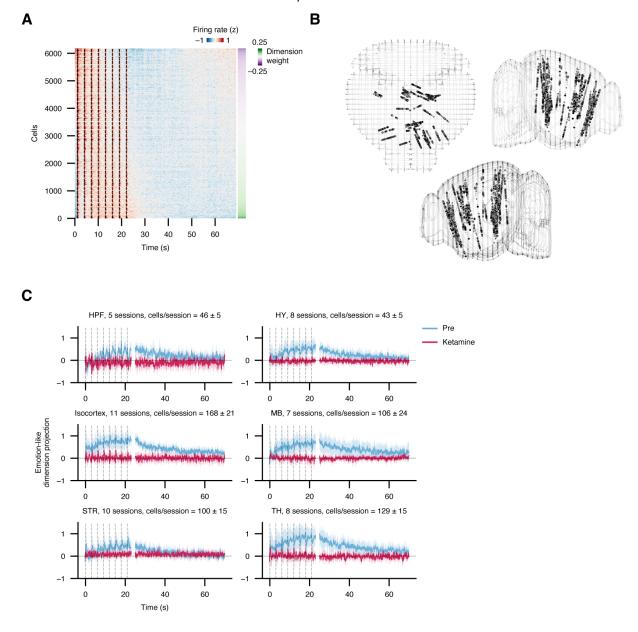


Fig. S32. Additional detail on emotion-like neural dimension. (A) Trial-averaged single cell responses, where a trial consists of a sequence of eight puffs. Cells sorted based on their weight in the emotion-like neural dimension, indicated on the right. (B) Spatial distribution of emotion-like neural dimension weights, showing broad anatomical distribution. (C) Emotion-like neural dimension projection computed by high-level brain area. Mean \pm 95% CI. All high-level brain areas (where there are sufficient numbers of simultaneously recorded neurons to perform population analyses) exhibit some degree of accumulating persistent dynamics during preinfusion that is abolished by ketamine.

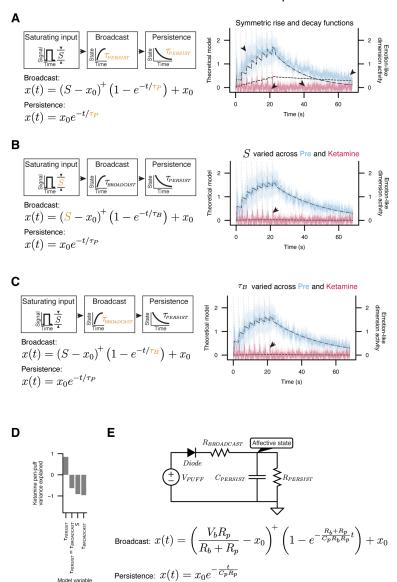


Fig. S33. Additional detail on first order system model of emotion-like state. (A) Alternative symmetric model in which there is only a single timescale across the broadcast and persistence phases. Forcing the rise and decay rates to be symmetric leads to an inability to accurately fit the preinfusion or infusion states. The leftmost and rightmost arrows indicates a poor preinfusion fit, and the middle two arrows indicate a poor infusion fit. (B) Alternative model in which the magnitude of the input signal S is varied across conditions instead of $t_{PERSIST}$. This leads to an inability to fit the transient peri-puff increases in emotion-like state during ketamine infusion, as indicated by the arrow. (C) A similar inability occurs in an alternative model where $t_{BROADCAST}$ is the only free parameter across conditions, as indicated by the arrow. (D) Variance explained (R^2) of the model evaluated specifically on the puff series during ketamine epoch. This suggests that a model where ketamine modulates $t_{PERSIST}$, as opposed to $t_{BROADCAST}$ or S, is a better fit to the experimental data. (E) An example of a physical system that exhibits the same first order dynamics as our model: an analog electrical circuit with passive components. The affective state is represented as the voltage at the node preceding the capacitor. The diode enforces asymmetry between a broadcast phase when V_{PUFF} is high, and a persistence phase when V_{PUFF} is low. The capacitor serves to store the input signal and causes the input signal to saturate. The persistence timescale is determined by $t_{PERSIST} = C_{PERSIST}R_{PERSIST}$, and the broadcast timescale is further controlled by $t_{PBROADCAST}$. The dynamics were solved using Kirchoff's Current Law.

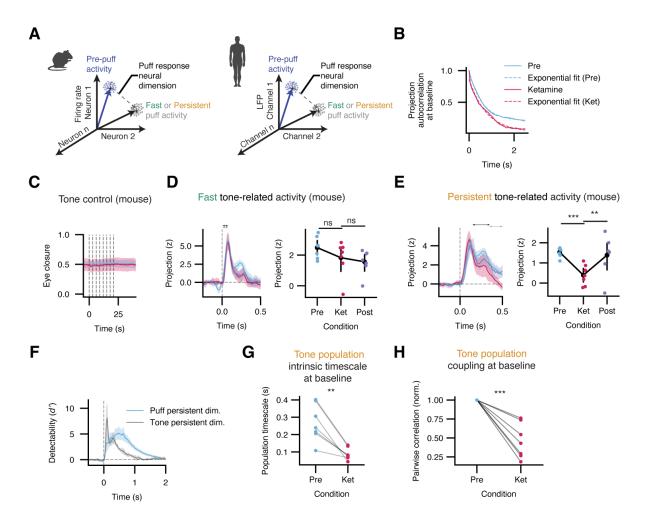


Fig. S34. Additional details on coding dimension analysis. (A) Schematic of analysis to identify neural dimensions encoding either fast or persistent puff-triggered population activity, relative to pre-puff activity. In mice, the dimension is in the space of neuron firing rates, whereas in humans it is in the space of channel LFPs. (B) Example autocorrelation of coding dimension projection during preinfusion or ketamine (solid lines), and exponential fits (dashed lines) used to compute the intrinsic timescale. Autocorrelation functions were fit to the exponential function f(x) = a*exp(-x/b)+c. The ketamine autocorrelation decays more quickly than the preinfusion autocorrelation, corresponding to a reduced intrinsic timescale (parameter b, above) with ketamine. (C) Mice do not exhibit substantial eye closure in response to a sequence of auditory tones during preinfusion or ketamine (dashed lines represent onset of each 250 ms tone). (D) Fast tone-related population activity is not significantly altered by ketamine (coding dimension and mean activity computed during 0 to 70 ms following tone onset). (E) Persistent tone-related population activity is reduced by ketamine relative to preinfusion and postinfusion (coding dimension computed during 150 to 350 ms following tone onset, and mean activity quantified 350 to 500 ms following tone onset). Black arrow, coding dimension window; gray arrow, mean activity window. See Methods for detail on choice of time windows. (F) Detectability (d') of the projection of the persistent neural dimension for either eyepuff or tone. The affectively salient eyepuff elicits substantially longer persistent dynamics than the neutral tone. C-F, mean ± 95% CI. (G) Intrinsic timescale at baseline of persistent tone population is reduced during ketamine relative to preinfusion. (H) Coupling at baseline within persistent tone population is reduced during ketamine relative to preinfusion. ns, P-value ≥ 0.05 . *, P-value < 0.05. **, P-value < 0.01. ***, P-value < 0.001. ***, P-value < 0.0001. See Supplementary table 3 for information on statistical analyses and sample sizes.

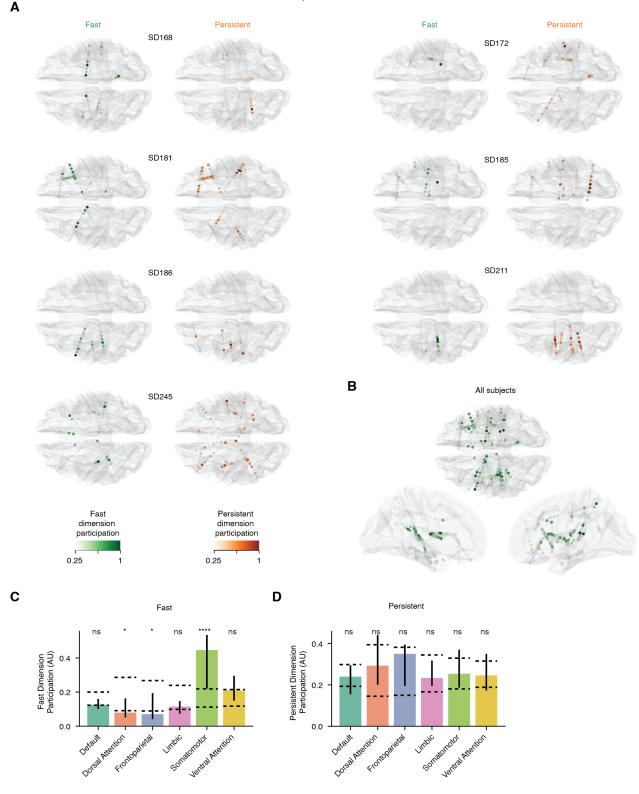


Fig. S35. Distributed human anatomical participation of puff-related coding dimensions. (A) Participation (loading) of each site for each subject for Fast and Persistent coding dimensions. (B) Sites from all subjects overlaid, for Fast coding dimension loadings. (C) Aggregate participation in Fast dimension for Yeo7 resting state networks. Participation is not uniformly distributed across networks, with somatomotor exhibiting over-participation. Dashed lines represent upper and lower thresholds signifying over- or under-participation in the Fast dimension relative to other networks (median \pm 95% confidence interval, thresholds determined with bootstrap shuffles of network label, 1000 shuffles). (D) Aggregate participation in Persistent dimension for Yeo7 resting state networks. Participation is global, and uniformly distributed

across networks. ns, P-value ≥ 0.05 . *, P-value < 0.05. **, P-value < 0.01. ***, P-value < 0.001. ***, P-value < 0.001. See Supplementary table 3 for information on statistical analyses and sample sizes.

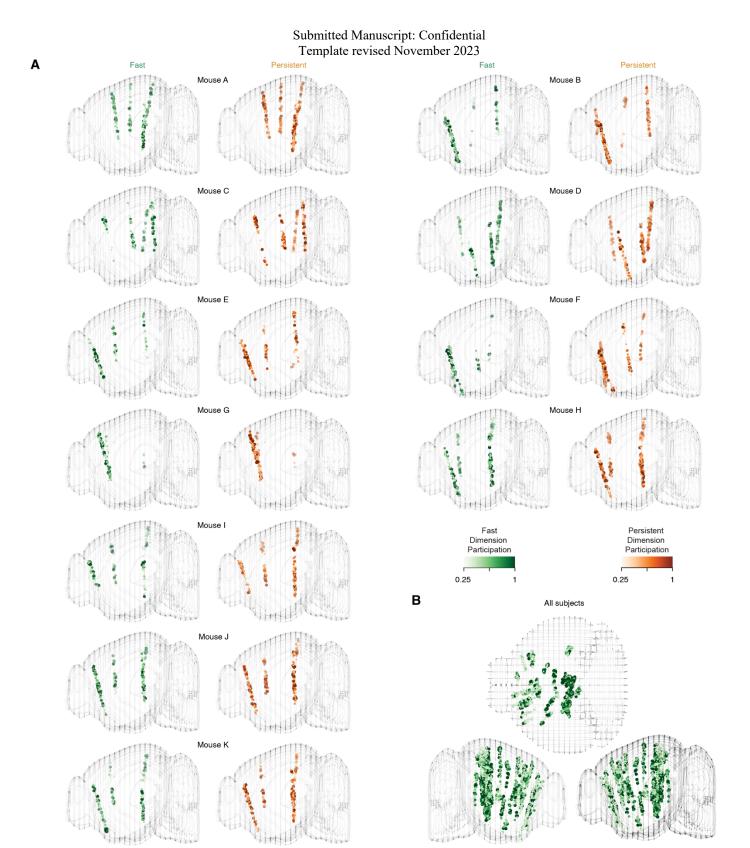


Fig. S36. Distributed mouse anatomical loading of puff-related coding dimensions. (A) Loading of each neuron for each mouse for Fast and Persistent coding dimensions. (B) Loading for all neurons from all mice for the Fast coding dimension.

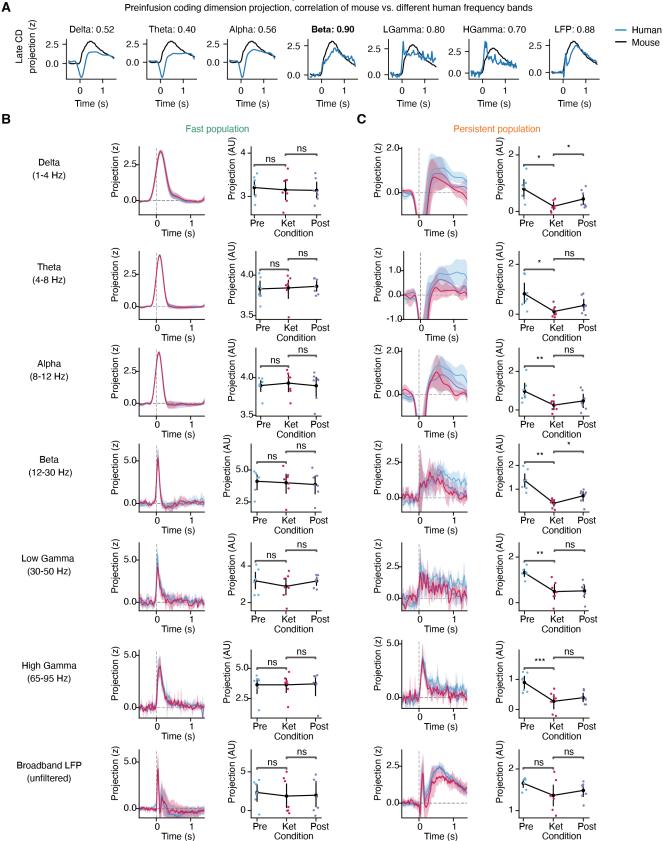


Fig. S37. Additional details on human coding dimension. (A) Preinfusion projection of late coding dimension (CD) computed using different spectral components of the electrical recording, and correlation with mouse firing rate-based CD. Beta power yields a CD projection that is most similar to that of the mouse. (B) Left: Early CD projections in humans for each spectral band during preinfusion, infusion, and postinfusion conditions (n=7 subjects, mean \pm s.e.m.). Right: No

significant differences across conditions of mean projection during early window. (C) Left: Late CD projections in humans for each spectral band (n=7 subjects, mean \pm s.e.m.). Right: All power bands (except for broadband LFP) exhibit significant decrease of persistent population projection from preinfusion to ketamine. Beta and delta further exhibit significant recovery during postinfusion. ns, P-value \geq 0.05. *, P-value < 0.05. **, P-value < 0.01. ****, P-value < 0.001. ****, P-value < 0.0001. See Supplementary table 3 for information on statistical analyses and sample sizes.

Fig. S38. Additional detail on human coupling changes across frequency bands and alterations by ketamine.

0.30

Pre-puff

ß phase locking

0.35

Affective

closure 0.75

0.50

0.2750.3000.325

Pre-puff

β phase locking

0.5

0.0

-0.5

regressior

Pre Ket

Single-trial,

Subject

2

0.0

2

0

10

20

Z-scored

0.0

Pre-puff

β phase locking (z)

Affective eye closure

(A) Preinfusion, out-of-task phase locking is not significantly different among edges between low puff-triggered persistence sites (low CD, denoted as L) vs. among edges between high persistence sites (high CD, denoted as H). (B) However, ketamine-induced decreases in coupling are greater in magnitude between high persistence sites, compared to between low persistence sites. Plot shows change (ketamine – preinfusion) in phase locking among low or high CD weight sites. Bipolar sites are classified as H or L by their weight assigned within the persistent peri-eyepuff neural activity dimension, or coding dimension, abbreviated CD. Low refers to sites in the bottom 10 percentile of CD weight and high refers to sites in the top 10 percentile of CD weight, for CDs computed using power in each spectral band. (C) Beta phase locking changes among high persistence sites show kinetics (mean over edges and patients) that resemble the pharmacokinetics of ketamine, and differ from the kinetics of low CD edges. This suggests a link between the dissociative state and coupling changes among the persistent sites. (D) Preinfusion power of low persistence sites is not significantly different from high persistence sites. (E) Ketamine alteration of power is also not significantly different between low and high persistence sites, and thus differences in ketamine-induced coupling changes between high and low persistence subnetworks (shown in (B)) are not necessarily due to differences in spectral power changes. Plot shows change (ketamine – preinfusion) in spectral power among low or high CD weight sites. (F) Temporal progression of beta spectral power around 40-minute ketamine infusion. Kinetics of spectral power do not markedly differ between high and low CD sites, in contrast to kinetics of phase locking (as shown in (C)). (G) Single-trial correlation of pre-puff network coupling with post-puff persistent activity for an example subject, SD168. Network coupling is measured as the beta phase locking value averaged across all recorded edges between

non-noise channels, averaged in the 0.5 seconds before a puff. Persistent activity is measured as the projection of population activity along the affective window beta coding dimension, i.e. the linear combination of all channel beta-filtered powers weighted by coding dimension weights. Left: single-trial scatter and best linear fit, comparing preinfusion and ketamine trials in different colors. Right: the progression of these metrics over trials during preinfusion (top) and ketamine (bottom) air puff sessions. Pre-puff network coupling is represented by a dotted line, and post-puff persistent activity is represented by a solid line. (H) Network coupling and persistent activity correlation as in (G) but represented across all patients. The two plots represent the best linear fits for all patients during preinfusion and ketamine, each patient represented in a different color. P-value is computed as the significance of the cross-patient correlation coefficient, from the hierarchical mixedeffects linear model with random intercept, grouping trials across patients. (I) Statistical significance of each patients' pertrial coupling to persistent activity correlation. Left: per-patient correlation coefficient (r) with bootstrapped confidence intervals. Right: cross-patient regression coefficient is significantly positive for preinfusion air puff sessions, but not significant for ketamine sessions. Cross-patient coefficients are computed using a hierarchical mixed linear model with random intercept grouped across patients. Error bars denote the 95% confidence interval of coefficient estimates. (J) Single trial correlation of pre-puff network coupling with post-puff affective eye closure for an example subject, SD211. Left and right are analogous to (G). Pre-puff network coupling is represented by a dotted line, and post-puff affective behavior is represented by a solid line. (K) Network coupling and affective behavior correlation as in (J) but represented across all patients. The two plots represent best linear fits for all patients during preinfusion and ketamine, each patient represented in a different color. (L) Statistical significance of each patients' per-trial coupling to eye closure correlation. Left: per-patient correlation coefficient (r) with bootstrapped confidence intervals. Right: cross-patient regression coefficient is significantly positive for preinfusion air puff sessions, but not significant for ketamine sessions. Cross-patient coefficients computed as above. ns, P-value ≥ 0.05. *, P-value < 0.05. **, P-value < 0.01. ***, P-value < 0.001. ****, P-value < 0.001. See Supplementary table 3 for information on statistical analyses and sample sizes.

			Num	ber of S	Significan	t Contacts	3	C 1C (To	tal Nun	nber of C	ontacts			Total	Total	Fraction
HCP Cortical Area	Region	SD168	SD172	SD181	SD185	SD186 SI	D211 S	SD245	SD168	SD172 S	SD181	SD185 S	SD186 S	SD211 SE)245	Significant	Contacts	Significant
R_RSC_ROI	PMC	0	0	0	0	0	0	1	2	0	0	0	0	0	3	1	5	0.2
R_PSL_ROI	SMG	0	2	0	0	0	0	0	0	2	0	0	0	0	1	2	3	0.7
R_PCV_ROI	PMC	0	1	0	0	0	0	0	0	2	0	0	0	0	0		2	0.5
R_STV_ROI	SMG	0	2	0	0	0	0	0	0	2	0	0	0	0	0	2	2	1.0
	PMC	0	0	0	0	0	0	2	0	1	0	1	0	0	3	2	5	0.4
R_5mv_ROI	PMC	1	0	3	0	0	0	0	2	0	3	0	0	0	0		5	0.8
R_23c_ROI R_24dd_ROI	PMC SMC	0	0	0	1	0	0	0	0	2	0	2	0	0	0	1 2	2	0.3 1.0
R_7Am_ROI	PMC	0	3	0	0	0	0	0	0	3	0	0	0	0	0	3	3	1.0
	IPS	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1	1.0
R_p32pr_ROI	ACC	0	0	2	0	0	0	1	0	0	3	0	0	0	1	3	4	0.8
R_47m_ROI	ORB	0	1	0	0	0	0	0	0	1	0	0	0	0	0		1	1.0
R_44_ROI	IFG	0	3	0	0	0	0	0	0	4	3	0	0	0	1	3	8	0.4
R_45_ROI	IFG	0	1	0	0	0	0	0	0	1	2	3	0	0	1	1	7	0.1
R_11I_ROI	ORB	0	2	0	0	0	0	0	0	3	0	0	0	0	0	2	3	0.7
R_13I_ROI	ORB	0	0	3	0	0	0	0	0	0	3	0	0	0	0	3	3	1.0
R_OFC_ROI	ORB	0	3	0	0	0	0	0	0	3	2	1	0	0	0	3	6	0.5
R_OP1_ROI	SMC	0	0	0	1	0	0	0	0	0	0	3	0	0	0	1	3	0.3
R_OP2-3_ROI	INS	0	2	0	0	0	0	0	0	3	0	0	0	0	0	2	3	0.7
R_RI_ROI	IFG	0	0	0	0	0	0	4	0	0	0	0	0	0	5	4	5	0.8
R_PFcm_ROI	SMG	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1	1.0
	INS	1	4	0	0	0	0	0	2	4	0	0	0	0	0	_	6	0.8
R_FOP4_ROI	IFG	0	1	2	0	0	0	1	0	1	3	0	0	0	1	4	5	0.8
	INS	0	3	0	0	0	0	0	0	5	3	0	0	0	0	-	8	0.4
	IFG	0	0	1	0	0	0	0	0	0	1	2	0	0	0	1	3	0.3
	INS	0	1	0	0	0	0	0	0	5	0	0	0	0	0	1	5	0.2
R_H_ROI	HIPP	4	0	0	0	0	0	0	5	2	5	3	0	0	2	4	17	0.2
R_A5_ROI	STS	0	2	0	0	0	0	0	1	3	0	0	0	0	1	2	5	0.4
	STS MTG	0	1	0	0	0	0	0	0	2	0	0 2	0	0	0	1	6	0.5
	FG	0	0	1	0	0	0	0	0	0	1	0	0	0	0	1	1	0.2 1.0
R_TPOJ3_ROI	AG	0	0	0	1	0	0	0	0	0	0	1	0	0	0	1	1	1.0
	PMC	2	0	0	0	0	0	0	2	0	0	0	0	0	1	2	3	0.7
R_pOFC_ROI	ORB	0	0	0	2	0	0	0	0	0	0	2	0	0	0		2	1.0
·	INS	0	1	0	0	0	0	5	0	3	0	0	0	0	6		9	0.7
	INS	0	2	0	1	0	0	0	0	2	0	1	0	0	0	3	3	1.0
	IFG	0	1	2	0	0	0	0	0	1	3	0	0	0	0	3	4	0.8
R_MBelt_ROI	STG	1	0	0	0	0	0	0	1	0	0	0	0	0	0		1	1.0
R_PI_ROI	STG	0	0	0	0	0	0	2	0	0	0	0	0	0	2	2	2	1.0
L_RSC_ROI	PMC	1	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1.0
L_PSL_ROI	SMG	0	0	0	0	0	0	2	0	0	0	0	0	0	2	2	2	1.0
L_PCV_ROI	PMC	0	0	0	0	0	0	1	0	0	0	0	0	0	2	1	2	0.5
L_d23ab_ROI	PMC	1	0	0	0	0	0	0	1	2	0	0	1	0	0	1	4	0.3
L_5mv_ROI	PMC	0	0	0	0	1	0	0	1	0	0	0	1	0	0	1	2	0.5
L_23c_ROI	PMC	0	0	0	0	1	0	1	1	1	0	0	2	0	2	2	6	0.3
L_33pr_ROI	MCC	0	0	0	0	0	0	3	0	0	0	0	0	0	3	3	3	1.0
L_9p_ROI	SFG	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1	1.0
L_44_ROI	IFG	0	0	1	0	0	0	0	0	2	2	0	0	0	0	1	4	0.3
L_47I_ROI	IFG	0	0	0	0	0	0	1	0	0	0	0	1	0	1	1	2	0.5
	IFG	0	0	1	0	0	0	0	0	2	1	0	0	0	0		3	0.3
	SFG	0	0	0	0	0	0	2	0	0	0	0	0	0	2			1.0
L_OP4_ROI	PRECG	0	0	0	0	0	1	0	1	0	0	0	0	4	0		5	0.2
L_OP1_ROI L 52 ROI	SMG STG	0	0	0	0	0	1	0	0	0	0	0	0	1	0	1	1 2	1.0 0.5
	INS	1 0	0	0	0	2	3	6	1	0	0	0	3	3	6		12	0.5
	INS	0	0	0	0	0	2	0	1	0	0	0	0	2	0			0.9
	INS	0	0	0	0	0	0	2	0	1	0	0	0	0	2			0.7
	IFG	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1	1.0
	INS	1	0	0	0	0	1	0	2	0	0	0	0	1	0			0.7
L H ROI	HIPP	4		1	0	0	0	1	12	5	4	0	5	5	2			0.2
	PHG	1	0	0	0	0	0	0	1	0	0	0	0	1	0		2	0.5
	HIPP	1	0	0	0	0	0	1	1	0	0	0	0	2	1	2		0.5
	STS	2		0	0	0	0	0		0	0	0	0	0	1	2		0.3
	ITG	0	0	0	0	0	2	0	2	0	0	0	0	7	0			0.2
	TP	0	1	0	0	0	0	0	0	1	0	0	0	9	1	1	11	0.1
	ITG	0	0	0	0	0	2	0	0	0	0	0	1	3	0	2	4	0.5
L_TPOJ1_ROI	MTG	1	0	0	0	0	0	0	1	0	0	0	0	0	0		1	1.0
	MTG	1	0	0	0	0	0	0	3	0	0	0	0	0	0	1	3	0.3
	PHG	1	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1.0
	PHG	1	0	0	0	0	0	0	1	0	0	0	0	0	0		1	1.0
	INS	0	0	0	0	0	0	7	0	0	0	0	0	0	9	7	9	0.8
	IFG	0	0	0	0	0	0	1	0	0	0	0	0	0	1	1	1	1.0
	STG	1	0	0	0	0	0	0	1	0	0	0	0	0	0		1	1.0
	MCC	0	0	0	0	0	0	1	0	0	0	0	0	0	2		2	0.5
All areas with signific	cant contacts	26	39	19	7	4	12	49	55	70	41	21	14	38	70	156	309	0.5

			Nur	nber of	Significa	nt Cont			Visca i v			oer of Co	ntacts			Total	Total	Fraction
HCP Cortical Area	Region	SD168	SD172					1 SD245	SD168 S					SD211 SE	245	Significant	Contacts	Significant
	MOT	0	0		0 0			0 0		0	0	1	0	0	0	0	1	0.0
R_v23ab_ROI	PMC	0	0	(0 0	C) (0 0	0	0	0	1	0	0	0	0	1	0.0
R_d23ab_ROI	PMC	0	0	(0 0	C) (0 0	0	1	0	1	0	0	1	0	3	0.0
	ACC	0	0	(0	C		0 0		0	3	0	0	0	0	0	3	
	SS	0	0	(0	C) (0 0	0	0	0	3	0	0	0	0	3	0.0
	MCC	0	0		0			0 0		0	0	0	0	0	3	0	3	
	ACC	0	0		0 0			0 0		0	0	1	0	0	0	0	1	0.0
	MSFG	0	0		0 0			0 0		0	0	0	0	0	2	0	2	
	MFG	0	0		0			0 0		0	0	2	0	0	0	0	2	
	MFG	0	0					0 0		0	0	2	0	0	1	0	3	
	MFC	0	0		0			0 0		0	0	1	0	0	0	0	1	0.0
	MFC	0	0		0 0			0 0		0	0	1	0	0	1	0	2	
	MFG	0	0		0 0			0 0		0	1	0	0	0	0	0	1	0.0
	IFG	0	0					0 0		1	0	1	0	0	0	0	2	
	IFG	0	0					0 0		1	0	0	0	0	0	0	1	0.0
																-	·	
	IFG	0	0					0 0		1	0	0	0	0	0	0	1	0.0
	INS	0	0		0			0 0		0	1	0	0	0	0	0	1	0.0
	HIPP	0	0					0 0		0	0	0	0	0	0	0	1	0.0
	STG	0	0		0			0 0		0	0	0	0	0	0	0	1	0.0
	PHG	0	0		0			0 0		1	0	0	0	0	0	0	1	0.0
	MTG	0	0		0			0 0		2	0	0	0	0	1	0	3	
	MTG	0	0		0 0			0 0		1	0	0	0	0	0	0	1	0.0
	TP	0	0		0 0			0 0		3	0	0	0	0	0	0	3	
	ITG	0	0		0 0			0 0		2	4	0	0	0	0	0	6	
	ITG	0	0		0 0	C) (0 0		1	0	0	0	0	0	0	1	0.0
R_IP0_ROI	MOG	0	0	(0 0	C) (0 0	0	0	0	0	0	0	1	0	1	0.0
	SS	0	0	(0 0	C) (0 0	0	0	0	1	0	0	0	0	1	0.0
R_PF_ROI	SMG	0	0	(0 0	C) (0 0	0	0	0	0	0	0	1	0	1	0.0
R_PGi_ROI	AG	0	0	(0 0	C) (0 0	0	0	0	4	0	0	4	0	8	0.0
R_A4_ROI	STG	0	0	(0 0	C) (0 0	1	0	0	0	0	0	0	0	1	0.0
R_STSva_ROI	MTG	0	0	(0 0	C) (0 0	0	3	0	2	0	0	0	0	5	0.0
R_TE1m_ROI	ITG	0	0	(0 0	C) (0 0	0	0	1	0	0	0	0	0	1	0.0
R_a32pr_ROI	ACC	0	0	(0 0	C) (0 0	0	0	1	1	0	0	0	0	2	0.0
R_p24_ROI	ACC	0	0	(0 0	C) (0 0	0	0	0	1	0	0	0	0	1	0.0
L_FFC_ROI	CS	0	0	(0 0	C) (0 0	0	0	0	0	0	1	0	0	1	0.0
	MCC	0	0		0 0	C		0 0	0	0	0	0	0	0	3	0	3	0.0
	SPL	0	0	. (0	C) (0 0	1	0	0	0	2	0	0	0	3	0.0
	SPL	0	0		0 0			0 0		0	0	0	1	0	0	0	1	0.0
	MCC	0	0		0 0			0 0		0	0	0	0	0	1	0	1	0.0
	ORB	0	0					0 0		0	0	0	2	0	0	0	2	
	IFG	0	0		0 0			0 0		0	0	0	4	0	5	0	9	
	IFG	0	0					0 0		1	0	0	2	0	0	0	3	
	ORB	0	0		0			0 0		0	0	0	2	0	1	0	3	
	OFC	0	0		0 0			0 0		0	0	0	2	0	2	0	4	0.0
	SMG	0	0		0 0			0 0		0	0	0	1	1	0	0	2	
	INS	0	0		0 0			0 0		0	0	0		0	0	0	1	
	IFG	0	0		0 0			0 0		0	0	0	1	0	0	0	1	0.0
		0	0										1			0		
	INS				0 0			0 0	-	1	1	0	0	0	0		2	
	TP	0	0		0			0 0		0	0	0	0	2	0	0	2	
	STG	0	0		0			0 0		0	0	0	0	0	0	0	1	0.0
	MTG	0	0		0			0 0		0	0	0	1	0	0	0	1	
	TP	0	0		0			0 0		0	3	0	0	1	0			
	MTG	0	0		0			0 0		0	1	0	0	2	0	0	5	
	MTG	0	0		0			0 0		1	2	0	2	2	2	0	9	
	ITG	0	0					0 0		0	0	0	0	1	0	0	1	
	ITG	0	0		0			0 0		0	0	0	0	2	0	0	2	
L_PF_ROI	SMG	0	0	(0 0	C) (0 0	0	0	0	0	0	2	0	0	2	0.0
	SMG	0	0		0 0			0 0		0	0	0	0	0	1	0	1	
	PMC	0	0	(0 0	C) (0 0	1	1	0	0	1	0	1	0	4	0.0
L_VVC_ROI	FG	0	0	(0 0	C)	0 0	0	0	0	0	0	3	0	0	3	0.0
	INS	0	0	(0 0	C) (0 0	0	0	0	0	0	0	1	0	1	0.0
	STG	0	0	(0 0	C) (0 0	1	0	0	0	0	0	0	0	1	0.0
	STG	0	0		0 0			0 0		0	0	0	1	0	0	0	1	
	MTG	0	0		0			0 0		0	0	0	3	0	1	0	4	
	MTG	0	0		0 0			0 0		2	0	0	0	3	0	0	5	
	INS	0			0 0			0 0		0	0	0	2	0	1	0	3	
L_PI_ROI																		

Table S1. Anatomical detail about human iEEG changes on ketamine, by contact, and parcellated by Human Connectome Project (HCP) area. For each area, the number of total recorded contacts are specified for each subject and overall, and the number of recorded contacts that exhibit a significant ketamine-elicited spectrotemporal change in the puff-triggered spectrogram, as determined by permutation cluster test across trials paired between preinfusion and infusion (from 12 to 39 trials depending on which subject a contact came from). Top table specifies regions with at least one recorded contact that exhibits significant changes, and the bottom table specifies regions with no significantly changing contacts.

Acronym			
	Full name	Unit count	Session count
AAA	Anterior amygdalar area	23	1
ACAd1	Anterior cingulate area, dorsal part, layer 1	9	1
	<u> </u>		
ACAd2/3	Anterior cingulate area, dorsal part, layer 2/3	39	4
ACAd5	Anterior cingulate area, dorsal part, layer 5	153	6
ACAd6b	Anterior cingulate area, dorsal part, layer 6b	3	1
ACAv1	Anterior cingulate area, ventral part, layer 1	43	4
ACAv2/3		77	4
	Anterior cingulate area, ventral part, layer 2/3		
ACAv5	Anterior cingulate area, ventral part, layer 5	77	3
ACAv6a	Anterior cingulate area, ventral part, 6a	4	1
ACB	Nucleus accumbens	317	8
Ald5	Agranular insular area, dorsal part, layer 5	153	2
Ald6a	Agranular insular area, dorsal part, layer 6a	70	2
Alv2/3	Agranular insular area, ventral part, layer 2/3	36	2
Alv5	Agranular insular area, ventral part, layer 5	171	2
AON	Anterior olfactory nucleus	89	2
BLAa	Basolateral amygdalar nucleus, anterior part	1	1
BLAv	Basolateral amygdalar nucleus, ventral part	1	1
BMAp	Basomedial amygdalar nucleus, posterior part	1	1
BST	Bed nuclei of the stria terminalis	35	3
CA1	Field CA1	146	7
CA2	Field CA2	5	2
CA3	Field CA3	57	6
CEAm	Central amygdalar nucleus, medial part	13	1
CLI	Central linear nucleus raphe	9	1
CM	Central medial nucleus of the thalamus	20	2
COApl	Cortical amygdalar area, posterior part, lateral zone	1	1
COApm	Cortical amygdalar area, posterior part, medial zone	26	2
CP	Caudoputamen	516	7
DG-mo	Dentate gyrus, molecular layer	87	6
		10	4
DG-po	Dentate gyrus, polymorph layer		
DG-sg	Dentate gyrus, granule cell layer	29	7
DP	Dorsal peduncular area	37	3
EPd	Endopiriform nucleus, dorsal part	135	1
Eth	Ethmoid nucleus of the thalamus	16	1
FF	Fields of Forel	59	4
FRP6a	Frontal pole, layer 6a	22	1
FS	Fundus of striatum	10	1
HPF	Hippocampal formation	9	3
HY	Hypothalamus	59	7
mr	nypothalamus		
IF	Interfascicular nucleus raphe	4	1
IF IG	Interfascicular nucleus raphe Induseum griseum		1
		4	
IG IGL	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex	4 13 15	1 2
IG IGL ILA1	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1	4 13 15 46	1 2 1
IG IGL ILA1 ILA2/3	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3	4 13 15 46 11	1 2 1 1
IG IGL ILA1	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1	4 13 15 46	1 2 1
IG IGL ILA1 ILA2/3	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3	4 13 15 46 11	1 2 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5	4 13 15 46 11 26	1 2 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core	13 15 46 11 26 1	1 2 1 1 1 1 1 1 3
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infrailmbic area, layer 1 Infrailmbic area, layer 2/3 Infrailmbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone	4 13 15 46 11 26 1 12	1 2 1 1 1 1 1 1 3 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infrailmbic area, layer 1 Infrailmbic area, layer 2/3 Infrailmbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell	4 13 15 46 11 26 1 12 12 13	1 2 1 1 1 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infrailmbic area, layer 1 Infrailmbic area, layer 2/3 Infrailmbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone	4 13 15 46 11 26 1 12	1 2 1 1 1 1 1 1 3 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infrailmbic area, layer 1 Infrailmbic area, layer 2/3 Infrailmbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell	4 13 15 46 11 26 1 12 12 13	1 2 1 1 1 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGv	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex.	4 13 15 46 11 26 1 12 12 13 28	1 2 1 1 1 1 1 3 1 1 1 3 3 1 1 1 3 3 1 1 1 3 3 1 1 1 3 3 1 1 1 3 3 1 1 1 1 1 3 3 1 1 1 1 1 3 3 1 1 1 1 1 1 3 3 1 1 1 1 1 3 3 1 1 1 1 1 3 3 1
IG IGL ILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGV LH LHA	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex complex Lateral habenula Lateral hypothalamic area	4 13 15 46 11 26 1 12 12 13 28 18	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus	4 13 15 46 111 26 1 12 12 13 28 18 25 102	1 2 1 1 1 1 1 1 1 3 3 1 1 1 1 3 3 2 2 5 5
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hyborhalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part	4 13 15 46 11 1 26 1 12 12 13 28 18 25 102 8	1 2 1 1 1 1 1 1 1 1 3 3 1 1 1 1 3 3 2 2 5 5 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus	4 13 15 46 111 26 1 12 12 13 28 18 25 102	1 2 1 1 1 1 1 1 1 3 3 1 1 1 1 3 3 2 2 5 5
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hyborhalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part	4 13 15 46 11 1 26 1 12 12 13 28 18 25 102 8	1 2 1 1 1 1 1 1 1 1 3 3 1 1 1 1 3 3 2 2 5 5 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSv	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infrailmbic area, layer 1 Infrailmbic area, layer 2/3 Infrailmbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part	4 13 15 46 11 1 12 12 12 13 28 18 25 102 8 521	1 2 1 1 1 1 1 1 1 1 3 3 1 1 1 1 3 3 3 2 2 5 5 1 1 7 5 5
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSc LSr LSv MB	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex complex Lateral habenula Lateral habenula Lateral hypothalamic area Lateral septal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral (rostroventral) part Lateral septal nucleus, ventral (rostroventral) part Midbrain	4 13 15 46 111 26 1 1 2 12 12 13 28 18 25 102 8 8 521 64 310	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IIMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSr LSv MB	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ispallateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral soptal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus	4 13 15 46 11 26 1 12 12 13 28 18 25 102 8 521 64 310 240	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSc LSr LSv MB	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex complex Lateral habenula Lateral habenula Lateral hypothalamic area Lateral septal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral (rostroventral) part Lateral septal nucleus, ventral (rostroventral) part Midbrain	4 13 15 46 111 26 1 1 2 12 12 13 28 18 25 102 8 8 521 64 310	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IIMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSr LSv MB	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ispallateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral soptal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus	4 13 15 46 11 26 1 12 12 13 28 18 25 102 8 521 64 310 240	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr LSv MB MD MEPO MGd	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Median preoptic nucleus Median preoptic nucleus	4 13 15 46 11 26 1 12 12 13 28 28 18 25 102 8 521 64 310 240	1 2 1 1 1 1 1 1 3 3 1 1 1 1 3 3 2 2 5 5 1 1 7 7 5 8 8 4 1 1 1 1 1 1
IG IGL IILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGv LH LHA LP LSc LSr LSv MB MD MEPO MGd MGm	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 2/3 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Midbrain Mediodorsal nucleus of thalamus Median prepoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, dorsal part	4 13 15 46 11 26 1 1 22 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 ILA6 IMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSc MB MD MEPO MGG MGG IGA1	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Lateral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral hypothalamic area Lateral septal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, costral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, medial part	4 13 15 46 11 12 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3	1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL IILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGv LH LHA LP LSc LSr LSv MB MD MEPO MGd MGm	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 2/3 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Midbrain Mediodorsal nucleus of thalamus Median prepoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, dorsal part	4 13 15 46 11 26 1 1 22 12 13 28 18 25 102 8 521 64 310 240 1	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 ILA6 IMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSc MB MD MEPO MGG MGG IGA1	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Infralimbic area, layer 5/3 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Lateral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral hypothalamic area Lateral septal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, costral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, medial part	4 13 15 46 11 12 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3	1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSv MB MD MEPO MGd MGd MGv MOp2/3	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral hypothalamic area Lateral hypothalamic area Lateral septal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, medial part Medial geniculate complex, medial part Primary motor area, Layer 2/3	4 13 15 46 11 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 4 1 1 1 1 2 1 1 1 1 1 1 1 1 1 1 1 1 1	1 2 1 1 1 1 3 3 3 2 2 5 5 1 1 7 7 5 8 8 4 1 1 1 1 3 3 3 3 3 3 3 3 3 3 3 3 3 3 3
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr LSv MB MD MEPO MGd MGm MGW MOp2/3 MOp5 MOp6a	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Median preoptic nucleus Medial geniculate complex, medial part Medial geniculate complex, medial part Medial geniculate complex, ventral part Primary motor area, Layer 5 Primary motor area, Layer 5	4 13 15 46 11 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 1 1 21 79 113	1 2 1 1 1 1 1 1 3 3 3 3 3 4 4 1 1 1 1 1 3 3 3 3
IG IGL IILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr MB MD MEPO MGd MGm MGW MOp2/3 MOp6a MOp6b	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Ventral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral hosterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, dorsal part Medial geniculate complex, ventral part Primary motor area, Layer 5 Primary motor area, Layer 6a Primary motor area, Layer 6b	4 13 15 46 11 26 1 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1 1	1 2 1 1 1 1 1 3 3 3 4 4 3 3
IG IGL ILA1 ILA2/3 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr LSv MB MD MEPO MGd MGm MGW MOp2/3 MOp5 MOp6a	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Median preoptic nucleus Medial geniculate complex, medial part Medial geniculate complex, medial part Medial geniculate complex, ventral part Primary motor area, Layer 5 Primary motor area, Layer 5	4 13 15 46 11 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 1 1 21 79 113	1 2 1 1 1 1 1 1 3 3 3 3 3 4 4 1 1 1 1 1 3 3 3 3
IG IGL IILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr MB MD MEPO MGd MGm MGW MOp2/3 MOp6a MOp6b	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex shell Ventral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral hosterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, dorsal part Medial geniculate complex, ventral part Primary motor area, Layer 5 Primary motor area, Layer 6a Primary motor area, Layer 6b	4 13 15 46 11 26 1 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1 1	1 2 1 1 1 1 1 3 3 3 4 4 3 3
IG IGL ILA1 ILA2/3 ILA5 ILA5 IMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSv MB MD MEPO MGG MGG MGPS MOpSa MOpSa MOpSa	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral habenula Lateral habenula Lateral septal nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, wentral part Primary motor area, Layer 2/3 Primary motor area, Layer 6 Primary motor area, Layer 6 Secondary motor area, layer 5	4 13 15 46 11 12 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1 1	1 1 2 1 1 1 1 3 3 3 2 5 5 1 1 7 7 5 8 4 4 1 1 1 1 3 3 3 3 4 4 3 3 5 5
IG IGL ILA1 ILA2/3 ILA5 ILA5 IMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSv MB MD MEPO MGd MGm MGV MOp2/3 MOp56 MOp66 MOp66 MCN	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, rostral (rostroventral) part Midbrain Mediodorsal nucleus of thalamus Medial geniculate complex, dorsal part Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, medial part Medial geniculate complex, wentral part Primary motor area, Layer 2/3 Primary motor area, Layer 6b Secondary motor area, Layer 6b Secondary motor area, layer 5 Secondary motor area, layer 6a Midbrain reticular nucleus	4 13 15 46 11 12 12 13 28 18 25 102 8 521 64 310 240 1 1 1 21 79 113 10 39 91 100	1 1 2 1 1 1 1 3 3 3 2 2 5 5 1 1 7 7 5 8 8 4 4 1 1 1 1 3 3 3 3 4 4 3 3 5 5 4 4 2 2
IG IGL IILA1 IILA2/3 IILA5 IIMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSr LSv MB MD MEPO MGd MGm MGy MOp5/3 MOp5a MOp6a MOp6b MOs5a MRN ND	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, pelateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral spetal nucleus, caudal (caudodorsal) part Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, wentral part Primary motor area, Layer 2/3 Primary motor area, Layer 6 Primary motor area, Layer 6 Secondary motor area, layer 6 Secondary motor area, layer 5 Secondary motor area, layer 6a Midbrain reticular nucleus Nucleus of Darkschewitsch	4 13 15 46 11 26 1 12 12 13 28 18 25 102 8 521 64 310 240 1 1 1 21 79 113 10 39 91 100 7	1 1 2 1 1 1 1 1 3 3 3 2 2 5 5 1 1 7 7 5 5 8 8 4 4 1 1 1 1 1 3 3 3 3 4 4 3 3 5 5 4 4 2 2 1 1
IG IGL IILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr MB MD MEPO MGd MGm MGC MOp6a MOp6a MOp6b MOs6a MOs6a MRN ND NLOT1	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, ventral part Primary motor area, Layer 2/3 Primary motor area, Layer 6/b Secondary motor area, Layer 6a Midbrain reticular nucleus Nucleus of Darkschewitsch Nucleus of the lateral olfactory tract, molecular layer	4 13 15 46 11 26 1 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 1 1 2 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1 1	1 2 1 1 1 1 1 3 3 3 4 4 3 3 5 4 4 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1
IG IGL IILA1 IILA2/3 IILA5 IIMD LGd-co LGd-lp LGd-sh LGV LH LHA LP LSc LSr LSv MB MD MEPO MGd MGm MGy MOp5/3 MOp5a MOp6a MOp6b MOs5a MRN ND	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, pelateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral hypothalamic area Lateral spetal nucleus, caudal (caudodorsal) part Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, rostral (rostroventral) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, wentral part Primary motor area, Layer 2/3 Primary motor area, Layer 6 Primary motor area, Layer 6 Secondary motor area, layer 6 Secondary motor area, layer 5 Secondary motor area, layer 6a Midbrain reticular nucleus Nucleus of Darkschewitsch	4 13 15 46 11 26 1 12 12 13 28 18 25 102 8 521 64 310 240 1 1 1 21 79 113 10 39 91 100 7	1 1 2 1 1 1 1 1 3 3 3 2 2 5 5 1 1 7 7 5 5 8 8 4 4 1 1 1 1 1 3 3 3 3 4 4 3 3 5 5 4 4 2 2 1 1
IG IGL IILA1 ILA2/3 ILA5 IIMD LGd-co LGd-ip LGd-sh LGV LH LHA LP LSc LSr LSr MB MD MEPO MGd MGm MGC MOp6a MOp6a MOp6b MOs6a MOs6a MRN ND NLOT1	Induseum griseum Intergeniculate leaflet of the lateral geniculate complex Infralimbic area, layer 1 Infralimbic area, layer 2/3 Infralimbic area, layer 5 Intermediodorsal nucleus of the thalamus Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, core Dorsal part of the lateral geniculate complex, ipsilateral zone Dorsal part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex, shell Ventral part of the lateral geniculate complex Lateral habenula Lateral habenula Lateral posterior nucleus of the thalamus Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, caudal (caudodorsal) part Lateral septal nucleus, ventral part Midbrain Mediodorsal nucleus of thalamus Median preoptic nucleus Medial geniculate complex, dorsal part Medial geniculate complex, medial part Medial geniculate complex, ventral part Primary motor area, Layer 2/3 Primary motor area, Layer 6/b Secondary motor area, Layer 6a Midbrain reticular nucleus Nucleus of Darkschewitsch Nucleus of the lateral olfactory tract, molecular layer	4 13 15 46 11 26 1 12 12 12 13 28 18 25 102 8 521 64 310 240 1 1 3 1 1 2 1 1 3 1 1 1 1 1 1 1 1 1 1 1 1 1	1 2 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1 1

		Template	e revise
Acronym	Full name	Unit count	Session coun
NPC	Nucleus of the posterior commissure	106	6
OLF	Olfactory areas	277	9
OP	Olivary pretectal nucleus	20	2
ORBI1	Orbital area, lateral part, layer 1	22	1
ORBI2/3	Orbital area, lateral part, layer 2/3	118	4
ORBI5	Orbital area, lateral part, layer 5	360	5
ORBI6a	Orbital area, lateral part, layer 6a	100	3
ORBm5	Orbital area, medial part, layer 5	50	1
ORBm6a	Orbital area, medial part, layer 6a	18	1
ORBvI2/3	Orbital area, ventrolateral part, layer 2/3	111	2
ORBvl5	Orbital area, ventrolateral part, layer 5	57	2
OT	Olfactory tubercle	91	2
PA	Posterior amygdalar nucleus	15	1
PAG	Periaqueductal gray	159	5
PCN	Paracentral nucleus	23	1
PF	Parafascicular nucleus	2	1
PIR	Piriform area	212	3
PL1	Prelimbic area, layer 1	49	1
PL2/3	Prelimbic area, layer 2/3	85	1
PL5	Prelimbic area, layer 5	128	3
PL6a	Prelimbic area, layer 6a	11	1
PO	Posterior complex of the thalamus	345	5
PPT	Posterior pretectal nucleus	7	1
PRC	Precommissural nucleus	13	2
PS	Parastrial nucleus	3	1
PSTN	Parasubthalamic nucleus	25	2
PVT	Paraventricular nucleus of the thalamus	1	1
PoT	Posterior triangular thalamic nucleus	95	2
RPF	Retroparafascicular nucleus	11	1
SPagl2/3	Retrosplenial area, lateral agranular part, layer 2/3	8	1
RSPagl5	Retrosplenial area, lateral agranular part, layer 5	20	1
SPagl6a	Retrosplenial area, lateral agranular part, layer 6a	7	1
SPagl6b	Retrosplenial area, lateral agranular part, layer 6b	2	1
RSPd2/3	Retrosplenial area, dorsal part, layer 2/3	3	1
RSPd5	Retrosplenial area, dorsal part, layer 5	11	3
RSPd6a	Retrosplenial area, dorsal part, layer 6a	25	2
RSPv1	Retrosplenial area, ventral part, layer 1	35	5
RSPv2/3	Retrosplenial area, ventral part, layer 2/3	51	5
RSPv5	Retrosplenial area, ventral part, layer 5	243	6
RSPv6a	Retrosplenial area, ventral part, layer 6a	89	3
RSPv6b	Retrosplenial area, ventral part, layer 6b	12	3
RT	Reticular nucleus of the thalamus	46	1
SCdg	Superior colliculus, motor related, deep gray layer	10	1
SCdw	Superior colliculus, motor related, deep white layer	8	1
SCig	Superior colliculus, motor related, intermediate gray layer	50	2
SCiw	Superior colliculus, motor related, intermediate white layer	25	2
SCop	Superior colliculus, optic layer	8	2
SCsg	Superior colliculus, superficial gray layer	15	2
SCzo	Superior colliculus, zonal layer	9	3
SF	Septofimbrial nucleus	40	1
SI	Substantia innominata	79	4
SNc	Substantia nigra, compact part	6	1
SNr	Substantia nigra, reticular part	3	1
SPA	Subparafascicular area	8	2
SPFp	Subparafascicular nucleus, parvicellular part	8	1
SSp-II6a	Primary somatosensory area, lower limb, layer 6a	35	1
	Primary somatosensory area, lower limb, layer 6b	3	1
SSp-II6b			
SSp-II6b SSp-m1	Primary somatosensory area, mouth, layer 1	4	1
SSp-II6b SSp-m1			1
SSp-II6b SSp-m1 Sp-m2/3	Primary somatosensory area, mouth, layer 1	4	
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3	7	1
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m5	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4	4 7 16	1
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m5 SSp-m6a	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5	4 7 16 17	1 1
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m5 SSp-m6a SSp-uI6a	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a	4 7 16 17 40	1 1 1
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m5 SSp-m6a SSp-u16a SSp-u16b	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a	4 7 16 17 40 8	1 1 1 1
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m5 SSp-m6a SSp-u16a SSp-u16b	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a	4 7 16 17 40 8 1	1 1 1 1 1
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m6 SSp-m6 SSp-m6 SSp-m6 SSp-m1 SSp-m6 SSp-u16b SSS6b	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b	4 7 16 17 40 8 1 1 1	1 1 1 1 1 1
\$\$p-ll6b \$\$p-m1 \$p-m2/3 \$\$p-m4 \$\$p-m6 \$\$p-m6 \$\$p-m6 \$\$p-m6a \$\$p-ul6a \$\$p-ul6b \$\$ss6b \$\$TN	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Subthalamic nucleus Striatum	4 7 16 17 40 8 1 1 1 19 59	1 1 1 1 1 1 1 1 1 1 9
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m4 SSp-m6 SSp-m6 SSp-m6a SSp-m6a SSp-ul6a SSp-ul6b STN STR SUB	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upuper limb, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Subthalamic nucleus Striatum Subiculum	4 7 16 17 40 8 1 1 1 19 59 67	1 1 1 1 1 1 1 1 1 9
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m5 SSp-m6 SSp-m6a SSp-m16a SSp-u16a SSp-u16b SS-s6b STN STR SUB TH	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Subthalamic nucleus Striatum Subiculum Thalamus	4 7 16 17 40 8 1 1 1 19 59 67	1 1 1 1 1 1 1 1 1 1 9 3
SSp-II6b SSp-m1 Sp-m2/3 SSp-m4 SSp-m4 SSp-m6 SSp-m6 SSp-m6 SSp-m6 SSp-m6 SSp-u16a SSp-u16b SSs6b STN STR SUB TH	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Subthalamic nucleus Striatum Subiculum Thalamus Taenia tecta, dorsal part	4 7 16 17 40 8 1 1 1 19 59 67 62	1 1 1 1 1 1 1 1 1 9 3 6
\$\$p.116b \$\$p.m12 \$\$p.m23 \$\$p.m23 \$\$\$p.m23 \$\$\$\$p.m4 \$\$\$\$p.m5 \$\$\$p.m65 \$\$\$p.m65 \$\$\$p.m65 \$\$\$\$p.m66 \$\$\$\$p.m66 \$\$\$\$\$p.m66 \$	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Supplemental somatosensory area, layer 6b Subthalamic nucleus Striatum Subiculum Thalamus Taenia tecta, dorsal part Ventral anterior-lateral complex of the thalamus	4 7 16 17 40 8 1 1 1 19 59 67 62 14	1 1 1 1 1 1 1 1 1 1 9 3 6
\$\$p.116b \$\$p.m17 \$\$p.m273 \$\$p.m273 \$\$p.m273 \$\$p.m273 \$\$p.m61 \$\$p.m61 \$\$p.m61 \$\$p.m61 \$\$p.m61 \$\$p.u16b \$\$p.u16b \$\$p.u16b \$\$TN \$\$TR \$\$UB \$\$TH \$\$TH \$\$TTT \$\$VAL \$\$VPLpc	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 5 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Supplemental somatosensory area, layer 6b Subtitaliamic nucleus Striatum Subiculum Thalamus Taenia tecta, dorsal part Ventral anterior-lateral complex of the thalamus Ventral posterolateral nucleus of the thalamus, parvicellular part	4 7 16 17 40 8 1 1 1 19 59 67 62 14 40	1 1 1 1 1 1 1 1 1 1 9 3 6 3 2 2
\$\$p.116b \$\$p.m1 \$\$p.m273 \$\$p.m273 \$\$\$p.m273 \$\$\$\$p.m4 \$\$\$\$p.m273 \$\$\$\$p.m4 \$\$\$\$p.m4 \$\$\$\$p.m61 \$\$\$\$p.m61 \$\$\$\$p.m61 \$\$\$\$\$p.m61 \$\$\$\$\$\$\$p.u160 \$	Primary somatosensory area, mouth, layer 1 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 2/3 Primary somatosensory area, mouth, layer 4 Primary somatosensory area, mouth, layer 6a Primary somatosensory area, mouth, layer 6a Primary somatosensory area, upper limb, layer 6a Primary somatosensory area, upper limb, layer 6b Supplemental somatosensory area, layer 6b Supplemental somatosensory area, layer 6b Subthalamic nucleus Striatum Subiculum Thalamus Taenia tecta, dorsal part Ventral anterior-lateral complex of the thalamus	4 7 16 17 40 8 1 1 1 19 59 67 62 14	1 1 1 1 1 1 1 1 1 1 9 3 6

Table S2. Anatomical details about mouse Neuropixels recordings. Areas parcellated according to Allen Brain Institute Reference Atlas (CCFv3).

Table S3.

Summary of statistical analyses

Fig. 1D

Analysis: Puff-triggered eye closure of an example human subject, averaged across trials, comparing saline and ketamine (0.5 mg/kg) conditions.

Condition	# trials	Plot
Saline	38	Mean \pm s.e.m.
Ketamine	38	Mean \pm s.e.m.

Fig. 1E

Analysis: Normalized late eye closure (late divided by early) in humans, saline vs. ketamine. Point estimates for each subject are mean \pm s.e.m. across trials, excluding the first two trials.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# subjects	# trials per subject	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	4	38	0.71 ± 0.08	Paired t-test	2 702	0.0342
Ketamine	4	38	0.43 ± 0.12	raired t-test	3.703	0.0342

Fig. 1J

Analysis: Puff-triggered eye closure of an example mouse, averaged across the first puff of each trial (trials consist of 8 closely-spaced puffs), comparing saline and ketamine (50 mg/kg) conditions.

Condition	# trials	Plot
Saline	15	Mean \pm s.e.m.
Ketamine	15	Mean \pm s.e.m.

Fig. 1K

Analysis: Normalized late eye closure (late divided by early) in mice for first puff of each trial, saline vs. ketamine.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	5	15	0.26 ± 0.06	D : 144 4	4.727	0.000
Ketamine	5	15	-0.07 ± 0.06	Paired t-test	4.737	0.009

Fig. 1L

Analysis: Normalized late eye closure (late divided by early) in mice for first puff of each trial, preinfusion vs. infusion for saline, ketamine (50 mg/kg), and PCP (20 mg/kg).

Null hypothesis: No difference between preinfusion and infusion.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value	
Pre-saline	5	15	0.29 ± 0.08		Pre-saline vs. saline	0.73	0.504	
Saline	5	15	0.26 ± 0.06	D: 144 4-14 C1 11	Fie-samile vs. samile	0.73	0.304	
Pre- ketamine	5	15	0.37 ± 0.06	Paired t-test with fdr-bh correction	Pre-ketamine vs.	9.53	0.002	
Ketamine	5	15	-0.07 ± 0.06		ketamine			

D DCD DCD	D.,, DCD DCD 2.41
Pre-PCP vs. PCP	Pre-PCP vs. PCP 3.41
	3.41

Fig. 1M

Analysis: Average eye closure across trials (a trial consists of 8 puffs each separated by 3 s), averaged across mice, comparing preinfusion and saline infusion conditions.

Condition	# mice	# trials per mouse	Plot
Preinfusion	5	15	mean \pm s.e.m.
Saline	5	15	mean \pm s.e.m.

Fig. 1N

Analysis: Average eye closure across trials (a trial consists of 8 puffs each separated by 3 s), averaged across mice, comparing preinfusion and ketamine (50 mg/kg) infusion conditions.

Condition	# mice	# trials per mouse	Plot
Preinfusion	5	15	mean \pm s.e.m.
Ketamine	5	15	mean \pm s.e.m.

Fig. 10

Analysis: Average eye closure across trials (a trial consists of 8 puffs each separated by 3 s), averaged across mice, comparing preinfusion and PCP (20 mg/kg) infusion conditions.

Condition	# mice	# trials per mouse	Plot	
Preinfusion	5	15	mean \pm s.e.m.	
PCP	5	15	mean \pm s.e.m.	

Fig. 1P

Analysis: Average eye closure across trials (a trial consists of 8 puffs each separated by 3 s), averaged across mice, comparing preinfusion and anesthetic dose of K/X (135 mg/kg ketamine with 15 mg/kg xylazine).

Condition	# mice	# trials per mouse	Plot
Preinfusion	5	15	mean \pm s.e.m.
K/X	5	15	mean \pm s.e.m.

Fig. 2C

Analysis: Average puff-triggered eye closure comparing preinfusion, infusion, and postinfusion conditions.

Condition	# human participants	# trials per subject	Plot
Preinfusion	7	mean: 30.6, min: 12, max: 39	mean \pm s.e.m.
Infusion	7	mean: 30.6, min: 12, max: 39	mean \pm s.e.m.
Postinfusion	7	mean: 30.6, min: 12, max: 39	mean \pm s.e.m.

Fig. 2D

Analysis: Clinician administered dissociative states score (CADSS) before, during, and after ketamine infusion across the subjects that met the eyepuff behavioral inclusion criteria.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Preinfusion	7	0.86 ± 0.46	D-i1 4 44i41- £1 1-1-			
Ketamine	7	22.43 ± 4.75	Paired t-test with fdr-bh correction	Preinfusion vs. ketamine	4.807	0.004

Postinfusion	7	0.86 ± 0.46		Postinfusion vs. ketamine	4.512	0.004
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Fig. 2E

Analysis: Preinfusion event related potential for example channels from insula (INS) and posteromedial cortex (PMC), both from subject SD172. Mean across trials shown in teal, individual trials in gray. All trials, excluding the first trial, are shown (39 trials).

Fig. 2G

Analysis: Factor projections, corresponding to matrix H in a 6-component non-negative matrix factorization D = WH. The procedure is schematized in 2F. The matrix D is of shape [channels x frequency x time], formed by stacking the mean spectrogram across a training set of half of the preinfusion and postinfusion trials for all channels from all subjects (458 total channels). W is [channels x 6], and H is [6 x frequency x time]. NMF is applied by first splitting D into positive and negative parts, and computing NMF with k=3 on each part. Only channels localized to the 21 regions that were sampled in at least two subjects were included (ORB, IFG, MFG, INS, ACC, MCC, PMC, MOT, SS, SMG, PHG, STG, STS, MTG, ITG, TP, HIPP, BG, AMY, THAL ANT, THAL POS). The total number of channels was 458, with a breakdown by subject as follows: SD168=86, SD172=97, SD181=58, SD185=43, SD186=47, SD211=44, SD245=83. The number of components was selected by using cross-validation on a test set of trials, and identifying an elbow in the test set explained variance, which occurred at K=6.

For subsequent analyses, factor loadings for each channel, corresponding to rows in the matrix W are computed for each condition (e.g. preinfusion or infusion) by projecting H on to the D corresponding to that condition. Here, D is formed by averaging the spectrogram across a test set of half of the trials of that condition (and that are distinct from the training set of trials used to compute the factorization). Loadings represent the extent to which a factor is 'active' in the neural response of that channel. For each factor, channels with a loading of less than 0.1 on the average across preinfusion and postinfusion test set trials were excluded from subsequent analysis for that factor.

Fig. 2H

Analysis: Factor loadings, corresponding to W in NMF factorization D = WH, where W is [channels x 6], averaged across channels grouped by Yeo7 network. Loadings are then normalized within each row (divided by the sum of the row).

Fig. 2I

Analysis: Factor loading before, during, and after infusion, mean \pm s.e.m. across channels. Mixed effect linear model, channels grouped by subject, [pre + post]/2 vs. ketamine (for channels with factor loading across the average of preinfusion and postinfusion test trials of at least 0.1).

Null hypothesis: No difference between infusion and average of preinfusion and postinfusion.

Condition	# human participants	# channels	Mean ± s.e.m.	Statistical test	Comparison	Coefficient	Corrected p- Value
Factor 0 Pre. Factor 0 Inf. Factor 0 Post.	7	312	0.48 ± 0.02 0.42 ± 0.02 0.45 ± 0.02	Mixed-	(Pre + Post)/2 vs. Infusion	-0.049	0.121
Factor 1 Pre. Factor 1 Inf. Factor 1 Post.	7	260	$0.51 \pm 0.02 \\ 0.32 \pm 0.02 \\ 0.38 \pm 0.02$	subject, with idr-bh correction across factors	(Pre + Post)/2 vs. Infusion	-0.125	2.56e-05
Factor 2 Pre.	7	185	0.35 ± 0.02			-0.115	5.51e-08

				inplace revised froveinger 2025		
Factor 2 Inf. Factor 2 Post.			$0.21 \pm 0.01 \\ 0.30 \pm 0.02$	(Pre + Post)/ Infusion		
Factor 3 Pre. Factor 3 Inf. Factor 3 Post.	5	158	0.28 ± 0.01 0.21 ± 0.01 0.23 ± 0.01	(Pre + Post)/ Infusion	-0.041	0.016^{\dagger}
Factor 4 Pre. Factor 4 Inf. Factor 4 Post.	6	370	0.32 ± 0.01 0.39 ± 0.01 0.42 ± 0.01	(Pre + Post)/ Infusion		0.406
Factor 5 Pre. Factor 5 Inf. Factor 5 Post.	7	378	0.39 ± 0.01 0.43 ± 0.01 0.44 ± 0.01	(Pre + Post)/ Infusion	1 (1)(1)(3	0.341

[†]Mixed-effect model, grouped by subject, did not converge for Factor 3. Linear model without grouping by subject does converge and yields a corrected p-value of 0.023.

As a post-hoc test, we further assessed the comparison between infusion and preinfusion and separately between infusion and postinfusion.

Null hypothesis: No difference between infusion and either preinfusion or postinfusion.

Condition	# human participants	# channels	Mean ± s.e.m.	Statistical test	Comparison	Coefficient	Corrected p- Value
Factor 0 Pre.	7	312	0.48 ± 0.02	Mixed- effect linear model, grouped by subject,	Pre vs. Infusion	0.065	0.036
Factor 0 Inf.			0.42 ± 0.02	with fdr-bh correction across factors	Post vs. Infusion	0.033	0.304
Factor 0 Post.			0.45 ± 0.02				
Factor 1 Pre.	7	260	0.51 ± 0.02		Pre vs. Infusion	0.190	1.80e-09
Factor 1 Inf.			0.32 ± 0.02		Post vs. Infusion	0.061	0.046
Factor 1 Post.			0.38 ± 0.02				
Factor 2 Pre.	7	185	0.35 ± 0.02		Pre vs. Infusion	0.140	1.25e-08
Factor 2 Inf.			0.21 ± 0.01		Post vs. Infusion	0.089	5.16e-05
Factor 2 Post.			0.30 ± 0.02				
Factor 3 Pre.	5	158	0.28 ± 0.01		Pre vs. Infusion	0.068	5.16e-04

Factor 3 Inf.			0.21 ± 0.01	•	Post vs. Infusion	0.014	0.394
Factor 3 Post.			0.23 ± 0.01				
Factor 4 Pre.	6	370	0.32 ± 0.01		Pre vs. Infusion	0.063	2.99e-04
Factor 4 Inf.			0.39 ± 0.01		Post vs. Infusion	0.037	0.046
Factor 4 Post.			0.42 ± 0.01				
Factor 5 Pre.	7	378	0.39 ± 0.01		Pre vs. Infusion	0.038	0.010
Factor 5 Inf.			0.43 ± 0.01		Post vs. Infusion	0.011	0.394
Factor 5 Post.			0.44 ± 0.01				

Fig. 2J

Analysis: Change in factor loadings, ketamine – preinfusion, corresponding to W in NMF factorization D = WH, where W is [channels x 6], averaged over the first dimension across channels grouped by Yeo7 network. Factorization H is computed using the average across preinfusion and postinfusion trials as described above, and used to compute W during ketamine and preinfusion.

Fig. 3B

Analysis: Permutation cluster test (mne.stats.permutation_cluster_test) comparing preinfusion and infusion puff-triggered spectrograms on each trial for each Yeo7 network (except for the Visual network, which was not present in our dataset). For each subject, all channels in a network were averaged into a single trace for each trial. All trials were stacked across subjects, yielding the total number of trials (which are indicated parenthetically for each network). The test statistic was a 1-sample t-test of the difference between each preinfusion and infusion trial (infusion – preinfusion). As the same eyepuff protocol was used during preinfusion and infusion (with the same ordering of interpuff interval), trials were paired to account for any within-puff-sequence trends. A pixel threshold of p <0.01 was used to identify clusters. At the cluster level, a threshold of p<0.01 was used to assign significance to a cluster. Fdr-bh multiple comparisons was used across the most significant (lowest p-value) cluster of each network. For results with additional test statistics and thresholds, see supplement.

Fig. 3C

Analysis: Same as previous, but with regions instead of Yeo7 networks. Regions were included if there were at least two subjects with representation of the region. Number of trials specified in parentheses.

Fig. 4B

Analysis: Average eye closure behavior on the first puff of each puff series, for pre and ketamine conditions. Traces are baseline subtracted with a 1s window before puffs. Mean and 95% confidence interval of the mean value is given (1.96 * s.e.m.), with trials averaged within session and variability taken across sessions.

Condition	# sessions	# trials	# subjects
Preinfusion	13	20	10
Ketamine	13	10	10

Fig. 4E

Analysis: Average cluster per-puff firing rate. Averages are baseline subtracted by average cluster activity 1s prior to airpuff onset. Cluster labels obtained in Fig. 3D. Mean cluster activity is z-scored across the peri-puff time window within cluster. Mean across all cells assigned to cluster, pooled across all datasets.

Condition	# cells	# airpuffs	# sessions	# subjects
Cluster 0	1070	160	13	10
Cluster 1	832	160	13	10
Cluster 2	803	160	13	10
Cluster 3	663	160	13	10
Cluster 4	639	160	13	10
Cluster 5	597	160	13	10
Cluster 6	547	160	13	10
Cluster 7	508	160	13	10
Cluster 8	473	160	13	10
Cluster 9	344	160	13	10
Cluster 10	317	160	13	10
Cluster 11	176	160	13	10

Fig 4G, fig. S31 A

Analysis: Average cluster per-puff firing rate compared between pre-infusion and ketamine infusion conditions. fig. S31 additionally shows post-infusion condition. Averages are baseline subtracted by average cluster activity 1s prior to airpuff onset. Cluster labels obtained in Fig. 4D. Mean cluster activity is z-scored across the peri-puff time window within cluster and condition. Mean across all cells assigned to cluster, pooled across all datasets. Mean \pm 95% confidence interval (1.96 * s.e.m.), with variability measured across all cells within cluster.

Condition	# cells	# airpuffs	# sessions	# subjects
Cluster 0 - Pre		160		
Cluster 0 - Ketamine	1070	64	13	10
Cluster 0 - Post		64		
Cluster 1 - Pre		160		
Cluster 1 - Ketamine	832	64	13	10
Cluster 1 - Post		64		
Cluster 2 - Pre		160		
Cluster 2 - Ketamine	803	64	13	10
Cluster 2 - Post		64		
Cluster 3 - Pre		160		
Cluster 3 - Ketamine	663	64	13	10
Cluster 3 - Post	003	64	13	10
Cluster 4 - Pre		160		
Cluster 4 - Ketamine	639	64	13	10
Cluster 4 - Post	037	64	13	10
Cluster 5 - Pre		160		
Cluster 5 - Ketamine	597	64	13	10
Cluster 5 - Post	371	64		10
Cluster 6 - Pre		160		
Cluster 6 - Ketamine	547	64	13	10
Cluster 6 - Post	517	64	15	10

Cluster 7 - Pre		160		
Cluster 7 - Ketamine	508	64	13	10
Cluster 7 - Post	300	64	13	10
Cluster 8 - Pre		160		
Cluster 8 - Ketamine	473	64	13	10
Cluster 8 - Post	175	64	13	10
Cluster 9 - Pre		160		
Cluster 9 - Ketamine	344	64	13	10
Cluster 9 - Post	511	64	13	10
Cluster 10 - Pre		160		
Cluster 10 - Ketamine	317	64	13	10
Cluster 10 - Post	317	64	13	10
Cluster 11 - Pre		160		
Cluster 11 - Ketamine	176	64	13	10
Cluster 11 - Post	170	64	13	10

Fig. 4H, I

Analysis: Rise and decay times of average regional firing rates. Peri-puff regional firing rates are averaged together within cluster identity and transformed to d' measurements on a per-time-bin basis. Time to rise is defined as the time post puff to reach 25% of the maximum d' value post puff. Time to decay is defined as the time to decay to 25% of the maximum d' value, following the peak. Regions with maximum d' less than 2 during either pre-infusion or ketamine-infusion conditions were excluded from the analysis. Regions with peak d' occurring 750ms following the puff were excluded from the analysis. After per-region per-cluster decay or rise times are obtained, they are combined per region into a single value using a weighted average, with weights determined by the fraction of cells in a given region assigned to each cluster. Point estimates given are median values across replicates of puffs. Bars indicate 95% confidence intervals of the median rise and decay values, obtained by bootstrap sampling of rise and decay times across puff replicates for each region.

Fig. 5B

Analysis: Average mouse eye closure behavior across puff series, for pre and ketamine conditions. Traces are baseline subtracted with the mean eye closure in a 1s window before each puff series. Mean and 95% confidence interval of the mean value is given (1.96 * s.e.m.), with trials averaged within session and variability taken across sessions.

Condition	# sessions	# trials	# subjects
Preinfusion	11	20	9
Ketamine	11	10	9

Fig. 5D

Analysis: Simultaneously recorded population neural activity along affective neural dimension, shown across puff series and subsequent inter-trial interval. Mean and 95% confidence interval of the mean value is given (1.96 * s.e.m.), with trials (each puff series) averaged within session and variability taken across sessions.

Condition	# sessions	# trials	# subjects
Preinfusion	11	20	9
Ketamine	11	10	9

Fig. 5E

Analysis: Mean activity along the affective neural dimension during the first puff of each puff-series, comparing

mean activity between pre-infusion and ketamine during a 0–0.25s window post-puff and during a 2–3s window post-puff. Point estimates of mean activity are obtained by averaging across trials within session. Statistics are computed on values across sessions. Error bars are given as 95% confidence intervals of the means.

Null hypothesis: No difference between each condition.

Condition	# mice	# sessions	# trials	Mean ± s.e.m.	Statistical test	Test statistic	Comparison	Corrected p- Value
0–0.25s	9	11	20 Pre, 10 Ket	Pre: 0.601 ± 0.28 ; Ket: 0.332 ± 0.138	Paired t-test with fdr-bh	1.209	Pre vs. Ketamine	0.254
2–3s	9	11	20 Pre, 10 Ket	Pre: 0.546 ± 0.174 ; Ket: 0.005 ± 0.043	correction	2.874	Pre vs. Ketamine	0.033

Fig. 5G

Analysis: Comparison of simultaneously recorded population neural activity along affective neural dimension and mathematical model. Model is fit to the average within condition across sessions. Variance explained is calculated as 1–MSE_model / MSE_null, where MSE_model is the mean squared error of the fitted model prediction—allowing only the tau_persistence parameter to vary across condition—and MSE_null is the mean squared error of the fitted model with no parameter allowed to vary across condition.

Condition	# sessions	# trials	# subjects
Preinfusion	11	20	9
Ketamine	11	10	9

Fig. 5H, I

Analysis: Comparison of model fit to behavioral data across drug doses, with fits performed separately for each drug. Variance explained is calculated as described above. Only the tau_persistence parameter of the model is allowed to vary across drug dose conditions.

Condition	# sessions	# trials	# subjects
Ketamine 35 mg/kg	5	10	5
Ketamine 25 mg/kg	5	10	5
Ketamine 13 mg/kg	5	10	5
Ketamine 6 mg/kg	5	10	5
PCP 10 mg/kg	5	10	5
PCP 5 mg/kg	5	10	5
PCP 1 mg/kg	5	10	5

Fig. 6C

Analysis: Left, peri-puff population activity along persistent dimension. Mean \pm 95% confidence interval (1.96 x s.e.m.). Right, average persistent population activity during 0.5 to 1.5 s after puff onset, compared between preinfusion, infusion, and postinfusion. Variation quantified across subjects, trials averaged per subject per condition.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p-Value
Preinfusion	7	1.36 ± 0.16		D V4	5.887	0.002
Ketamine	7	0.41 ± 0.07	Paired t-test with fdr-bh correction	Pre. vs. Ket.	3.887	0.002
Postinfusion	7	0.72 ± 0.11		Post vs. Ket.	3.171	0.019

Fig. 6D

Analysis: Left, peri-puff population activity along persistent dimension. Mean \pm 95% confidence interval (1.96 x s.e.m.). Right, average persistent population activity during 1-2 s after puff onset, comparing preinfusion, infusion, and postinfusion. Variation quantified across sessions, trials averaged per session per condition.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Preinfusion	9	11	20	0.86 ± 0.05	D : 1 :1 01 11	D., V.4	5.660	0.0004
Ketamine	9	11	10	0.02 ± 0.13	Paired t-test with fdr-bh correction	Pre. vs. Ket.	5.662	0.0004
Postinfusion	9	11	6	0.52 ± 0.21	correction	Post vs. Ket.	2.657	0.024

Fig. 6E

Analysis: Left, peri-puff population activity along fast dimension. Mean \pm 95% confidence interval (1.96 x s.e.m.). Right, average fast population activity during 0.05 to 0.10 s after puff onset, compared between preinfusion, infusion, and postinfusion, using beta band power. Variation quantified across subjects, trials averaged per subject per condition (12 to 40 trials per subject per condition).

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p-Value
Preinfusion	7	4.12 ± 0.31				
Ketamine	7	4.00 ± 0.41	Paired t-test with fdr-bh correction	Pre. vs. Ket.	0.432	0.681
Postinfusion	7	3.86 ± 0.38	i ancu i-test with ful-bil confection	Post vs. Ket.	0.576	0.585

Fig. 6F

Analysis: Left, peri-puff population activity along fast dimension. Mean \pm 95% confidence interval (1.96 x s.e.m.). Right, average fast population activity during 0 to 70 ms after puff onset, compared between preinfusion, infusion, and postinfusion. 70 ms is chosen based on the median time to peak signal per region. Variation quantified across sessions, trials averaged per session per condition.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Test statistic	Comparison	Corrected p- Value
Preinfusion	9	11	20	4.48 ± 0.129	Paired t-test with fdr-bh	-0.548	Pre. vs. Ket.	0.595
Ketamine	9	11	10	4.59 ± 0.185	correction	-0.546	Tic. vs. Rct.	0.575
Postinfusion	9	11	6	4.32 ± 0.184	zon zonon	2.468	Post vs. Ket.	0.066

Fig. 6G

Analysis: Intrinsic timescale of persistent population activity during the 10 minutes preceding eyepuff assay during preinfusion and infusion conditions.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	p-Value
Preinfusion	7	0.11 ± 0.01	Paired t-test	Preinfusion vs. ketamine	4.74	0.0032
Ketamine	7	0.09 ± 0.01				

Fig. 6H

Analysis: Intrinsic timescale of persistent population activity during 30s preceding each puff series, during preinfusion and infusion conditions. Statistics across session replicates. Values averaged within session across trials for each condition.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	p-Value
Preinfusion	9	11	10	0.45 ± 0.07	Paired t-test	Preinfusion vs. ketamine	3.357	0.0073
Ketamine	9	11	10	0.14 ± 0.06				

Fig. 6I

Analysis: Phase locking of persistent channels (defined as channels with greater than 90th percentile weight in the persistent population dimension), during preinfusion and infusion conditions, in the 10 minutes preceding eyepuff assay. Values are divisively normalized to preinfusion.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	p-Value
Preinfusion	7	1 ± 0.00	Paired t-test	Preinfusion vs. ketamine	3.344	0.0155
Ketamine	7	0.979 ± 0.0061				

Fig. 6J

Analysis: Pairwise correlation between persistent neurons (selected as neurons with greater than 90th percentile weight in persistent population dimension), during 30s preceding each puff series, for preinfusion and infusion conditions. Statistics across session replicates. Values averaged within session across trials for each condition. Values are divisively normalized to preinfusion.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	p-Value
Preinfusion	9	11	10	1.0 ± 0.0	Paired t-test	Preinfusion vs. ketamine	7.463	2.2 x 10 ⁻⁵
Ketamine	9	11	10	0.49 ± 0.07				

fig. S1 B

Analysis: Reflexive eye closure (sum of eye closure during 0.1 to 0.2 s after puff initiation), ketamine vs. saline. Point estimates for each subject are mean \pm s.e.m. across trials, excluding the first two trials.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# human participants	# trials per subject	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	4	38	0.84 ± 0.01	Paired t-test	1.622	0.2032
Ketamine	4	38	0.78 ± 0.04	Paired t-test		

fig. S1 C

Analysis: Affective eye closure (sum of eye closure during 0.3 to 0.8 s after puff initiation), ketamine vs. saline. Point estimates for each subject are mean \pm s.e.m. across trials, excluding the first two trials.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# human participants	# trials per subject	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	4	38	0.59 ± 0.07	D-:1444	2 (52	0.0254
Ketamine	4	38	0.34 ± 0.10	Paired t-test	3.653	0.0354

fig. S1 D

Analysis: Clinician administered dissociative states scale (CADSS), saline vs. ketamine.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# human participants	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	4	0.50 ± 0.29	Paired t-test	7.024	0.006
Ketamine	4	9.75 ± 1.25	Paired t-test	7.034	0.006

fig. S1 E

Analysis: Puff-triggered eye closure of each human subject, averaged across trials excluding the first two trials, comparing saline and ketamine conditions.

Human subject	Condition	# trials	Plot
1	Saline	38	Mean \pm s.e.m.
1	Ketamine	38	Mean \pm s.e.m.
2	Saline	38	Mean \pm s.e.m.
2	Ketamine	38	Mean \pm s.e.m.
3	Saline	38	Mean \pm s.e.m.
3	Ketamine	38	Mean \pm s.e.m.
4	Saline	38	Mean \pm s.e.m.
4	Ketamine	38	Mean \pm s.e.m.

fig. S2 B

Analysis: Reflexive eye closure in mice during for first puff of each trial, saline vs. ketamine.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	5	15	0.78 ± 0.05	Paired t-test	6.854	0.002
Ketamine	5	15	0.45 ± 0.06	Paired t-test	0.834	0.002

fig. S2 C

Analysis: Affective eye closure in mice during for first puff of each trial, saline vs. ketamine.

Null hypothesis: No difference between ketamine and saline conditions.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Test statistic	p-Value
Saline	5	15	$0.34 \pm\ 0.07$	Paired t-test	4.951	0.000
Ketamine	5	15	-0.02 ± 0.02	Paired t-test	4.931	0.008

fig. S2 EAnalysis: eye closure persistence after final puff of each trial, assessed as Kaplan Meier survival time.

Condition		Number of trials	Statistical test	Test statistic
Preinfusion	5	20 trials per mouse, 100 trials total	Kaplan Meier median survival time	15.16 seconds

fig. S2 G

Analysis: Scaling of normalized affective eye closure scales with puff number or puff intensity, after the eighth puff of each trial.

Null hypothesis: No difference between first and eighth puff at fixed puff intensity or between the eighth puff of two different puff intensities.

Condition	# mice	# trials per mouse	mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Puff 1, 8 PSI	8	15	0.256 ± 0.067		8 PSI: Puff 1 vs. Puff 8	4.834	0.0038
Puff 8, 8 PSI	8	15	0.517 ± 0.058		18 PSI: Puff 1 vs. Puff 8	9.243	0.0001
Puff 1, 18 PSI	8	15	0.239 ± 0.032		28 PSI: Puff 1 vs. Puff 8	11.633	0.0001
Puff 8, 18 PSI	8	15	0.515 ± 0.037	Paired t-test with fdr-bh correction	Puff 8: 8 PSI vs. 18 PSI	0.024	0.982
Puff 1, 28 PSI	8	15	0.286 ± 0.068		Puff 8: 8 PSI vs. 28 PSI	1.106	0.458
Puff 8, 28 PSI	8	15	0.447 ± 0.061		Puff 8: 18 PSI vs. 28 PSI	0.718	0.596

fig. S2 H

Analysis: Effect of different doses of ketamine or PCP on normalized affective eye closure, for the eighth puff of each trial.

Null hypothesis: No difference between preinfusion and infusion at a given dose.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	0.66 ± 0.07		Pre vs. Saline	0.980	0.425
Saline	5	15	0.70 ± 0.09		Pre vs. Saime	0.980	0.423
Pre-ketamine 50 mg/kg	5	15	0.66 ± 0.03		Pre vs. Ketamine 50 mg/kg	0 100	0.011
Ketamine 50 mg/kg	5	15	-0.04 ± 0.08		Pre vs. Ketamine 30 mg/kg	8.488	0.011
Pre-ketamine 35 mg/kg	8	15	0.45 ± 0.06	Paired t-test with fdr-bh	Pre vs. Ketamine 35 mg/kg	3.843	0.0159
Ketamine 35 mg/kg	8	15	0.17 ± 0.07	correction			
Pre-ketamine 25 mg/kg	5	15	0.61 ± 0.07		Pre vs. Ketamine 25 mg/kg	1.232	0.357
Ketamine 25 mg/kg	5	15	0.44 ± 0.19		rie vs. Ketanine 23 mg/kg	1.232	0.337
Pre-ketamine 13 mg/kg	5	15	0.57 ± 0.03		Pre vs. Ketamine 13 mg/kg	1.270	0.357

Ketamine 13 mg/kg	5	15	1.00 ± 0.34				
Pre-ketamine 6 mg/kg	5	15	0.70 ± 0.04		Pre vs. Ketamine	0.329	0.759
Ketamine 6 mg/kg	5	15	0.68 ± 0.07		6 mg/kg	0.329	0.739
Pre-PCP 20 mg/kg	5	15	0.71 ± 0.06		Pre vs. PCP 20 mg/kg	6.080	0.016
PCP 20 mg/kg	5	15	-0.01 ± 0.07				
Pre-PCP 10mg/kg	5	15	0.72 ± 0.05		Pre vs. PCP	5.404	0.016
PCP 10 mg/kg	5	15	0.20 ± 0.13		10 mg/kg	3.404	
Pre-PCP 5 mg/kg	5	15	0.64 ± 0.06		Pre vs. PCP	2.682	0.110
PCP 5 mg/kg	5	15	0.51 ± 0.07		5 mg/kg	2.082	0.110
Pre-PCP 1 mg/kg	5	15	0.61 ± 0.08		Pre vs. PCP	2064	0.100
PCP 1 mg/kg	5	15	0.40 ± 0.13		1 mg/kg	2.064	0.180

fig. S2 IAnalysis: Effect of different doses of ketamine or PCP on reflexive eye closure, for the eighth puff of each trial.

Null hypothesis: No difference between preinfusion and infusion at a given dose.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	0.78 ± 0.05		Pre vs. Saline	0.103	0.923
Saline	5	15	0.78 ± 0.05		Tie vs. Saine	0.103	0.923
Pre-ketamine 50 mg/kg	5	15	0.76 ± 0.08		Pre vs. Ketamine 50 mg/kg	3.042	0.064
Ketamine 50 mg/kg	5	15	0.44 ± 0.06		Fie vs. Retainine 50 mg/kg	3.042	0.004
Pre-ketamine 35 mg/kg	8	15	0.69 ± 0.05		Pre vs. Ketamine 35 mg/kg	2 525	0.032
Ketamine 35 mg/kg	8	15	0.43 ± 0.06		Fre vs. Ketamine 55 mg/kg	3.525	0.032
Pre-ketamine 25 mg/kg	5	15	0.62 ± 0.11		Pre vs. Ketamine 25 mg/kg	2.299	0.119
Ketamine 25 mg/kg	5	15	0.54 ± 0.10	Paired t-test with fdr-bh	Tre vs. recumme 25 mg/kg		
Pre-ketamine 13 mg/kg	5	15	0.73 ± 0.05		Pre vs. Ketamine 13 mg/kg	5.989	0.032
Ketamine 13 mg/kg	5	15	0.34 ± 0.07	correction		3.969	0.032
Pre-ketamine 6 mg/kg	5	15	0.83 ± 0.05		Pre vs. Ketamine	3.585	0.058
Ketamine 6 mg/kg	5	15	0.55 ± 0.11		6 mg/kg	3.363	0.038
Pre-PCP 20 mg/kg	5	15	0.63 ± 0.03		Pre vs. PCP 20 mg/kg	4.842	0.032
PCP 20 mg/kg	5	15	0.43 ± 0.01		20 mg/kg		
Pre-PCP 10mg/kg	5	15	0.61 ± 0.03		Pre vs. PCP	3.220	0.064
PCP 10 mg/kg	5	15	0.44 ± 0.05		10 mg/kg	3.220	0.004
Pre-PCP 5 mg/kg	5	15	0.51 ± 0.02		Pre vs. PCP	1.511	0.228
PCP 5 mg/kg	5	15	0.57 ± 0.04		5 mg/kg	1.311	0.220
Pre-PCP 1 mg/kg	5	15	0.67 ± 0.05		Pre vs. PCP	2.002	0.122
PCP 1 mg/kg	5	15	0.47 ± 0.09		1 mg/kg	2.082	0.132

fig. S2 J
Analysis: Effect of different doses of ketamine or PCP on affective eye closure, for the eighth puff of each trial.

Null hypothesis: No difference between preinfusion and infusion at a given dose.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	0.51 ± 0.05		D C 1	0.000	0.4100
Saline	5	15	0.55 ± 0.08		Pre vs. Saline	0.898	0.4199
Pre-ketamine 50 mg/kg	5	15	0.49 ± 0.04		Pre vs. Ketamine 50	7.947	0.0068
Ketamine 50 mg/kg	5	15	-0.01 ± 0.03		mg/kg	7.547	0.0008
Pre-ketamine 35 mg/kg	8	15	0.31 ± 0.04		Pre vs. Ketamine 35	5.397	0.0068
Ketamine 35 mg/kg	8	15	0.08 ± 0.03		mg/kg	3.391	0.0008
Pre-ketamine 25 mg/kg	5	15	0.38 ± 0.07		Pre vs. Ketamine 25	1.716	0.2017
Ketamine 25 mg/kg	5	15	0.20 ± 0.08		mg/kg	1.710	0.2017
Pre-ketamine 13 mg/kg	5	15	0.42 ± 0.05		Pre vs. Ketamine 13	3.789	0.0386
Ketamine 13 mg/kg	5	15	0.27 ± 0.05	Paired t-test with fdr-bh correction	mg/kg		
Pre-ketamine 6 mg/kg	5	15	0.58 ± 0.06		Pre vs. Ketamine	2.252	0.1250
Ketamine 6 mg/kg	5	15	0.39 ± 0.09		6 mg/kg	2.232	0.1230
Pre-PCP 20 mg/kg	5	15	0.45 ± 0.05		Pre vs. PCP	5.750	0.0113
PCP 20 mg/kg	5	15	0.00 ± 0.03		20 mg/kg		
Pre-PCP 10mg/kg	5	15	0.44 ± 0.05		Pre vs. PCP	6 104	0.0113
PCP 10 mg/kg	5	15	0.11 ± 0.07		10 mg/kg	6.194	0.0113
Pre-PCP 5 mg/kg	5	15	0.33 ± 0.04		Pre vs. PCP	1 171	0.2406
PCP 5 mg/kg	5	15	0.29 ± 0.04		5 mg/kg	1.171	0.3406
Pre-PCP 1 mg/kg	5	15	0.42 ± 0.07		D DCD		
PCP 1 mg/kg	5	15	0.23 ± 0.09		Pre vs. PCP 1 mg/kg	2.491	0.1123

fig. S2 K

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, comparing preinfusion and saline infusion conditions for five different sessions per mouse (one session per day).

Condition	# mice	# trials per mouse	Plot
Preinfusion	5	15	$mean \pm s.e.m.$
Saline	5	15	mean \pm s.e.m.

fig. S2 L

Analysis: Reflexive, affective, and normalized affective preinfusion eye closure across five sessions (one per day) with the same subject, for the eighth puff of each trial.

Null hypothesis: No difference between the first session and each subsequent session.

Normalized affective

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Day 1	5	15	0.42 ± 0.06				
Day 2	5	15	0.47 ± 0.03		Day 1 vs. Day 2	0.74	0.6569
Day 3	5	15	0.31 ± 0.08	Paired t-test with fdr-bh correction	Day 1 vs. Day 3	1.38	0.5638
Day 4	5	15	0.50 ± 0.05	Correction	Day 1 vs. Day 4	1.24	0.5638
Day 5	5	15	0.48 ± 0.07		Day 1 vs. Day 5	0.48	0.6569

Reflexive

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Day 1	5	15	0.39 ± 0.02				
Day 2	5	15	0.36 ± 0.03		Day 1 vs. Day 2	0.91	0.7425
Day 3	5	15	0.41 ± 0.05	Paired t-test with fdr-bh correction	Day 1 vs. Day 3	0.35	0.7425
Day 4	5	15	0.68 ± 0.08	correction	Day 1 vs. Day 4	3.31	0.119
Day 5	5	15	0.43 ± 0.04		Day 1 vs. Day 5	0.63	0.7425

Affective

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Day 1	5	15	0.17 ± 0.03				
Day 2	5	15	0.17 ± 0.01		Day 1 vs. Day 2	0.02	0.9879
Day 3	5	15	0.12 ± 0.03	Paired t-test with fdr-bh correction	Day 1 vs. Day 3	1.13	0.6418
Day 4	5	15	0.34 ± 0.05	correction	Day 1 vs. Day 4	3.19	0.1324
Day 5	5	15	0.21 ± 0.04		Day 1 vs. Day 5	0.63	0.7475

fig. S2 M

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, comparing preinfusion conditions for three different airpuff pressures.

Condition	# mice	# trials per mouse	Plot
8 PSI	8	15	mean \pm s.e.m.
18 PSI	8	15	mean \pm s.e.m.
28 PSI	8	15	mean \pm s.e.m.

fig. S2 N

Analysis: Left, average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, comparing preinfusion condition for male and female mice.

Condition	# mice	# trials per mouse	Plot
Female	5	15	mean \pm s.e.m.
Male	5	15	mean \pm s.e.m.

Analysis: Right, normalized affective eye closure for male and female mice for the eighth puff of each trial. Paired t-test for within-subject comparisons, and independent t-test for across subject.

Null hypothesis: No difference between male and female mice.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Male Pre-saline	5	15	0.44 ± 0.08		Male Pre vs. Saline	1.080	0.3827
Male saline	5	15	0.57 ± 0.11	T-test with fdr-bh	Female Pre vs. Saline	0.980	0.3827
Female Pre- saline	5	15	0.66 ± 0.07	correction	Pre Male vs. Female	2.079	0.2851
Female saline	5	15	0.70 ± 0.09		Saline Male vs. Female	0.983	0.3827

fig. S2 O

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, on trials in which either the 6th or 7th puff was skipped.

Condition	# mice	# trials per mouse	Plot
Preinfusion skip 6th puff	5	10	mean \pm s.e.m.
Infusion skip 6th puff	5	10	mean \pm s.e.m.
Preinfusion skip 7th puff	5	10	mean \pm s.e.m.
Infusion skip 7th puff	5	10	mean \pm s.e.m.

fig. S5 A

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, comparing preinfusion and drug infusion conditions.

Condition	# mice	# trials per mouse	Plot
Preinfusion	5	15	mean \pm s.e.m.
Buprenorphine	5	15	mean \pm s.e.m.

fig. S5 B

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, comparing preinfusion and drug infusion conditions.

Condition	# mice	# trials per mouse	Plot
Preinfusion	5	15	$mean \pm s.e.m.$
Diazepam	5	15	mean \pm s.e.m.

fig. S5 C

Analysis: Normalized affective eye closure (affective divided by reflexive) in mice for eighth puff of each trial, preinfusion vs. infusion for saline, ketamine (50 mg/kg), PCP (20 mg/kg), buprenorphine (2 mg/kg), diazepam (2 mg/kg). K/X is not shown here because loss of reflexive response makes normalized affective measure not applicable.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	0.66 ± 0.03		Pre-saline vs. saline	0.9797	0.4784

Saline	5	15	0.70 ± 0.09				
Pre- ketamine	5	15	0.66 ± 0.03		Pre-ketamine vs.	8.4882	0.0053
Ketamine	5	15	$\textbf{-0.04} \pm 0.08$		ketamine		
Pre-PCP	5	15	0.71 ± 0.06	Paired t-test with fdr-bh	Pre-PCP vs. PCP	6.0804	0.0092
PCP	5	15	-0.01 ± 0.07	correction	PIE-PCP VS. PCP	0.0804	0.0092
Pre-bupr.	5	15	0.65 ± 0.09		Day 1, 1,	0.0878	0.9342
Bupr.	5	15	0.64 ± 0.08		Pre-bupr. vs. bupr.	0.0878	0.9342
Pre-diaz.	5	15	0.75 ± 0.04		Pre-diaz, vs. diaz.	1.6120	0.3038
Diaz.	5	15	0.62 ± 0.05		Pre-diaz. vs. diaz.	1.0120	0.3038

fig. S5 D

Analysis: Reflexive eye closure in mice for eighth puff of each trial, preinfusion vs. infusion for saline, ketamine (50 mg/kg), PCP (20 mg/kg), buprenorphine (2 mg/kg), diazepam (2 mg/kg), K/X (135 mg/kg ketamine with 15 mg/kg xylazine).

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	0.78 ± 0.05		Pre-saline vs. saline	0.1026	0.9232
Saline	5	15	0.78 ± 0.05		Pre-same vs. same	0.1026	0.9232
Pre- ketamine	5	15	0.76 ± 0.08		Pre-ketamine vs.	3.0416	0.0575
Ketamine	5	15	0.44 ± 0.06		ketamine		
Pre-PCP	5	15	0.63 ± 0.03		Pre-PCP vs. PCP	4.8422	0.0168
PCP	5	15	0.43 ± 0.01	Paired t-test with fdr-bh	Pre-PCP vs. PCP	4.8422	0.0108
Pre-bupr.	5	15	0.54 ± 0.07	correction	Day 1, 1,	1.2745	0.2259
Bupr.	5	15	0.47 ± 0.07		Pre-bupr. vs. bupr.	1.2743	0.3258
Pre-diaz.	5	15	0.78 ± 0.02		Pre-diaz. vs. diaz.	6.2356	0.0101
Diaz.	5	15	0.48 ± 0.04		Pre-diaz. vs. diaz.	0.2330	0.0101
Pre-K/X	5	15	0.77 ± 0.09		Pre-K/X vs. K/X	8.6017	0.0060
K/X	5	15	0.03 ± 0.02		ric-n/A VS. N/A	8.001/	0.0060

fig. S5 E

Analysis: Affective eye closure in mice for eighth puff of each trial, preinfusion vs. infusion for saline, ketamine (50 mg/kg), PCP (20 mg/kg), buprenorphine (2 mg/kg), diazepam (2 mg/kg), K/X (135 mg/kg ketamine with 15 mg/kg xylazine).

Null hypothesis: No difference between preinfusion and infusion conditions.

71			1				
Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	0.51 ± 0.05		Pre-saline vs. saline	0.8982	0.4199
Saline	5	15	0.55 ± 0.08		Pre-saline vs. saline	0.8982	0.4199
Pre- ketamine	5	15	0.49 ± 0.04		Pre-ketamine vs.	7.9465	0.0041
Ketamine	5	15	-0.01 ± 0.03		Ketamine		
Pre-PCP	5	15	0.45 ± 0.05	Paired t-test with fdr-bh	n non non	5 7504	0.0001
PCP	5	15	0.00 ± 0.03	correction	Pre-PCP vs. PCP	5.7504	0.0091
Pre-bupr.	5	15	0.37 ± 0.09		Day 1, 1,	1.5276	0.2416
Bupr.	5	15	0.29 ± 0.04		Pre-bupr. vs. bupr.	1.32/6	0.2416
Pre-diaz.	5	15	0.58 ± 0.04		D., 4: 4:	2.0525	0.0252
Diaz.	5	15	0.30 ± 0.04		Pre-diaz. vs. diaz.	3.9535	0.0252

Pre-K/X	15	0.47 ± 0.03	D V	D.,, V/V V/V	D., V/V V/V 12.0602
K/X	15	0.02 ± 0.01	Pre-K.	Pre-K/X vs. K/X	Pre-K/X vs. K/X 12.0602

fig. S5 F

Analysis: Exponential decay rate of eye closure after the last puff of each trial in mice, preinfusion vs. infusion for saline, ketamine (50 mg/kg), PCP (20 mg/kg), buprenorphine (2 mg/kg), diazepam (2 mg/kg).

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Pre-saline	5	15	3.33 ± 0.13		Pre-saline vs. saline	0.8204	0.7634
Saline	5	15	3.96 ± 0.77		Pre-saline vs. saline	0.8204	0.7034
Pre- ketamine	5	15	3.53 ± 0.28		Pre-ketamine vs.	4.5198	0.0266
Ketamine	5	15	11.89 ± 1.74		ketamine		
Pre-PCP	5	15	3.04 ± 0.46	Paired t-test with fdr-bh	Pre-PCP vs. PCP	5 2524	0.0266
PCP	5	15	13.20 ± 1.59	correction	Pre-PCP vs. PCP	5.3534	0.0266
Pre-bupr.	5	15	4.15 ± 1.15		D 1 1 .	0.4065	0.0242
Bupr.	5	15	3.60 ± 0.84		Pre-bupr. vs. bupr.	0.4065	0.8243
Pre-diaz.	5	15	3.29 ± 0.78		Pre-diaz, vs. diaz.	0.2370	0.8243
Diaz.	5	15	3.03 ± 0.43		rre-diaz. Vs. diaz.	0.2370	0.8243

fig. S5 G

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, zoomed in on the time surrounding the eighth puff of a trial, comparing preinfusion and drug infusion conditions.

Condition	# mice	# trials per mouse	Plot
Saline	5	15	mean \pm s.e.m.
Ketamine	5	15	mean \pm s.e.m.
PCP	5	15	mean \pm s.e.m.
Buprenorphine	5	15	mean \pm s.e.m.
Diazepam	5	15	mean \pm s.e.m.
Ketamine/Xylazine	5	15	mean \pm s.e.m.

fig. S5 H

Analysis: Example video frames immediately preceding, during and after an airpuff after infusion with different drugs.

fig. S6 B

Analysis: Average eye closure across trials (trials consist of 8 closely-spaced puffs), averaged across mice, comparing no injection (intact), saline injection, and VPM muscimol injection conditions. The conditions are paired across mice.

Condition	# mice	# trials per mouse	Plot
Intact	5	15	mean \pm s.e.m.
Saline	5	15	mean \pm s.e.m.
VPM Muscimol	5	15	mean \pm s.e.m.

fig. S6 C

Analysis: Normalized affective eye closure (affective divided by reflexive) in mice for eighth puff in trial, comparing no injection, saline injection, and muscimol injection bilaterally in ventral posteromedial (VPM) thalamus.

Null hypothesis: No difference between each condition.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
No injection	5	15	0.63 ± 0.03		No injection vs. saline	0.45	0.6727
Saline	5	15	0.60 ± 0.05	Paired t-test with fdr-bh	Saline vs. muscimol	3.85	0.0289
Muscimol	5	15	0.11 ± 0.12	correction	No injection vs. muscimol	3.79	0.0289

fig. S6 D

Analysis: Reflexive eye closure in mice for eighth puff in trial, comparing no injection, saline injection, and muscimol injection bilaterally in ventral posteromedial (VPM) thalamus.

Null hypothesis: No difference between each condition.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
No injection	5	15	0.45 ± 0.09		No injection vs. saline	0.37	0.7179
Saline	5	15	0.41 ± 0.07	Paired t-test with fdr-bh	Saline vs. muscimol	2.48	0.0614
Muscimol	5	15	0.20 ± 0.06	correction	No injection vs. muscimol	2.43	0.0614

fig. S6 E

Analysis: Affective eye closure in mice for eighth puff in trial, comparing no injection, saline injection, and muscimol injection bilaterally in ventral posteromedial (VPM) thalamus.

Null hypothesis: No difference between each condition.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
No injection	5	15	0.29 ± 0.06		No injection vs. saline	0.48	0.6423
Saline	5	15	0.25 ± 0.05	Paired t-test with fdr-bh	Saline vs. muscimol	3.96	0.0063
Muscimol	5	15	0.03 ± 0.03	correction	No injection vs. muscimol	4.07	0.0063

fig. S6 F

Analysis: Early reflexive (within 40 ms of eyepuff onset) eye closure in mice for eighth puff in trial, comparing no injection, saline injection, and muscimol injection bilaterally in ventral posteromedial (VPM) thalamus.

Null hypothesis: No difference between each condition.

Condition	# mice	# trials per mouse	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
No injection	5	15	0.08 ± 0.01		No injection vs. saline	0.14	0.8898
Saline	5	15	0.08 ± 0.03	Paired t-test with fdr-bh	Saline vs. muscimol	0.37	0.8898
Muscimol	5	15	0.09 ± 0.02	correction	No injection vs. muscimol	0.43	0.8898

fig. S6 G

Analysis: Average eye closure across trials around the first puff of a trial, averaged across mice, comparing no injection (intact), saline injection, and VPM muscimol injection conditions.

Condition	# mice	# trials per mouse	Plot
Intact	5	15	mean \pm s.e.m.

Saline	5	15	mean \pm s.e.m.
VPM Muscimol	5	15	mean \pm s.e.m.

fig. S6 H

Analysis: Average eye closure across trials zoomed in after the first puff of a trial, averaged across mice, comparing no injection (intact), saline injection, and VPM muscimol injection conditions.

Condition	# mice	# trials per mouse	Plot
Intact	5	15	mean \pm s.e.m.
Saline	5	15	mean \pm s.e.m.
VPM Muscimol	5	15	mean \pm s.e.m.

fig. S6 I

Analysis: Average eye closure across trials around the eighth puff of a trial, averaged across mice, comparing no injection (intact), saline injection, and VPM muscimol injection conditions.

Condition	# mice	# trials per mouse	Plot
Intact	5	15	mean \pm s.e.m.
Saline	5	15	mean \pm s.e.m.
VPM Muscimol	5	15	mean \pm s.e.m.

fig. S6 J

Analysis: Average eye closure across trials zoomed in after the eighth puff of a trial, averaged across mice, comparing no injection (intact), saline injection, and VPM muscimol injection conditions.

Condition	# mice	# trials per mouse	Plot
Intact	5	15	mean \pm s.e.m.
Saline	5	15	mean \pm s.e.m.
VPM Muscimol	5	15	mean \pm s.e.m.

fig. S8 A

Analysis: Comparison of normalized affective eye closure during preinfusion, ketamine infusion, and postinfusion for each human subject that met the behavioral inclusion criteria (significantly decreased eye closure during infusion relative to preinfusion and relative to postinfusion). Affective eye closure is normalized by the reflexive eye closure on each trial.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Subject ID	Condition	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p-Value
	Preinfusion		0.94 ± 0.01		Pre vs. Ketamine	5.8087	1.29e-05
SD168	Ketamine	24	0.83 ± 0.02	Paired t-test with fdr-bh correction	Dogt va Vatamina	3.2763	0.0033
	Postinfusion		0.90 ± 0.01		Post vs. Ketamine	3.2703	0.0055
	Preinfusion		0.53 ± 0.03		Pre vs. Ketamine	5.6281	1.84e-06
SD172	Ketamine	39	0.35 ± 0.02	Paired t-test with fdr-bh correction	Dogt va Vatamina	11.7475	6.47e-14
	Postinfusion		0.75 ± 0.03		Post vs. Ketamine	11.7473	0.4/6-14
	Preinfusion		0.85 ± 0.04		Pre vs. Ketamine	10.5245	4.09e-11
SD181	Ketamine	30	0.21 ± 0.04	Paired t-test with fdr-bh correction	Dogt va Vatamina	5 9904	2.21e-06
	Postinfusion		0.53 ± 0.04		Post vs. Ketamine	5.8804	2.216-00
	Preinfusion		0.60 ± 0.03		Pre vs. Ketamine	4.7462	4.77e-05
SD185	Ketamine	31	0.45 ± 0.03	Paired t-test with fdr-bh correction	Dogt va Vatamina	6.8624	2.58e-07
	Postinfusion		0.67 ± 0.02		Post vs. Ketamine	0.8024	2.386-07
	Preinfusion		0.54 ± 0.05		Pre vs. Ketamine	2.7912	0.0236
SD186 F	Ketamine	12	0.35 ± 0.04	Paired t-test with fdr-bh correction	Dogt va Vataria	2.6265	0.0226
	Postinfusion		0.55 ± 0.06		Post vs. Ketamine		0.0236

	Preinfusion		0.31 ± 0.02		Pre vs. Ketamine	7.1858	1.38e-08	
SD211	SD211 Ketamine	39	0.15 ± 0.02	Paired t-test with fdr-bh correction	Dogt va Votomino	10.4531	1.96e-12	
	Postinfusion		0.36 ± 0.01	Į.	Post vs. Ketamine	10.4331	1.96e-12	
	Preinfusion		0.56 ± 0.02		Pre vs. Ketamine	5.2553	1.19e-05	
SD245	Ketamine	39	0.40 ± 0.02	Paired t-test with fdr-bh correction	Post vs. Ketamine	2,9258	0.0058	
	Postinfusion		0.50 ± 0.02		Post vs. Ketamine	2.9238	0.0038	

fig. S8 B

Analysis: Comparison of normalized affective eye closure during preinfusion, ketamine infusion, and postinfusion for each human subject that did not meet the behavioral inclusion criteria. Affective eye closure is normalized by the reflexive eye closure on each trial.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Subject ID	Condition	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
	Preinfusion		0.67 ± 0.03	D: 1	Pre vs. Ketamine	0.3794	0.7065
SD163	Ketamine	39	0.68 ± 0.03	Paired t-test with fdr-bh correction	Post vs.	1.7575	0.1738
	Postinfusion		0.78 ± 0.05	correction	Ketamine	1.7575	0.1736
	Preinfusion		0.13 ± 0.03	D: 144 4-14 C1 11	Pre vs. Ketamine	0.1429	0.8871
SD170	Ketamine	39	0.12 ± 0.05	Paired t-test with fdr-bh correction	Post vs.	1.1008	0.5558
	Postinfusion		0.05 ± 0.03	correction	Ketamine	1.1008	0.5556
	Preinfusion		0.93 ± 0.02	D: 144 4-14 C1 11	Pre vs. Ketamine	1.7376	0.0941
SD178	Ketamine	27	0.98 ± 0.02	Paired t-test with fdr-bh correction	Post vs.	3.6936	0.0021
	Postinfusion		0.90 ± 0.01	correction	Ketamine		0.0021
	Preinfusion		0.40 ± 0.02	D: 144 4-14 C1 11	Pre vs. Ketamine	0.9389	0.3555
SD184	Ketamine	30	0.43 ± 0.02	Paired t-test with fdr-bh correction	Post vs.	2.4481	0.0413
	Postinfusion		0.50 ± 0.03	correction	Ketamine	2.4461	0.0413
	Preinfusion		0.52 ± 0.03	D: 1	Pre vs. Ketamine	4.8338	0.0014
SD207	Ketamine	11	0.27 ± 0.04	Paired t-test with fdr-bh correction	Post vs.	2.1669	0.0555
	Postinfusion		0.17 ± 0.03	correction	Ketamine	2.1009	0.0555
	Preinfusion		0.33 ± 0.04	D: 1	Pre vs. Ketamine	1.2251	0.2281
SD215	Ketamine	39	0.21 ± 0.11	Paired t-test with fdr-bh correction	Post vs.	2 2002	0.0449
	Postinfusion		0.45 ± 0.04	correction	Ketamine	2.3802	0.0448
	Preinfusion		0.75 ± 0.02		Pre vs. Ketamine	4.8557	4.16e-05
SD242	Ketamine	39	0.63 ± 0.01	Paired t-test with fdr-bh	Post vs. Ketamine		0.0204
	Postinfusion		0.62 ± 0.02	correction		0.2170	0.8294

fig. S12 A

Analysis: Magnitude of event related potential (average of absolute value of local field potential for puff-triggered blinks) for each channel. Sorted by peak time on average across a train-set of trials. Plotted using average across held-out set of trials.

fig. S12 B

Analysis: Channels ordered according to (A), but with magnitude of event related potential triggered on non-puff blinks.

fig. S12 C

Analysis: Average event related potential across all channels. Event related potential magnitude is average absolute value of local field potential triggered on either onset of a puff-triggered blink or a natural (non-puff-triggered) blink. Mean traces are baseline-subtracted using the pre-blink interval.

Condition	# human participants	Total # blinks	Total # channels	Plot
Puff blink	7	52	773	mean \pm s.e.m.
Non-puff blink	7	52	773	mean \pm s.e.m.

fig. S12 D

Analysis: Average eye closure across blinks, triggered on either onset of a puff-triggered blink or a natural (non-puff-triggered) blink. Mean traces are baseline-subtracted using the pre-blink interval.

Condition	# human participants	Total # blinks	Plot
Puff blink	7	52	mean \pm s.e.m.
Non-puff blink	7	52	mean \pm s.e.m.

fig. S12 E

Analysis: Rise and decay times of average regional firing rates. Peri-puff local field potentials are transformed to d' measurements on a per-time-bin basis. The d' traces were smoothed with a 4-th order sos butterworth filter (cutoff=12.5 Hz). Time to rise is defined as the time after puff onset to reach a d' of either 0.75, 1.0, or 1.5. Regions with peak d' occurring 750ms following the puff were excluded from the analysis. Point estimates given are median values across replicates of puffs. Bars indicate 95% confidence intervals of the rise and decay values, obtained by bootstrap sampling of rise times across puff replicates for each region.

fig. S13

Analysis: Mean puff-triggered spectrogram by region. For each subject for each trial, a regional spectrogram was calculated as the mean across all channels in that region. Spectrograms were then averaged across all trials across all subjects that sampled a given region. Different regions were sampled in different subjects, so the number of trials differed.

fig. S14 G

Analysis: Spearman correlation between change in normalized affective eye closure (ketamine - preinfusion) versus Total CADSS score.

Null hypothesis: No correlation between change in normalized affective eye closure and CADSS.

Comparison	# human participants	Statistical test	Correlation coefficient	p- Value
Total CADSS vs. Change in normalized affective eye closure (ketamine - preinfusion)	10	Spearman rank-order correlation coefficient	0.744	0.014

fig. S16

Analysis: Mean puff-triggered spectrogram by region by subject.

fig. S21

Analysis: Comparison of permutation cluster test results using various parameters. Statistical test: paired t-test, independent t-test, and mixed-effect linear model with trials grouped by patient. Pixel threshold for cluster formation: p=0.05, p=0.01, p=0.001.

fig. S22

Analysis: Comparison of permutation cluster test results using various parameters. Statistical test: paired t-test,

independent t-test, independent t-test with hat variance adjustment, and mixed-effect linear model with trials grouped by patient. Pixel threshold for cluster formation: p=0.05, p=0.01, p=0.001, and threshold-free cluster enhancement (tfce) with start=0 and step=0.1.

fig. S23

Analysis: Comparison of permutation cluster test results using various parameters. Statistical test: paired t-test, independent t-test, and mixed-effect linear model with trials grouped by patient. Pixel threshold for cluster formation: p=0.05, p=0.01, p=0.001.

fig. S24

Analysis: Permutation cluster test for each channel individually, comparing infusion – preinfusion. Statistical test: paired t-test with p=0.01 pixel threshold for cluster formation. Each channel had from 12 to 39 trials per session depending on which subject the channel came from. Using the lowest cluster p-value from each channel, the p-values were fdr-bh corrected for multiple comparisons across channels. A channel was considered to have a significant change if the lowest corrected cluster p-value was < 0.05.

fig. S31 B

Analysis: fraction of recorded cells of a given cluster in each brain region, given for all regions with greater than 30 cells. P-value is calculated from a two-sided tail statistic given a per-region null distribution and corrected for multiple hypotheses using a fdr-bh correction. Regions above a corrected P-value threshold of 0.01 are labeled with the region acronym.

Null hypothesis: a given region has a uniform categorical distribution of clusters, computed by 20,000 bootstrapped cluster fractions from randomly assigned cluster labels within region.

fig. S31 F, G

Analysis: Distribution of changes (on ketamine, from pre-infusion) in rise times and decay times for all regions. Red line indicates the median of each distribution.

fig. S34 B

Analysis: Autocorrelation of puff response neural dimension of an example mouse subject for preinfusion and ketamine conditions, with exponential fit, f(t) = a*exp(-t/b)+c), where the intrinsic timescale is operationalized as the decay rate, b.

fig. S34 C

Analysis: eye closure across tone-series, for preinfusion and ketamine conditions. Mean \pm 95% CI (1.96 x s.e.m.). Variation quantified across sessions, trials averaged within session.

Condition	# subjects	# sessions	# trials
Preinfusion	7	7	10
Ketamine	7	7	5
Postinfusion	7	7	3

fig. S34 D

Analysis: Left, peri-tone population activity along tone fast dimension. Mean \pm 95% confidence interval (1.96 x s.e.m.). Right, average tone fast population activity during 0 to 70 ms after tone onset, compared between preinfusion, infusion, and postinfusion. Variation quantified across sessions, trials averaged per session per condition.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Preinfusion	7	7	10	2.49 ± 0.245	Paired t-test with fdr-bh	Pre. vs. Ket.	1.088	0.318
Ketamine	7	7	5	1.83 ± 0.399	correction	Tic. vs. Kct.	1.088	0.318
Postinfusion	7	7	3	1.56 ± 0.275		Post vs. Ket.	1.438	0.318

fig. S34 E

Analysis: Left, peri-tone population activity along tone persistent dimension (difference between average activity in the window [150 ms, 350 ms] and baseline). Mean \pm 95% confidence interval (1.96 x s.e.m.). Right, average tone persistent population activity during 350 ms to 500 ms after tone onset, comparing preinfusion, infusion, and postinfusion. Variation quantified across sessions, trials averaged per session per condition.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
Preinfusion	7	7	10	1.531 ± 0.081	Paired t-test with fdr-bh	Pre. vs. Ket.	6.942	0.0009
Ketamine	7	7	5	0.392 ± 0.177	correction	11c. vs. Rct.	0.742	0.0009
Postinfusion	7	7	3	1.385 ± 0.338		Post vs. Ket.	-3.753	0.009

fig. S34 F

Analysis: Comparison of timecourse of detectability of activity along the persistent dimension for eyepuff vs. tone. Detectability is calculated as d-prime. Mean \pm 95% CI.

fig. S34 G

Analysis: Intrinsic timescale of tone persistent population activity during 30s preceding each tone series, during preinfusion and infusion conditions. Statistics across session replicates. Values averaged within session across trials for each condition.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	p-Value
Preinfusion	7	7	10	0.268 ± 0.037	Paired t-test	Preinfusion vs. ketamine	5.479	0.0015
Ketamine	7	7	5	0.089 ± 0.012				

fig. S34 H

Analysis: Pairwise correlation between tone persistent neurons (selected as neurons with greater than 90th percentile weight in tone persistent population dimension), during 30s preceding each tone series, for preinfusion and infusion conditions. Statistics across session replicates. Values averaged within session across trials for each condition.

Null hypothesis: No difference between preinfusion and infusion conditions.

Condition	# subjects	# sessions	# trials	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	p-Value
Preinfusion	7	7	10	1.0 ± 0.0	Paired t-test	Preinfusion vs. ketamine	6.217	0.0008
Ketamine	7	7	5	0.460 ± 0.080				

fig. S35 A

Analysis: Participation (loading) of each channel in the Fast coding dimension (left) or Persistent coding dimension (right). For each plot, the loadings are normalized by the maximum value.

fig. S35 B

Analysis: Participation (loading) of each channel in the Fast coding dimension across all subjects. The loadings for each subject are normalized by the maximum value of that subject.

fig. S35 C

Analysis: Aggregate participation (loading) of each Yeo7 resting state network in the Fast coding dimension. Median across all channels in a network, with 95% confidence interval errorbars. To determine whether a network has over- or under-represented in the coding dimension, bootstrap shuffling is used, in which the network label assigned to each channel is shuffled, with 1000 shuffles. Dashed lines represent the upper and lower cutoffs for two-sided significance at p=0.05. P-values are computed based on the percentile of the true value among the shuffles (p=0.05 corresponds to 97.5 or 2.5 percentile), and are corrected across networks with fdr-bh.

Null hypothesis: A network does not have more or less loading on the Fast coding dimension than other networks.

Yeo7 network	# subjects	# channels	Mean ± s.e.m.	Statistical test	Corrected p- Value
Default	7	88	0.126 ± 0.014		0.1056
Dorsal Attention	1 Attention 5		0.080 ± 0.027		0.0440
Frontoparietal	5	23	0.070 ± 0.027	Bootstrap shuffle (network labels) with fdr-bh correction across	0.0180
Limbic	7	34	0.115 ± 0.018	networks	0.1720
Somatomotor	7	7 50 (0.446 ± 0.045		0.0000
Ventral Attention	7		0.209 ± 0.032		0.0960

fig. S35 D

Analysis: Same as previous panel but for Persistent coding dimension.

Null hypothesis: A network does not have more or less loading on the Persistent coding dimension than other networks.

Yeo7 network	# subjects	# channels	Mean ± s.e.m.	Statistical test	Corrected p- Value
Default	Default 7 88 0.239 ± 0.021		0.239 ± 0.021		0.9740
Dorsal Attention	sal Attention 5	18	0.292 ± 0.060		0.9740
Frontoparietal	5	23	0.350 ± 0.037	Bootstrap shuffle (network labels) with fdr-bh correction across	0.7920
Limbic	7	34	0.233 ± 0.035	networks	0.9740
Somatomotor	7	50	0.253 ± 0.030		0.9740
Ventral Attention	7	63	0.246 ± 0.032		0.9740

fig. S37 A

Analysis: Pearson correlation coefficient between mean preinfusion persistent coding dimension projection (across subjects) for mouse firing and human bandpower, for each frequency band.

fig. S37 B

Analysis: Left, peri-puff population activity along persistent dimension. Mean \pm s.e.m. Right, average persistent population activity during 0.05 to 0.10 s after puff onset, compared between preinfusion, infusion, and postinfusion, using different frequency bands.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Frequency band	Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
	Preinfusion		3.20 ± 0.10				
Delta (1-4 Hz)	Ketamine	7	3.15 ± 0.13	Paired t-test with fdr-bh correction	Pre. vs. Ket.	0.817	0.859
,	Postinfusion		3.14 ± 0.10		Post vs. Ket.	0.183	0.859
TI 4	Preinfusion		3.83 ± 0.05	Paired t-test with fdr-bh	Pre. vs. Ket.	0.274	0.793
Theta (4-8 Hz)	Ketamine	7	3.84 ± 0.06	correction	FIE. VS. Ket.	0.274	0.793
(4-0 HZ)	Postinfusion		3.86 ± 0.04	correction	Post vs. Ket.	0.428	0.793
A 11	Preinfusion		3.89 ± 0.04	Paired t-test with fdr-bh	Pre. vs. Ket.	0.848	0.508
Alpha (8-12 Hz)	Ketamine	7	3.93 ± 0.06	correction	Tic. vs. Kct.	0.040	0.508
(0 12 112)	Postinfusion		3.89 ± 0.07	Correction	Post vs. Ket.	0.703	0.508
Beta	Preinfusion	7	4.12 ± 0.31	Paired t-test with fdr-bh	Pre. vs. Ket.	0.432	0.681
(12-30 Hz)	Ketamine	7	4.00 ± 0.41	correction	Tic. vs. Kct.	0.432	0.001
(12 30 112)	Postinfusion	7	3.86 ± 0.38	correction	Post vs. Ket.	1.063	0.657
Gamma	Preinfusion	7	3.19 ± 0.24	Paired t-test with fdr-bh	Pre. vs. Ket.	1.245	0.273
(30-50 Hz)	Ketamine	7	2.89 ± 0.26	correction	Tic. vs. Kct.	1.243	0.273
(50 50 112)	Postinfusion	7	3.18 ± 0.12	correction	Post vs. Ket.	1.207	0.273
II:-1. C	Preinfusion	7	3.64 ± 0.37	Paired t-test with fdr-bh	Pre. vs. Ket.	0.053	0.960
High Gamma (65-95 Hz)	Ketamine	7	3.64 ± 0.35	correction	Tic. vs. Kct.	0.055	0.900
(03-93 112)	Postinfusion	7	3.71 ± 0.46	correction	Post vs. Ket.	0.371	0.960
	Preinfusion	7	2.35 ± 0.59	D-i14 44i4l- £1 11	Pre. vs. Ket.	0.816	0.761
Broadband LFP	Ketamine	7	1.88 ± 0.88	Paired t-test with fdr-bh correction	116. vs. Ket.	0.010	0.701
	Postinfusion	7	1.99 ± 0.81	Correction	Post vs. Ket.	0.319	0.761

fig. S37 C

Analysis: Left, peri-puff population activity along persistent dimension. Mean \pm s.e.m. Right, average persistent population activity during 0.5 to 1.5 s after puff onset, compared between preinfusion, infusion, and postinfusion, using different frequency bands.

Null hypothesis: No difference between infusion and preinfusion or postinfusion.

Frequency band	Condition	# human participants	Mean ± s.e.m.	Statistical test	Comparison	Test statistic	Corrected p- Value
D-14-	Preinfusion		0.79 ± 0.16	Paired t-test with fdr-bh	Pre. vs. Ket.	3.930	0.015
Delta (1-4 Hz)	Ketamine	7	0.18 ± 0.07	correction	Tic. vs. Kct.	3.930	0.013
,	Postinfusion		0.44 ± 0.12		Post vs. Ket.	2.502	0.046
Theta	Preinfusion		0.82 ± 0.22	Paired t-test with fdr-bh	Pre. vs. Ket.	3.246	0.035
(4-8 Hz)	Ketamine	7	0.11 ± 0.08	correction	TIC. VS. KCt.	3.240	0.033
(4-8 112)	Postinfusion		0.35 ± 0.13	Correction	Post vs. Ket.	1.935	0.101

Alpha	Preinfusion Ketamine	7	0.97 ± 0.21 0.25 ± 0.12	Paired t-test with fdr-bh	Pre. vs. Ket.	4.620	0.007
(8-12 Hz)	Postinfusion	,	0.47 ± 0.18	correction	Post vs. Ket.	1.777	0.126
D-4-	Preinfusion	7	1.36 ± 0.16	Daine 14 4-4 i41 Cl. 1-1	Pre. vs. Ket.	5.887	0.002
Beta (12-30 Hz)	Ketamine	7	0.41 ± 0.07	Paired t-test with fdr-bh correction	1 1c. vs. Kct.	3.867	0.002
(12-30 112)	Postinfusion	7	0.72 ± 0.11	correction	Post vs. Ket.	3.171	0.019
Low Gamma	Preinfusion	7	1.32 ± 0.08	D: 1 1.1.01.11	Pre. vs. Ket.	6.252	0.002
(30-50 Hz)	Ketamine	7	0.49 ± 0.17	Paired t-test with fdr-bh correction	Pre. vs. Ket.	0.232	0.002
(30-30 112)	Postinfusion	7	0.52 ± 0.14	Correction	Post vs. Ket.	0.383	0.715
н: 1 С	Preinfusion	7	0.90 ± 0.11	D: 1 1.1.01.11	Pre. vs. Ket.	6.870	0.001
High Gamma (65-95 Hz)	Ketamine	7	0.27 ± 0.13	Paired t-test with fdr-bh correction	Fie. vs. Ket.	0.870	0.001
(03-93 112)	Postinfusion	7	0.39 ± 0.08	correction	Post vs. Ket.	1.189	0.279
	Preinfusion	7	1.66 ± 0.05	D: 1 11.01.11	D., V.4	2.625	0.079
Broadband LFP	Ketamine	7	1.37 ± 0.14	Paired t-test with fdr-bh correction	Pre. vs. Ket.	2.023	0.079
	Postinfusion	7	1.49 ± 0.08	Correction	Post vs. Ket.	0.790	0.460

fig. S38 A

Analysis: Mixed effects linear model, grouped by subject, of preinfusion phase locking ~ high vs. low CD edge identity (categorical).

Null hypothesis: No correlation between preinfusion phase locking and an edge's identity as being between high vs. low CD channels.

Frequency band	# human participants	# inter- channel edges	Statistical test	Comparison	Coefficient	Confidence interval 95%	R ²	Corrected p-value
Delta (1-4 Hz)	7	716		Low vs. High	0.0017	[-0.005, 0.008]	1.3e-4	0.76
Theta (4-8 Hz)	7	716	Mixed-effects	Low vs. High	0.0026	[-0.004, 0.009]	-7.4e-4	0.76
Alpha (8-12 Hz)	7	716	linear model with fdr-bh	Low vs. High	0.0017	[-0.006, 0.009]	-7.6e-4	0.76
Beta (12-30 Hz)	7	716	correction	Low vs. High	0.0016	[-0.005, 0.008]	-3.7e-4	0.76
Gamma (30-50 Hz)	7	716		Low vs. High	0.0018	[-0.004, 0.007]	3.7e-4	0.76
High Gamma (65-95 Hz)	7	716		Low vs. High	-0.00060	[-0.006, 0.005]	4.6e-5	0.83

fig. S38 B

Analysis: Mixed effects linear model, grouped by subject, of phase locking change on ketamine \sim high vs. low CD edge identity (categorical).

Null hypothesis: No correlation between ketamine-induced change in an edge's phase locking and an edge's identity as being either high or low CD channels.

Frequency band	# human participants	# inter- channel edges	Statistical test	Comparison	Coefficient	Confidence interval 95%	\mathbb{R}^2	Corrected p-value
Delta (1-4 Hz)	7	716	Mixed-effects linear model with fdr-bh	Low vs. High	-0.0038	[-0.024, 0.016]	-0.0093	0.70
Theta (4-8 Hz)	7	716	correction	Low vs. High	-0.016	[-0.038, 0.005]	-0.00068	0.16

Alpha (8-12 Hz)	7	716	Low vs. High	-0.028	[-0.050, -0.007]	-0.0063	0.021
Beta (12-30 Hz)	7	716	Low vs. High	-0.038	[-0.057, -0.020]	0.020	0.00026
Gamma (30-50 Hz)	7	716	Low vs. High	-0.023	[-0.040, -0.006]	0.0080	0.019
High Gamma (65-95 Hz)	7	716	Low vs. High	-0.012	[-0.028, 0.003]	0.0031	0.16

fig. S38 D

Analysis: Mixed effects linear model, grouped by subject, of preinfusion power ~ high vs. low CD channel identity (categorical).

Null hypothesis: No correlation between a channel's pre-infusion power and a channel's identity as being high vs. low CD.

Frequency band	# human participants	# channels	Statistical test	Comparison	Coefficient	Confidence interval 95%	R ²	Corrected p-value
Delta (1-4 Hz)	7	104		Low vs. High	-2.36	[-4.6, -0.094]	0.0092	0.24
Theta (4-8 Hz)	7	104		Low vs. High	-1.28	[-3.49, 0.91]	0.0011	0.48
Alpha (8-12 Hz)	7	104	Mixed-effects linear model with fdr-bh	Low vs. High	-0.43	[-2.80, 1.93]	-0.0025	0.71
Beta (12-30 Hz)	7	104	correction	Low vs. High	-0.74	[-2.82, 1.33]	-0.0011	0.57
Gamma (30-50 Hz)	7	104		Low vs. High	-0.99	[-2.97, 0.98]	-0.0014	0.48
High Gamma (65-95 Hz)	7	104		Low vs. High	-1.63	[-3.53, 0.25]	0.0015	0.26

fig. S38 E

Analysis: Mixed effects linear model, grouped by subject, of power change from preinfusion to ketamine ~ high vs. low CD channel identity (categorical).

Null hypothesis: No correlation between a channel's change in power on ketamine and a channel's identity as being high vs. low CD.

Frequency band	# human participants	# channels	Statistical test	Comparison	Coefficient	Confidence interval 95%	R ²	Corrected p- value
Delta (1-4 Hz)	7	104		Low vs. High	-0.19	[-0.47, 0.075]	0.0041	0.46
Theta (4-8 Hz)	7	104		Low vs. High	-0.11	[-0.39, 0.15]	0.0043	0.79
Alpha (8-12 Hz)	7	104	Mixed-effects linear model with fdr-bh	Low vs. High	-0.047	[-0.38, 0.28]	0.00063	0.80
Beta (12-30 Hz)	7	104	correction	Low vs. High	0.079	[-0.40, 0.56]	0.00022	0.80
Gamma (30-50 Hz)	7	104		Low vs. High	0.87	[0.11, 1.63]	0.032	0.14
High Gamma (65-95 Hz)	7	104		Low vs. High	0.081	[-0.56, 0.73]	0.00044	0.80

fig. S38 H

Analysis: Mixed effects linear model, grouped by subject, of **single-trial** post-puff affective (late) window CD projection ~ pre-puff network-wide beta phase locking.

Null hypothesis: No correlation on single trials between the post-puff late projection and pre-puff beta phase locking.

Condition	# human participants	# total trials	Statistical test	Coefficient	Coeff 95% CI	R ²	p-value
Preinfusion	7	214	Mixed-effects linear model	12.12	[5.19, 19.04]	0.057	6.04e-4
Infusion	7	214	winxed-effects fifical filoder	-2.15	[-6.78, 2.47]	0.00083	0.361

fig. S38 K

Analysis: Mixed effects linear model, grouped by subject, of **single-trial** post-puff affective eye closure ~ prepuff network-wide beta phase locking.

Null hypothesis: No correlation on single trials between the post-puff eye closure behavior and pre-puff beta phase locking.

Condition	# human participants	# total trials	Statistical test	Coefficient	Confidence interval 95%	R ²	p-value
Preinfusion	7	214	Mixed-effects linear model	3.35	[1.26, 5.45]	0.034	0.0017
Infusion	7	214		-0.065	[-2.83, 2.70]	-0.0080	0.96

References and Notes

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