

Cerebellar Activation Bidirectionally Regulates Nucleus Accumbens Shell and Core

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Abstract

Traditionally viewed as a motor control center, the cerebellum (CB) is now recognized as an integral part of a broad, long-range brain network that serves limbic functions and motivates behavior. This diverse CB functionality has been at least partly attributed to the multiplicity of its outputs. However, relatively little attention has been paid to CB connectivity with subcortical limbic structures, and nothing is known about how the CB connects to the nucleus accumbens (NAc), a complex striatal region with which the CB shares functionality in motivated behaviors. Here, we report findings from in vivo electrophysiological experiments that investigated the functional connectivity between CB and NAc. We found that electrical microstimulation of deep cerebellar nuclei (DCN) modulates NAc spiking activity. This modulation differed in terms of directionality (excitatory vs. inhibitory) and temporal characteristics, in a manner that depends on NAc subregions: in the medial shell of NAc (NAc_{Med}), slow inhibitory responses prevailed over excitatory ones, whereas the proportion of fast excitatory responses was greater in the NAc core (NAc_{Core}) compared to NAc_{Med}. Slow inhibitory modulation of NAc_{Core} was also observed but it required stronger CB inputs compared to NAc_{Med}. Finally, we observed shorter onset latencies for excitatory responses in NAc_{Core} than in NAc_{Med}, which argues for differential connectivity. If different pathways provide signal to each subregion, the divergence likely occurs downstream of the CB because we did not find any response-type clustering within DCN. Because there are no direct monosynaptic connections between CB and NAc, we performed viral tracing experiments to chart disynaptic pathways that could potentially mediate the newly discovered CB-NAc communication. We identified two anatomical pathways that recruit the ventral tegmental area and intralaminar thalamus as nodes. These pathways and the functional connectivity they support could underlie CB's role in motivated behaviors.

Keywords

Cerebellum, nucleus accumbens, ventral striatum, connectivity, mouse, electrophysiology, anatomy, circuit, thalamus, intralaminar, VTA, motivated behavior, non-motor

Introduction

The cerebellum (CB) exploits the parallel, modular organization of its circuitry to integrate information and perform complex computations. Most CB research sheds light on this complexity in the context of CB's role in predicting and updating motor movements¹. However, accumulating evidence supports CB involvement in high-order non-motor functions as well²⁻⁵. In humans, the CB is consistently activated during decision making, particularly during risk- or reward-based tasks^{6,7}, and during aversive experiences and emotional learning^{8,9}. Further support for non-motor CB roles stems from clinical translational studies, which have linked CB dysfunction with neurodevelopmental disorders, posttraumatic stress disorder, generalized anxiety disorder, addiction, and cognitive and emotional disturbances known as cerebellar cognitive affective syndrome^{8,10-18}. These findings are further corroborated by evidence from animal studies, which solidify a role for the CB in the processing of valence, reward, reward anticipation and omission¹⁹⁻²³; emotional learning and aggression²⁴⁻³⁰; and motivation³¹⁻³³.

The ability of the CB to derive a complex repertoire of non-motor functions from its relatively invariant cellular organization is largely attributed to its diverse outputs^{34,35}. In addition to the heavily emphasized non-motor cerebellar influences on cortical regions³⁶⁻⁴², functional and/or anatomical CB connections with subcortical structures critical for cognition and emotion have also been documented⁴³⁻⁴⁸. Here, we focused on the nucleus accumbens (NAc), a complex limbic structure that shares reward, motivation and affective functionality with the CB^{49,50}. Stimulation of CB cortex or deep cerebellar nuclei (DCN) modulates levels of NAc dopamine- an important, but not exclusive, regulator of NAc functions⁵¹⁻⁵⁷. However, how the CB connects to NAc is unknown.

Here we used in vivo electrophysiology in anesthetized mice to examine the effects of CB stimulation on spiking activity in the NAc medial shell (NAc_{Med}) and core (NAc_{Core}). These regions exhibit distinct input-output organization, anatomy, and function⁵⁸⁻⁶³. We provide the first evidence, to our knowledge, of an electrophysiological connection between CB and NAc, which shows NAc subregion-dependent specificity. Using viral tracing approaches, we offer first insights on the anatomical blueprint of CB-NAc connectivity, which includes nodal neurons in ventral tegmental area (VTA) and limbic thalamus. These findings may expound upon the CB's involvement in limbic functions such as motivation, reward learning, and affective processing.

Materials and Methods

In vivo electrophysiology

Mice

C57Bl/6J mice of both sexes (N = 29) were used in accordance with the National Institutes of Health guidelines and the policies of the University of California Davis Committee for Animal Care. Acute in vivo recordings were performed at postnatal days P21-P29. All animals were maintained on a light/dark cycle (light 7AM-7PM), and experiments were performed during the light cycle.

Surgery

Anesthesia was initially induced by brief inhalation of 4-5% isoflurane followed by an intraperitoneal injection of anesthetic cocktail (100 mg/kg Ketamine; 10 mg/kg xylazine; 1 mg/kg acepromazine), and was maintained with periodic injections of the anesthetic cocktail (20-50 mg/kg ketamine), as needed. Animals were placed in a stereotactic frame after confirmation of anesthesia depth using a toe pinch response test. Breathing rate and toe pinch responses were monitored to ensure maintenance of anesthesia. A small craniotomy and durectomy were performed over the DCN (lateral/interposed n.: relative to lambda, in mm: -2.1 to -2.6 AP; +/-2.1 to +/-2.3 ML), and over NAc targeting the medial shell and/or core (relative to bregma: +1.7 to +1.54 AP; +/- 0.4 to +/-1.15 ML).

Stimulation

A custom stimulating stereotrode (~200 μ m distance between tips) was lowered 1.95 – 2.45 mm below the brain surface (in DV axis) through the cerebellar craniotomy to reach the DCN. Ten trials of bipolar constant-current electrical microstimulation were delivered at each location at a 15-s inter-trial interval. Each stimulation trial consisted of a 200-Hz burst of five 0.5-ms monophasic square-waveform pulses at 100 μ A. In a subset of experiments (N = 11 mice), the stimulation intensity was varied between 30, 100, and 300 μ A in interleaved blocks of 5 trials per intensity. The use of stereotrode-delivered electrical stimuli in these experiments was instrumental in allowing us to 1) probe changes in NAc spiking activity in response to broad variations in current intensity; 2) exercise tight control over a small localized area; 3) simultaneously stimulate all neurons in that area.

Data Acquisition

For electrophysiological recordings, a 12-channel (platinum; 0.2-2 M Ω impedance) axial multi-electrode array (FHC; 150 μ m distance between channels), dipped in fluorescein dextran (3000 MW), was lowered into the NAc at a depth of 3.75 – 5.25 mm below the brain surface (DV axis). Electrode signals were fed to a digital headstage (RHD 2132, Intan Technologies) for amplification (x20), filtering (0.7-7500 Hz) and digitization (30 kS/s with 16-bit resolution), before transfer to an open-source acquisition system (OpenEphys) for display and storage.

Histology for Verification of Electrode Placement

Positioning of electrodes was initially guided by atlas-based stereotactic coordinates (Paxinos & Franklin) and, upon completion of experiments, histologically verified through electrolytic lesions (for location of DCN microstimulation electrodes; single 10-s cathodal pulse of 300 μ A) and fluorescence imaging (NAc fluorescein dye track). Coordinates for subsequent animals of the same litter were further adjusted accordingly. Briefly, at the end of experiments electrodes were retracted and animals were perfused with 4% (w/v) paraformaldehyde (PFA) in 0.1 M phosphate buffer (PB). Brains were dissected and post-fixed in 4% PFA, sliced in phosphate buffered saline (PBS) and inspected under a fluorescence stereoscope (Olympus SZX2). NAc slices containing the dye track from the recording array were identified under 488 nm light, and CB slices with electrolytically lesioned tissue were identified under brightfield illumination. Slices of interest were subsequently stained with DAPI (1 : 20,000, Thermo Fisher Scientific), mounted on slides, and imaged using a VS120 Olympus slide scanner. Images were manually registered to the Paxinos and Franklin Mouse Brain Atlas and the location of the recording electrode tip was mapped. Only channels along the recording array that were determined to be within NAc_{Med} and/or NAc_{Core} were included in analysis. Similarly, experiments in which cerebellar electrolytic lesions were localized outside the lateral and/or interposed DCN were excluded from analysis.

Quantification and Statistical Analysis

Data processing and quantification

Custom-written MATLAB scripts (Mathworks) were used for data processing and analysis. Separation of multi-unit spikes from local field potentials was achieved through a 4th-order Butterworth high-pass filter (cutoff at 300 Hz). The filtered signal was thresholded at 3-3.5 standard deviations below the average voltage of a 10-s baseline (taken before stimulation

onset), and neural firing rates were computed from the spike counts in consecutive 10-ms time bins (peri-stimulus time histogram; PSTH). Putative spikes with time differences less than 10 ms from stimulus artifacts or 2 ms from the previous spike were discarded.

To quantify the DCN stimulation-induced modulation of NAc spiking activity we first averaged the PSTH across all trials of the same intensity at the same recording and stimulation sites, and randomly permuted a trial-averaged 10-s pre-stimulus baseline 1,000 times. The observed trial-averaged PSTH was then z-transformed based on the mean and standard deviation of the randomization distribution (z-transformed firing rate). A change of firing rate from baseline was considered a response to DCN stimulation only if it (a) took place within the first 3 s following stimulus onset (response window), and (b) exceeded either a positive threshold of $z = 4$ (for excitatory responses) or a negative threshold of $z = -3.2$ (for inhibitory responses). The same thresholds were employed to identify responses of both NAc subregions and to all stimulation intensities that were tested in this study. Recordings with responses were categorized as responders (vs. non-responders). Extensive analyses of candidate thresholds (an example subset of the candidate thresholds we tested appears in Fig. 2S1) allowed us to employ response thresholds that (a) kept the relative frequency of threshold crossings during a 3-s pre-stimulus baseline window (i.e., same duration as the response window) less than 5% (false positive responses; typically around 2-3%), and (b) the relative distribution of excitatory/inhibitory responses was robust to small changes in threshold. Onset latency of responses was quantified as the time of threshold crossing relative to stimulus onset.

Statistics

Unless otherwise noted, we performed all statistical comparisons using the random permutation method^{64,65}. We randomly shuffled the data between groups 1,000 times and estimated the probability to find a difference greater than or equal to the observed difference by chance alone. We corrected for multiple comparisons with False Discovery Rate (FDR)⁶⁶ set at 10%.

Anatomical Tracing

Surgery and viral injections

Mice of both sex (P35 - P49) were anesthetized with isoflurane (4 % - 5 % induction; 1.5 % maintenance) and secured to a stereotactic frame. After exposing the top of the skull, the head was leveled and small craniotomies were drilled over areas of interest. Injections were made using a Micro4 controller and UltraMicroPump 3 (VWR). Glass needles were made from 1-mm outer diameter glass pipettes (Wiretrol II, Drummond) pulled to a fine tip (20 - 50 μ m tip diameter, 3 - 4 mm tip length) using a pipette puller (P-97, Sutter). Needles were left in place for 7-10 min following injections to minimize diffusion. Following surgery and analgesia administration (0.1 mg/kg buprenorphine, 5 mg/kg meloxicam), mice were allowed to recover on a warm heating pad before being transferred back to the vivarium. Mice remained in the colony to allow for recovery and retrograde labeling/virus expression for 2-3 weeks prior to euthanasia and tissue collection/processing.

For co-localization experiments (Fig. 6): AAV9-CAG-GFP (2×10^{12} viral particles/ml, UNC viral core) was injected in DCN (from bregma, in mm: medial n.: -2.55 AP, \pm 0.75 ML, -2.1 DV, 50 nl; interposed n. : -2.5 AP, \pm 1.55 ML, -2.1 DV, 50 nl; lateral n.: -2.2 AP, \pm 2.3 ML, -2.12 DV, 50 nl; and -1.8, \pm 2.35 ML, -2.12 DV, 50 nl). Cholera toxin subunit B (ctb)- 640 or -568 (5 mg/ml, Biotium) was injected in NAc shell and core (from bregma, in mm: 1.8 AP, \pm 0.8 ML, -4.2 DV, 200 nl). For anterograde transsynaptic tracing (Fig. 7): AAV1-hSyn-Cre-WPRE-hGH (10^{13} gc/ml, Addgene; 1:10 dilution) was injected in DCN (coordinates as above). AAV5-CAG-FLEX-tdTomato (7.8×10^{12} viral particles/ml, UNC viral core; 1 : 2 dilution) or AAV5-pCAG-FLEX-EGFP-WPRE (1.1×10^{13} gc/ml, Addgene; 1 : 2 dilution) was injected in VTA (from bregma, in mm: 2.8 AP, \pm 0.35 ML, -4.2 DV, 100 nl; and -2.85 AP, \pm 0.6 ML, -4.2 DV, 100 nl) or thalamus (from bregma, in mm: -0.85 AP, \pm 0.3 ML, -3.3 DV, 300 nl; and -1.2 AP, \pm 0.5 ML, -3.5 DV, 300 nl; 1:5 dilution).

Histology and fluorescence microscopy

Two to three weeks following tracer/virus injections, mice were anesthetized with an anesthetic cocktail (100 mg/kg ketamine, 10 mg/kg xylazine, 1 mg/kg acepromazine) and perfused transcardially with 4 % (w/v) PFA in PB. Brains were post-fixed in 4 % PFA for 6 h and transferred to 30 % sucrose in PBS for overnight incubation at 4 °C. Brains were coronally sectioned (60 μ m) on a sliding microtome, stained with DAPI, mounted to slides, coverslipped with Mowiol-based antifade solution and imaged. VTA sections were immunostained for

tyrosine hydroxylase (TH) prior to mounting, as follows: slices were first incubated with blocking solution [10 % normal goat serum (NGS) in PBS supplemented with 0.3 % Triton-X100; PBST] for 1 h at room temperature, then with mouse anti-TH (clone LNC11, Millipore Sigma; 1 : 1000) in blocking solution with 2 % NGS overnight at 4 °C. Sections were washed with PBST (3 x 20 min) and incubated for 1 h at room temperature with goat anti-mouse Alexa fluor 488 secondary antibody (1 : 1000), washed with PBS (3 x 20 min), mounted, coverslipped and imaged. Epifluorescence image mosaics were acquired on an Olympus VS120 slide scanner with a 10x air objective. High magnification confocal images were taken sequentially with different laser lines and a 60x oil-immersion objective on a Zeiss LSM800 microscope with Airyscan. Image brightness/contrast was adjusted using ImageJ (NIH) for display purposes.

Results

DCN microstimulation bidirectionally alters NAc spiking activity

To examine functional connectivity between CB and NAc, we recorded ongoing spiking activity from NAc_{Med} and NAc_{Core} *in vivo* and electrically microstimulated the DCN (Figure 1A,B). We primarily targeted the lateral DCN, activation of which has been shown to modulate levels of dopamine in NAc⁵³. In a subset of experiments, post-hoc analysis localized the bipolar stimulating electrode in the interposed DCN, and these data were included in our analysis. Activation of DCN with 100- μ A current pulses evoked two distinct types of responses in NAc: inhibitory (Fig. 1C) and excitatory (Fig. 1D). We define a response as the positive and/or negative threshold-crossing spiking activity that occurred within 3 s from stimulus presentation. The time course of all positive (*blue*) and negative (*red*) threshold-crossing spiking activity throughout the 10 s trial can be seen in the peri-stimulus time histograms in Fig. 2 (Fig. 2A1 and 2A2). Analysis of NAc responses revealed differences in the prevalence of each response type (i.e., inhibitory vs. excitatory), based on NAc subregion. In NAc_{Med}, DCN microstimulation elicited a significant number of inhibitory responses compared to pre-stimulus baseline (Fig. 2B,C)(proportion of inhibitory responses in NAc_{Med}: 16.7 %, comparison to pre-stimulus baseline activity: $p = 0.0$), as well as some excitatory responses (6.1 %)(Fig. 2B); however, the proportion of the latter did not differ significantly from that expected by chance alone ($p = 0.18$)(Fig. 2C). By contrast, DCN microstimulation elicited a significant number of both inhibitory and excitatory responses in NAc_{Core} (Fig. 2B,C; 17.1 % inh.: $p = 0.0$; 13.7 % excit.: $p = 0.003$). There was no significant difference between the prevalence of all (pooled excitatory

and inhibitory) responses between the two subregions ($p = 0.11$), or between the proportion of excitatory and inhibitory responses in NAc_{Core} ($p = 0.14$). Finally, there was no difference in the proportion of inhibitory responses between NAc_{Med} and NAc_{Core} ($p = 0.55$) (Table 1A-E).

To ensure that the observed relationships between excitatory and inhibitory responses in NAc_{Med} and NAc_{Core} did not hinge on response thresholds, we quantified proportions of responses for different threshold values (Fig. 2S1). We chose values that maintained a maximum of 5 % threshold crossings during pre-stimulus baseline (i.e., false positive response rate of 5 %). We found that the relative distribution of excitatory and inhibitory responses was robust to small changes in threshold. Together, these results suggest that (a) activation of DCN can modulate ongoing spiking activity in NAc and (b) this modulation is both inhibitory and excitatory. Interestingly, (c) excitatory modulation of spiking activity is more prevalent in NAc_{Core} than in NAc_{Med} ; and (d) in NAc_{Med} , inhibitory modulation is more prevalent than excitatory.

Temporal profiles of NAc responses

We evaluated the temporal profiles of NAc responses. In both NAc_{Med} and NAc_{Core} , excitatory responses appeared to precede inhibitory ones (Fig. 2A1,A2). This can be seen more clearly in 3D color plots (Fig. 3A,B), in which all excitatory and inhibitory responders are aligned to the time of stimulus onset and ordered by response-onset latency, with the z-transformed firing rate represented by color (corresponding color plots of all non-responders ordered by latency of peak activity are shown in Fig. 3S1). To quantitatively describe these differences, we compared onset latencies for excitatory and inhibitory responses using multiple-comparison-corrected random permutation tests (Fig. 3C and Table 1F,G). We found that excitatory responses were significantly faster (i.e., had shorter onset latencies) than inhibitory ones both in NAc_{Med} (mean \pm SEM: excit.: 0.63 ± 0.33 s, inh.: 1.24 ± 0.15 s, $p = 0.03$) and in NAc_{Core} (mean \pm SEM: excit.: 0.12 ± 0.02 s, inh.: 1.08 ± 0.14 s, $p = 0.0$). Moreover, excitatory responses were faster in NAc_{Core} than in NAc_{Med} ($p = 0.048$), whereas the onset latency of inhibitory responses did not differ significantly between the two subregions ($p = 0.22$).

To evaluate the time period over which NAc spiking activity remained supra-threshold, we plotted the cumulative probability of threshold crossings in NAc_{Med} and NAc_{Core} excitatory and

inhibitory responders as a function of time from stimulus onset (Fig. 3D1,D2). These plots are also informative of how fast, and presumably synchronous, the initial rise in activity was. This approach does not impose a theoretical model on the data, which may or may not be fitted adequately by the model, in order to extract information on temporal characteristics. We found that excitatory responses were indeed faster and shorter-lived than inhibitory responses in both NAc subregions [50 % rise-time (in s): NAc_{Med}: excit.: 0.22; inh.: 1.795, permutation test, $p = 0.01$; NAc_{Core}: excit.: 0.175, inh.: 1.195, permutation test, $p = 0.0$]. The 50 % rise-times of excitatory responses in NAc_{Med} and NAc_{Core} were not significantly different ($p = 0.36$), despite the few threshold crossings observed at later time points (Fig. 3D1). Finally, the small difference between NAc_{Med} and NAc_{Core} inhibitory responses (Fig. 3D2) did not reach significance ($p = 0.06$) (Table 1H). Collectively, these results suggest that DCN microstimulation reliably evokes fast, short-lived excitatory responses in NAc_{Core}, as well as slower and temporally protracted inhibitory responses in NAc_{Med} and NAc_{Core}.

Spatial considerations for electrical stimulation

To maintain fine control of electrical current spread at DCN stimulation sites, we used bipolar stimulation electrodes. To confirm that the stimulation was indeed localized, we performed current-spread analysis (Fig. 4D), which took advantage of the fact that we probed individual NAc recording sites with stimuli delivered at multiple DCN sites. We quantified the likelihood that stimulation of a DCN site would evoke a significant excitatory or inhibitory response in a NAc site and plotted it as a function of the Euclidean distance from the DCN site that evoked the strongest response in that NAc site (most effective DCN stimulation site, at $d = 0 \mu\text{m}$). We reasoned that if the electrical current spread over a large radius from the most effective DCN site, stimulation of other sites within that radius would also evoke NAc responses with likelihood greater than expected by chance. We found that even at close distances (100 μm) from the most effective site, the probability of a DCN stimulation site evoking a significant response was not different from the marginal probability (i.e., the probability to evoke a response across all recordings, regardless of distance from the most effective site) ($p = 0.45$) (Table 1I). This result confirms that our stimulation was localized.

Given that DCN stimulation was localized, could the observed differences in NAc response types arise from regional differences within DCN? To address this question, we

examined whether there was clustering of DCN sites with respect to their effectiveness to evoke significant excitatory or inhibitory responses in NAc_{Med} and/or NAc_{Core}. We did not find indications of such clustering (Fig. 4B,C), which argues against topographical organization of lateral DCN with respect to NAc response types. Using a similar approach, we examined whether there was topographical clustering of excitatory and/or inhibitory responder sites within NAc subregions, but we did not find evidence for any such organization either (not shown).

Differential and non-linear dependence of NAc responses on DCN stimulation intensity

To further probe the functional connectivity between DCN and NAc, in a subset of experiments we assessed the effectiveness of DCN microstimulation to modulate spiking activity in NAc_{Med} and NAc_{Core} at three different intensities: 30 μ A, 100 μ A, and 300 μ A. We found that the roughly 3-fold changes in stimulation intensity had (a) differential effects based on response type and NAc subregion, and (b) non-linear effects on the prevalence of response types in both NAc subregions.

Specifically, weak (30 μ A) DCN stimulation effectively evoked a small but significantly greater-than-baseline number of only inhibitory responses and only in NAc_{Med} (Fig. 5A,B) (proportion of responses: NAc_{Med}: inh.: 7.5 %, compared to pre-stimulus baseline, $p = 0.03$; exc.: 10.5 %, compared to pre-stimulus baseline, $p = 0.07$; excit. vs. inh: $p = 0.61$; NAc_{Core}: inh.: 1.5 %, compared to pre-stimulus baseline, $p = 0.5$; exc.: 8.7 %, compared to pre-stimulus baseline, $p = 0.14$) (Table 2A-B,D). Even after increasing intensity to 100 μ A, only inhibitory responses could be evoked reliably above baseline in NAc_{Med} (proportion of responses: inh.: 15.8 %, compared to pre-stimulus baseline, $p = 0.0$; exc.: 3.2 %, compared to pre-stimulus baseline, $p = 0.64$; exc. vs. inh.: $p = 0.02$). In contrast, in NAc_{Core} 100- μ A DCN stimulation evoked significant proportions of both inhibitory and excitatory responses (inh.: 19.8 %, compared to pre-stimulus baseline, $p = 0.0$; exc.: 11.9 %, compared to pre-stimulus baseline, $p = 0.01$) (exc. vs. inh.: $p = 0.01$). The differential effect of 100- μ A microstimulation on NAc subregions was statistically significant: more excitatory responses were evoked in NAc_{Core} than in NAc_{Med} ($p = 0.02$), whereas the prevalence of inhibitory responses was similar between subregions ($p = 0.29$). Finally, the 300- μ A DCN microstimulation evoked significant proportions of both inhibitory and excitatory responses in NAc_{Med} (inh.: 13.1 %, compared to pre-stimulus

baseline, $p = 0.0$; exc.: 19.1 %, compared to pre-stimulus baseline, $p = 0.003$; exc. vs. inh.: $p = 0.4$), as well as in NAC_{Core} (inh.: 17.3 %, compared to pre-stimulus baseline, $p = 0.004$; exc.: 24 %, compared to pre-stimulus baseline, $p = 0.0$)(exc. vs. inh.: $p = 0.0$), with no significant difference between the two subregions ($p = 0.28$ for excit. in NAC_{Med} vs. NAC_{Core}; $p = 0.29$ for inh. in NAC_{Med} vs. NAC_{Core}) (Table 2A-E).

The non-linear effects of stimulation intensity can be appreciated if we consider the change in the prevalence of response types across different intensities (Fig. 5B; Table 2F). In both NAC_{Med} and NAC_{Core}, increasing stimulation intensity from 30 μA to 100 μA did not significantly increase excitatory responses. In fact, in NAC_{Med} we observed a strong trend toward fewer responses at 100 μA (30 vs. 100 μA : excit.: NAC_{Med}: $p = 0.05$; NAC_{Core}: $p = 0.26$). Further increasing stimulation intensity from 100 μA to 300 μA significantly increased the proportions of excitatory responses in both subregions (100 vs. 300 μA : excit.: NAC_{Med}: $p = 0.0$; NAC_{Core}: $p = 0.004$). By contrast, for inhibitory responses a 30-to-100 μA stimulus intensity change was sufficient to significantly increase their prevalence in both NAC_{Med} and NAC_{Core} (30 vs. 100 μA : inh.: NAC_{Med}: $p = 0.04$; NAC_{Core}: $p = 0.0$). Further increase to 300 μA did not significantly affect inhibitory response prevalence in either subregion (100 vs. 300 μA : inh.: NAC_{Med}: $p = 0.27$; NAC_{Core}: $p = 0.29$). Notably, the localized nature of DCN stimulation was not compromised at 300 μA (comparison to marginal probability at 100 μm from most effective DCN site: $p = 0.2$; Table 2J-K). Collectively, these results suggest that DCN stimulation more effectively (i.e., at lower intensities) modulates inhibitory responses than excitatory ones, with NAC_{Med} being more responsive than NAC_{Core}. Modulation of excitatory responses requires stronger inputs, and this is particularly applicable to NAC_{Med}.

Response onset latencies are largely unaffected by DCN stimulation intensity

To investigate effects of DCN input strength on temporal characteristics of NAc responses, we examined the dependence of response onset latencies on DCN stimulation intensity. After correction for multiple comparisons, we found that increasing stimulation intensity did not significantly affect onset latencies of either excitatory or inhibitory responses, in either NAC_{Med} or NAC_{Core} (Table 2G-I). In agreement with our previous results (Fig. 3C), we found that, overall, excitatory responses tended to be faster than inhibitory ones. This is apparent in the 3-D color plots of z-transformed spiking activity for inhibitory and excitatory

responders (Fig. 5B) (see Fig. 5S1 for non-responders), as well as the frequency histograms of threshold crossings across all recorded sites (Fig. 5S2). Further quantification indicated that excitatory responses of NAc_{Med} had significantly shorter onset latencies than inhibitory ones at 300- μ A stimulation [onset latency (in s): mean \pm SEM: NAc_{Med}: 300 μ A: excit. 0.32 ± 0.17 , inh.: 1.24 ± 0.20 , $p = 0.0$]. The 100- μ A dataset did not reach significant difference ($p = 0.25$), probably due to sample size differences. In NAc_{Core}, excitatory responses were faster than inhibitory ones at 100- μ A and 300- μ A stimulation [onset latency (in s): NAc_{Core}: 100 μ A: excit. 0.11 ± 0.03 , inh.: 1.08 ± 0.14 , $p = 0.0$; 300 μ A: excit. 0.32 ± 0.15 , inh.: 1.29 ± 0.18 , $p = 0.0$] (Fig. 5D) (Table 2G,H). Comparing between subregions, our previous observation that excitatory responses in NAc_{Core} were faster than in NAc_{Med} (Table 1G) was also confirmed here (100 μ A: $p = 0.03$). There was no significant difference between the two subregions in the onset latencies of excitatory responses evoked by 30- μ A or 300- μ A stimulation (excit.: NAc_{Med} vs. NAc_{Core}: 30 μ A: $p = 0.06$; 300 μ A: $p = 0.57$); or in the latencies of inhibitory responses evoked by any stimulation intensity (inh.: NAc_{Med} vs. NAc_{Core}: 30 μ A: $p = 0.98$; 100 μ A: $p = 0.18$; 300 μ A: $p = 0.59$). The overall picture emerging from these analyses points to NAc receiving fast excitatory and slow inhibitory inputs in response to DCN stimulation. NAc_{Med} appears more attuned to inhibitory inputs, with significant excitation recruited only with very strong stimulation, whereas NAc_{Core} receives faster and more prevalent excitation compared to NAc_{Med}.

Anatomical blueprint of CB-NAc connectivity

There are no direct, monosynaptic connections between CB and NAc (Allen Brain Atlas, and our own observations); we therefore hypothesized the existence of at least disynaptic anatomical pathways between the two regions. To test this hypothesis, we adopted a 2-prong approach. First, we combined injection of a retrograde tracer (cholera toxin subunit B (ctb) - 640 or -568) in NAc with injection of an anterograde viral tracer (AAV9-CAG-GFP) in DCN to identify areas of overlap (nodes) in a putative disynaptic CB-NAc circuit (Fig. 6A,B). Histological processing and confocal imaging of brain sections ($n = 3$ mice with successful NAc and DCN injections, without spill to neighboring regions) revealed two regions of overlap between ctb-filled neurons that project to NAc and GFP-labelled DCN axonal projections: the VTA and limbic thalamus (Fig. 6C,D). In VTA, most areas of overlap localized medially and caudally and involved both TH+ (dopaminergic) neurons and TH- neurons (Fig. 6C). In

thalamus, we found areas of overlap in parafascicular (PF) and anterior centromedial (CM) intralaminar nuclei (Fig. 6D). These results are consistent with the existence of a disynaptic CB-NAc anatomical circuit, and point to VTA and intralaminar thalamic nuclei as putative nodes.

To independently confirm these observations, we performed AAV1-mediated anterograde transsynaptic tracing experiments⁶⁷ via stereotactic injections of AAV1-Cre in DCN and AAV-FLEX-tdTomato in VTA or limbic thalamus (Fig. 7A,B). The approach relies on the transsynaptic transfer of Cre to postsynaptic neurons, which -once infected with a floxed fluorophore- will become fluorescently labeled. With this method, we confirmed the existence of neurons receiving CB input in VTA (Fig. 7C1,C2) and limbic thalamus (Fig. 7E1,E2). Importantly, we were able to localize their labeled axonal projections in NAc, in both NAc_{Med} and NAc_{Core} (Fig. 7D,F). Similar projection patterns were observed when AAV1-Cre was injected in lateral DCN only (not shown). These findings chart the blueprint of disynaptic CB-NAc connectivity and provide an anatomical foundation for our newly discovered CB-NAc functional connection.

Discussion

In this study we examined the uncharted functional connectivity between CB and NAc through in vivo electrophysiology. We found that electrical microstimulation of DCN modulates spiking activity in NAc and elicits responses in both NAc_{Med} and NAc_{Core}. In both subregions, these responses manifested as inhibitory and/or excitatory modulations of spiking activity and exhibited NAc subregion-specificity with respect to response-type prevalence, latency, and dependence on stimulus intensity.

First, we found that weak-to-moderate DCN stimulation reliably evoked inhibitory, but not excitatory, responses in NAc_{Med}. Excitatory responses in NAc_{Med} could be reliably evoked only with the strongest DCN stimulation. In contrast, even moderate stimulation could reliably induce both inhibitory and excitatory responses in NAc_{Core}. Second, we found differences between the two subregions with respect to onset latency: excitatory responses were faster to start and shorter-lasting than inhibitory ones in both subregions. In NAc_{Core} in particular, excitatory responses tended to be faster than in NAc_{Med}. The differences in response profiles

between NAc_{Med} and NAc_{Core} are consistent with the fact that the anatomy and functions of these NAc subregions also differ^{55,60,61,63,68,69}.

The differences in prevalence, temporal characteristics and induction threshold of responses between NAc_{Med} and NAc_{Core} suggest that the two subregions do not receive the same copy of the CB signal. These differences are unlikely to arise from within DCN, because our analyses of the topographical organization of effective DCN stimulation sites (Fig. 4) indicated widespread and overlapping distributions of sites that could evoke excitatory or inhibitory responses in NAc_{Med} and/or NAc_{Core}. Even sites as close as 100 μm from each other could evoke quite different responses. If these different response profiles in NAc arise from distinct sources within DCN, these sources are likely organized at a scale below the resolution of our study (e.g., at the cellular level). Moreover, even if distinct pools of DCN neurons transmit signals to the two NAc subregions, the onset latency differential between responses suggests at least some divergence in the downstream routes that DCN signals take to arrive at their NAc destinations.

The CB's influence on NAc spiking activity is probably served by disynaptic (and/or possibly polysynaptic) pathways. This is supported by the finding that the onset of both excitatory and inhibitory responses in NAc_{Med}, as well as the onset of at least the inhibitory responses in NAc_{Core}, occurred well past the onset of stimulation and its artifact, which placed a lower bound on the latencies we were able to detect confidently. Our electrophysiological observations are also consistent with tracing experiments (Allen Brain Atlas; our own preliminary work) that have not found direct anatomical connections between CB and NAc, and are further corroborated by our viral tracing experiments (Figs. 6,7), which point to VTA and intralaminar nuclei as nodes in CB-NAc communication routes.

Anatomical and functional connections between DCN and VTA, and DCN and intralaminar n. nodes, have been described previously^{20,44,70-73}. VTA and thalamic projections to NAc are also well established^{62,74-81}. It might therefore not be surprising that the CB recruits these nodes to communicate with NAc. However, the VTA and thalamus are exquisitely complex areas with multiple output streams. For example, the VTA projects not only to NAc but also to hippocampus, hypothalamus, lateral habenula, entorhinal cortex, etc.⁸²⁻⁸⁴. None of these downstream target regions appears to be disynaptically connected to DCN, which points

to specificity in circuit wiring. Our study is the first one, to our knowledge, to map a CB-VTA-NAc circuit. A very recent study provided anatomical evidence for a disynaptic medial DCN-NAc circuit⁴³. Here we extend these observations to lateral DCN and also provide evidence for the centromedial and parafascicular nuclei as parts of the thalamic node.

The cell types and properties of nodal neurons that receive DCN input remain incompletely understood. In the VTA, for example, glutamatergic DCN projections target both dopaminergic and non-dopaminergic neurons^{20,73,85}. Given the known heterogeneity of VTA neural populations^{86,87}, DCN inputs could activate dopaminergic, GABAergic, glutamatergic, and/or dual neurotransmitter-releasing neurons. This diversity in neural types, which are known to project to NAc with some subregion-specificity^{57,81,88–92}, could at least partly explain the differences in responses and onset latencies we observed in our experiments. For example, dopamine's action on G protein-coupled receptors to modulate neuronal activity could be responsible for the long latencies and/or duration in NAc_{Med} inhibitory responses. Thalamic neurons also show at least partial NAc subregion specificity^{61,74,93}, which could explain the faster excitatory responses in NAc_{Core} compared to NAc_{Med}.

It is tempting to speculate on network-wide implications of the different CB-NAc response profiles. The fast excitatory profiles would be well poised to support the rapid communication of information critical to the control of motivated behavior, such as prediction or prediction-error signals, which are well established in the cerebellum^{19,21,22,94}. The slower, less synchronous and longer-lasting inputs may be suggestive of a regulatory function, e.g. gain control of the communication between NAc and other brain regions (Buzsáki et al., 2007). If the DCN signals do in fact reach NAc_{Med} and NAc_{Core} through distinct pathways, this leaves room for their distinct multiplexing with signals originating elsewhere. Future investigations into the cellular basis of CB-NAc communication as well as its behavioral contributions are clearly in order. Here, we have broken new ground by providing the first evidence of functional connectivity between CB and NAc, identifying its NAc subregion-dependence, and offering an anatomical blueprint that could serve as its foundation.

Conflict of Interest

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

Author contribution

AD, EA and DF designed the research; AD and SJJ performed the research; AD and EA analyzed the data; SG assisted with tissue slicing and registration; SJJ assisted with preparation of anatomy figures; AD, EA and DF wrote the paper.

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Figure legends

Figure 1. Cerebellar stimulation elicits excitatory and inhibitory responses in NAc. **A**, Schematic diagram of recording setup. Recording electrode array was lowered into NAc and bipolar stimulating electrode was lowered into deep cerebellar nuclei. Stimulation protocol consisted of 10 trials (inter-trial interval: 15 s) of five 100 μ A, 0.5-ms pulses at 200 Hz. **B**, Recording sites in NAc_{Med} and NAc_{Core}. Numbers at bottom of slices indicate distance from bregma. **C-D**, Examples of inhibitory and excitatory responses in NAc_{Med} (C) and NAc_{Core} (D), respectively, as a function of time. *Top*, 4-s raster plots of spiking activity; *Middle*, Peri-stimulus time histograms of firing rate (10-ms bin); *Bottom*, Average firing rate, normalized to baseline (z-score). Red dotted lines indicate response threshold, set at -3.2 for inhibitory responses and +4.0 for excitatory responses. Stimulus artifacts have been masked for clarity (teal bars).

Figure 2. The distribution of response types differs between NAc shell and core. **A**, Percent of recordings crossing positive (*blue*) or negative (*red*) threshold at each 10-ms time bin for NAc_{Med} (A1) and NAc_{Core} (A2) regions, as a function of time. Shaded area indicates 3-s response window. **B**, Pie charts of distribution of response types in NAc_{Med} and NAc_{Core}. Recordings that cross positive threshold within the response window are classified as excitatory

responders (*blue*); recordings that cross negative threshold: inhibitory (*red*); recordings that do not cross either threshold: non-responders (*gray*). **C**, Comparison of % responses vs. pre-stimulus baseline, by type (excitatory, inhibitory) and NAc region (shell, core). * $p < 0.05$, corrected for multiple comparisons (FDR 10%).

Figure 2- figure supplement 1. Robustness of response distributions to small fluctuations in threshold. Percent threshold crossings during response window (solid lines) and pre-stimulus baseline (dotted lines) for different types of responses (*red*: inhibitory; *blue*: excitatory) in NAc_{Med} and NAc_{Core}, as a function of threshold. Selected threshold values (vertical dashed line) maintain a maximum of 5% crossings during baseline (horizontal dashed line) for each response type and NAc subregion. **B**, Same as in A, but for subset of data that compared three CB stimulation intensities (30, 100 and 300 μ A).

Figure 3. Excitatory and inhibitory responses show different temporal profiles. A,B, Heat maps of firing rate (z-score) of excitatory (A) and inhibitory (B) responders in NAc shell and core, as a function of time. Responders were ordered on the y-axis by response onset latency. Black triangle denotes time of stimulation; white vertical bars mask stimulus artifact for clarity. **C**, Comparison of onset latencies for excitatory and inhibitory responses within each NAc subregion. * $p < 0.05$, corrected for multiple comparisons (FDR 10%). **D1,D2**, Cumulative probability histograms of threshold crossings for excitatory (D1) and inhibitory (D2) responders in NAc shell and core, with most crossings falling within 3 seconds of stimulus onset (shaded region).

Figure 3- figure supplement 1. Heat maps of non-responders at 100 μ A stimulation. Firing rate (z-score) of non-responders, ordered on the y-axis by latency of peak response amplitude, as a function of time. Gray box outlines the response window.

Figure 4. Differential distribution of excitatory and inhibitory responses in NAc is not due to topographical specialization within CB. A, The lateral or interposed DCN was electrically stimulated. Abbreviations: m: medial n., ip: interposed n.; lat: lateral n. **B,C**, Stereotactic coordinates of CB stimulation sites in ML-DV (B1,C1), ML-AP (B2,C2), and AP-DV (B3,C3)

planes. Colored dots denote sites that evoked excitatory (*blue*), inhibitory (*red*), or no responses (*gray*) in NAc_{Med} (B) and NAc_{Core} (C). Abbreviations: ML: medio-lateral, DV: dorso-ventral, AP: antero-posterior, excit.: excitatory, inh.: inhibitory, non-resp.: non-responding. **D**, Spatial resolution of stimulation (current spread). Plot of the probability that a stimulation site would elicit a NAc response at various distances from the CB site that elicited the strongest response in each experiment (most effective site at distance $d = 0 \mu\text{m}$). Black horizontal line shows marginal response probability.

Figure 5. Varying CB stimulation intensity changes the relative distribution of response types within NAc, but leaves temporal profiles mostly unaffected. **A**, Distribution of response types in NAc shell (A1) and core (A2) evoked by 30 μA , 100 μA and 300 μA bipolar CB stimulation. **B**, Comparison of % responses vs. pre-stimulus baseline, by type (excitatory, inhibitory), stimulation intensity (30 μA , 100 μA and 300 μA), and NAc subregion (shell, core). * $p < 0.05$, corrected for multiple comparisons (FDR 10%)(see also Table 2). **C**, Firing rate (z-score) heat maps of excitatory and inhibitory responders in NAc shell and core, as a function of time. Responders were ordered on the y-axis by latency of response onset. Black triangle denotes time of stimulation; white vertical bars mask stimulus artifacts for clarity. **D**, Boxplots of onset latencies for excitatory and inhibitory responses in NAc shell (*top*) and core (*bottom*), elicited by 30 μA , 100 μA and 300 μA CB stimulation. Figure legend in A also applies to panels B and D. Abbreviations: non-resp.: non-responding, inh.: inhibitory, excit.: excitatory.

Figure 5- figure supplement 1. Profiles of NAc population activity at different CB stimulation intensities. **A,B**, Time histograms of % recordings in NAc shell (A) and core (B) that cross positive (*blue*) or negative (*red*) threshold at each 10-ms time bin, for CB stimulation intensity of 30 μA (A1,B1), 100 μA (A2,B2) and 300 (A3,B3) μA . Shaded region designates 3-s response window.

Figure 6. Co-localization of NAc-projecting neurons with DCN projections in VTA and intralaminar thalamus. **A**, Schematic diagrams of stereotactic injections in DCN and NAc. **B1**, Expression of GFP at DCN injection sites. **B2**, Ctb-568 injection site in NAc. **C**, Overlap of ctb-labeled NAc projectors and DCN axons in VTA. **C1**, VTA identification through TH

immunostaining. **C2**, TH+ neurons. **C3**, Ctb-labeled NAc projectors. **C4**, GFP-expressing DCN axons. **C5**, C2-C4 merged. **D**, Overlap of ctb-labeled NAc projectors (red) and GFP-expressing DCN axons (green) in intralaminar thalamic nuclei. **D1-D2**, Overlap in parafascicular n. **D3-D4**, Overlap in centromedial n. Yellow boxes in C1,D1,D3 denote zoom-in areas depicted in C2-C5, D2 and D4, respectively. Scale bars: B1-2,C1,D1,D3: 200 μm ; C2-C5: 20 μm ; D2,D4: 10 μm . Blue: DAPI. Numbers denote distance from bregma.

Figure 7. Disynaptic CB-NAc connectivity is confirmed via anterograde transsynaptic viral tracing. **A**, Schematic diagram of AAV1-mediated transsynaptic labeling approach. **B**, Schematic diagrams of stereotactic injections of floxed fluorophore in VTA and thalamic nodes. **C1**, The VTA is visualized via TH immunostaining (green). Yellow box denotes zoom-in area depicted in C2. Rn: red nucleus; ml: medial lemniscus. **C2**, Neurons that receive DCN input are labeled with tdTomato. Green: TH+ neurons (arrow). Red: tdTomato+ neurons (white arrowhead). Orange: TH+, tdTomato+ neuron (yellow arrowhead). **D**, Projections of CB-VTA neurons in NAc shell (NAc_{Med}) and core (NAc_{Core}). **E1**, Neurons that receive DCN input in thalamus are labeled with tdTomato (red). Blue: NeuN. CM: centromedial n.; PC: paracentral n.; CL: centrolateral n.; VL: ventrolateral n. **D2**, Same as D1, but for parafascicular n. (PF). fr: fasciculus retroflexus. **F**, Projections of CB-thalamic neurons in NAc_{Med} and in NAc_{Core}. Scale bars below images: 500 μm , except for C2: 100 μm . Numbers denote distance from bregma.

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TABLE 1							
A	<i>NUMBER OF OBSERVATIONS</i>						
		BASELINE			RESPONSE		
	TOTAL	POSITIVE	NEGATIVE	POSITIVE	NEGATIVE	(threshold crossings)	
	NAc SHELL	132	4 (3.0%)	0 (0%)	8 (6.1%)	22 (16.7%)	
	Nac CORE	117	3 (2.6%)	4 (3.4%)	16 (13.7%)	20 (17.1%)	
B	<i>COMPARISONS OF RELATIVE FREQUENCY OF RESPONSES VS. PRE-STIMULUS BASELINE</i>						
		NAc SHELL		NAc CORE			
		POSITIVE	NEGATIVE	POSITIVE	NEGATIVE		
	p-value*	0.176	0.0	0.003	0.0		
	* bold red if significant after FDR correction						
C	<i>COMPARISON OF PERCENT RESPONDERS IN CORE VS. SHELL</i>						
	NAc SHELL	NAc CORE	p-value				
	22.727273	30.769231	0.111				
D	<i>COMPARISONS OF FREQUENCY OF INHIBITORY VS. EXCITATORY RESPONSES</i>						
		NAc SHELL	NAc CORE				
	p-value	0.0	0.142				
E	<i>COMPARISONS OF RESPONSE FREQUENCY IN SHELL VS. CORE</i>						
		EXCITATORY	INHIBITORY				
	p-value	0.036	0.548				
F	<i>RESPONSE LATENCIES (sec; mean ± SEM)</i>						
		EXCITATORY	INHIBITORY				
	NAc SHELL	0.63 ± 0.33	1.24 ± 0.15				
	NAc CORE	0.12 ± 0.02	1.08 ± 0.14				

TABLE 2							
A							
<i>NUMBER OF OBSERVATIONS</i>							
				BASELINE		RESPONSE	
		TOTAL	POSITIVE	NEGATIVE	POSITIVE	NEGATIVE	
	NAc SHELL	30 μ A	67	2 (3.0%)	0 (0%)	7 (10.5%)	5 (7.5%)
		100 μ A	95	3 (3.2%)	0 (0%)	3 (3.2%)	15 (15.8%)
		300 μ A	84	4 (4.8%)	0 (0%)	16 (19.1%)	11 (13.1%)
	NAc CORE	30 μ A	69	2 (2.9%)	1 (1.5%)	6 (8.7%)	1 (1.5%)
		100 μ A	101	2 (2.0%)	4 (4.0%)	12 (11.9%)	20 (19.8%)
		300 μ A	75	1 (1.3%)	3 (4.0%)	18 (24%)	13 (17.3%)
B							
<i>COMPARISONS OF RELATIVE FREQUENCY OF RESPONSES VS. PRE-STIMULUS BASELINE</i>							
			NAc SHELL		NAc CORE		
			POSITIVE	NEGATIVE	POSITIVE	NEGATIVE	
	p-value*	30 μ A	0.066	0.028	0.14	0.50	
		100 μ A	0.64	0.0	0.005	0.0	
		300 μ A	0.003	0.0	0.0	0.004	
	* bold red only if significant after FDR correction						
C							
<i>COMPARISON OF PERCENT RESPONDERS IN CORE VS. SHELL</i>							
		NAc SHELL	NAc CORE	p-value			
	30 μ A	17.9104478	10.144928	0.15			
	100 μ A	18.9473684	31.683168	0.028			
	300 μ A	32.1428571	41.333333	0.13			
D							
<i>COMPARISONS OF FREQUENCY OF INHIBITORY VS. EXCITATORY RESPONSES</i>							
			NAc SHELL	NAc CORE			
	p-value	30 μ A	0.61	0.029			
		100 μ A	0.0	0.011			
		300 μ A	0.38	0.0			
E							
<i>COMPARISONS OF RESPONSE FREQUENCY IN SHELL VS. CORE</i>							
			EXCITATORY	INHIBITORY			
	p-value	30 μ A	0.5	0.11			
		100 μ A	0.022	0.287			
		300 μ A	0.28	0.285			
F							
<i>COMPARISONS OF RESPONSE FREQUENCY ACROSS STIMULATION INTENSITY</i>							
			EXCITATORY		INHIBITORY		
			NAc SHELL	NAc CORE	NAc SHELL	NAc CORE	
	30 μ A vs. 100 μ A	0.05	0.257	0.037	0.0		
	100 μ A vs. 300 μ A	0.0	0.004	0.268	0.29		
	30 μ A vs. 300 μ A	0.023	0.004	0.12	0.001		

TABLE 2- continued					
G	<i>RESPONSE LATENCIES (sec; mean ± SEM)</i>				
		NAc SHELL		NAc CORE	
		EXCITATORY	INHIBITORY	EXCITATORY	INHIBITORY
	30 μA	1.0 ± 0.29	0.89 ± 0.41	0.41 ± 0.22	2.58 ± 0.0
	100 μA	0.88 ± 0.69	1.27 ± 0.20	0.11 ± 0.03	1.08 ± 0.14
	300 μA	0.32 ± 0.17	1.24 ± 0.20	0.32 ± 0.15	1.29 ± 0.18
H	<i>Random permutation tests on response latencies</i>				
		p-value			
		30 μA	100 μA	300 μA	
	Excitatory v Inhibitory	0.13	0.0	0.0	
	E v I in Core	0.13	0.0	0.0	
	E v I in Shell	0.59	0.25	0.0	
	Core v Shell	0.27	0.021	0.55	
	Core v Shell in Excit.	0.06	0.029	0.57	
	Core v Shell in Inhib.	0.98	0.18	0.59	
I		30vs100 μA	100vs300 μA	30vs300 μA	
	Core/Excitatory	0.13	0.13	0.24	
	Core/Inhibitory	0.034	0.16	0.052	
	Shell/Excitatory	0.44	0.16	0.045	
	Shell/Inhibitory	0.15	0.44	0.21	
J	<i>SPATIAL SPREAD OF CURRENT (responder probability)</i>				
		stimulation intensity			
		30 μA	100 μA	300 μA	
	distance from most effective site	100 μm	0.13	0.17	0.27
		200 μm	0.06	0.28	0.24
		300 μm	0.0	0.25	0.0
		400 μm	0.0	0.20	0.25
	marginal prob	0.14	0.26	0.37	
K	<i>Comparison to marginal (p-values of random permutation test)</i>				
		stimulation intensity			
		30 μA	100 μA	300 μA	
	distance from most effective site	100 μm	0.662	0.284	0.211
		200 μm	0.094	0.699	0.057
		300 μm	0.637	0.720	0.027
		400 μm	0.371	0.383	0.277

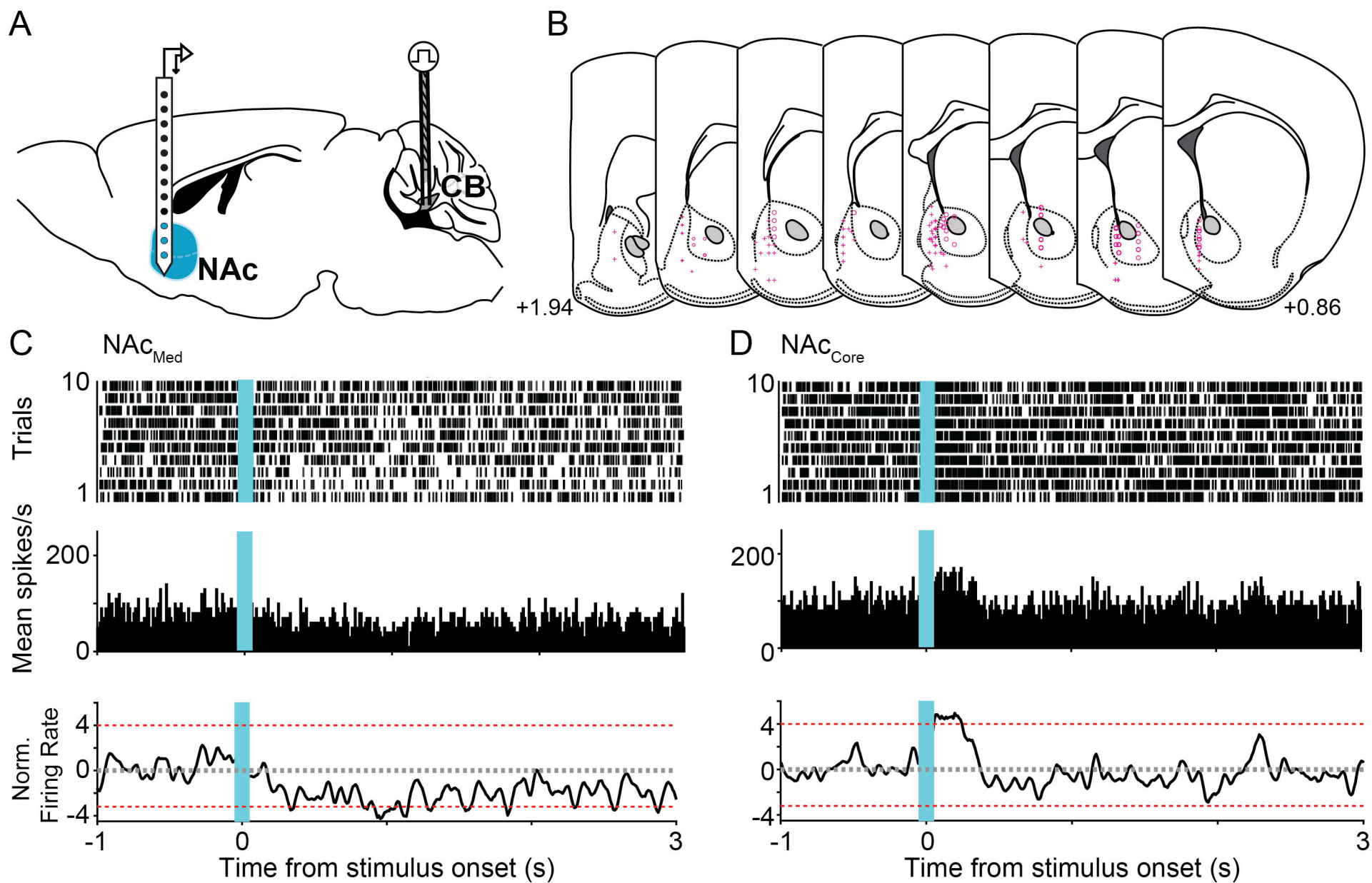


Figure 1

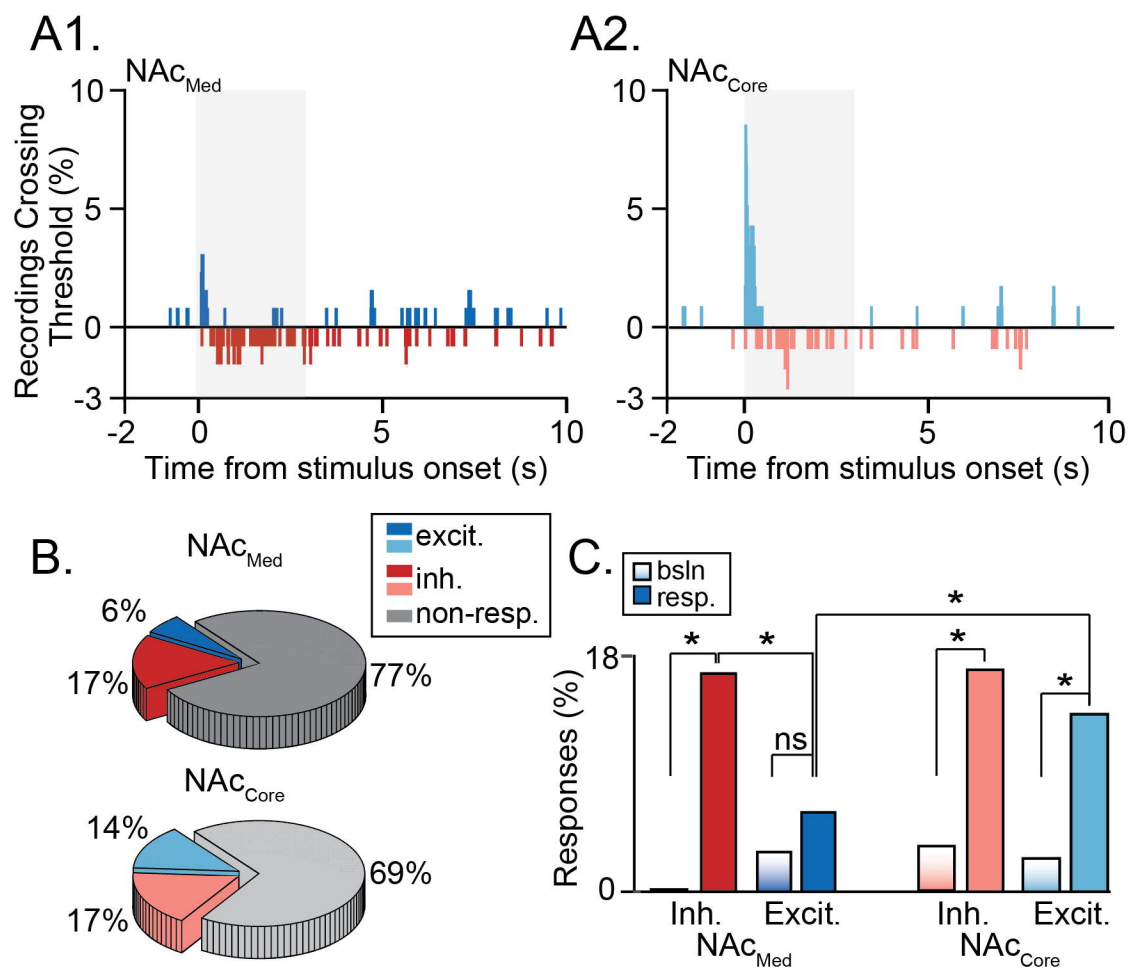


Figure 2

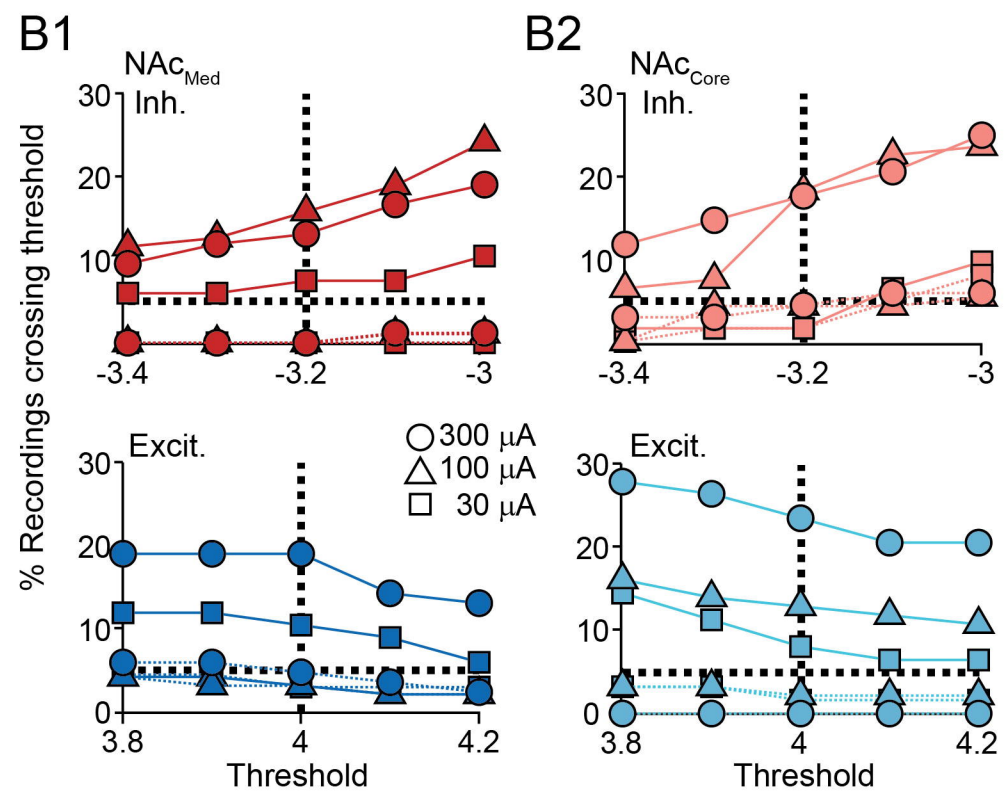
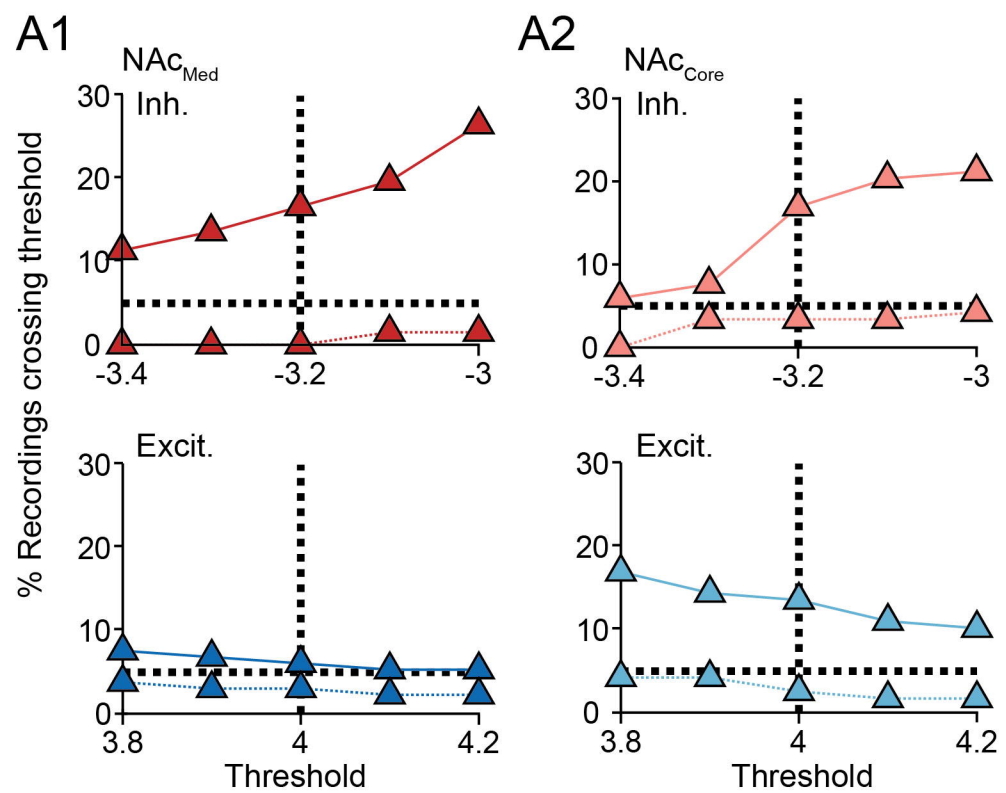
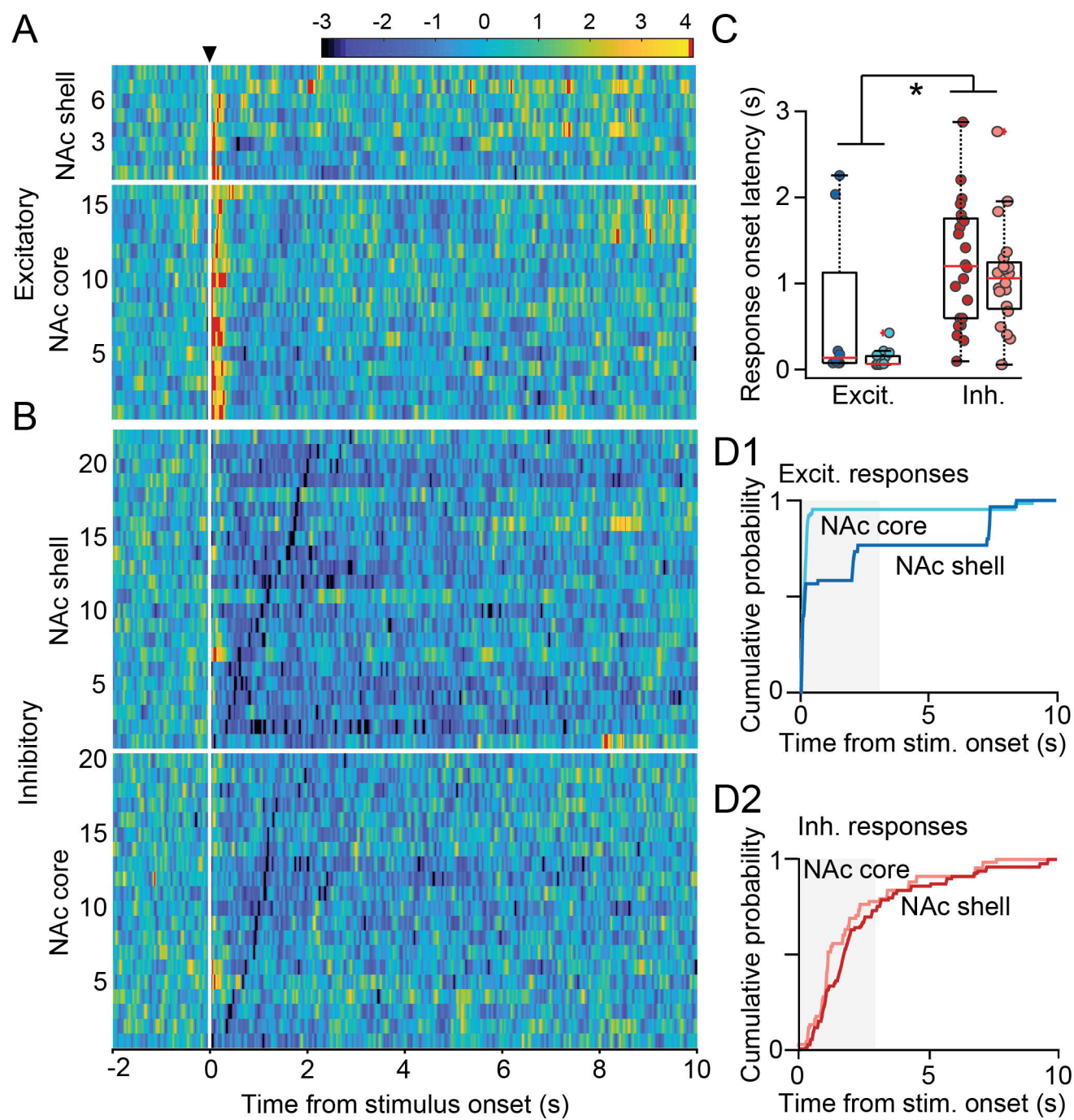


Figure 2S1



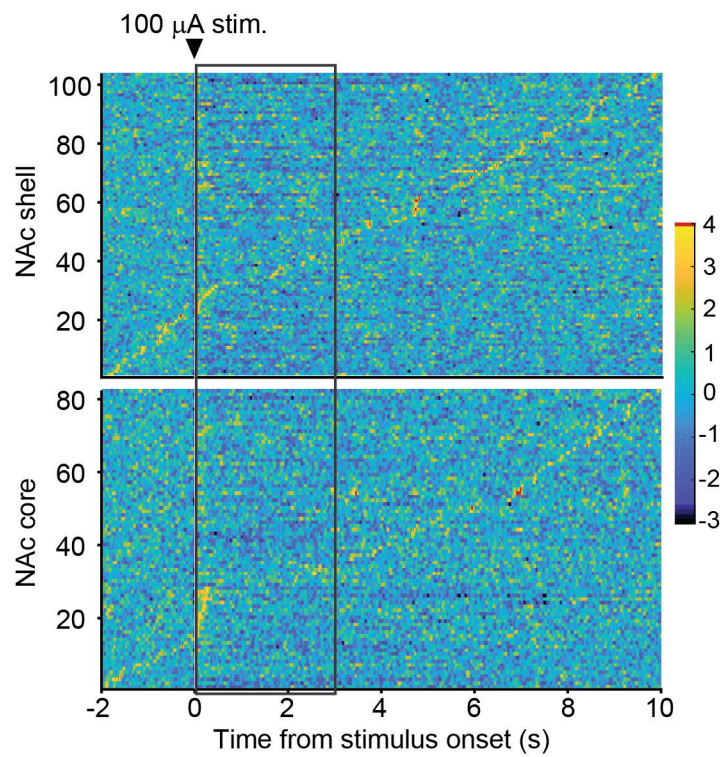


Figure 3S1

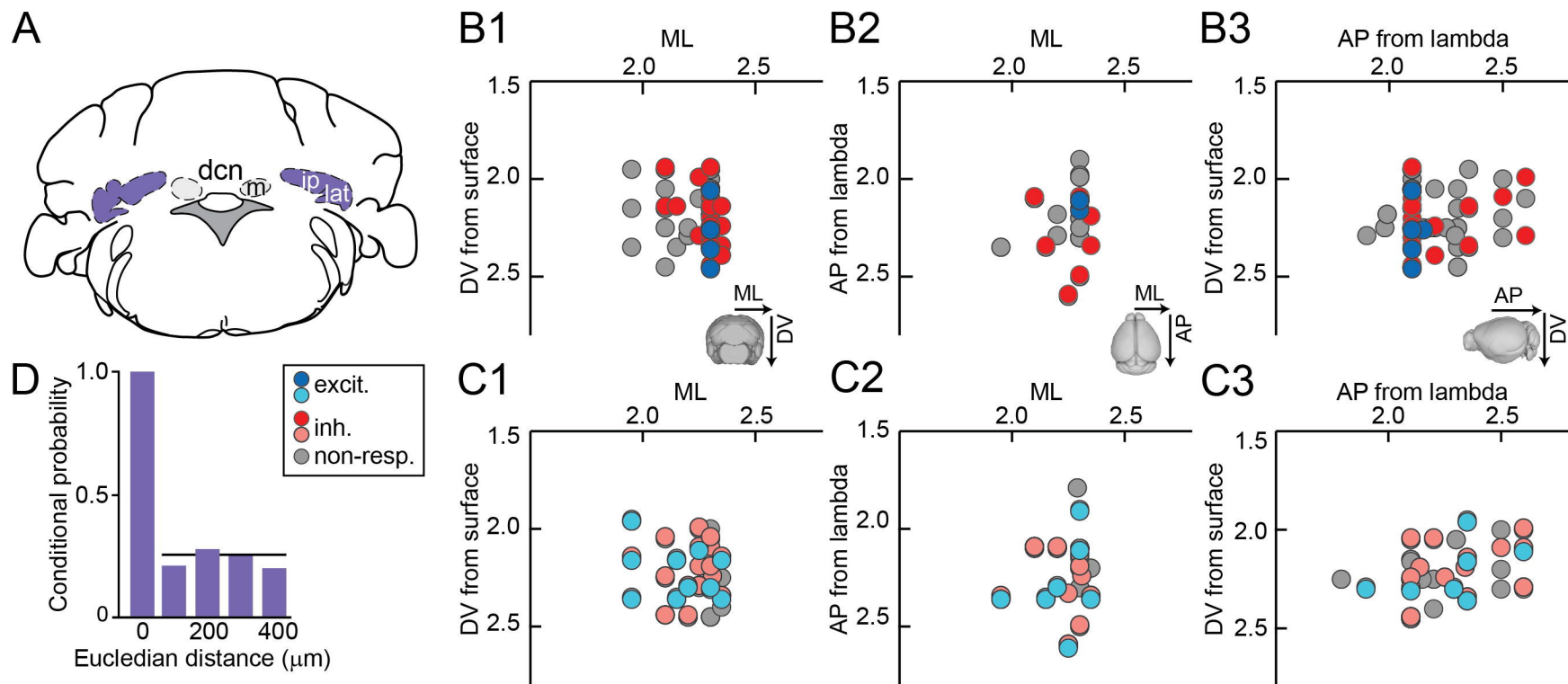


Figure 4

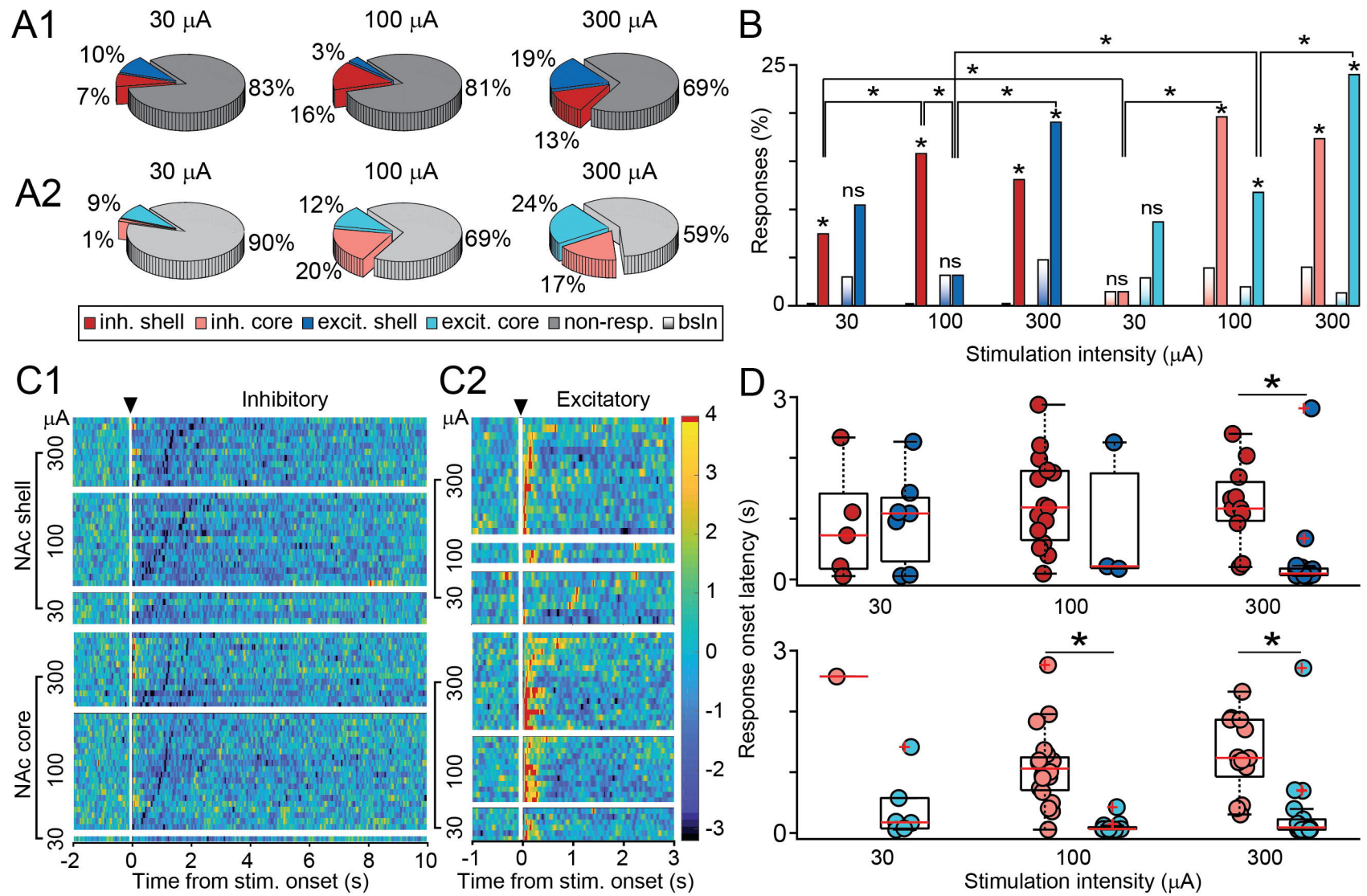


Figure 5

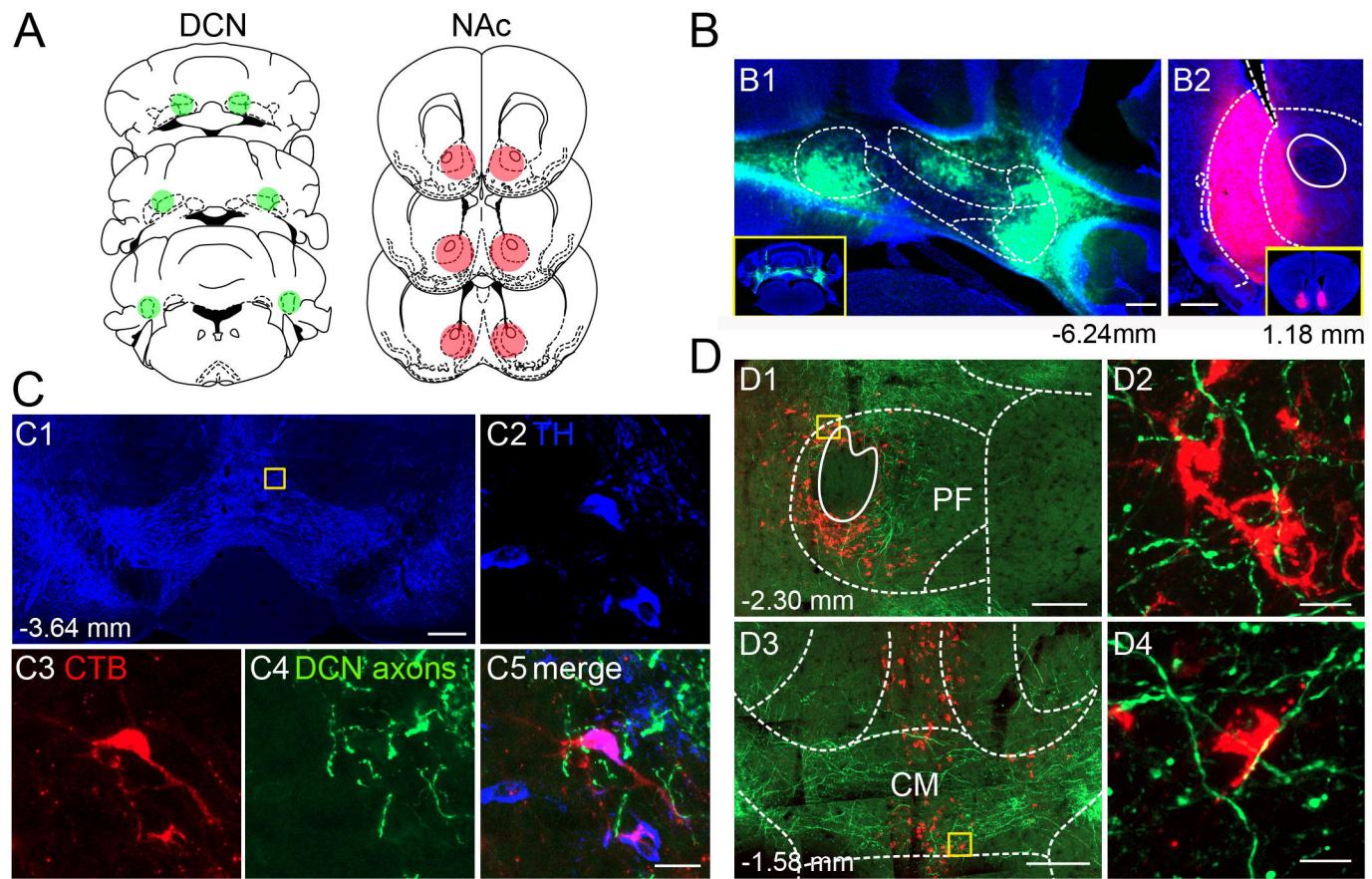


Figure 6

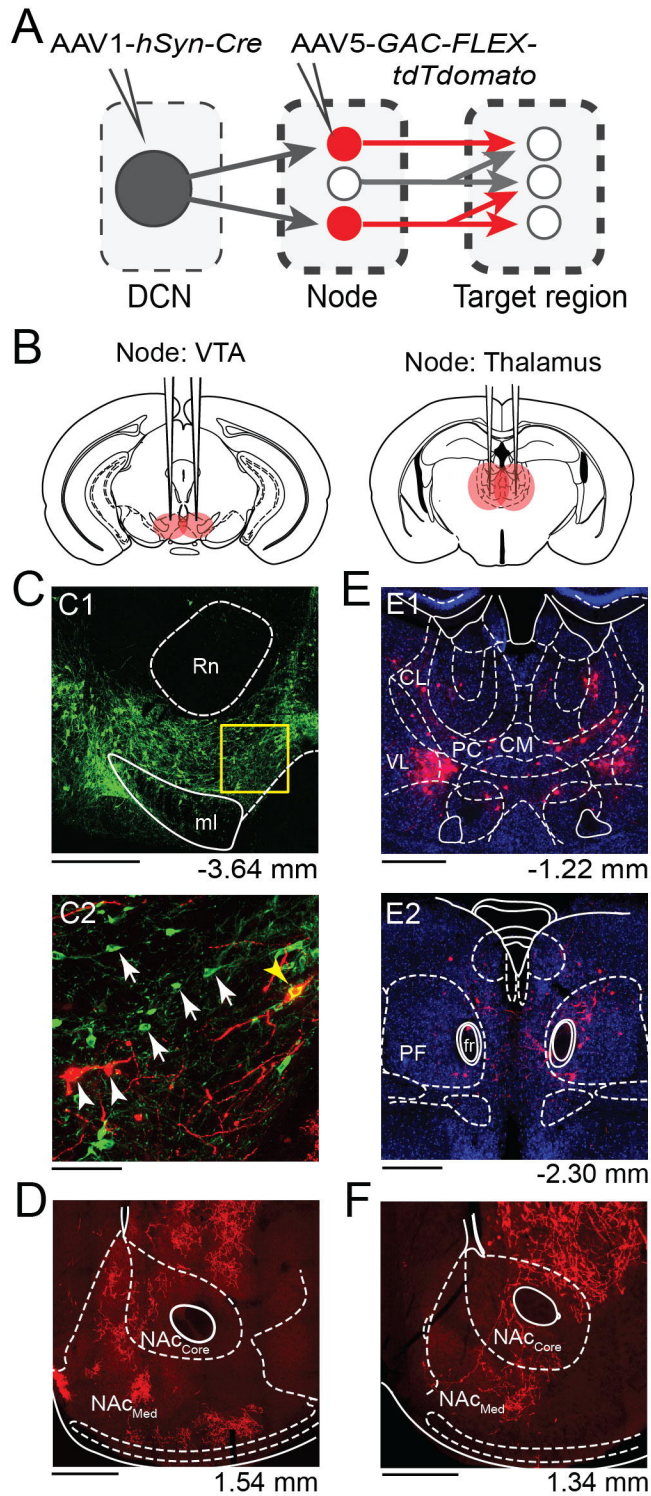


Figure 7