Thalamic regulation of switching between cortical representations enables cognitive flexibility

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Interactions between the prefrontal cortex (PFC) and mediodorsal thalamus are critical for cognitive flexibility, yet the underlying computations are unknown. To investigate frontothalamic substrates of cognitive flexibility, we developed a behavioral task in which mice switched between different sets of learned cues that guided attention toward either visual or auditory targets. We found that PFC responses reflected both the individual cues and their meaning as task rules, indicating a hierarchical cueto-rule transformation. Conversely, mediodorsal thalamus responses reflected the statistical regularity of cue presentation and were required for switching between such experimentally specified cueing contexts. A subset of these thalamic responses sustained context-relevant PFC representations, while another suppressed the context-irrelevant ones. Through modeling and experimental validation, we find that thalamic-mediated suppression may not only reduce PFC representational interference but could also preserve unused cortical traces for future use. Overall, our study provides a computational foundation for thalamic engagement in cognitive flexibility.

ognitive flexibility, the ability to mentally switch between different thoughts and action plans, is critical for survival in a rapidly changing environment¹⁻³. This important process allows us to flexibly switch attention among competing inputs⁴⁻⁷. A lack of cognitive flexibility is a hallmark of many mental illnesses, such as schizophrenia^{8,9}. Furthermore, a key limiting factor to artificial general intelligence is the inability of deep learning algorithms to perform multiple tasks without interference^{10,11}. Therefore, elucidating the circuit and computational principles underlying cognitive flexibility will have a broad multidisciplinary impact.

The PFC plays a central role in cognitive flexibility^{5,12,13}, including the differential allocation of attentional resources based on learned cues^{14–16}. Multiple recent studies have also demonstrated that PFC function is highly dependent on its interactions with the mediodorsal thalamus (MD)^{17–22}. In particular, the MD sustains task-relevant representations in the PFC by augmenting effective connectivity between cortical neurons¹⁷. However, because previous studies have not included a controllable switching component, the role of MD–PFC interactions in cognitive flexibility remain unclear.

In this study, we examined the substrates of cognitive flexibility through a series of behavioral manipulations, temporally precise optogenetic perturbations, and multisite multielectrode recordings. We found that PFC responses reflected both the individual cues and their meaning as task rules, indicating a hierarchical cue-to-rule transformation in this cortical area. In contrast, MD responses reflected the statistical regularity of the cue presentation, which we refer to as the cueing context. Using causal perturbations, we found that in addition to stabilizing context-relevant representations, MD neurons also suppress context-irrelevant PFC representations. These processes impart to the PFC the flexibility to dynamically switch between different contexts with minimal interference. Altogether, our work clarifies how MD neurons regulate prefrontal representational switching and provides a computational foundation for thalamic engagement in cognitive flexibility.

Results

Prefrontal neurons display mixed selectivity during attentional switching. To examine how mouse PFC ensembles operate when cognitive flexibility is required, we expanded an attentional control task^{17,23} to incorporate a cue-switching component (Fig. 1a). At the core of the task is sensory selection, where freely behaving mice select between spatially conflicting visual and auditory targets. On each trial, a mouse selects between the two targets based on one of two 100-ms-long learned cues, a high-pass or a low-pass noise burst. These cues correspond to two rules: attend to audition and attend to vision. Mice were required to hold a pseudorandomly presented cue in mind for up to 1 s by maintaining snout fixation in the initiation port before the simultaneous presentation of the two targets. The targets corresponded to the spatial location of the reward being delivered through the right or left reward port (for example, left light-emitting diode (LED) flash on an attend-to-vision trial signaled a response on the left reward port). Correct performance was rewarded with 10 µL of condensed milk, while incorrect performance was punished with a timeout. Logistic regression modeling of behavior across all mice used in this study revealed that they used the cue to guide their choices (Supplementary Fig. 1a-d and see Methods).

Once mice became proficient at using the auditory noise cues, we introduced two visual cues, an ultraviolet flash and a green LED flash, which corresponded to the same rules (Fig. 1b). To assess how mice switched from using one cue set to another, we trained mice to perform this task in blocks (Fig. 1b). Mice had equivalent performance across both blocks regardless of their presentation order (Supplementary Fig. 1e,f), suggesting that they used different cue sets equivalently. Critically, this demonstrates an ability to flexibly switch their attention when the cueing context changes.

Given the well-known role for the PFC in cognitive flexibility^{1,20,24,25}, we asked how PFC neurons (prelimbic cortex; see Methods) engage in this task. Using unbiased trial-selection (Supplementary Fig. 1g) and spike-waveform-clustering analysis^{26–28}, we classified

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recorded neurons into two categories: regular spiking (RS, putative excitatory) and fast spiking (FS, putative inhibitory; Supplementary Fig. 2a-d). As previously reported¹⁷, a subset of RS neurons showed a brief increase in spike-timing reliability during the delay period. We refer to these cells as transient (see Supplementary Fig. 2e-g for classification of cells). As a population, these transient cells tiled time in the delay period with distinct neurons responding at different timepoints (Supplementary Fig. 2h). Notably, these cells could be further categorized into two groups. One group responded selectively to one of the four learned cues (Fig. 1c,d; cue-selective, 233 of 1,789 cells in 5 mice), while another responded equivalently to two cues that had the same meaning, which may be interpreted as a single task rule (Fig. 1e,f and Supplementary Fig. 2i; cue-invariant, 102 of 1,789 cells in 5 mice). For example, a cue-invariant PFC neuron selective to the attend-to-vision rule responded with transient elevation in spiking reliability at the same delay-period time in both low-pass noise and green-LED trials. To our knowledge, this type of cue-invariance, indicative of rule-selectivity^{29,30}, has not been previously reported in mouse PFC.

In stark contrast to the transient RS neural responses, putative inhibitory FS neurons showed broad changes in spike rate that distinguished between the two cueing contexts but not rule meaning (Fig. 1g,h and Supplementary Fig. 2j; 418 neurons 5 mice). In fact, RS and FS populations encoded distinct cognitive variables, with the rule signal being more readily decodable from RS neurons while the context signal was more readily decodable from FS neurons (Fig. 1i,j). Further analysis confirmed that RS neurons encoded cues and rules through changes in spiking reliability (Supplementary Fig. 3a,b). On the other hand, FS populations encoded context through broader (persistent) changes in spike rates across both the intertrial interval and the delay period. Note that in addition to transient PFC RS cells, we observed another RS group that showed persistent spike rate changes that also correlated with context (Supplementary Fig. 3c). However, it was much harder to decode context from these cells than from PFC FS neurons (Supplementary Fig. 3d,e).

To further test whether transient PFC responses were indeed due to the sensory cue, we omitted the cue on 20% of the trials (Supplementary Fig. 4a). In addition to a decrease in behavioral performance, we observed a significant decrease in tuning strength of both PFC cue-selective and cue-invariant neurons (P < 0.0001; Supplementary Fig. 4b-d). Taken together, these results indicate that transient PFC responses were due to the learned cue, with

cue-selective cells representing a physical cue and cue-invariant cells representing its meaning as a task rule.

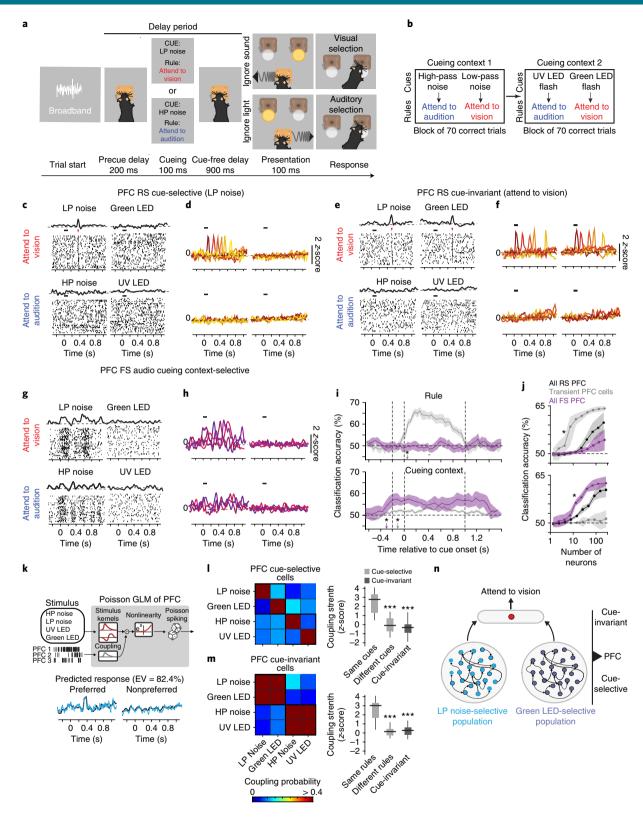
Is it possible that the observed mixed selectivity among cue-invariant neurons reflected differential sampling of local cueselective neurons in a context-dependent manner? To address this question, we constructed a multineuronal generalized linear model (GLM; Fig. 1k) to predict the spike rate of each PFC neuron^{31,32}. In addition to external sensory variables, our GLM model also included coupling terms to capture the dependencies between neurons (see Methods). These coupling terms were in the form of a coupling filter that allowed us to explain the effect that spiking in other simultaneously recorded neurons had on the unit being modeled. Notably, we used a rigorous cross-validation approach to statistically evaluate the predicted coupling and to constrain the parameters of the model (see Methods). On average, the GLM was able to explain close to 75% of the response variance of both cue-selective and cue-invariant PFC neurons on each trial (Supplementary Fig. 5a-c).

By analyzing the strength of the coupling filters, we were able to make inferences about the functional connectivity between the different classes of PFC neurons. This analysis revealed that cue-selective neurons were strongly coupled among themselves only when they encoded the same cue, and they did not receive substantial reciprocal coupling from cue-invariant neurons (Fig. 11). Additionally, cue-invariant neurons received strong functional inputs from cue-selective neurons across both contexts (Fig. 1m). As such, based on this pattern of functional connectivity, we reasoned that task-relevant PFC computations were hierarchically organized, with cue-invariant neurons gaining their rule selectivity from cue-selective neurons across contexts (Fig. 1n).

Mediodorsal thalamus encodes cueing context. The MD projects extensively to the PFC and has a critical role in maintaining task-relevant activity in this cortical region^{18,20–22,33}. Furthermore, a recent study found that the MD might play a role in cognitive flexibility by recruiting cortical inhibitory neurons³⁴. Given our finding that FS neurons were contextually selective (Fig. 1h,i), we next asked how MD neurons responded in our context-switching task (Fig. 2a). In agreement with our previously published results¹⁷, we found that a subset of MD neurons exhibited a transient increase in spiking reliability during the delay and that distinct MD neurons tiled the delay period (Fig. 2b). Unlike PFC RS neurons, these transient MD

Fig. 1 | Prefrontal neurons display selectivity indicative of a hierarchical cue to rule transformation during attentional switching. a, Schematic of task design. b, Mice were trained to associate four cues with two rules. These cues were presented in two blocks, each containing two cues. An animal had to achieve at least 70 correct trials in a block before moving on to the next block. For details, see Methods. UV, ultraviolet. c, Example peristimulus time histogram (PSTH) and raster plot (number of trials vs. time) for an RS PFC neuron that is selective to a low-pass (LP) noise. The black bar above the raster marks the cueing period, and the red arrowhead indicates the transient increase in spiking reliability. HP, high-pass. d, Transient responses tile the duration of the delay period. Each color is a different cue-selective neuron. e.f., As in c,d but for PFC cue-invariant cells. g,h, As in c,d but for PFC FS cells. Unlike RS cells, these neurons have persistent changes in firing rate over the delay period. Representative examples in \mathbf{d} , \mathbf{f} , and \mathbf{h} drawn from n=5 mice (independent samples). i, Classification accuracy over time relative to cue onset for a decoder trained to predict either rule (top) or cue context decoding (bottom) for PFC RS and FS neuronal populations. Asterisks denote the timepoints at which classification accuracy is significantly (i.e., P < 0.05, permutation test from n = 5 mice) above chance (50% classification accuracy). j, Classification accuracy (within delay period) scales with the number of neurons. As in i, the asterisk indicates the number of neurons at which classification accuracy is significantly above chance levels (P < 0.05, permutation test from n = 5mice). k, Top: schematic of Poisson GLM. Bottom: model prediction (gray) of the PSTH (black) for one example PFC neuron. EV, explained variance. I, Left: heatmap showing coupling probability between the four cue-selective cell PFC cell types. Right: box-and-whisker plots comparing the coupling strengths of inputs to PFC cue-selective neurons from cue-selective neurons preferring the same or different cues (light gray; $P=1.23\times10^{-4}$) or cue-invariant neurons (dark gray; $P = 0.18 \times 10^{-4}$; Bonferroni-corrected Kruskal-Wallis ANOVA with post hoc rank-sum test relative to neurons preferring the same cues; n = 5 mice). **m**, As in **I** but characterizing the inputs to cue-invariant PFC neurons from cue-selective neurons preferring the same or different rules $(P=1.89\times10^{-6})$ or cue-invariant neurons $(P=1.42\times10^{-6})$; Bonferroni-corrected Kruskal-Wallis ANOVA with post hoc rank-sum test relative to neurons preferring the same rules; n = 5 mice). **n**, Cartoon schematic showing how cue-invariant neurons gain their selectivity by pooling from cue-selective neurons across both cueing contexts. Data is shown as mean ± 95% confidence interval (shaded error bars). Boxplots: center line, median; box edges, 95% confidence interval; whiskers, range.

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neurons did not display cue-selective responses but had equivalent responses to both cues within the same block. Additionally, another subset of MD neuron showed the same selectivity toward the cueing context but through persistent changes in spike rate over the delay period (see Fig. 2c, d for classification of transient and persistent MD neurons). At the population level, the cueing context was much more decodable from both persistent and transient MD neurons than from PFC RS neurons (Fig. 2e, f). Therefore, across

the PFC-MD network, MD and PFC FS neurons were the most informative of the cueing context, whereas PFC transient neurons were most informative of the rule (Fig. 2f).

Are these thalamic responses reflective of sensory input (i.e., the modality of the cues) or of something more cognitive? To answer this question, we required mice to perform the task using blocks with cues of different modalities. MD neuronal activity reflected these heteromodal cueing blocks (Supplementary Fig. 6a–d).

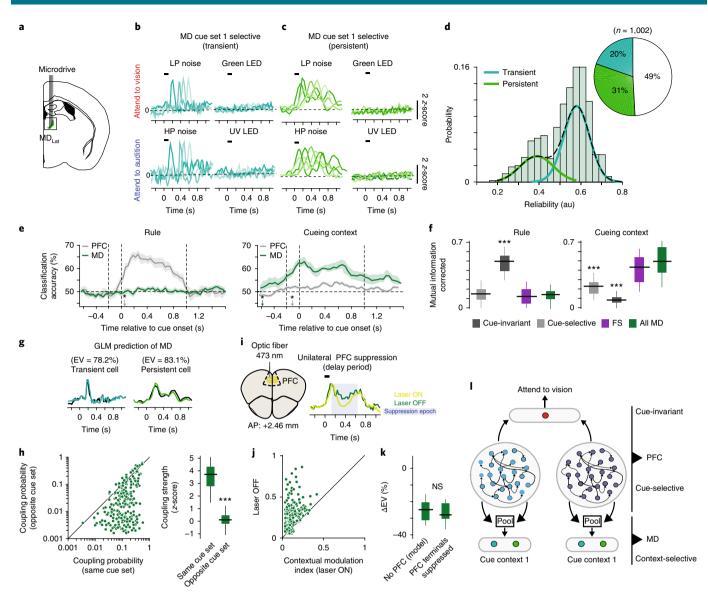


Fig. 2 | MD responses reflect the cueing context. a, Schematic for MD recordings. MDLat, lateral subdivision of the MD. b,c, Example PSTHs of MD neurons with transient (b) and persistent (c) responses to both cues within the auditory cueing context. Each color indicates a distinct neuron. Representative examples drawn from n=5 mice (independent samples). **d**, Histogram of intertrial spiking reliability scores from all task-modulated MD neurons. A Gaussian mixture model (dashed lines) was used to classify cells into either persistent (low reliability) or transient (high reliability). Inset: pie chart quantifying the fraction of each MD neuron type. e, Classification accuracy over time relative to cue onset for a decoder trained to predict either rule (top) or cue-context decoding (bottom) trained to classify rule (left) and cue context (right) from PFC RS and MD populations (5 mice). Asterisks denote the timepoints at which classification accuracy is significantly (i.e., P < 0.05, permutation test, from n = 5 mice) above chance (50% classification accuracy). f, Comparison of rules decoding and cueing context decoding accuracy, measured as mutual information (see Methods), between all recorded cell types (***P < 0.001, Bonferroni-corrected Kruskal-Wallis ANOVA with post hoc rank-sum test relative to MD neurons; n = 5 mice). \mathbf{g} , GLM of MD neurons can accurately predict the responses of both transient (right) and persistent cells (left). EV, explained variance. Black lines, data; colored lines, predictions. h, Left: comparison of coupling probability between PFC cue-selective neurons and MD neurons within the same cue set and the opposite cue set. Right: boxplot comparing coupling strengths between MD and PFC cells selective to cues in the same cue context or the opposite context (***P=0.15 x 10⁻⁴; Bonferroni-corrected Kruskal-Wallis test; n=345 MD neurons, 5 mice). i, Left: schematic illustrating unilateral PFC suppression. Right: example MD neuron with suppressed firing rate following PFC suppression. Shaded blue area marks time during which the laser was turned on. AP, anteroposterior. i, Comparison of contextual modulation index on laser ON and laser OFF trials. k, Change in GLM prediction, measured as Δ EV, when PFC filters are excluded from the model (no PFC) compared to a model fit to MD responses following PFC suppression (nonsignificant (NS), P = 0.42; Bonferroni-corrected Kruskal-Wallis test, n=186 MD neurons, 3 mice). I, Schematic of proposed PFC-MD connectivity. Data in d-h is from 5 mice; data in j,k is from 3 mice. Data is shown as mean ± 95% confidence interval (shaded error bars). Boxplots: center line, median; box edges, 95% confidence interval; whiskers, range.

Furthermore, when we presented all four cues in a randomized manner, we found MD neurons that responded to all four cues, suggesting that their combination was encoded as a single context (Supplementary Fig. 6e-h). Altogether, MD activity reflected the statistical regularity of cue presentation over a multitrial timescale, which we refer to as the cueing context (Supplementary Fig. 6i,j).

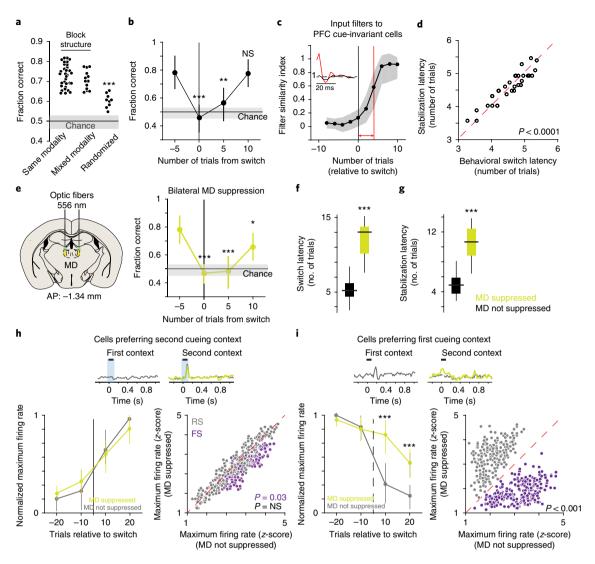


Fig. 3 | Flexible switching between contexts is associated with MD-dependent changes in PFC activity. a, Comparison of fraction correct trials between the different task conditions. Each data point is one session per mouse from 3 mice in total (***P = 0.023 × 10⁻³; Bonferroni-corrected Kruskal-Wallis ANOVA). Note that each session is treated as an independent sample. b, Change in behavioral performance (fraction correct) relative to switch (**P < 0.01, ***P < 0.001; one-way rank-sum test relative to 5 trials before the switch; n = 33 independent sessions from 3 mice). Data is shown as mean ± s.e.m. c, Change in filter similarity index of coupling filters from cue-selective to cue-invariant neurons in the PFC relative to switch. Insets: example coupling filter changing between the point of switch (black) to its final stable value (red). Red line marks the filter stabilization latency. Shaded area, 95% CI. For details, see Methods. d, Scatter plot relating behavioral switch latency with the filter stabilization latency for inputs to cue-invariant PFC neurons. Each data point is a session ($P = 0.0068 \times 10^{-6}$, two-way rank-sum test, n = 33 independent sessions from 3 mice). **e**, Left: schematic illustrating bilateral MD suppression. Right: change in behavioral performance (fraction correct) relative to switch for sessions with bilateral MD suppression. Statistics and plotting as in **b. f.g.** Boxplots comparing the effect of MD suppression on (**f**) behavioral switching latency (*** $P = 0.78 \times 10^{-4}$, Kruskal-Wallis ANOVA) and (g) cue-invariant input filter stabilization latency (*** $P = 0.19 \times 10^{-4}$, Kruskal-Wallis ANOVA; n = 33 sessions with no suppression, 31 sessions with MD suppression, from 3 mice). h, Top: example PSTH of a PFC neuron selective to cues in the second cueing context. Left: time course of the change in normalized maximum firing rate (relative to stable behavior) relative to the switch. No significant difference (P = 0.92 between suppressed and nonsuppressed conditions). Right: scatter plot comparing the maximum firing rate of PFC cue selective neurons (gray) and PFC FS neurons (purple) 10 trials after switch. i, As in h but showing the effect of MD suppression on PFC cells that are selective for cues in the first cueing context (left: ***P < 0.001, one-way rank-sum test for each timepoint between suppressed and nonsuppressed conditions, n=3 mice; right, P<0.001, one-way rank-sum test for each timepoint between suppressed and nonsuppressed conditions; n=3 mice). Data is shown as mean \pm s.e.m.

What factors could explain this contextual selectivity in the MD? We have previously shown that transient responses in the MD are dependent on PFC inputs¹⁷. Therefore, one possibility is that the MD gains contextual selectivity from PFC inputs, either from persistent RS neurons or from cue-selective ones (Supplementary Fig. 7a,b). To test these two models, we fit GLMs to MD neurons and analyzed how different PFC cell types contributed to their selectivity (Fig. 2g). PFC RS persistent cells did not contribute to

spiking of either MD transient or persistent cells (Supplementary Fig. 7c–f). In contrast, more than 75% of the variance in the delay-period activity of both transient and persistent MD neurons could be explained by inputs from cue-selective PFC neurons, with MD neurons more likely to receive inputs from the context-congruent cortical cue set (Fig. 2h). Also, consistent with the fact that PFC FS neurons do not project to the MD, we found that they exerted no causal influence on MD spiking, further validating the power of the

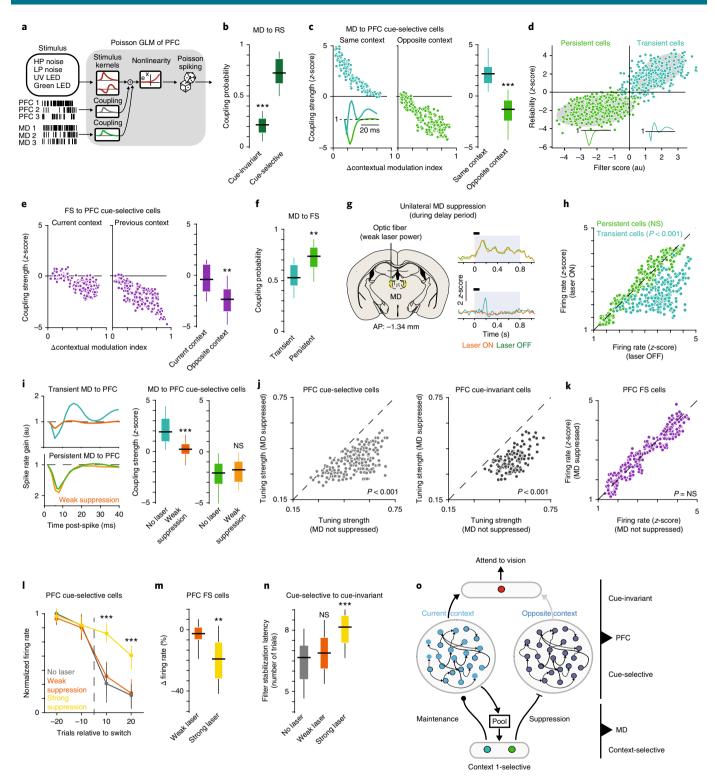
GLM to infer biologically plausible circuit models (Supplementary Fig. 7c,f). Therefore, these findings suggest that MD neurons gained their contextual selectivity, at least partly, by pooling context-specific cue inputs from the PFC. MD transient cells pooled from PFC cells in a temporally precise manner, while MD persistent cells pooled from PFC cells over a broader temporal window (Supplementary Fig. 7g,h).

To further verify these model predictions, we expressed the inhibitory channelrhodopsin iC++ in PFC to directly inactivate its neurons or their terminals in MD (PFC $_{\rm MD}$; Fig. 2i and Supplementary Fig. 8a-c) in a temporally precise manner. We reasoned that if MD neurons derived contextual selectivity from the PFC, then suppressing PFC_{MD} inputs should decrease its selectivity. To mitigate the detrimental effect that bilateral PFC suppression has on behavior in this task¹⁷, we suppressed the PFC unilaterally once the animal achieved stable performance in each block (Supplementary Fig. 8b and see Methods). This method allowed us to dissociate neural selectivity in the MD from changes in behavioral performance. Suppressing PFC neurons themselves, or their terminals in MD, diminished contextual signals in the MD (Fig. 2j and Supplementary Fig. 8d,e). Notably, removing PFC input filters in the MD GLM had an effect on response predictability similar to that of suppressing PFC_{MD} terminals, providing further experimental validation of our GLM (Fig. 2k). Finally, to test the idea that PFC inputs were causally involved in generating and not just modulating contextual selectivity in the MD, we suppressed PFC_{MD} inputs on every trial during the cueing period (Supplementary Fig. 9a). When performed unilaterally, this manipulation significantly decreased MD contextual selectivity (P<0.0001; Supplementary Fig. 9b,c), and when performed bilaterally, it significantly impaired behavioral performance (P < 0.0001; Supplementary Fig. 9d-f). Taken together, these results suggest that by pooling from cue-specific PFC neurons, the MD encodes the cueing context (Fig. 21).

Encoding a cueing context is critical for behavioral performance and flexibility. Our results so far suggest that both MD and PFC FS explicitly encode cueing context, while PFC RS neurons encode other cognitive variables, such as cue identity and rule. We wondered whether behavioral performance benefitted from such a behavioral encoding scheme, and if so, how? We noticed that mice performed much better on sessions in which the four cues were separated into two cueing contexts compared to sessions in which the cues were equiprobable (Fig. 3a). This performance advantage occurred regardless of how cues in the block were grouped with respect to modality. Critically, although performance within either of the two blocks was higher than when cues were fully randomized across the session, there was a clear and consistent behavioral detriment for 4–8 trials upon switching from one block to another (Fig. 3b). This decline in behavioral performance at the switch suggested that, despite previous learning, mice had to readjust to using cues in the new cueing set.

This behavioral switching dynamic was associated with several neural ones. We found that cue-selective PFC neurons showed reliable spiking earlier than cue-invariant ones upon first exposure to the new context (Supplementary Fig. 10a). This suggests that the decrease in behavioral performance at the point of context switch could be due to a remapping of inputs onto the shared cue-invariant cells. To test this hypothesis, we analyzed the temporal evolution of coupling filters from cue-selective to cue-invariant PFC cells on a trial on a trial-by-trial basis at the point of the switch (Supplementary Fig. 10b and see Methods). Notably, the time taken for these coupling filters to stabilize followed broadly similar dynamics to that of the behavioral performance (Fig. 3c) and could explain close to 87% of the variance in the behavioral switch latency: sessions in which the mouse took longer to switch were also associated with a longer time to stabilize cue-selective inputs into cue-invariant neurons (Fig. 3d). This would be expected if PFC cue-invariance was the source of cognitive control signals (attendto-vision versus attend-to-audition). In contrast, although the coupling between cue-selective PFC neurons and context-selective MD neurons followed broadly similar dynamics, they were too fast to correlate with behavioral performance (Supplementary Fig. 10c-g). Therefore, the output of the PFC cue-invariant cells, and not the MD or PFC cue-selective cells, were most likely used for controlling sensory selection and successful task performance.

Fig. 4 | Distinct MD neurons augment and suppress context-relevant PFC representations. a, Schematic of Poisson GLM used to model PFC neurons including MD interactions. **b**, Boxplot comparing the coupling probability between MD and PFC cue-selective (n = 230 neurons from 5 mice) or cueinvariant neurons (n = 86 neurons from 5 mice; *** $P = 0.58 \times 10^{-5}$; two-way rank-sum test relative to cue-selective neurons). **c**, Left: scatter plot relating the coupling strength between MD and PFC cue-selective cell with the difference in contextual selectivity for both cells that prefer the current context or the previous context. Inset: two different coupling filters between MD and PFC cells. Each data point is one PFC cell from 5 mice. Right: boxplot comparing the difference in coupling strength between MD and PFC neurons preferring the same context (n=141 MD neurons from 5 mice) or the opposite context $(n=211 \text{ MD neurons from 5 mice}; ***P=0.89\times10^{-4}; \text{two-way rank-sum test}).$ **d**, Clustering analysis relating MD-PFC coupling strength to MD spiking reliability. Each data point is one MD cell (n = 352 neurons from 5 mice). Shaded gray area is the 95% confidence interval ellipse of a Gaussian mixture model. Inset: median filter shape from each cluster. e, Right: scatter plot relating FS to PFC cue-selective coupling strength with difference in contextual modulation. Each data point is one PFC cell from 5 mice. Left: boxplot comparing the difference in coupling strength between FS and PFC neurons preferring the same context (n=141 neurons from 5 mice) or the opposite context (n=211 neurons from 5 mice; ** $P=0.31\times10^{-2}$; two-way rank-sum test). f, Boxplot of coupling probabilities between MD and PFC FS cells (**P=0.78×10⁻²; two-way rank-sum test relative to transient MD neurons; n=410 PFC FS neurons, 5 mice). g, Left: method for unilaterally suppressing the MD. Right: PSTHs of two example MD neurons. Persistent MD cells (top) are less affected by weak MD suppression than transient MD cells (bottom). Shaded blue area marks the duration of the laser; black bar marks the cueing period. h, Scatter plot comparing the effect of weak MD suppression on the firing rates of transient (n = 260 neurons from 3 mice, $P = 0.89 \times 10^{-3}$) and persistent MD cells (n = 247 neurons from 3 mice, NS, P = 0.22, Friedman test between laser ON and laser OFF trials). i, Left: example MD-PFC coupling filters with (orange) and without (green) MD suppression. Right: boxplot comparing the effect of MD suppression on the coupling strength between transient and persistent MD cells and PFC cue-selective neurons (n=177 neurons, 3 mice; *** $P=1.02\times10^{-4}$ and P=0.12 (NS), two-way rank-sum text). **j**, Scatter plot showing the change in tuning strength of cue selective (left, n=177 neurons) and cue-invariant (right, n=127 neurons, 3 mice) neurons caused by weak MD suppression. Friedman test between laser ON and laser OFF trials. k, As in j, but showing no significant effect of weak MD suppression on PFC FS neurons (264 neurons, 3 mice, P = 0.81). I, Time course of the change in normalized maximum firing rate relative to switch of PFC cells selective for cues in the first cue set (n=3 mice; ***P < 0.0001, one-way rank-sum test relative to laser OFF condition). Colors indicate various levels of MD suppression; data shown as mean \pm s.e.m. **m**, Boxplot comparing the change in firing rate of PFC FS neurons with weak and strong MD suppression (n = 264 and 212 neurons, respectively, from 3 mice; $**P = 0.71 \times 10^{-2}$, one-way Kruskal-Wallis ANOVA). **n**, Boxplot comparing cue-selective to cue-invariant filter stabilization latency (laser OFF, n = 33 sessions; weak laser, n = 31 sessions; strong laser = 18 sessions from 3 mice; *** $P = 0.063 \times 10^{-4}$, one-way Kruskal-Wallis ANOVA with post hoc rank-sum test). o, Cartoon summarizing the distinct effect that MD transient and persistent cells exert on the PFC. All boxplots: center line, median; box edges, 95% confidence interval; whiskers, range.



These results also suggest that the contextual selectivity of the MD may be required for the generation of rule signals in the PFC by adjusting the functional connectivity between cue-selective and cue-invariant PFC neurons in a context specific manner. To causally test this model, we designed and executed a series of optogenetic perturbation experiments. Our previous study demonstrated that bilateral suppression of MD neurons during the duration of the delay period diminished task-relevant activity in the PFC¹⁷. In addition, in tasks lacking a delay period, MD suppression via halorhodopsin (see Methods) during the cueing period (100 ms) had

minimal effect on behavioral performance¹⁷. As such, we first asked whether cue-specific, interleaved, bilateral MD suppression had a measurable impact on behavioral performance. Notably, once an animal achieved stable performance within a block, such manipulation had no impact (Fig. 3e). Instead, the biggest behavioral deficit that we observed was the prolonged time taken to achieve stable performance in the new context (Fig. 3f and Supplementary Fig. 11). Consistent with the idea that MD contextual signals are relevant for establishing PFC task-relevant connectivity patterns, this MD manipulation also increased the number of trials taken to stabilize

cue-invariant representations in the PFC (Fig. 3g). Notably, this laser manipulation had no unwanted effects on the MD, as the same laser power and duration had no effect in control mice that expressed GFP in the MD (data not shown).

MD neurons regulate PFC representational switching, likely through cortical inhibition. In addition to the effects on behavioral performance and PFC representational stability, we also noted that temporally precise MD suppression during the cueing period impacted cue-selective PFC neural spiking. Specifically, although the increase in spiking of cells preferring the second context was unaffected (Fig. 3h), we observed that cells preferring the first context continued to fire even though their sensory cue was no longer present (Fig. 3i). Suppressing MD terminals in the PFC resulted in a similar 'out-of-context' spike rate elevation (Supplementary Fig. 12). Critically, these MD-dependent changes in PFC RS spiking activity were contrasted by changes in PFC FS firing; MD suppression attenuated the normal elevation of FS neural spiking associated with the second context (Fig. 3i). Therefore, at least a subset of MD neurons may regulate representational switching by suppressing out-of-context activity in the PFC through cortical inhibitory mechanisms.

To more directly probe this process, we turned to our multineuronal GLM to assess the impact of MD neurons on PFC targets (Fig. 4a). We found that, in contrast to cue-invariant PFC neurons (Fig. 4b), cue-selective PFC neurons received substantial MD inputs, which varied according to context (Fig. 4c). These functional inputs could be broadly segregated into two types, one predominantly inhibitory and another predominantly excitatory (Fig. 4c). Notably, these functional inputs originated from the two distinct MD functional subgroups; persistent MD neurons were more likely to provide inhibitory functional inputs, while transient MD neurons predominantly provided excitatory ones (Fig. 4d).

Similar to MD neurons, PFC FS neurons also exerted a context-dependent inhibitory effect on cue-selective neurons, with FS neurons having a larger inhibitory effect on PFC neurons that preferred cues of the opposite context (Fig. 4e). Consistent with the idea that MD cell types may be exerting part of their effects on the PFC through local inhibitory circuits^{34,35}, we found that MD inputs could explain more of the variance of PFC FS neuron firing than PFC cue-selective neurons could (Supplementary Fig. 13a–c). MD persistent neurons were more coupled to PFC FS neurons than MD

transient neurons were (Fig. 4f). Also, in contrast to FS neurons, PFC RS persistent neural responses were poorly explained by inputs from either the MD or cue-selective PFC neurons (Supplementary Fig. 13d), reinforcing the notion that they may be part of a distinct functional circuit other than the one under study. Altogether, these results support a model in which the MD controls contextual switching by suppressing PFC neurons of the irrelevant context through mechanisms that involving cortical inhibition.

To further test this model causally, we needed to gain a degree of selectivity over the two identified functional MD subtypes (Fig. 4g). MD neurons in vitro may have a bimodal resting-membrane-potential distribution³⁶, suggesting different degrees of excitability. Because our analysis suggested that two MD populations are driven by different degrees of cortical engagement (Supplementary Fig. 7), we reasoned that this might also be due to differential excitability that may impart differential susceptibilities to optogenetic inhibition. Specifically, the less-excitable MD population (likely transient MD cells) would require stronger or more-coincident PFC inputs to fire and hence would be more susceptible to weak suppression. Conversely, persistent MD neurons may be more excitable and would require weaker and less-coincident PFC inputs to fire.

By parametrically controlling laser power on an animal-by-animal basis without influencing behavior (Supplementary Fig. 14a,b), we found that MD transient cells were far more susceptible than MD persistent cells to low levels of yellow laser power (556 nm, power at fiber tip: 0.6-1.1 mW; Fig. 4h and Supplementary Fig. 14c,d). These laser powers did not have an appreciable effect on the spiking properties of MD persistent cells (Supplementary Fig. 14e-g). Higher laser powers (power at fiber tip: 2.1–3.5 mW) affected both transient and persistent MD neurons (Supplementary Fig. 14h). In support of the predictions made by our GLM, selectively suppressing MD transient cells with weak laser powers selectively eliminated excitatory functional inputs to the PFC but had no impact on the inhibitory functional inputs from transient MD neurons (Fig. 4i). Suppressing MD terminals in the PFC had a similar effect on the PFC without affecting the firing rates of these neurons (Supplementary Fig. 14i-k). This manipulation also revealed a selective effect on the response properties of both transient PFC RS neuron subtypes (Fig. 4j and Supplementary Fig. 15a), but not on PFC FS neurons (Fig. 4k). In agreement with our earlier studies¹⁷, temporally limited MD suppression had a stronger effect on the maintenance of these peaks than on their initiation in the PFC, confirming that the

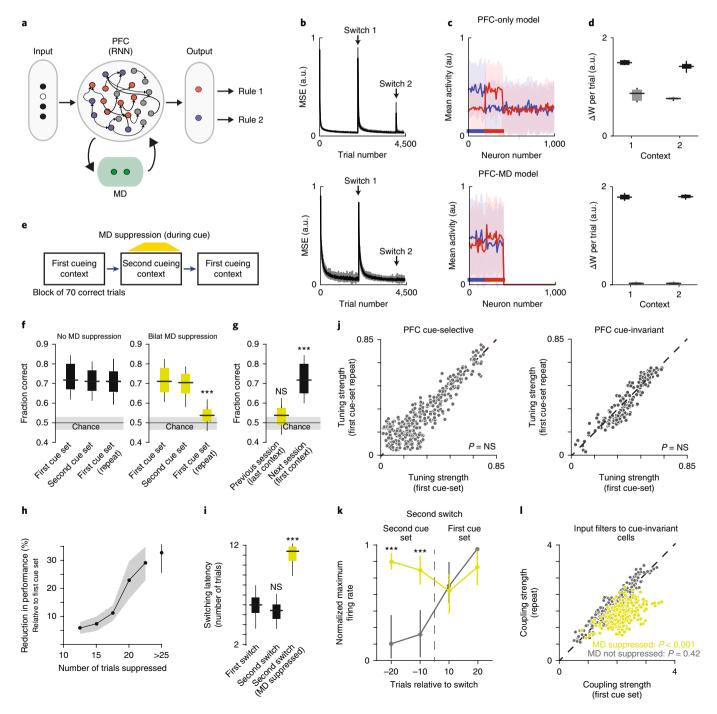
Fig. 5 | Benefit of PFC-MD over PFC-only architecture on switching contexts. a, Recurrent neural network (RNN) model of the PFC-MD network. The drawing depicts neural activation in a single context; gray RNN neurons represent the currently irrelevant context. b, The mean squared error (MSE) in decoding the desired output from the PFC over two context switches (indicated by the arrows). For details, see Methods. c, Trial-averaged responses of 1,000 neurons in the PFC to LP noise (blue) and HP noise (red). Horizontal lines below the plots indicate the sets of neurons activated by the input cues. Shaded area shows s.e.m. **d**, Trial averaged change (n = 200 trials) in connection weights (ΔW) per trial, from current-context neurons (black) and from other-context neurons (gray) to rule-selective output neurons during context 1 and context 2 presentations. Each box extends from lower to upper quartiles, the middle line marks the median, and the whiskers represent the range (from 10 network instances). e, Schematic of the three-block switching task that mice were required to complete. f, Boxplots showing the effect of bilateral MD suppression in the second context on behavioral performance (fraction correct). Shaded area indicates 95% confidence interval of chance behavioral performance derived from a probabilistic model $(n=12 \text{ independent sessions without MD suppression and } 12 \text{ sessions with MD suppression from } 3 \text{ mice}; ***P = 0.05 \times 10^{-4}, Bonferroni-corrected rank$ sum test). \mathbf{g} , Comparison of performance on the consecutive sessions (separated by 1 d; one-way rank-sum test relative to chance; n=10 independent sessions each; *** $P = 0.08 \times 10^{-4}$). **h**, Relationship between the reduction in performance and the number of MD suppression trials. Data shown as mean \pm 95% confidence interval (shaded error bar); n=3 mice. i, Bilateral MD suppression significantly increases the latency of switch back to the first context (n=12 independent sessions each from 3 mice; *** $P=0.09\times10^{-4}$, Bonferroni-corrected Kruskal-Wallis ANOVA with post hoc rank-sum test). j, Scatter plot relating the tuning strengths of PFC cue-selective (left, n = 236 neurons) and cue-invariant (right, n = 158 neurons from 3 mice) cells in the first block with their tuning strengths in the third block. NS, nonsignificant by Friedman test. k, Change in normalized maximum firing rate relative to the second switch of PFC cells selective for cues in the first cue set showing an increase in maximal spiking for out-of-context neurons when the MD is ontogenetically suppressed (n=3 mice; ***P < 0.0001, one-way rank-sum test relative to no MD suppression group). Data shown as mean $\pm 95\%$ confidence interval. I, Scatter plot of the coupling strength between first context PFC cue-selective neurons and cue-invariant ones averaged over trials 10-20 following the switch (gray dots, n=223 neurons). Unilateral optogenetic MD suppression substantially diminishes the size of these functional connections (yellow dots, n=150 neurons from 3 mice). P values calculated using Friedman test between first and third cueing contexts. All boxplots: center line, median; box edges, 95% confidence interval; whiskers, range.

MD was not the source of PFC cue information (Supplementary Fig. 15b). Although the congruence between terminal and somatic inactivation is surprising, the relatively large volumes of MD terminals may result in a larger impact on somatic excitability than would otherwise be expected.

Consistent with the idea that persistent MD neurons provide cross-contextual PFC suppression, strong laser suppression significantly increased out-of-context cue-selective PFC spiking (Fig. 4l) and concomitantly decreased PFC FS neural spiking (Fig. 4m). Because of this, inputs from cue-selective PFC neurons onto cue-invariant neurons took a longer time to stabilize (Fig. 4n). Weak laser suppression, which targeted only transient cells, did not have a similar effect. Taken together, our findings strongly suggest that the MD has two distinct computational functions: (i) transient MD cells maintain the context-relevant representation in the PFC, permitting

cue information to be held in working memory; and (ii) persistent MD cells suppress context-irrelevant representations in the PFC by recruiting FS neurons in a context-dependent manner (Fig. 40).

MD-dependent suppression of context-irrelevant representations protects them for near-future use. Our data thus far suggest a model in which persistent MD neurons suppress PFC representations when they are no longer relevant for the current context. What computational advantage could this architecture impart? Recent theoretical work³⁷ has shown that a context-dependent gating mechanism, which suppresses task-irrelevant nodes in deep neural network, can increase flexibility by allowing the network to learn more tasks sequentially. We wondered whether MD-mediated inhibition could impart such a benefit onto PFC, allowing it to flexibly switch between the different cueing contexts.



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To test this idea, we used a reservoir network of rate neurons as a model for PFC function^{38,39} and incorporated an MD-like node³⁸ that suppressed context-irrelevant reservoir neurons (Fig. 5a). The network was trained to perform a classification task in which it had to classify four cues into two rules, which was analogous to the task that mice were trained to do. Notably, the PFC-MD model outperforms a PFC-only model in being able to flexibly switch between cueing contexts (Fig. 5b). Without an MD, weights relevant to context 1 change in the second context (Fig. 5c,d), which in turn increase errors when context 1 is required again. Incorporating the MD limits the spread of recurrent excitation to the context-relevant PFC neurons, making the two contextual representations practically disjoint and disabling weight changes involving context-irrelevant neurons. Notably, this weight-protection benefit generalizes to a more computationally demanding exclusive-or classification task that is (by design) not linearly separable^{40,41} (Supplementary Fig. 16). Overall, the computational benefits imparted by an MD-like node are even more relevant in the exclusive-or task, suggesting broad benefits of cross-contextual suppression in cognitive flexibility.

We reasoned that we could test these theoretical weight-protection benefits experimentally if we employed a three-block switching framework (Fig. 5e and see Methods), in which mice were re-exposed to the first cueing context in the third block. In this experiment, PFC neurons selective for the first cue-set should be suppressed in the second block, but perhaps reactivated again in the third, as it would be computationally efficient to simply re-engage the same functional ensemble rather than generate a new one de novo. We found that mice performed this three-block switching task well, with no significant difference in performance across blocks (Fig. 5f). Notably, cue-specific and bilateral MD suppression after behavior stabilized in the second block significantly impaired performance when the animal was re-exposed to the first block (Fig. 5f). Although performance was close to chance (as if the mouse had forgotten the first context), this manipulation did not have any long-lasting effects, as performance returned to normal the following day (Fig. 5g). This reduction in behavioral performance on reexposure to the first block parametrically varied with the number of trials suppressed in the second block; suppressing a larger number of trials in the second block resulted in a larger behavioral deficit (Fig. 5h; inflection point, 20 trials). Notably, although the switching latency was marginally shorter when the animal moved from the second context back to the first cueing context, MD suppression in the second block prolonged this switch (Fig. 5i and Supplementary Fig. 17). This effect was stronger than what we show in Fig. 3f, because unlike in the two-block switching task, mice were now required to reactivate representations for the first cueing context in the PFC.

To examine the neural substrates of this behavioral detriment, we again aimed at dissociating behavioral from neural manipulations and therefore employed a unilateral optogenetic suppression condition, in which we suppressed MD neurons during the cue, once behavior stabilized in the second block. In sessions in which no such optical manipulation was deployed, both PFC cue-selective and cue-invariant neurons were largely shared between the first cue-set and their repeat in the third block (Fig. 5j). As expected, unilateral cue-specific MD suppression resulted in out-of-context spiking of the first cue-set neurons during the second block (Fig. 5k and Supplementary Fig. 18). Although this was not associated with a delay in how these neurons were recruited in the third context, their functional inputs onto cue-invariant neurons were much weaker upon the switch (Fig. 51). Therefore, our data suggest that in addition to suppressing context-irrelevant cortical representations, such that context-relevant functional connections rapidly stabilize, such processes may protect recently engaged but currently irrelevant connectivity patterns for near-future use (see Supplementary Fig. 19 for summary model).

Discussion

In this study, we expanded on a behavioral experiment we had previously developed^{17,23} by nesting it in a cognitive hierarchy. Specifically, while our previous studies explored the neural correlates of crossmodal sensory selection based on two learned cues, the current design nested the selection process within multiple cueing contexts. Crucially, these contexts were under complete experimental control and could be arbitrarily constructed on a session-by-session basis.

This allowed us to make multiple observations. First, we identified a prefrontal neural hierarchy that matches the cognitive one; neurons that reflected the meaning of the cue (the rule) derived their representations from local cue-selective inputs. Similar hierarchies are seen in sensory areas⁴², potentially speaking to broadly similar cortical organization principles. Notably, we found only 5% of cells in the mouse PFC to be rule-selective, a contrast to higher species, which have substantially larger fractions of such cells^{29,43}. This difference may explain certain cross-species differences in generalization and cognitive capacity.

Second, unlike structures like the lateral geniculate nucleus or thalamic circuits that primarily drive excitatory responses in the cortex⁴⁴, we found that the MD exerts effects on cognitive switching through local inhibitory cortical interneurons. This builds on similar recent studies^{22,34,35}, but also provides a computational framework linking thalamic output to cortical inhibitory microcircuits. For example, transient MD neurons could recruit disinhibitory motifs⁴⁵ to maintain activity in the PFC, while persistent MD neurons could target soma-targeting interneurons³⁵. Without further evidence however, we can only speculate that the diversity of thalamocortical computations may match the diversity of cortical interneurons⁴⁶.

Third, the unique connectivity patterns of the lateral MD are consistent with our physiological results; convergence of individual small cortical terminal inputs onto single MD neurons^{20,47} may explain their lack of selectivity to categorical information that originates in cortex. Instead, our model shows that this convergence of PFC inputs may facilitate the emergence of contextual signals in the MD²⁰. Additionally, the lack of thalamic lateral connectivity may allow MD neurons to multiplex incoming signals, a process that would be harder for cortical circuits to implement given their extensive recurrence.

Fourth, the experiments involving multiple switches point to a variety of plasticity rules governing cortical function, as has been recently shown through recurrent neural network simulations⁴⁸. Within this framework, our data suggest a unique role for the thalamus in generating contextual representations that may regulate cortical plasticity. The exact nature of cortical inhibitory neurons involved in cross-contextual suppression is still unknown, and our study provides a starting point for such detailed exploration.

Lastly, it is worth mentioning that recent progress in artificial intelligence research has shown that incorporating context-specific gating mechanisms in convolutional networks is beneficial for the ability to perform multiple tasks and the mitigation of 'catastrophic forgetting' 10,37. The key idea in these studies is the generation of non-overlapping, task-specific representations in a context-specific manner 19. We envision that the MD imparts a similar computational benefit for task-specific PFC representations: rapid separation of potentially overlapping representations such that they are more easily decoded. Overall, our findings may not only be relevant to future research in neuroscience but may also lead to the generation of artificial networks that exhibit more stable learning and robust performance.

Online content

Any methods, additional references, Nature Research reporting summaries, source data, statements of data availability and associated accession codes are available at https://doi.org/10.1038/s41593-018-0269-z.

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Author contributions

R.V.R. conceived and performed experiments, analyzed and interpreted data, and wrote the paper. A.G. developed, simulated, and analyzed the thalamocortical computational model. M.M.H. conceived and supervised experiments, analyzed and interpreted the data, and wrote the paper. M.M.H. also acquired funding.

Competing interests

The authors declare no competing interests.

Additional information

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Methods

Mice. All experiments were carried out under protocols approved by MIT's Committee on Animal Care and conformed to NIH guidelines. With the exception of one mouse that had the Sst-IRES-Cre (Jax: 013044) genotype, all mice were C57/BL6 (Taconic Biosciences). Only male mice older than 8 weeks old were used in this study. Please refer to the Nature Research Reporting Summary for further details. Mice were housed in the vivarium on a standard 12-h light/dark cycle and were singly housed throughout the experimental period. Experiments were performed during the light portion of the cycle.

Behavioral setup. Behavioral training and testing took place in gridded floormounted, custom-built enclosures made of sheet metal covered with a thin layer of antistatic coating for electrical insulation (dimensions in cm: length, 15.2; width, 12.7; height, 24). All enclosures contained custom-designed operant ports, each of which was equipped with an IR LED/IR phototransistor pair (Digikey) for nosepoke detection. Trial initiation was achieved through an 'initiation port' mounted on the grid floor 6 cm away from the 'response ports' located at the front of the chamber. Task rule cues and auditory sweeps were presented with millisecond precision through a ceiling-mounted speaker controlled by an RX8 Multi I/O processing system (Tucker-Davis Technologies). Visual stimuli were presented by two dimmable, white-light-emitting diodes (Mouser) mounted on each side of the initiation port and controlled by an Arduino Mega microcontroller (Ivrea). Similarly, the visual cues were delivered through a pair of a wall-mounted 5-mm LEDs (UV: 320-380 nm, green: 495-510 nm, 100-mW, 25° viewing angle Mouser). These LEDs were bright enough to illuminate the whole arena. Two response ports were mounted at the angled front wall 7.5 cm apart, respectively. Milk reward ($10\,\mu$ L evaporated milk, Carnation) was delivered by a single syringe pump (New Era Pump Systems) when mice made a correct choice. Access to the response ports was restricted by vertical sliding gates which were controlled by a servo motor (Tower Hobbies). The TDT Rx8 sound production system (Tucker Davis Technologies) was triggered through Matlab (MathWorks), interfacing with custom-written software running on an Arduino Mega (Ivrea) for trial logic control.

Multielectrode array construction and implantation. Custom multielectrode array scaffolds (drive bodies) were designed using 3D CAD software (SolidWorks) and printed in Accura 55 plastic (American Precision Prototyping) as described previously 17,23,50 . Prior to implantation, each array scaffold was loaded with 12–18 independently movable microdrives carrying 12.5-µm nichrome (California Fine Wire Company) stereotrodes or tetrodes. Electrodes were pinned to custom-designed, 96-channel electrode interface boards (EIB, Sunstone Circuits), along with a common reference wire (A-M systems). For combined optogenetic manipulations and electrophysiological recordings of the PFC, optic fibers delivering the light beam lateral (45° angled tips) were embedded adjacent to the electrodes.

During implantation, mice were deeply anaesthetized with 1% isoflurane and mounted on a stereotaxic frame. A craniotomy was drilled centered at AP $-2\,\mathrm{mm}$, ML $0.6\,\mathrm{mm}$ for PFC recordings and at AP $1\,\mathrm{mm}$, ML $1.2\,\mathrm{mm}$ for mediodorsal recordings. The range of coordinates covered in our recordings for the lateral MD are: AP: -1 to $-1.5\,\mathrm{mm}$, ML: 0.3 to $0.8\,\mathrm{mm}$ relative to bregma. Similarly, for the PFC, the range of coordinates covered in our recordings are: AP: 2.1 to $2.7\,\mathrm{mm}$, ML: 0.25 to $0.6\,\mathrm{mm}$ relative to bregma.

The dura was carefully removed, and the drive implant was lowered into the craniotomy using a stereotaxic arm until stereotrode tips touched the cortical surface. Surgilube (Savage Laboratories) was applied around electrodes to guard against fixation through dental cement. Stainless-steel screws were implanted into the skull to provide electrical and mechanical stability and the entire array was secured to the skull using dental cement.

Optogenetic manipulation. We used a dual-wavelength, optical-silencing method to independently suppress neurons in the PFC and MD. Specifically, we virally expressed the inhibitory channelrhodopsin iC++ in the PFC (AAV-CaMKIIA-iC++-eYFP) $^{\rm SI}$, which is selective to blue-shifted wavelengths (473 nm), and expressed halorhodopsin (AAV-CaMKIIA-eNpHR3.0-eYFP) in the MD $^{\rm S2}$. Since the peak spectrum of eNpHR is red-shifted (peak at $\sim\!550$ nm), we could independently suppress both populations without affecting their terminals in either structures. Light was delivered to these structures using optic fibers that were part of the microdrive (as described above). We used a 473-nm laser and a 556-nm laser (OptoEngine) to activate iC++ and eNpHR, respectively.

Behavioral training. Mice were trained to perform this task in subsequent stages. First, $10~\mu L$ of evaporated milk (reward) was delivered randomly to each reward port for shaping and reward habituation. Next, the location of the rewarded port was signaled by a white LED (same used as the visual target) to establish an association between the location of the visual target and the location of the reward port. Following this, mice learned the association between the auditory targets—up-sweep, 10-15~kHz, and down-sweep, 16-12~kHz—with the left and right ports, respectively. An individual trial was terminated 20~s after reward collection, and a new trial became available 5~s later. As soon as mice achieved

criterion performance in this block (>60% correct), visual and auditory targets were randomly interleaved.

Second, mice learned to poke (i.e., break the IR barrier in each reward port) in order to receive reward. All other parameters remained constant. An incorrect poke had no negative consequence. By the end of this training phase, all mice collected at least 20 rewards per 30-min session.

Third, mice were trained to initiate trials. Initially, mice had to briefly (50 ms) break the infrared beam in the initiation port to trigger target stimulus presentation and render reward ports accessible. Trial rule (attend to vision or attend to audition) was indicated by 10-kHz low-pass-filtered white noise (vision) or 11-kHz high-pass-filtered white noise (audition) sound cues. Stimuli were presented in blocks of six trials consisting of single-modality stimulus presentation (no conflict). An incorrect response immediately rendered the response port inaccessible. Rewards were available for 15 s following correct response, followed by a 5-s intertrial interval (ITI). Incorrect responses were punished with a timeout, which consisted of a 30-s ITI. During an ITI, mice could not initiate new trials. During this stage, the duration of the initiation time was gradually increased from 50 ms to 800 ms. Mice progressed to the next stage only when they were able to maintain snout fixation for at least 800 ms.

Fourth, conflict trials were introduced, in which auditory and visual targets were co presented indicating reward at opposing locations. Four different trial types were presented in repeating blocks: (i) three auditory-only trials; (ii) three visual-only trials; (iii) six conflict trials with auditory target; and (iv) six conflict trials with visual target. The time that mice had to break the IR barrier in the initiation port was continuously increased over the course of this training stage (1–2 weeks) until it reached 0.5 s. At the same time, duration of the target stimuli was successively shortened to a final duration of 0.1 s. Once mice performed successfully on conflict trials, single-modality trials were removed, and block length was reduced to three trials.

Fifth, during the final stage of training, trial availability and task rule were dissociated. Broadband white noise indicated trial availability, which prompted a mouse to initiate a trial. Upon successful initiation, the white noise was immediately replaced by either low-pass or high-pass filtered noise for 0.1 s to indicate the rule. This was followed by a delay period (variable, but for most experiments it was 0.4 s) before target stimuli presentation. All block structure was removed, and trial type was randomized. Particular steps were taken throughout the training and testing periods to ensure that mice used the rules for sensory selection.

Once mice were fully familiarized with the main structure of the task and achieved consistent performance on the final stage of training, they were exposed to the visual cueing condition. After achieving 40 correct responses, mice were moved to an association block in which the LEDs were paired with the congruent auditory cues (LP with green LED and HP with UV LED). The volume of the sound decayed linearly over the course of trials, with full volume for the first 10 trials up to 1/5 of the volume for the last few trials. The sound volume was changed only after the mouse made two consecutive correct responses (an indication that the mouse understands the task). At the end of 70 trials, and depending on performance, mice progressed to the visual-only block, in which no auditory cues were played. In the following session, the length of the association block was gradually reduced. Once mice were able to achieve a consistent performance of > 60% on three consecutive sessions in the visual-only block, the association block was removed completely. At this point, mice were considered experts on the task.

Behavioral testing. In the double block cueing experiment, mice were required to complete 70 trials in each block. Blocks were constructed based on either cues of the same modality (HP–LP and UV–green) or cues from both modalities (HP–green and UV–LP). In each block, cues were drawn pseudorandomly and the order of blocks was randomized from session to session. Sessions in which mice did not perform > 60% overall in each block were discarded and were not analyzed further.

In the randomized cueing experiment, mice were required to complete a total of 200 trials per session. On each trial, one cue of a possible four cues (HP, LP, UV, and green) was drawn at random. To further ensure that the cues appeared in random order (i.e., without any regularity), we imposed the additional constrained that no more than three draws could be from the same modality. That is, after three UV–green draws, the next cue had to be either HP or LP. A new random seed was used each day. Mice that were previously trained on the block design took approximately a week to adjust to this new cueing condition. Although average performance was low, mice had brief periods in which their local performance was close to 80%.

In the three-block switching experiment, mice were required to complete a total of 70 trials in the first two blocks and 90 trials in the third block. The identity of the first block (i.e., visual or auditory) was pseudorandomized from day to day. In total, we mice completed four sessions per mouse (two auditory-visual-auditory, two visual-auditory-visual) on the standard version of the task, and four sessions per mouse with MD suppressed in the second block. We did not notice a difference in the effect of MD suppression in the visual block compared to the auditory cueing block, and hence have pooled these sessions for analysis.

Behavioral analysis. To quantify the behavior, we carried out regression analysis to weigh the contributions of rule and history of choice and reward on the animal's choice on the current trial⁵³. To do so, we concatenated data from multiple sessions for each mouse and fit the animal's choice with a logistic regression model of the form

$$P(V/vision) = lapse + \frac{1 + 2lapse}{1 + e^{-A}}$$

$$A = \beta_0 + \sum_{t=1}^{T} \beta_{rule}(t) R(t) + \beta_{success}(t) S(t) + \beta_{failure}(t) F(t)$$

where T is the number of trials in the past. For this model, we calculated that the model explained behavioral variance best when we included up to 10 trials in the past. In this equation, $S(t),F(t)\in[+1,-1]$ if a trial is a success or a failure, respectively. Similarly, $R(t)\in[+1,-1]$ if the rule on that trial is attend-to-vision or attend-to-audition, respectively. This model was fit using a custom-written ridge regression routine in Matlab. The hyperparameter value for ridge regression was calculated using fivefold cross-validation.

To assess the effect that choice history had on the probability of success on the next trial, we developed a probabilistic model. Given the 2-AFC structure of our task, we assumed that on each trial, the mouse makes a coin-flip choice with a bias ($q = P_{\text{success}}$) that depends on the animal's success on past trials. Therefore, the likelihood of a success given s successes and f failures in the past 10 trials is given by the binomial distribution:

$$p(s;q) = {s+f \choose s} q^{s} (1-q)^{f}$$

Knowing that the conjugate prior of the binomial distribution is the beta distribution, we can calculate the posterior distribution using Bayes' theorem

$$p(q) = \frac{q^{\alpha-1}(1-q)^{\beta-1}}{B(\alpha,\beta)}$$

Posterior =
$$\frac{p(q)p(s;q)}{\int p(q)p(s;q)dq}$$

$$\sim Beta((\alpha+s),(\beta+f))$$

Using this, the expected probability of success (E(q)) on the next trial is then

$$E(q) = \frac{B(\alpha + s + 1, \beta + f)}{B(\alpha + s, \beta + f)}$$

All models were fit separately for each mouse (n=5) using 1,000 runs of fivefold cross-validation. For each run, we computed the log-likelihood for the test dataset for the mean value if P(vision) or P(success). Model fit quality was assessed by computing the deviance statistic. We also used this model to assess overall behavioral performance. Trials in which behavior was close to the predicted chance levels were ignored and the overall fraction correct was computed from trials in which actual behavior deviated substantially from the model (we define these periods as 'stable' behavior). As such, changes in performance accuracy reported throughout are only calculated over the period of stable behavior and do not include switch trials.

Trial selection. By comparing the two models (a model that includes rule-dependent components and a reward-history model) we were able to determine trials over which the animal was making either an informed choice based on the current cue or a biased choice based on history of reward. We used this to select trials. Specifically, for each trial, we computed the deviance between the full model and the history model and selected a trial if deviance was above a threshold value that was calculated using cross-validation for each session. Unexpectedly, tuning peaks (described in the following section) were much more apparent in the selected trials than in the rejected trials. Please refer to the Nature Research Reporting Summary for further details on omitted data.

Electrophysiological recordings and spike sorting. Signals were acquired using a Neuralynx multiplexing digital recording system (Neuralynx) through a combination of 32- and 64-channel digital multiplexing headstages plugged into the 96-channel EIB of the implant. Signals from each electrode were amplified, filtered between 0.1 Hz and 9 kHz, and digitized at 30 kHz. For thalamic recordings, tetrodes were lowered from the cortex into the mediodorsal thalamus over the course of 1–2 weeks where recording depths ranged from –2.8 to –3.2 mm DV. For PFC recordings, adjustments accounted for the change of depth of PFC across

the AP axis. Thus, in anterior regions, unit recordings were obtained -1.2 to -1.7 mm DV, whereas for more posterior recordings, electrodes were lowered -2 to -2.4 mm DV.

Spike sorting was done automatically using MountainSort⁵⁴. Following sorting, each cluster was manually inspected for quality. Only well-isolated clusters with biologically plausible waveforms were selected for further analysis.

Identification of FS and RS cells. For each spike waveform, we extracted four metrics: (i) peak-to-trough time; (ii) peak-to-trough ratio; (iii) spike width; and (iv) spike amplitude. We combined this four-dimensional feature vector with the overall firing rate of the neuron to form a five-dimensional feature vector for each cell. We applied k-means clustering (k++ algorithm, 1,000 runs with randomly initialized seed) and determined the optimal number of clusters using the Calinski–Harabasz criterion Cluster separability was assessed statistically by calculating the ratio of between-cluster variance to within-cluster variance. For most sessions, the waveforms clustered reliably into two clusters, corresponding to FS and RS waveforms. Approximately 10% of all recorded spike waveforms could not be reliably classified into either subtype (based on 1,000 runs) and hence were not included in further analysis.

Analysis of firing rate. For all PFC and mediodorsal neurons, changes in firing rate-associated task performance were assessed using peristimulus time histograms (PSTHs). PSTHs were computed using 10-ms bins for individual neurons in each recording session, convolved with a Gaussian kernel (25 ms full-width at half-maximum) to create a spike-density function (SDF) which was then converted to a z-score by subtracting the mean firing rate in the baseline (500 ms before event onset) and dividing by the variance over the same period. For comparison of overall firing rates across conditions, trial number and window size were matched between groups. Except for switching analysis, we analyzed firing rates only in trials in which local performance deviated significantly from the probabilistic model (Supplementary Fig. 1).

Computing reliability and tuning strength. For each recorded neuron, we computed trial-to-trial reliability using a 150-ms sliding window. Reliability was simply the correlation in spike times between each pairwise combination of trials, such that a neuron with perfect reliability had no spike time variation and a correlation coefficient of 1. Only neurons with responses on 15 trials or more were selected for this analysis.

To determine whether the observed level of reliability was significantly different from chance, we used a randomization test where the time period of analysis was randomly picked in the range [–2.5, 1.5], the trials randomly shuffled, and the reliability score recalculated. By repeating this process 1,000 times, a null distribution of the reliability time series was constructed. A neuron was reliable if the unshuffled reliability time series exceeded the null distribution by 1.5 s.d. (*z*-score > 1.5). Using this method, we were able to calculate a significant reliability trace for each neuron and for each stimulus condition.

Classification of cells into persistent and reliable. The method described above allowed us to extract reliability scores (max in the delay period). In the PFC RS and MD populations, we noticed a bimodal distribution of reliability scores: some neurons responded with high trial-to-trial spiking in the delay period (transient) and others responded with low reliability (persistent). To formally classify these cells, we used the expectation-maximization algorithm (Python sklearn package) to fit a Gaussian mixture model to the reliability histogram. This procedure was run separately for PFC RS, FS, and MD neurons. The goodness of fit of the Gaussian was assessed using the Bayes information criterion (BIC). Separability of the resulting Gaussians was assessed by ROC analysis. PFC RS and MD populations had separable Gaussians and passed the Hartigan's dip-test for bimodality. In these populations, we classified cells as transient if they were within 95% CI of the mean of the high-reliability Gaussian model. Cells were classified as persistent if they were within the 95% CI of the mean of the low-reliability Gaussian model. This method allowed us to robustly classify neurons without the need to define an arbitrary threshold.

Classification of cells into cue-selective or cue-invariant. Using the reliability time series across the delay period, we computed a cross-correlogram for all pairs of conditions (six-way comparison). Neurons with significant correlation with lags within $\pm\,50$ ms were scored. A neuron was classified as cue-selective if it had a significant reliability event for only one of the four stimulus conditions. A neuron was classified as mixed-selective if it had a significant reliability event for two of the four stimulus conditions. We found no neurons in either the MD or PFC with significant reliability in more than two conditions (except for randomized cueing experiments).

To further assess the tuning strength of PFC neurons, we first sampled trials with replacement to calculate an estimate of d-prime for either cues or rules for each cell. Second, we also computed a bootstrapped value of the reliability of that cell. We defined the tuning strength as the slope of the regression line between reliability and d-prime. As such, a neuron with high reliability and high d-prime had a higher tuning-strength value, indicating that this neuron was strongly

selective to a cue. We used this tuning-strength metric to better define cueinvariant neurons. Cue-invariant, rule-selective neurons should respond to both cues that map onto the same rule. Hence, we calculated the selectivity angle for each pair of cues corresponding to the same rule using the formula⁶

$$\theta = \tan^{-1} \! \left(\frac{Tuning \ strength \ to \ cue \ 1 \ of \ rule \ A}{Tuning \ strength \ to \ cue \ 2 \ of \ rule \ A} \right)$$

Neurons with selectivity angle of 45° had the same tuning strength for both cues in the same rule and were hence classified as cue-invariant. Therefore, we used a hierarchical selection process to classify cue-invariant cells: (i) the significant reliability time series for cues 1 and 2 of rule A must be correlated within a lag of 50 ms and (ii) the selectivity angle must be close to 45°. Since MD and PFC FS neurons were weakly reliable, we calculated their selectivity using trial-averaged firing rates instead. In this way, MD and FS neurons were classified as context selective when (i) they had correlated responses for both cues within a context and (ii) they had a within-context selectivity angle was also close to 45°. Each of these measures was tested for significance using a permutation test in which hybrid data were created by shuffling trial labels.

Calculating contextual modulation index. We assessed the contextual modulation index (CMI) of the trial-averaged firing rate of a neuron using the following formula:

$$CMI = \frac{Rate_{context1} - Rate_{context2}}{Rate_{context1} + Rate_{context2}}$$

As such, because the firing rate is non-negative, $CMI \in [+1,-1]$. To determine significance, we calculated the CMI for two hybrid spike trains created by randomly shuffling trial labels from 1,000 iterations. This created a null distribution. A cell was considered significantly contextually modulated if the unshuffled CMI was outside the 95% confidence interval of the shuffled CMI (P < 0.05, two-tailed Student's t test).

Decoding analysis. Trial-by-trial classification analysis was performed using a support vector machine (SVM) implemented through LIBSVM and the Matlab neural decoding toolbox 56 . The firing rates of neurons on each trial from the entire population (pooled across sessions) were first smoothed using a 20-ms-wide Gaussian filter. The SVM classifier with a Gaussian radial basis function kernel was then trained on 60% of the data (randomly selected) while 40% of the data were used for prediction. This classifier works by first constructing an optimal hyperplane based on labeled training data and then generating predictions of the labels on testing data. Accuracy of the decoding was assessed by comparing the predicted labels to the actual labels. Classification accuracy was also quantified by computing the mutual information via the following equation

$$MI = \sum_{i=1}^{S} \sum_{j=1}^{S} p_{ij} \log \frac{p_{ij}}{p_i p_j}$$

where p_{ij} is the probability of observing label i (cue, rule, or context) given that the original label is j. This classification process was repeated 1,000 times to obtain and accurately estimate the error of the classification accuracy.

To test the dependence of the number of neurons on classification, we used a Monte Carlo sampling technique (repeated 500 times) to pick n neurons (range: 1 to population size) at random from the population with replacement. The single-trial responses from these n neurons were compared to the template as described above

Generalized linear model (GLM). We modeled the spike trains of neurons using a generalized linear model (GLM) 31,32,57 . The spike trains were discretized (Δ) into 5-ms bins. As explained elsewhere 58 , the log-likelihood for a single neuron (up to an additive constant) is given by the formula

$$\log L(\varphi, r) = \sum_{t} r(t) \log(\Delta \varphi(t)) - \Delta \varphi(t)$$

where $\varphi(t)$ is the instantaneous spike rate (conditional intensity) of the fully coupled GLM

$$\varphi(t) = \exp(\mathbf{k}x(t) + \mathbf{h}r(t-1) + \mathbf{c}s(t) + \mathbf{b})$$

In this equation, k is the weights on the stimulus covariates (akin to a receptive field), h is the postsynaptic weights that integrate the neuron's own spiking history, and c is the coupling weights (filters) on other simultaneously recorded spikes (s). In the uncoupled model, we ignored this coupling term. To avoid overfitting, regression weights were fit with a maximum a posteriori estimate with an L2 penalty. Matlab scripts used to build the GLM can be found here: https://github.com/pillowlab/neuroGLM.

These coupling filters are analogous to the positive lag of a cross-correlogram, with the additional benefit of accounting for the response variance that is not already explained by the cue and other task-relevant variables. In other words, each neuron produces a coupling filter that, when convolved with the spike train of that neuron, explains part of the variance of the neuron being modeled. Mathematically, this operation can be written as

$$\sum_{i=1}^{m} \sum_{j=1}^{n} c_{ij} f_{j} (s_{i}(t-\tau : t))$$

where m is the number of simultaneously recorded neurons and f_i are temporal basis functions that we assumed to be nonlinearly time-scaled, raised cosine functions³². In each session, the GLM was constructed using a median of 25 PFC and 18 MD neurons with well-isolated units.

To statistically validate these coupling filters, we randomized both neuronal labels and trial order and used leave-one-neuron-out cross-validation. This allowed us to determine the probability of a coupling being significant above chance levels. We also calculate the coupling strength as the integral (area under the curve) of each coupling filter. Because each neuron can receive many coupling filters, we used a dimensionality reduction (SVD) to determine the most common filter shapes (i.e., those that explained the largest fraction of variance). We note here that we fit each GLM in an unbiased manner and determined the most significant couplings based on shuffling. When we tested the effect of removing certain filters on predicting the firing rate of neurons, we first fit a model to 80% of the trials, set the necessary filter components to 0, and then used that model to predict the remaining 20% of the trials. We computed explained variance (EV) using the following formula

$$EV = 1 - \frac{\sum_{i} (y_{i} - Model_{i})^{2}}{\sum_{i} (y_{i} - \hat{y}_{i})^{2}}$$

We repeated this procedure 100 times. In this way, we do not bias the other terms of the model by removing terms before performing the regression.

We derived a filter similarity index to determine how inputs to a neuron changed as the animal switched from one context to another. First, we used the behavioral model (see section on trial selection above) to determine trials in which choice behavior was stable in each context. Using these trials, we derived coupling filters (i) between PFC cue-selective cells, (ii) from PFC cue-selective cells to MD cells, and (iii) from PFC cue-selective cells to PFC cue-invariant cells. We refer to these filters as stable input filters. Next, we refit the GLM on a trial-to-trial basis from 10 trials before the switch to 10 trials after the switch and extracted single-trial input filters. The filter similarity index is the Pearson's correlation coefficient between the single-trial coupling filter and the stable coupling filter in each context. In particular, for cells preferring the second context, we report the correlation coefficients between the single-trial input filters and the stable filter in the second context. This analysis allowed us to visualize the remapping of intracortical and corticothalamic inputs as mice switched from one cueing context to another. We defined the filter stabilization latency as the trial number at which the correlation coefficient between the single-trial coupling filter and the stable coupling filter in each context is significantly above chance levels.

For the clustering analysis in Fig. 4d, we quantified the shape of the filter using a filter score. For filters with a larger inhibitory magnitude, the filter score was the signed area under the curve of the inhibitory component. For filters with a larger excitatory magnitude, the filter score was the area under the curve of the positive component. In this way, negative filter scores correspond to MD neurons that exert an inhibitory effect on their targets, while positive filters scores correspond to MD neurons that exert an excitatory effect on their targets.

Model to explain MD responses. We constructed a simple model to determine whether and how MD neurons derive their contextual selectivity from PFC cueselective neurons (Supplementary Fig. 7). To do so, we first generated a population of 1,000 Poisson spiking units with transient elevations that spanned the duration of the delay period (50-ms peak spacing). These model neurons mimicked, for example HP and LP selective neurons in the PFC, with the aim of predicting the responses of the auditory cueing context-selective MD cells. For each model PFC neuron, we computed a PSTH. Each PSTH was then convolved with the PFC-MD input kernel (described above). This convolved output was then weighted and summed. We then used a least-squares method to determine the best fit model that could explain the trial-averaged firing rate of either persistent or transient MD neurons. For persistent MD neurons, weights were almost uniformly distributed over all PFC inputs, suggesting that their inputs were not temporally selective. In contrast, transient MD neurons weighted inputs from co-tuned PFC neurons more strongly, suggesting that they receive temporally selective inputs from these cotuned cue-selective neurons.

Computational modeling. We use a recurrently connected reservoir of 1,000 rate neurons to model the PFC. The rate of each neuron, indexed by i, is given as a

function of its input I_i as r_i = tanh(I_i) if I_i > 0, and 0 otherwise. The input consists of cue input, recurrent input, and MD gating together, filtered with a decaying exponential synapse with time constant τ = 20 ms, as

$$\begin{split} \tau \frac{dI_i(t)}{dt} = -I_i &+ \sum_k w_{ik}^{in} cue_k(t) \\ &+ \sum_j \left(1 + \mu_i(t)\right) w_{ij} r_j(t) + \sum_l w_{il}^{MD+} r_l^{MD}(t) \end{split}$$

where cue_k is a vector of length equal to the number of possible cues (corresponding to HP and LP noise and UV and green LED flash). It has entries 1 for cues that are on at the current time and 0 for those off. The input weights w_{ik}^{in} are set such that each cue k stimulates a set of 200 neurons, disjoint with the sets for other cues, with each weight chosen uniformly between 0.75 and 1.5. w_{ij} is set as a Gaussian-distributed variable with mean = 0 and s.d. =0.75/ $\sqrt{400}$, and then the mean is subtracted across each row of the matrix r_l^{MD} is a vector representing the activity of MD neurons with dimensionality equal to the number of contexts. We set the entry for the current context to 1 and the rest to 0. w_{il}^{MD+} is set to -10 for those neurons that are not stimulated by cues belonging to context l and to 0 for those that are, effectively suppressing activity of context-irrelevant neurons. μ_l mediates the multiplicative effect of the MD on the total recurrent input to neuron i and is given by

$$\mu_i = \sum_m w_{im}^{MD \times} r_m^{MD}$$

 $w_{im}^{MD\times}$ is set to 8 if neuron *i* is one of the neurons stimulated by cues belonging to context *m*, else it is set to 0, effectively enhancing the recurrent input for context-relevant neurons. Note that all sums run over all the full range of the summed indice

relevant neurons. Note that all sums run over all the full range of the summed indices. When simulating the PFC-only network, we set all w_{il}^{MD+} to zero and all $w_{im}^{MD\times}$ to 2, effectively removing all context-specific suppression and enhancement, yet ensuring enough recurrent input to sustain activity in the delay period. The model has two output neurons: the first corresponding to attend to audition and the second to attend to vision, receiving input from the PFC as

$$\tau \frac{dI_n^{out}}{dt} = -I_n^{out} + \sum w_{ni}^{out} r_i$$

with output r_n^{out} = tanh (I_n^{out}) if I_n^{out} > 0, and 0 otherwise. w_{ni}^{out} is initialized to zero, and is plastic, evolving as

$$\tau_{w} \frac{dw_{ni}^{out}}{dt} = -r_{i}(r_{n}^{out} - r_{n}^{target}) \equiv r_{i} \varepsilon_{n}$$

where $\tau_w = 200 \, \mathrm{s}$, and the instantaneous error \in_n in output n is defined in terms of the target output r_n^{target} which is cue-specific as below. Learning on the output weights is on throughout the simulations.

Each task was simulated as a run of 1,000 cycles of context 1 (block 1), followed by 1,000 cycles of context 2 (block 2), and then again followed by 200 cycles of context 1 (block 3). Each cycle consists of two trials of cue = (1,0,0,0) and cue = (0,1,0,0) during context 1 and two trials of cue = (0,0,1,0) and cue = (0,0,0,1) during context 2, in random order within each cycle, for the experimental linearly separable task, representing high-pass and low-pass noise and UV and green LED flash, respectively. The target output r_n^{target} values for these cues are (1,0), (0,1), (1,0), and (0,1) respectively. The two longer blocks allowed the network to learn the two contextual tasks sequentially, while the shorter third block served to test the ability of the network to recall the first context.

Similarly, for the XOR task, each cycle consists of 4 trials of *cue* equal to (0,0,0,0), (0,1,0,0), (1,0,0,0), and (1,1,0,0) during context 1 and (0,0,0,0), (0,0,0,1), (0,0,1,0), and (0,0,1,1) during context 2, in random order. These must map to target output r_n^{target} equal to (1,0) if only one of the cues in a context is active and to (0,1) if none or both are active.

Each trial consists of a 100-ms-long cue presentation followed by a 100-ms delay period when (0,0,0,0) is presented. The target output is maintained throughout the trial for plasticity of the output weights, and the mean squared error is computed over the full trial and across the two outputs.

Statistical testing. All data in this paper are pooled from 5 mice (except for optical perturbation, for which we used 3 mice). No statistical tests were done to determine the sample size, but our sample sizes are similar to those reported in previous publications^{17,23}. Note that data collection and analysis were not performed blind to the conditions of the experiments.

Data were first tested for normality using the Shapiro–Wilk test. All data presented in this paper are non-normally distributed; thus, all statistical tests were conducted using nonparametric statistics. Our experiments involved testing the influence of different conditions (cues, optical manipulations, etc.) on the same population of neurons; thus, all comparisons were performed using nonparametric repeated-measures ANOVA (Friedman test) with Bonferroni's correction for multiple comparisons. Comparisons between independent measures were performed using the nonparametric Kruskal–Wallis ANOVA. For Bonferroni corrections, the significance value was set to 0.05. Post hoc tests were performed using two-tailed signed-rank tests (for repeated measures) or Wilcoxon rank-sum tests for independent measures. All other statistical tests that were performed are described in the text. The 95% CIs were computed by bootstrapping.

Reporting Summary. Further information on research design is available in the Nature Research Reporting Summary linked to this article.

Code availability. All Matlab and python scripts used to analyses the data will be deposited on GitHub at https://github.com/toxine4610/ThalamusContextSwitchingCode and https://github.com/adityagilra/PFC_MD_weights_stability.

Data availability

All data are available from the corresponding author upon reasonable request.

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Reporting Summary

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Statistical parameters

		Methods section).
n/a	Cor	nfirmed
	\boxtimes	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement
	\boxtimes	An indication of whether measurements were taken from distinct samples or whether the same sample was measured repeatedly
	\boxtimes	The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.
	\boxtimes	A description of all covariates tested
	\boxtimes	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons
	\boxtimes	A full description of the statistics including <u>central tendency</u> (e.g. means) or other basic estimates (e.g. regression coefficient) AND <u>variation</u> (e.g. standard deviation) or associated <u>estimates of uncertainty</u> (e.g. confidence intervals)
	\boxtimes	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted <i>Give P values as exact values whenever suitable.</i>
\times		For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings
\times		For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes
\boxtimes		Estimates of effect sizes (e.g. Cohen's d, Pearson's r), indicating how they were calculated
	\boxtimes	Clearly defined error bars State explicitly what error bars represent (e.g. SD, SE, CI)

Our web collection on <u>statistics for biologists</u> may be useful.

Software and code

Policy information about availability of computer code

Data collection

Electrophysiological data was collected using the Neuralynx (Cheetah 6.3.2) system. Behavioral data was collected using custom MATLAB (Mathworks, version: 2016a) scripts that interfaced with and Ardunio Uno using custom-written code.

Data analysis

All analysis was conducted using custom written code in MATLAB (Mathworks, version: 2016a) using standard toolboxes. Links to non-standard toolboxes (eg. GLM) are provided in the Methods section of the paper. All statistical analysis was performed in MATLAB using standard toolboxes. Code used for analysis can be found in the Github link provided in the Methods section of the paper.

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors/reviewers upon request. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Research guidelines for submitting code & software for further information.

Data

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A list of figures that have associated raw data
- A description of any restrictions on data availability

All data that support the findings presented in this study are available from the corresponding author upon request.

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For a reference copy of the document with all sections, see <u>nature.com/authors/policies/ReportingSummary-flat.pdf</u>

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

Sample size

No sample-size calculation was performed. However, sample sizes, such as the number of neurons or animals, were similar to, or exceeded those reported in recently published papers from our laboratory (Schmitt et al. 2017, Nature) and others (Runyan et al. 2017, Nature).

Data exclusions

Behavioral sessions in which mouse performance was close to chance (as assessed by regression models, described in the Methods) were excluded from further analysis. Neurons with spiking on less that 25% of trials were also excluded from analysis. For all neural Poisson GLM data analysis, only cells with explained variance greater than 70% were used

Replication

Reproducibility of all modeling results was ensured via 200 runs of five-fold cross-validation. For electrophysiological results, reproducibility was ensured by sampling from a similar number of neurons across 3-4 different mice. Most experiments also included data collected by different experimenters. Optical perturbations of both the MD and PFC resulted in similar effects in each session, and all attempts at replicating the behavioral results reported in the paper were successful.

Randomization

Due to the design of our task, we did not have different experimental groups. Each mouse was trained to complete all the tasks described in the paper. The order of cues in each context block, as well as the order of the context blocks, were pseudo-randomized from session to session with a new random seed for each session. Optogenetic manipulations were also pseudorandomized.

Blinding

Due to the nature of our experiments, the experimenters were not blind to the type of optogenetic manipulation being performed. This is because the laser intensity had to be carefully calibrated for each mouse to ensure weak suppression of MD neurons. This is discussed further in the Methods section of the paper.

Reporting for specific materials, systems and methods

Materials & experimental systems

n/a Involved in the study

Unique biological materials

Antibodies

Eukaryotic cell lines

Palaeontology

Animals and other organisms

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n/a Involved in the study

ChIP-seq

Flow cytometry

MRI-based neuroimaging

Animals and other organisms

Human research participants

Policy information about studies involving animals; ARRIVE guidelines recommended for reporting animal research

Laboratory animals

This study involved a total of 5 mice. All mice were male and were 8 weeks of age at the time of surgery. One mouse had a SOM-Cre genotype (Jax: 013044, Sst-IRES-Cre). Four mice had the C57BL/6J genotype (Taconic Biosciences).

Field-collected samples

No field-collected samples were used in this study.

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