



Supplementary Materials for

Hierarchical reasoning by neural circuits in the frontal cortex

Morteza Sarafyazd and Mehrdad Jazayeri*

*Corresponding author. Email: mjaz@mit.edu

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Materials and Methods

Inferred rule experiment

A trial was initiated when the animal fixated a gray central fixation spot (diameter = 0.5 deg). After a random delay (400-800 ms, uniform hazard), a blue and a red visual target (diameter = 1 deg) were presented above and below the fixation point (distance = 7 deg). The animal had to report its belief about the rule by making a saccade to one of the targets within 1000 ms of their appearance (blue for C_1 and red for C_2). After the rule report epoch, the animal had to shift its gaze back to the fixation point to initiate the interval discrimination epoch (trial was aborted if re-fixation took longer than 500 ms). The trial proceeded regardless of whether the rule was reported correctly. After another random delay (400-900 ms, uniform hazard), two 50-ms flashes (diameter = 0.75 deg) were presented sequentially, a flash at the center of the screen followed by a flash to the left or right of the fixation point (distance = 9 deg). These two flashes demarcated the sample interval, t_s , which was drawn randomly from a 9-point discrete uniform distribution ranging between 530 and 1170 ms. In less than 700 ms after the second flash, the animal had to make a prosaccade toward the second flash or an antisaccade away from it to report whether its estimate of t_s was shorter or longer than a criterion of 850 ms, which corresponds to the mean of t_s distribution. We limited the reaction time to 700 ms to motivate the monkey to control the preparatory prepotent response inhibition dynamically during the interval measurement epoch. At the end of each trial, reward was provided (with a green visual feedback at the target location) if both the rule report in the first epoch and the interval discrimination in the second epoch were correct. No visual feedback or reward was given in error trials. When t_s was equal to the criterion (850 ms) in which case there was no correct answer for the interval discrimination, the animal got reward on a random 50% of the trials if it reported the rule correctly. The trial was aborted if the animal broke fixation before the second flash.

Instructed rule experiment

This task was identical to the rule inference experiment task with the only difference that the rule was cued explicitly and therefore the animal did not need to infer the rule based on history of errors. To specify the rule, we presented a red or blue disc around the fixation spot during the first 3-5 trials after every rule switch and 50-80% of trials until the next rule switch. The color of the disc served as an explicit cue for the current rule. To make sure the structure of the experiments was identical, in the instructed rule experiment, similar to the inferred rule experiment, animals had to first report the rule. On trials that the disc was not presented, the animal could infer it from memory of the previous trial since all rule switches were explicitly cued. The colored disc around the fixation spot disappeared immediately after the animal made a saccade to report the rule so that the interval discrimination epoch of the two tasks were identical.

Model for behavior

We developed a model with two hierarchically organized stages to capture behavior. The model is illustrated schematically in Fig. S3 and S4. The first stage captures the psychometric function describing the probability of Anti (Pr(Anti)) within a trial as a function of rule and t_d in that trial (Fig. S3). The second stage captures the probability of

switch ($\text{Pr}(\text{Sw})$) across trials as a function of feedback, t_d and rule in the preceding trials (Fig. S4). 1 and 2 refer to the parameters associated with the first and second stages, respectively. We use simple letter names for either experimentally controlled variables (e.g., t_s) or observable behavioral variables (e.g., action: A), and letter names with superscript (^) for behaviorally unobservable variables (e.g., internal belief about rule, \hat{C} , or internal estimate of discriminant interval, \hat{t}_d) and intended actions (e.g., intended pro or antisaccade, \hat{A}).

Modeling the first level of the decision hierarchy

We assumed that the observer makes a noisy measure of t_s (\hat{t}_s). Consistent with scalar variability of timing (93, 94), we modeled the noise by a zero-mean Gaussian whose standard deviation scales with t_s with the constant of proportionality w_m , as follows:

$$p(\hat{t}_s | t_s, w_m) \sim \mathcal{N}(t_s, w_m t_s)$$

The observer compares \hat{t}_s to an internal criterion, \hat{t}_c ($\hat{t}_d = \hat{t}_s - \hat{t}_c$), to categorize the interval as “Short” or “Long”. We treated \hat{t}_c as a parameter that was fixed throughout each behavioral session but was allowed to change across sessions. Moreover, we included a lapse rate, Γ_t to accommodate trials in which the animal categorized t_s randomly (50% “Short” and 50% “Long”). With this formulation, the probability of an intended antisaccade can be written as follows:

$$p(\hat{A} = \text{Anti} | \hat{C}, t_s) = \frac{\Gamma_t}{2} + (1 - \Gamma_t) \int_{\hat{t}_s} p(\hat{A} = \text{Anti} | \hat{C}, \hat{t}_s, \hat{t}_c) p(\hat{t}_s | t_s, w_m) d\hat{t}_s$$

where the first term in the integrand is determined by task contingencies as follows:

$$p(\hat{A} = \text{Anti} | \hat{C}, \hat{t}_s, \hat{t}_c) = \begin{cases} 1, & \text{if } \hat{C} = C_1 \text{ and } \hat{t}_s > \hat{t}_c \\ 0, & \text{if } \hat{C} = C_1 \text{ and } \hat{t}_s < \hat{t}_c \\ 0, & \text{if } \hat{C} = C_2 \text{ and } \hat{t}_s > \hat{t}_c \\ 1, & \text{if } \hat{C} = C_2 \text{ and } \hat{t}_s < \hat{t}_c \end{cases}$$

We assumed that the observer uses \hat{A} to generate a Pro or Anti action (A). In decision making studies, typically there is no need to distinguish between intended and executed actions. In our work, this distinction is warranted because animals occasionally fail to execute an intended antisaccade (39) when the inhibition to suppress the competing prepotent prosaccade is not sufficiently strong (25, 39, 40). To account for this asymmetry, the model was augmented with a parameter, Γ_{Anti} , to account for the small percentage of trials in which an intended antisaccade resulted in an erroneous prosaccade. The full model thus used 4 parameters ($\Psi_1 = \{w_m, \hat{t}_c, \Gamma_t, \Gamma_{\text{Anti}}\}$) to capture the probability of antisaccades as a function of t_s and C :

$$p(A = \text{Anti}|\hat{C}, t_s) = (1 - \Gamma_{\text{Anti}}) \left[\frac{\Gamma_t}{2} + (1 - \Gamma_t) \int_{\hat{t}_s} p(\hat{A} = \text{Anti}|\hat{C}, \hat{t}_s, \hat{t}_c) p(\hat{t}_s|t_s, w_m) d\hat{t}_s \right]$$

Finally, we assumed that actions across trials were conditionally independent and used the sum of the logarithm of individual conditional probabilities to derive the maximum likelihood estimate (MLE) of the model parameters:

$$\Psi_1(\text{MLE}) = \arg \max_{\Psi} \left[\sum_{i=1}^{N_{\text{Anti}}} \log p(A^i = \text{Anti}|\hat{C}^i, \hat{t}_s^i, \Psi) + \sum_{i=1}^{N_{\text{Pro}}} \log(1 - p(A^i = \text{Anti}|\hat{C}^i, \hat{t}_s^i, \Psi)) \right]$$

where indices i and j count over N_{Anti} antisaccade and N_{Pro} prosaccade trials, respectively. We fit this model to behavioral data in individual sessions of both the inferred rule (subjective rule) and instructed rule experiments.

Modeling the second level of the decision hierarchy

As a first step, we developed an ideal observer model (IOM) whose switch behavior on trial k is governed by posterior odds of rule change compared to no rule change (3), denoted Q_{IOM}^k . If we index the last rewarded trials by k_0-1 , Q_{IOM}^k has to be computed based on the consecutive error trials k_0, \dots, k (denote $[k_0:k]$) following the last rewarded trials:

$$Q_{IOM}^k = \frac{Pr(C^{k+1} \neq \hat{C}^k | t_s^{[k_0:k]}, \hat{C}^{[k_0:k]})}{Pr(C^{k+1} = \hat{C}^k | t_s^{[k_0:k]}, \hat{C}^{[k_0:k]})}$$

The denominator is the probability of no rule change despite negative feedback across all $[k_0:k]$ trials. For every trial i , the probability of being incorrect without a rule change is the product of the probability that no objective rule change occurred multiplied by the probability that the interval was discriminated incorrectly. If we denote the probability of correctly discriminating t_s in trial i by y^i , and the probability of objective covert rule change in trial i by its hazard function, denoted λ^i , the probability of an incorrect discrimination in the absence of rule change becomes $(1-\lambda^i)(1-y^i)$. The denominator therefore is the product of this probability across all $[k_0:k]$ trials. Using a similar logic, the numerator is the sum of multiple probabilities including the probability that the rule change occurred on trial k_0 , that it occurred on trial k_0+1 but not on trial k_0 , and the all the other terms until the last terms, which corresponds to a rule change on trial k with no rule change in the preceding $[k_0:k-1]$ trials. Accordingly, Q_{IOM}^k can be written as follows:

$$Q_{IOM}^k = \frac{\lambda^{k_0} + \sum_{i=k_0+1}^k \lambda^i \prod_{j=k_0}^{i-1} (1-\lambda^j)(1-y^j)}{\prod_{i=k_0}^k (1-\lambda^i)(1-y^i)}$$

Since y^i is a measure of expected accuracy (i.e., the probability of being correct), we quantified it based on our model of the subjective psychometric function after marginalizing over t_s (which the ideal observer does not know). For λ^i , to avoid over-parameterizing the model, we did not attempt to model the animal's subjective hazard

function and instead, assumed a constant hazard rate consistent with our experimentally imposed block transition scheme. With y^i and λ^i specified, we can fully derive Q_{IOM}^k with no additional free parameters.

Due to its history-dependent nature, direct use of IOM to characterize animal's behavior is not feasible (i.e., would need far more data that we can collect). As an approximation, we developed a simpler confidence-based model (CBM). The behavior of CBM is governed by a graded latent variable representing trial-by-trial evidence for a covert switch, which we denote by X_Σ . In this model, X_Σ is incremented after each error with a magnitude that depends on the previous trial's expected accuracy, and leads to a switch when X_Σ breaches a threshold (θ). To match CBM as closely as possible to IOM, we assumed that X_Σ on each trial has a Gaussian distribution. We set the mean and standard deviation (denoted by μ_{X_Σ} and σ_{X_Σ}) of X_Σ such that Q_{CBM}^k (probability that X_Σ is above θ over the probability it is below θ) was equal to Q_{IOM}^k for 1B-Er. Moreover, we assumed that switch evidence is subject to diffusive noise across trials; i.e., that variance grows linearly with the number of error trials (i.e., $\sigma_{X_\Sigma}^2$ for N errors = $N \sigma_{X_\Sigma}^2$ for 1 error). We also considered a nonlinear relationship between variance and number of trials (Fig. S5D-F) but the simple diffusive (linear) model was able to capture animals' switch behavior more accurately. Note that in the main manuscript, we denote latent representation of subjective switch evidence as \hat{X}_Σ , however, here we use X_Σ (which is the actual switch evidence) for simplicity.

$$p(X_\Sigma) \sim \mathcal{N}(\mu_{X_\Sigma}, \sigma_{X_\Sigma})$$

$$Q_{CBM}^k = \frac{P(X_\Sigma \geq \theta)}{P(X_\Sigma < \theta)} = \frac{\int_{\theta}^{\text{inf}} \mathcal{N}(\mu_{X_\Sigma}, \sigma_{X_\Sigma}) dX_\Sigma}{1 - \int_{\theta}^{\text{inf}} \mathcal{N}(\mu_{X_\Sigma}, \sigma_{X_\Sigma}) dX_\Sigma} = Q_{IOM}^k$$

Rearranging this equation, we can write:

$$\int_{-\text{inf}}^{\theta} \mathcal{N}(\mu_{X_\Sigma}, \sigma_{X_\Sigma}) dX_\Sigma = 1 - \frac{Q_{IOM}^k}{1 + Q_{IOM}^k}$$

which can be solved analytically and numerically to derive μ_{X_Σ} in terms of Q_{IOM}^k and σ_{X_Σ} . This formulation enabled CBM to closely approximate IOM. To fit CBM to animal's behavior, we made two simplifying assumptions: $X_\Sigma = \theta$ after a rewarded trial, and $\theta = 1$. These are inconsequential because their effect on behavior can be fully subsumed by μ_{X_Σ} and σ_{X_Σ} .

To account for the deviation of each animal's behavior from the ideal observer model, we augmented μ_{X_Σ} by a less-than-one multiplicative 'perseverance factor', α . This factor reduces the magnitude of increments in X_Σ and causes an overall drop in the probability of switch. The full model thus has three parameters, μ_{X_Σ} , σ_{X_Σ} , and α . However because of the relationship between μ_{X_Σ} , σ_{X_Σ} , and the known Q_{IOM}^k , μ_{X_Σ} is not a free parameter. Therefore, our model for rule inference had only two free parameters

($\Psi_2 = \{\sigma_{X\Sigma}, \alpha\}$). In this formulation, the probability of switch ($X_{y/n}=1$) can be simplified as follows:

$$p(X_{y/n} = \text{Sw}|\Psi_2) = \int_1^{\text{inf}} \mathcal{N}(\alpha\mu_{X\Sigma}, \sigma_{X\Sigma})dX_\Sigma$$

Finally, we assumed that switch decisions were conditionally independent across trials and used the sum of the logarithm of individual conditional probabilities to derive the maximum likelihood estimate (MLE) of the model parameters. The model fits were based on data in the 1B-Er and 2B-Er trials.

$$\Psi_2(\text{MLE}) = \arg \max_{\Psi} \left[\sum_{i=1}^{N_{\text{Sw}}} \log p(X_{y/n} = \text{Sw}|\Psi) + \sum_{j=1}^{N_{\text{No Sw}}} \log(1 - p(X_{y/n} = \text{Sw}|\Psi)) \right]$$

We performed several additional analyses to validate our assumptions about CBM and to verify that it was a suitable model: (1) *In-silico* model lesioning: we assessed the necessity of $\sigma_{X\Sigma}$ and α by testing the capacity of the model to capture the observed switch behavior when either of those parameters were fixed. Results in Fig. S5G-L show two examples, one in which $\sigma_{X\Sigma}$ was set to 0.5, and α was fitted, and another when α was set to 1 and $\sigma_{X\Sigma}$ was fitted. However, our conclusions were general and applied to other fixed values of $\sigma_{X\Sigma}$ and α so long as the fixed values were not identical to those recovered by the full model. (2) Model identification: we applied the model fitting procedure to simulated data generated by CBM to ensure that parameters could be recovered. Results are shown in Fig. S5C. (3) 2-fold cross-validation: to ensure that the success of the model was not due to overfitting, we repeatedly fit the model to a random half of each dataset and test the model on the remaining half. Results are shown in Fig. S5B. (4) Predictive validity: we simulated the model fit to the subjective psychometric function in the presence of experimentally imposed covert rule switches, and asked whether it could predict the observed objective psychometric function without additional parameterization and/or fitting. Results are shown in Fig. 1F.

Model free analysis of switch behavior

We also evaluated whether switch probability depended on trial difficulty (indexed by $|t_d|$) and the number of consecutive trials using a simple logistic regression:

$$P(X_{y/n}^{i+1} = \text{Sw}|t_d^i, n\text{B-Er}) = \frac{1}{1 + e^{-(\beta_0 + \beta_{|t_d|}|t_d^i| + \beta_{n\text{B-Er}}[n\text{B-Er}])}} \quad \text{Equation 1}$$

where $P(X_{y/n}^{i+1})$ represents the probability of switch in trial $i+1$, t_d^i , the discriminant interval in trial i , and $n\text{B-Er}$, the number of consecutive error before trial $i+1$. The parameters of the logistic model ($\beta_0, \beta_{|t_d|}, \beta_{n\text{B-Er}}$) capture the baseline, dependence on $|t_d^i|$, and dependence on $n\text{B-Er}$ of $P(X_{y/n}^{i+1})$, respectively. We used MLE to fit the model to behavior, and used fits to $\beta_{|t_d|}$ and $\beta_{n\text{B-Er}}$ to assess the statistical significance of the effect of $|t_d^i|$ and $n\text{B-Er}$ on switch probability, respectively.

Analysis of single neurons

We estimated firing rates of each neuron per condition by computing average spike counts in 25 ms time bins followed by convolution with a Gaussian boxcar filter ($\sigma = 50$ ms). For each neuron, we compared average firing rates during the inter-trial-interval (ITI) after rewarded and unrewarded trials (Fig. 2). We also compared responses for both rewarded and unrewarded trials further broken down according to whether the preceding trial was relatively “easy” or “difficult. We defined “easy” and “difficult” as trial in which $|t_d| \geq 160$ ms and $|t_d| < 160$ ms, respectively. Collapsing the full range of t_d to a small number of categories was necessary to overcome the limited number of error trials per sessions, but the results were qualitatively the same regardless of the criterion for defining easy versus difficult trials (we also did all the analyses for $|t_d| \geq 80$ ms and $|t_d| < 80$ ms). We also compared average firing rates as a function of number of preceding consecutive errors either across all values of t_d or after further dividing trials of the last error according to trial difficulty ($|t_d| \geq 160$ ms versus $|t_d| < 160$ ms).

Retrospective and prospective information in DMFC and ACC

We used a time-dependent linear regression model to examine the time course of the representations of trial outcome (Rw/Er) and switch behavior ($X_{y/n}$) across individual neurons in DMFC and ACC. We z-scored the firing rates of each neuron across all trials and times ($r(t)$), and formed a regression model that aimed to explain the firing rates as a function of trial outcome, $X_{y/n}$, and other experimental and behavioral variables as follows:

$$r(t) = \beta_{outcome} I_{Rw/Er} + \beta_{Sw} I_{y/n} + \beta_{Target} I_{left/right} + \beta_A I_{Pro/Anti} + \beta_{td} t_d + \beta_{Rule} I_{C1/C2} + \beta_0$$

Equation 2

In this regression, $I_{Rw/Er}$ was set to 1 (Er) or 0 (Rw), $I_{y/n}$ was set to 0 (no switch) or 1 (switch), $I_{left/right}$, to -1 or 1 depending on the saccade direction, $I_{Pro/Anti}$, to -1 or 1 depending on the animal’s action, $I_{C1/C2}$ to -1 or 1 depending on rule, and t_d was the experimentally imposed variable. Solving this equation yielded $\beta_{outcome}$ and β_{Sw} as a function of time for each neurons. Since our interest was to evaluate the relative strength of sensitivity across neurons and areas, we converted $\beta_{outcome}$ to a selectivity index by 1) taking the absolute value of each $\beta_{outcome}$ and 2) normalizing all $\beta_{outcome}$ at each time point by the average $\beta_{outcome}$ across all neurons, areas, and time during ITI. We did the same for β_{Sw} .

$$SI_{outcome} = \frac{|\beta_{outcome}|}{\langle |\beta_{outcome}| \rangle_{neurons, areas}}$$

$$SI_{Sw} = \frac{|\beta_{Sw}|}{\langle |\beta_{Sw}| \rangle_{neurons, areas}}$$

Finally, we averaged the selectivity index associated with $\beta_{outcome}$ and β_{Sw} across significantly sensitive neurons in each area and compared the time course of these two selectivity indices across the two areas (Fig. 3).

Sensitivity of population activity in DMFC and ACC to X_{Σ} and $X_{y/n}$

We employed a targeted dimensionality reduction (63) to assess the encoding of information about X_{Σ} and $X_{y/n}$ across the population in DMFC and ACC. This approach has been carefully described before (63). Here, we briefly describe the procedure. First, we z-scored firing rates of each neuron across all trials and times, and formed a regression model that aimed to explain the entire vector of firing rates across the population (Z) at each point in time as a function of various experimental and behavioral variables as follows:

$$Z = \beta_{outcome} I_{Rw/Er} + \beta_{X_{\Sigma}} X_{\Sigma} + \beta_{S_w} I_{y/n} + \beta_{Target} I_{left/right} + \beta_A I_{Pro/Anti} + \beta_{td} t_d + \beta_{Rule} I_{C1/C2} + \beta_0$$

Equation 3

The regressors are the same as described for Equation 2, with the addition of X_{Σ} which was inferred on a trial-by-trial basis from the fit of the confidence-based model to behavior. Second, we solved the regression model to derive the regression coefficients. Third, we denoised the coefficients by finding their projection in the subspace spanned by the first 10 principal components (PCs) of data that explained ~70% (or more) of the total variance in both areas and animals. To compute the PCs, we applied PCA to data resorted to a matrix of dimensions $[N_{units}, N_C \times T]$ (N_{units} : number of neurons, N_C : number of conditions, T : number of time bins). Fourth, for each regressor, we found the time at which the coefficient has the largest norm (i.e., the encoding of the corresponding variable of interest was strongest). Fifth, we orthogonalized the coefficients. To do so, we applied QR-decomposition to a matrix whose columns were the coefficients, and used the columns of the matrix Q as the new orthogonalized coefficients.

Next, we used the orthogonalized coefficients, $\beta_{X_{\Sigma}}$ and β_{S_w} , to examine how DMFC and ACC encoded X_{Σ} and $X_{y/n}$ at the population level. We first describe the procedure of $\beta_{X_{\Sigma}}$: 1) we found the projection of data onto $\beta_{X_{\Sigma}}$, which we refer to as $Z_{X_{\Sigma}}$; 2) we divided $Z_{X_{\Sigma}}$ into two groups, one containing all the trials that did not lead to a switch and another for all the trials that did lead to a switch; 3) we divided trials in each group to 6 overlapping bins depending on the value of X_{Σ} (i.e., low X_{Σ} , intermediate X_{Σ} , and high X_{Σ}). The bin and overlapping window were chosen so that the number of trials in each bin were nearly equal. 4) we used d' as a distance metric to quantify the separation between the distributions of $Z_{X_{\Sigma}}$ in each bin to the distribution of $Z_{X_{\Sigma}}$ associated with rewarded trials (i.e., when $X_{\Sigma}=0$). This distance metric was used to assess the sensitivity of DMFC and ACC to X_{Σ} . To examine the statistics, the population analysis method was cross-validated by resampling dataset into equally-sized training and testing sets (bootstrapping method, 100-times). The procedure for β_{S_w} was identical except that we used the projections of data onto β_{S_w} (instead of $\beta_{X_{\Sigma}}$), which we refer to as Z_{S_w} . Analysis of Z_{S_w} with respect to X_{Σ} enables us to assess the extent to which DMFC and/or ACC encoded $X_{y/n}$ as a binary variable (i.e., independent from X_{Σ}). In addition, as a control analysis, we conditioned the same subsets of trials on anti/pro-saccades, and separately on left/right saccades, and the projections into $\beta_{X_{\Sigma}}$ were measured to quantify the action dependency of X_{Σ} neural state (Fig. S10).

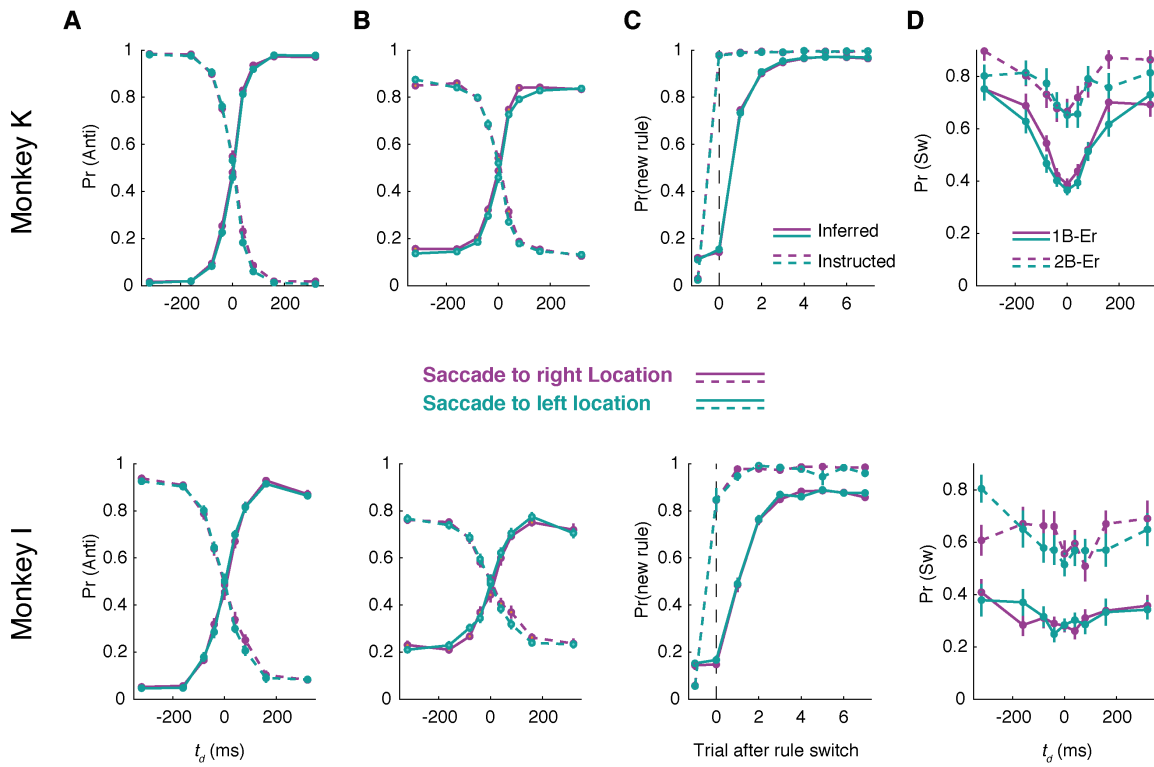


Fig. S1. Switch behavior of the two monkeys as a function of saccade location. (A-D) Results are shown with the same ordering and format as in Fig. 1E-H. Behavior is largely independent of the direction of saccade. There were no significant difference between corresponding points on each plot (point-by-point Wilcoxon rank sum test; $P > 0.05$).

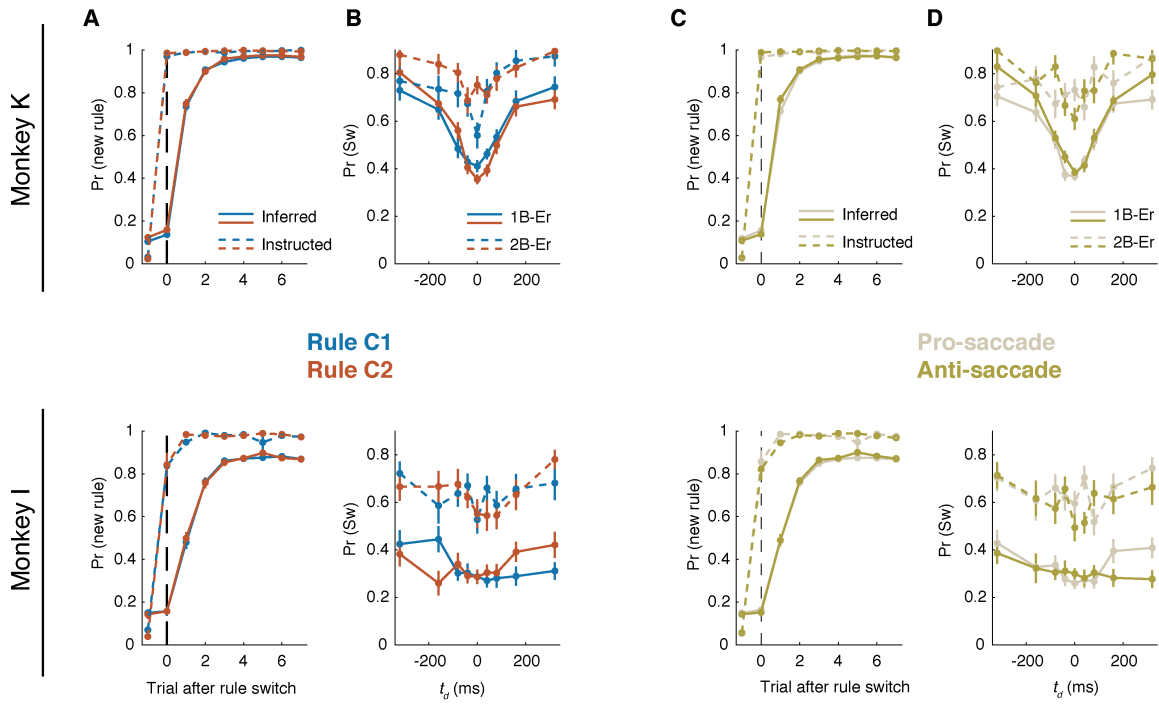


Fig. S2. Switch behavior of the two monkeys as a function of rule (C_1 versus C_2) and action (Pro versus Anti). Results are shown with the same ordering and format as in Fig. 1G-H. Behavior is largely independent of the direction of saccade. There were no significant difference between corresponding points on each plot (point-by-point Wilcoxon rank sum test, $P > 0.05$).

Time interval discrimination (Action within a trial)

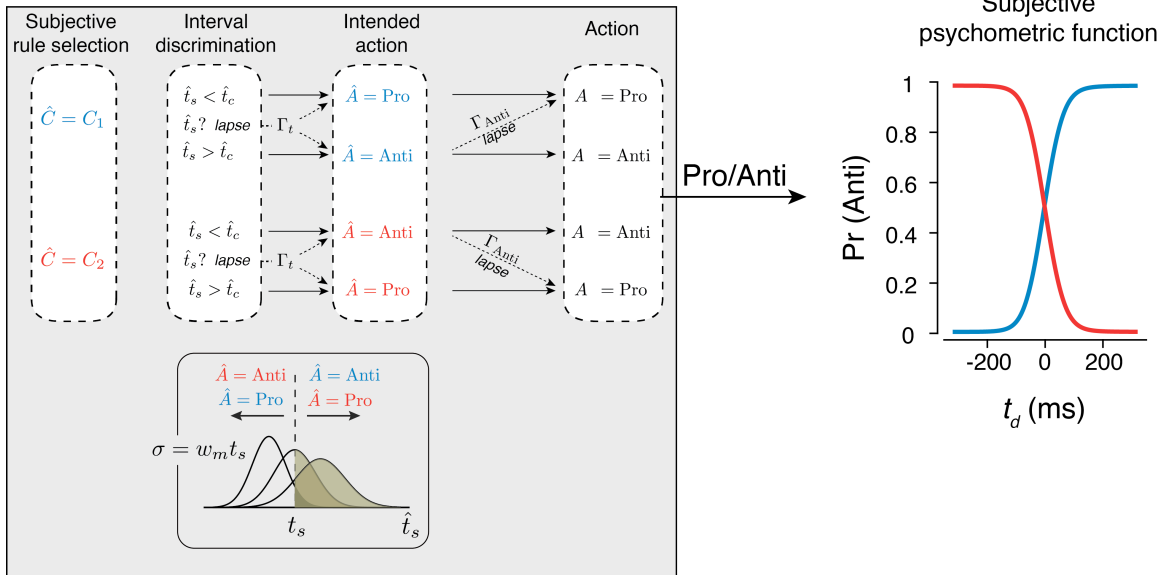


Fig. S3. Description of the observer model of time interval discrimination. The observer model makes a noisy measurement of the sample interval, t_s , denoted \hat{t}_s . The measurement noise has a zero-mean Gaussian distribution and its standard deviation (σ) scales with t_s with constant of proportionality, w_m ($\sigma = w_m t_s$). The model determines the intended action, \hat{A} based on rule \hat{C} ($\hat{C} = C_1$ or $\hat{C} = C_2$) and \hat{t}_s relative to an internal criterion (\hat{t}_c). When $\hat{C} = C_1$ and $\hat{t}_s > \hat{t}_c$, \hat{A} is an antisaccade (Anti), and when $\hat{C} = C_1$ and $\hat{t}_s < \hat{t}_c$, \hat{A} is a prosaccade (Pro). For $\hat{C} = C_2$, these response contingencies are reversed. The model also allows for a percentage of ‘lapse’ trials (Γ_t), in which \hat{A} is randomly assigned to Pro or Anti (shown as $\hat{t}_s?$). The model’s action, A usually follows from \hat{A} , but it also allows for a percentage of additional lapses parametrized by Γ_{Anti} in which $A = \text{Pro}$ despite $\hat{A} = \text{Anti}$ due to a failure to inhibit the prepotent prosaccade response. In sum, the model has 4 parameters ($\Psi_1 = \{w_m, \hat{t}_c, \Gamma_t, \Gamma_{\text{Anti}}\}$) to capture the probability of antisaccades as a function of t_s and \hat{C} . The two psychometric functions on the right are examples of the model’s behavior for the two rules shown in terms of proportion of antisaccades ($\text{Pr}(\text{Anti})$) as a function of the discriminant interval, t_d .

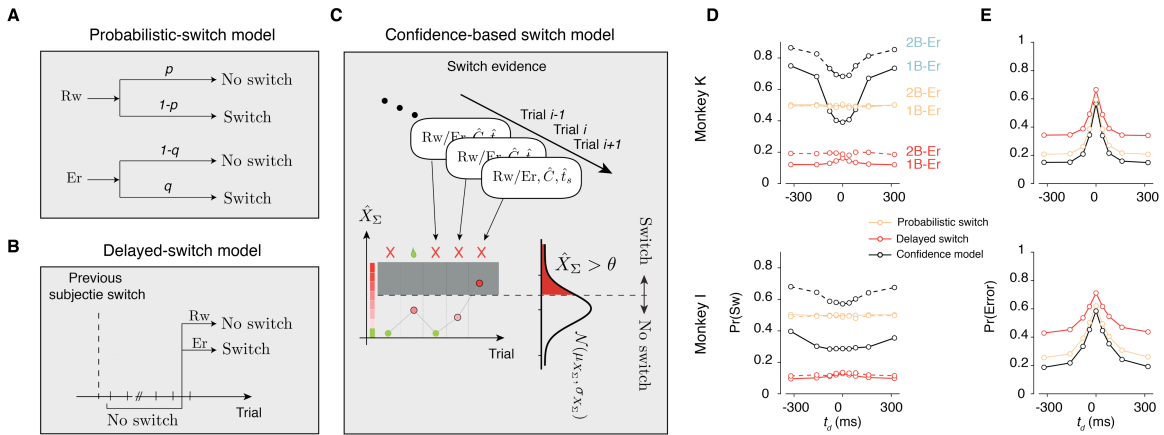


Fig. S4. Alternative models for switching behavior. (A) Probabilistic switch model. This model implements a variant of the win-stay lose-switch strategy in which the animal switches the rule with different fixed probabilities (p and q) depending on trial outcome (Rw: rewarded trial, Er: error trial). (B) Delayed switch model. This model switches after an error only after a certain number of trial since the last successful switch. The model emulates an observer that exploits knowledge about the block-structure in the experiment. (C) Confidence-based switch model. This model computes a graded variable, \hat{X}_Σ , whose value is determined by the outcome (Rw/Er) and difficulty (indexed by \hat{t}_s) of preceding trials. \hat{X}_Σ is reset to zero after each trial with positive outcome, and is incremented by a sample of Gaussian distribution after error trials. The mean and standard deviation of the distribution (μ_{X_Σ} and σ_{X_Σ}) are set so that the model's behavior approximates the behavior of an ideal observer (see Methods). When the updated \hat{X}_Σ breaches a threshold, θ , the model switches its belief about the rule (\hat{C}). The model also uses a perseveration factor (α) to account for the deviation of each animal's behavior from optimality (not shown in C but see Methods). Since μ_{X_Σ} was determined by the ideal observer, the behavior of this stage of the model was fully explained by two parameters ($\Psi_2 = \{\sigma_{X_\Sigma}, \alpha\}$). (D) Switch probability (Pr(sw)) based on simulations of three models. Each model was constructed such that it matched as closely as possible to each animal's subjective psychometric function. The data from the probabilistic switch model was generated assuming no switch after rewarded trials and 50% switch after error trials irrespective of confidence. The data for the delay switch model was generated by assuming that no switch for the first 10 trials (consistent with the experimentally imposed hazard) after each successful switch, and 50% switch for each error trial afterwards. The data for the confidence-based switch model are the same as those shown in the main manuscript. For each model, the results are shown for 1-back error (1B-Er; solid) and 2-back error (2B-Er; dashed) conditions. (E) Percentage error as a function of the discriminant interval, t_d , for each of the three models with the same specifications as in D. The Confidence-based switch model has the lowest probability of error in comparison to the the other two models.

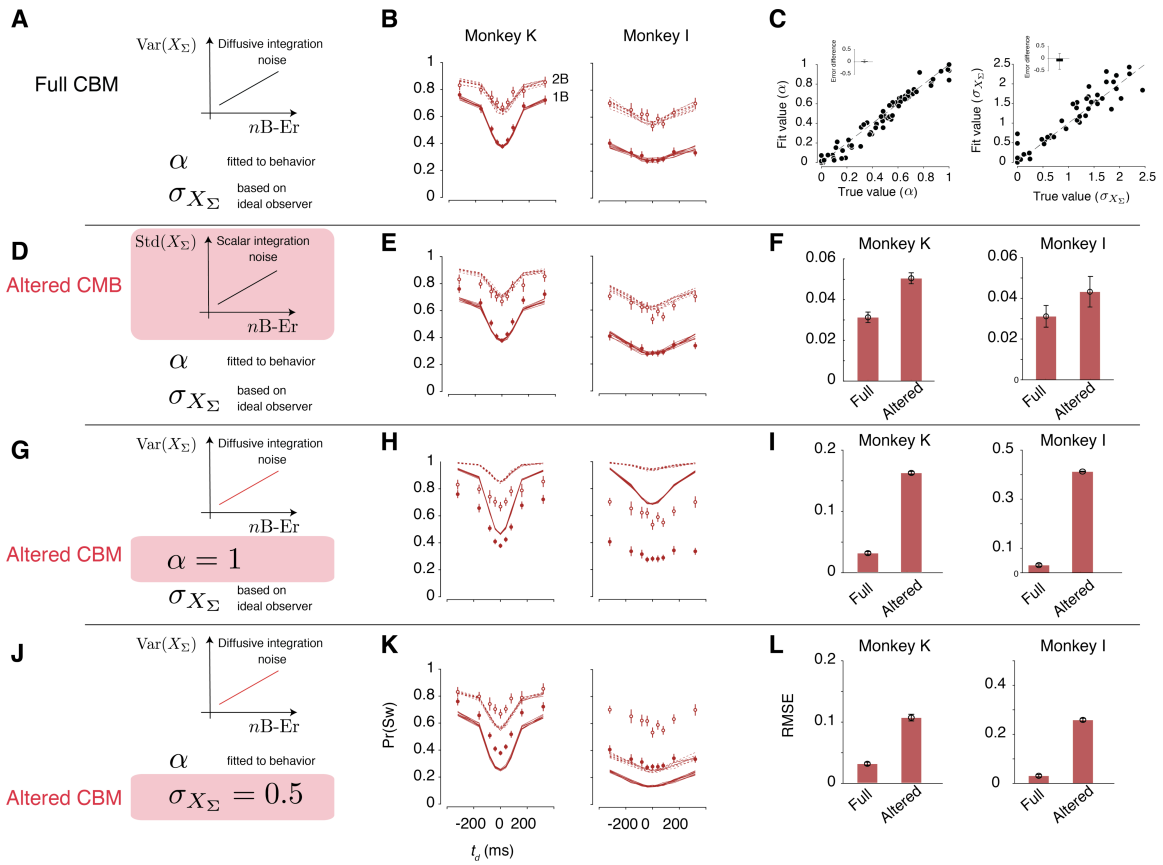


Fig. S5. Cross-validation, identification and lesion studies of the confidence-based belief update model. (A) Full CBM model with two parameters α and $\sigma_{X\Sigma}$. (B) Validation of the full CBM model. Fit of CBM (line) to switch behavior of two monkeys (symbols) in the same format as Fig. 1H. The model fits were cross-validated and repeated 10 times for random train and validation set. (C) Model identifiability. Fits of CBM to data generated from CBM with random choices of α and $\sigma_{X\Sigma}$. Inset: Difference between fitted and actual value across simulations. There is not significant difference between the fit and true values ($P > 0.05$). (D) Altered CBM in which the standard deviation (instead of variance) of noise increases with consecutive trials. The altered component (the one that differs from the full CBM) is highlighted by the shaded box. (E) Same as panel B for alternate model in D. (F) Comparison of the full and alternate CBM by using RMSE across 10 cross-validated fits to behavioral data (error bar: SEM). (G) Altered CBM in which α is set to 1 (no perseverance) while $\sigma_{X\Sigma}$ is fitted to behavior. (H,I) Same as E,F for the altered CBM in G. (J-L) Same as panels (G-I) for an altered CBM in which $\sigma_{X\Sigma}$ is set to 0.5 (instead of being inferred from the ideal observer) while α is fit.

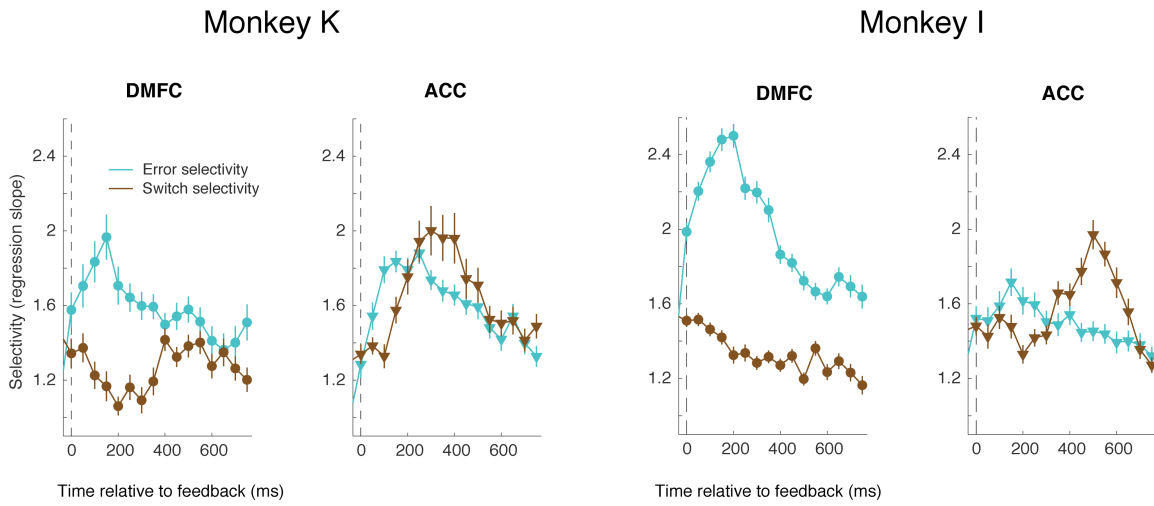


Fig. S6. Retrospective and prospective computations in DMFC and ACC during ITI, separately for the two animals. Results are shown in the same format as in Fig. 3A,B, separately for the two animals.

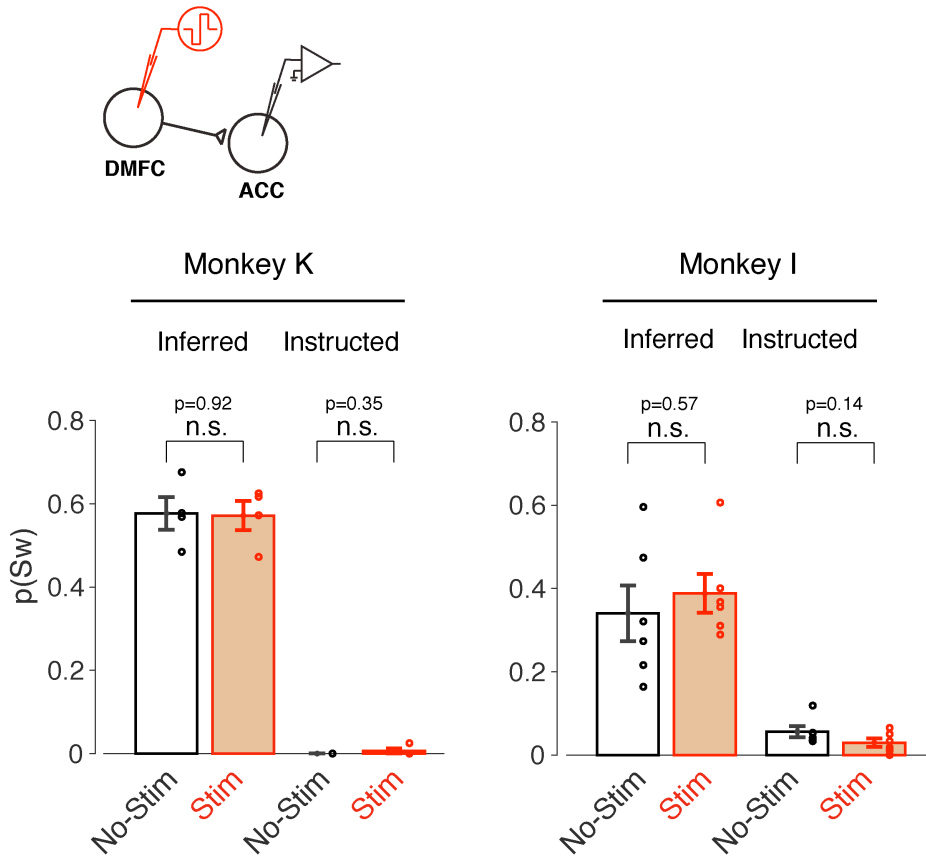


Fig. S7. No effect of DMFC microstimulation on switch probability. The ordinate shows average switch probability ($\text{Pr}(\text{Sw})$) across all sample intervals after 1-back error trials separately for trials when DMFC was and was not stimulated. There was no significant change in $\text{Pr}(\text{Sw})$ in the inferred or instructed experiments as a result of DMFC microstimulation (n.s.: $P > 0.05$). For the instructed experiment, $\text{Pr}(\text{Sw})$ is only shown for trials where the subjective switch was not cued.

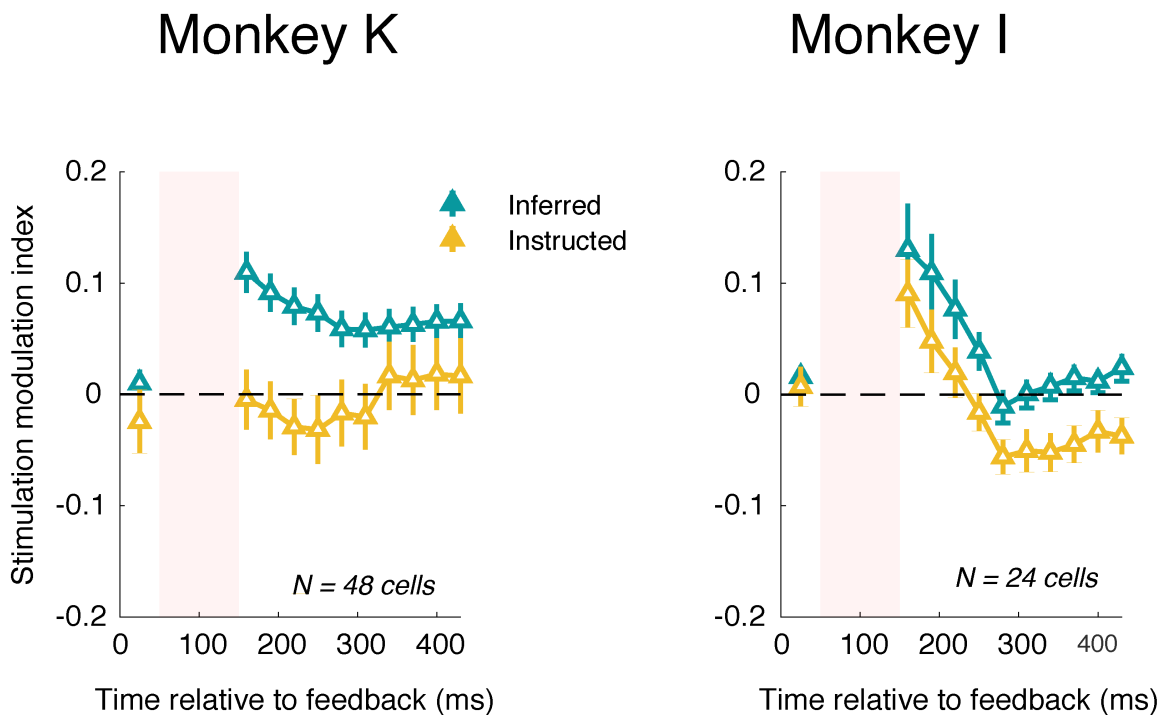


Fig. S8. Modulation of ACC activity due to DMFC microstimulation shown for the two animals separately. Results are shown in the same format as Fig. 3D. We examine the statistical significance of the stimulation modulation index in the inferred and instructed experiments independently. In each case, we compared the distribution of firing rates in the stimulated trials to a null distribution created by random labeling of trials as stimulated and unstimulated. We used a shuffling procedure to avoid making arbitrary assumptions about the sampling distributions. In the inferred experiment, DMFC microstimulation significantly increased the stimulation modulation index in ACC within 150-300 ms relative to feedback (Monkey K: mean=0.091, $P < 0.001$; Monkey I: mean=0.102, $P < 0.01$;). In contrast, there was no significant change in ACC activity in the same window after DMFC microstimulation in the instructed experiment (Wilcoxon rank sum test, $P > 0.05$). Since the results for the instructed experiment were not significant, we did not perform a direct statistical comparison of the inferred and instructed experiments. Although the results for both animals were statistically significant, we wish to highlight the difference between the two animals as this difference may provide stronger support for our hypothesis. The animal for which the effect of stimulation in the instructed task was stronger (“monkey I”), was less flexible at alternating between the inferred and instructed tasks. This is evident from a careful examination of the behavior. In monkey I, who occasionally did not switch following an instructed switch (compare yellow traces in Fig. 1G), DMFC stimulation influenced ACC during the instructed experiment. In contrast, in monkey K who followed the rule instruction nearly perfectly (Fig. 1G), the DMFC-stimulation had nearly no effect on ACC in the instructed experiment. In other words, the stimulation effects, when considered in relation to each animal’s behavior, seem to suggest that when rule inference via feedback is at play, DMFC stimulation has a more prominent effect on ACC, which is consistent with our hypothesis.

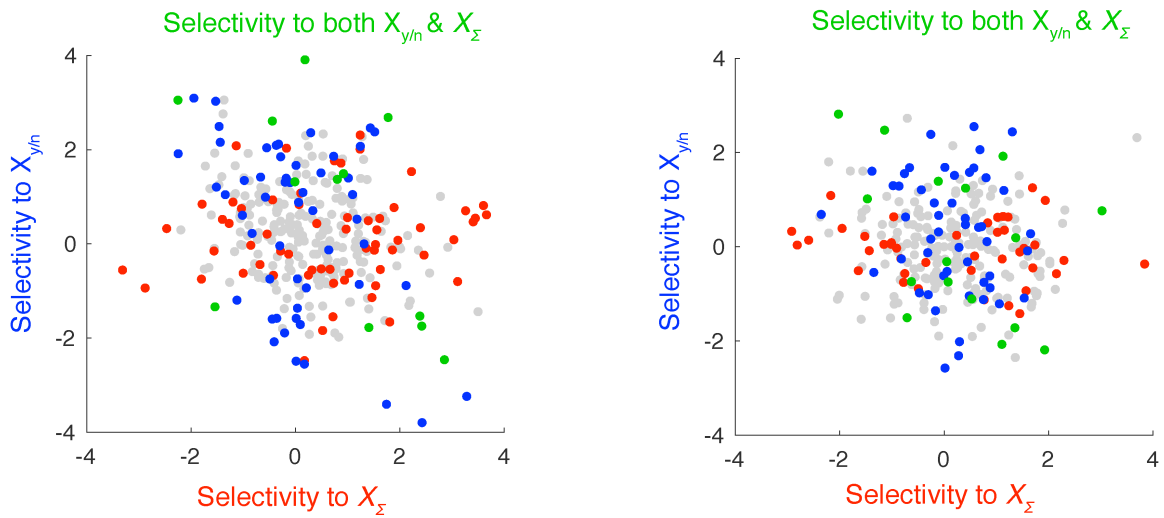


Fig. S9. Mixed representation of \hat{X}_{Σ} and $X_{y/n}$ at the level of individual neurons in ACC. The selectivity of each neuron to switch evidence and switch choice was measured using linear regression (see Methods). Colored dots represent neurons with significant selectivity (permutation test, 100 times, $P < 0.01$) to \hat{X}_{Σ} (red) and $X_{y/n}$ (blue) or both (green).

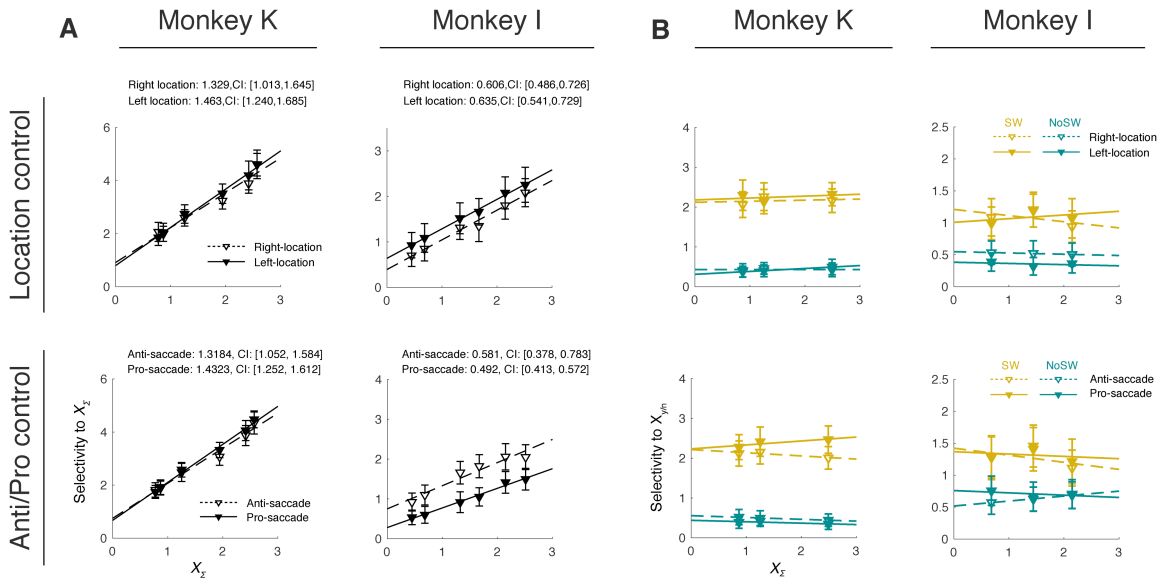


Fig. S10. Sensitivity of ACC to switch evidence was robust to nuisance parameters. (A) After fitting the regression model to ACC (Fig. 4), we separated trials based on saccade direction (left versus right) and response type (prosaccade versus antisaccade) to characterize the extent to which the encoding of switch evidence (X_{Σ}) was dependent on these nuisance variables. Results indicated that representation of X_{Σ} in ACC was present for both saccade directions and both response types. All regression slopes were significantly larger than zero ($P < 0.01$). (B) Same as panel (A) for the switch choice selectivity ($X_{\Sigma/n}$). Results indicated that representation of switch choice in ACC was present for both saccade directions and both response types ($P < 0.01$).

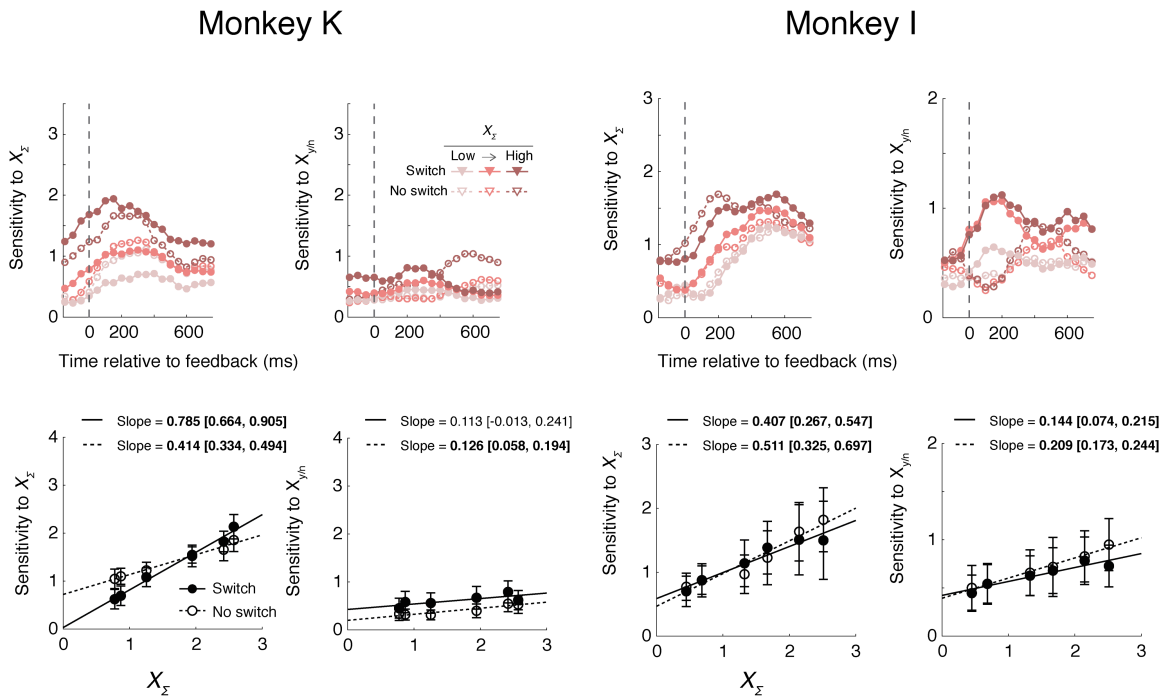


Fig. S11. Representations of switch evidence and switch choice in DMFC. Results are shown in the same format as in Fig. 4A,B, separately for the two animals. For the bottom panel, the slope of the regression line and the corresponding confidence intervals are reported. Regression slopes that were significantly different from zero are in bold.

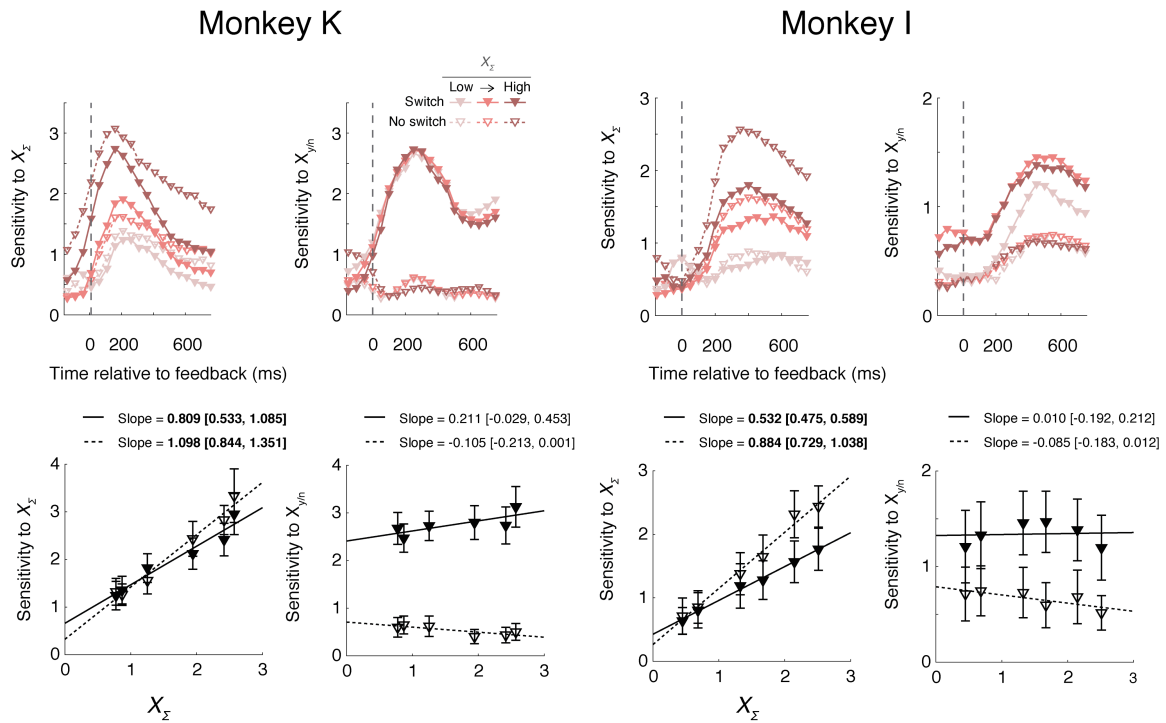


Fig. S12. Representations of switch evidence and switch choice in ACC. Results are shown in the same format as in Fig. 4A,B, separately for the two animals. For the bottom panel, the slope of the regression line and the corresponding confidence intervals are reported (slope [confidence interval]). Regression slopes that were significantly different from zero are in bold. Note that in ACC, selectivity to switch is largely independent of X_z (bottom row, second and fourth panels from left), and that neural responses strongly predict animals' switch behavior in the next trial (Table below shows the baseline of the regression line as a function of $X_{y/n}$ for the two animals). These results support our hypothesis that the decision to switch (i.e., threshold crossing) is likely made in ACC.

	$X_{y/n}$ (Sw)	Sensitivity to $X_{y/n}$ b_0 [CI]
Monkey K	Yes	2.408 , [1.977, 2.840]
	No	0.707 , [0.514, 0.900]
Monkey I	Yes	1.325 , [0.993, 1.657]
	No	0.788 , [0.627, 0.950]

* CI: 95% Confidence interval. Significant estimates ($P < 0.01$) are in bold.

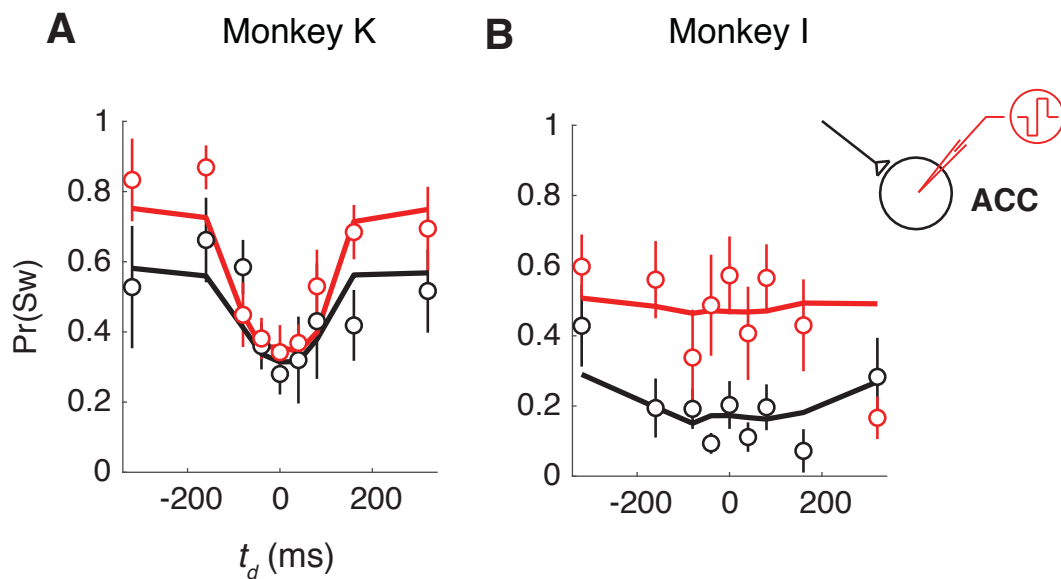


Fig. S13. The effect of ACC microstimulation on switch probability of the two animals separately. The effect of ACC microstimulation (red) compared to stimulation (black) is shown as a function of t_d in the preceding trial for monkey K (A) and monkey I (B). Stimulation was performed after a randomly selected 50% of 1-back error trials. The lines represent the fit of the confidence-based model (see Methods) to each animal for each condition.

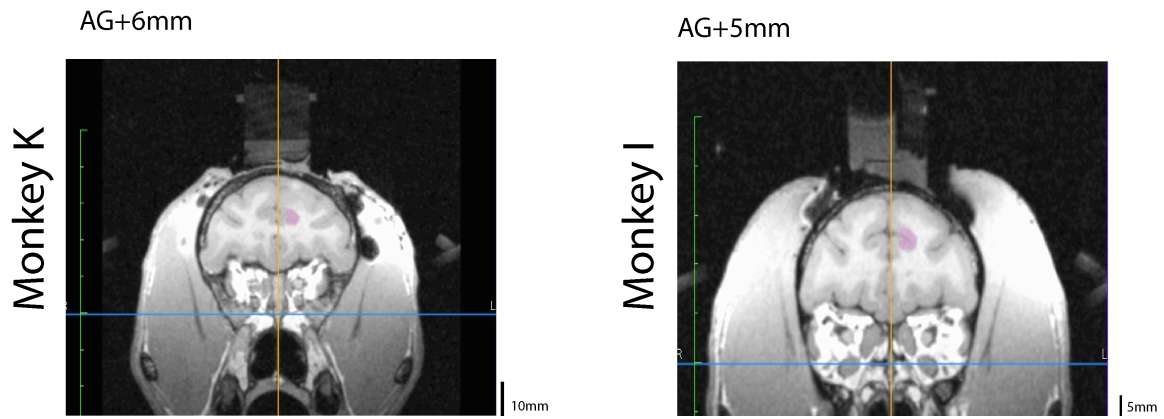


Fig. S14. Coronal section containing ACC from the structural MRI scan of the two animals. Magenta shows the recording site in ACC (AG: anterior to the genu of arcuate sulcus).

Parameter	Inferred rule experiment		Instructed rule experiment	
	Monkey K	Monkey I	Monkey K	Monkey I
w_m	0.0581 \pm 0.0042	0.1024 \pm 0.0107	0.0596 \pm 0.0043	0.1083 \pm 0.0093
\hat{t}_c (ms)	850.6 \pm 129.7 (C_1) 846.8 \pm 129.1 (C_2)	853.3 \pm 164.2 (C_1) 846.0 \pm 162.8 (C_2)	848.4 \pm 129.4 (C_1) 856.0 \pm 130.5 (C_2)	856.6 \pm 164.8 (C_1) 838.7 \pm 161.4 (C_2)
Γ_{Anti}	0.0084 \pm 0.0013	0.0108 \pm 0.0021	0.0080 \pm 0.0012	0.0095 \pm 0.0018
Γ_t	0.0122 \pm 0.0019	0.0150 \pm 0.0029	0.0096 \pm 0.0015	0.0240 \pm 0.0046

Table S1. Model parameters for the psychometric functions ($M \pm SEM$). Since the responses were asymmetric (prosaccade versus antisaccade), we fit the \hat{t}_c for the two rules (C_1 and C_2) separately. However, results indicated that there was not significant difference in \hat{t}_c between the two rules ($P > 0.05$). There was no significant difference between fitted model parameters in inferred and instructed rule experiments ($P > 0.05$).

Parameters	Monkey K	Monkey I
α	0.72 \pm 0.03	0.44 \pm 0.07
$\sigma_{X\Sigma}$	1.56 \pm 0.30	1.09 \pm 0.11

Table. S2. Model parameters for switch probability (M \pm SEM)

	Monkey K		Monkey I	
Area	DMFC	ACC	DMFC	ACC
Stereotactic coordinates	ML 2-6mm AG 0-6 mm	ML 2-5mm AG 0-8 mm	ML 2-6mm AG 0-7 mm	ML 2-5mm AG 0-8 mm
Recording sessions/trials	18/32148	16/27463	14/18008	17/24672
Stimulation sessions/trials	4/5738	4/6362	6/8685	8/10062

Table S3. DMFC/ACC electrophysiology. ML: mediolateral relative to midline; AG: Anterior-posterior relative to the genu of arcuate sulcus. DMFC stimulation pertains to experiments in which we simultaneously recorded from ACC. Fig. S14 shows a coronal section of the two animals' frontal cortex containing a sample recording site in ACC.

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