

1 Thunderstruck: The ACDC model of flexible sequences and rhythms in 2 recurrent neural circuits

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7
8 **Abbreviated title:** Action sequence flexibility

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Abstract

34 Adaptive sequential behavior is a hallmark of human cognition. In particular, humans can learn to
35 produce precise spatiotemporal sequences given a certain context. For instance, musicians can not only
36 reproduce learned action sequences in a context-dependent manner, they can also quickly and flexibly
37 reapply them in any desired tempo or rhythm without overwriting previous learning. Existing neural
38 network models fail to account for these properties. We argue that this limitation emerges from the fact
39 that order information (i.e., the position of the action) and timing (i.e., the moment of response
40 execution) are typically stored in the same neural network weights. Here, we augment a biologically
41 plausible recurrent neural network of cortical dynamics to include a basal ganglia-thalamic module
42 which uses reinforcement learning to dynamically modulate action. This “associative cluster-dependent
43 chain” (ACDC) model modularly stores order and timing information in distinct loci of the network. This
44 feature increases computational power and allows ACDC to display a wide range of temporal properties
45 (e.g., multiple sequences, temporal shifting, rescaling, and compositionality), while still accounting for
46 several behavioral and neurophysiological empirical observations. Finally, we apply this ACDC network
47 to show how it can learn the famous “Thunderstruck” song and then flexibly play it in a “bossa nova”
48 rhythm without further training.

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Introduction

51 Learning and manipulating sequential patterns of motor output are essential for virtually all domains of
52 human behavior. For instance, musicians can learn multiple precise spatiotemporal sequences each with
53 their own rhythm. They can modify the rhythm within each sequence, i.e. speed up or slow down the
54 tempo; or apply different rhythms on a previously learned sequence, i.e. perform a rock song with a
55 bossa nova rhythm. This implies that musicians can quickly and flexibly manipulate action timing in
56 action sequences. Similar capabilities abound in many other domains, such as language production and
57 athletics.

58 Precisely timed action sequences are thought to emerge from dynamical neural patterns of activity. In
59 particular, sparse sequential activity patterns observed in basal ganglia (Jin et al., 2009; Gouvêa et al.,
60 2015; Mello et al., 2015; Bakhurin et al., 2017; Dhawale et al., 2017), hippocampus (Pastalkova et al.,
61 2008; MacDonald et al., 2013; Eichenbaum, 2014) and the cortex (Luczak et al., 2007; Harvey et al.,
62 2012; Remington et al., 2018) are thought to provide a temporal (ordinal) signal for these action
63 sequences to emerge. However, the mechanistic and dynamic principles by which these neural patterns
64 afford sequential flexibility remain unknown. While several neural network models of corticostriatal
65 circuit exist, these are typically applied to single shot stimulus-action pairings rather than sequential
66 choices, despite extensive evidence that basal ganglia is implicated in such sequential behaviors
67 (Graybiel, 1998).

68 In this paper, we sought to develop a biologically plausible neural computational model of cortico-basal
69 ganglia circuitry sufficiently powerful to learn arbitrary sequences (e.g., scales) and easily adjust their
70 timing and expression on the fly. In particular, we aimed for the network to be able to learn multiple
71 arbitrary sequences and to allow for temporal asynchrony, shifting, rescaling, and compositionality. We
72 define these terms more precisely below.

73 Existing neurocomputational models of sequence production can be broadly categorized in three
74 classes, each with their advantages and disadvantages in computational power and their ability to
75 account for behavioral and neural features of action sequences.

- 76 • In associative chain models (also termed synfire chain; e.g., Fiete et al., 2010), activation flows
77 sequentially from one neuron (or neuronal population) to another through feedforward
78 connections (e.g., Cone and Shouval, 2021). The sequence emerges from the hard-wired

79 structure of the chain. Associative chain models naturally produce sequential but also
80 persistent neural activity, both of which are observed empirically (Veliz-Cuba et al., 2015;
81 Pereira and Brunel, 2020). They can also deal with inherent compression of sequential activity,
82 and thereby learn to produce each action in the sequence at any desired precise time (Cone and
83 Shouval, 2021). However, these models are not equipped to facilitate *temporal rescaling*: the
84 finding that learned action sequences can be sped up (compressed) or slowed down (dilated)
85 without the need to overwrite previous learning (Goodbody and Wolpert, 1998; Shmuelof et al.,
86 2012). Indeed, a musician who has learned a novel rhythm can directly speed up or slow down
87 this tempo without any additional learning. Moreover, it is unclear how these models
88 implement *temporal shifting*: the ability to start the action sequence earlier or later in time,
89 without modifying the action sequence structure. Chain models also do not straightforwardly
90 allow networks to encode more than a single sequence, given their hard-wired nature.

91

- 92 • Cluster-based models also involve a chained sequence of activation, but this sequence is learned
93 via cell assemblies (i.e. clusters) that form within a recurrent neural network (RNN) through, for
94 instance, spike timing dependent plasticity (Murray and Escola, 2017; Maes et al., 2020).
95 Depending on the timing of the sequential input to the distinct subsets of the RNN, connectivity
96 may emerge within and between clusters. Once this connectivity matrix is learned, input to the
97 RNN induces a sequential activation whereby activation flows from one cluster to another. In
98 contrast to associative chain models, cluster-based models allow temporal rescaling (Murray
99 and Escola, 2017) while also producing sequential and persistent patterns of activity (Maes et
100 al., 2020). Furthermore, they provide a simple mechanism allowing a network to encode
101 multiple sequences. By selectively activating a specific cluster within the RNN, only the cluster
102 “in line” (i.e., connected to the previous cluster) will be activated sequentially (and so forth).
103 Therefore, the RNN can encode multiple sequential behaviors by learning (and selectively
104 activating) distinct cluster chains encoded in the RNN connectivity matrix (Murray and Escola,
105 2017). Yet, it is unclear how these models could facilitate action sequences with *temporal*
106 *asynchrony*: the ability to learn, and flexibly manipulate, motor sequences with varying inter
107 action intervals (an advantage of associative chain models, see Cone and Shouval, 2021). Indeed,
108 cluster-based models can flexibly manipulate sequences; however, these sequences are typically
109 iso-synchronous (see Murray and Escola, 2017).

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- 111 • State-space models (e.g., Hardy et al., 2018) do not assume a chaining structure at all. Based on
112 a sparsely connected RNN structure, these models are able to learn and reproduce (in the
113 presence of a noise) a neural trajectory represented in high-dimensional space (Sussillo and
114 Abbott, 2009; Laje and Buonomano, 2013; Rajan et al., 2016). This neural trajectory acts as a
115 travelling wave which can then be decoded by downstream neurons to produce sequential
116 orders. State-space models have the ability to learn highly complex and flexible motor
117 sequences. However, unlike the other models, state-space models typically require highly
118 supervised (i.e. continuous teaching signal) and non-biological learning mechanisms. Moreover,
119 they do not provide a potential mechanism for encoding multiple sequences and fail at
120 implementing temporal rescaling (unless resorting to very specific learning regimes, see Hardy
121 et al., 2018).
- 122
- 123 • Finally, none of the models have tackled how a learned sequence at a particular tempo can be
124 executed with a completely different tempo which may have been learned for a different
125 sequence (e.g., applying a bossa nova rhythm to a rock song). We refer to this ability as
126 *temporal compositionality*.

127 In sum, all models can account for distinct functionalities in sequence production, but fail to provide a
128 plausible neurocomputational mechanism from which most fundamental abilities – temporal
129 asynchrony, shifting, rescaling, compositionality – can emerge and interact. These limitations arise from
130 a property common to all action sequence models: action identity, timing and order are represented
131 jointly within the recurrent weights of the network¹. In related sequential decision-making contexts in

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In associative chain models, action timing is principally controlled by the strength of recurrent connections of each excitatory neuron pool, which controls delay times for the next action in line (Cone and Shouval, 2021). Similarly, in cluster models, order and timing information (i.e., when activation “jumps” from one cluster to the other) also depends on the RNN weights (Maes et al., 2020). Finally, state-space models contain both order and time information within the same sets of RNN weights, which define both the neural trajectory and the speed at which the trajectory unfolds (Rajan et al., 2016; Remington et al., 2018).

132 the reinforcement learning domain, such joint coding of task features facilitates only very rigid forms of
133 generalization and transfer, whereas the ability to code task features compositionally facilitates more
134 robust transfer (Franklin and Frank, 2018) that can better account for human behavior (Franklin and
135 Frank, 2020). However, the mechanisms for such compositionality in neural networks remains unknown.

136 Here, we develop a biologically plausible RNN called the associative cluster-dependent chain (ACDC)
137 model. By combining strengths of the associative chain and cluster-based models, ACDC accounts for
138 biological data. To increase computational flexibility, the network factorizes order and timing
139 information by storing them separately in a premotor cortical RNN which is dynamically gated by a basal
140 ganglia-thalamus module. This modularity thereby affords independent (and flexible) manipulation of
141 sequence order and action timing. For instance, once an action sequence has been learned, temporal
142 rescaling can be accomplished by targeting the locus representing time, while still allowing the network
143 to produce the same desired action order sequence.

144 In the remainder of the paper we first present the architecture of the model, and ground it within the
145 context of neurophysiological observations on the premotor cortex (PMC) and the BG. Second, we
146 describe how the model *learns* to produce precisely timed, temporally asynchronous, action sequences.
147 Third, we demonstrate how the mechanistic properties of the model can account for temporal
148 properties: temporal shifting, rescaling, compositionality, and sustained motor activation. Fourth, we
149 simulate both empirical and neurophysiological observations. Finally, we discuss the characteristics and
150 abilities of the ACDC model in light of neurophysiological evidence and alternative neurocomputational
151 models.

152 **Methods**

153 ***The associative cluster-dependent chain (ACDC) model for flexible motor timing***

154 In this section, we provide the reader with an intuitive functioning of the ACDC model in simplified and
155 more detailed forms (Fig. 1); we refer to Appendix A for a comprehensive mathematical formulation. A
156 context module encodes the sequences to be executed (e.g., which song is to be played), and is
157 provided as input to an RNN, which learns to encode sequence order via Hebbian learning. Order is
158 encoded as a sequence of attractor states represented by persistent activation in distinct excitatory RNN
159 unit clusters. Cell assemblies (or clusters) learn to project to the appropriate action identity (again via
160 Hebbian learning), represented topographically in the BG. In turn, the BG project to the corresponding

161 motor thalamus to control action execution. To optimize precise action timing, the weights between
162 action identity and execution are learned via supervised learning (i.e. Delta rule), perhaps summarizing
163 the role of cerebellum in error corrective learning. Finally, thalamic activity about the executed action is
164 fed back to the RNN, ultimately creating a cortico-basal ganglia loop. Each loop subtends the
165 appropriate action order, identity and timing execution, allowing precisely timed action sequences to
166 unfold.

167 In a more detailed manner, our ACDC model contains four main modules (Fig. 1 right panel): an input
168 layer (Fig. 1A), an RNN (representing premotor cortex; Fig. 1B) and a BG-thalamus unit (Fig. 1C).

169 The input layer (Fig. 1A) consists of a vector of neurons, of which a subset is activated, representing
170 sensory or other context that would signal the identity of the sequence to be produced or learned.

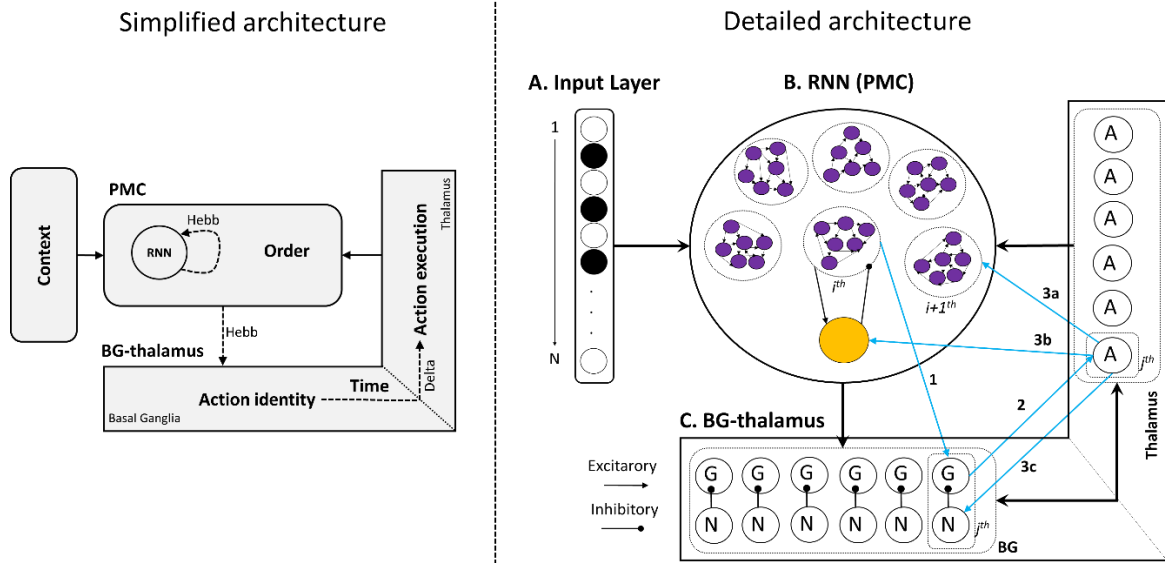
171 Crucially, the dynamics within the ACDC model evolve as a sequential unfolding of RNN-BG-thalamus-
172 RNN (i.e., cortico-basal ganglia) loops, depicted by the light blue arrows in figure 1. The sequence starts
173 with the activation of a cluster (i.e., densely interconnected) of excitatory RNN neurons (Fig. 1B). Each
174 cluster will come to encode the i^{th} element in the action sequence. As opposed to single unit, clustered
175 neurons provide a biologically plausible mechanism for supporting persistent activation within the
176 cluster given a phasic input (i.e., an attractor; Amit, 1988; Durstewitz et al., 2000). In prefrontal cortical –
177 BG models, such clusters are referred to as “stripes” based on their anatomical existence, and are
178 independently gated by BG (O’Reilly and Frank, 2006). Once a cluster is activated, the RNN temporarily
179 settles on an attractor state indicating the ordinal position (order or rank) in the sequence, analogous to
180 how distinct PFC stripes code for ordinal positions in phonological loop tasks(O’Reilly and Frank, 2006).
181 However, in ACDC such clusters emerge naturally via learning rather than hard-coded anatomical
182 entities. Moreover, attractor states are maintained via E-I balance: each excitatory neuron projects to a
183 common single inhibitory neuron (orange circle in Fig. 1B) which reciprocally inhibits all excitatory RNN
184 neurons. As long as the E-I balance is not perturbed by another input (see below), activation in the
185 cluster will persist and the RNN will continue representing the i^{th} order in the sequence.

186 In turn, each excitatory RNN cluster projects to its corresponding “Go” unit in the BG (blue arrow 1 from
187 i^{th} cluster in Fig. 1B to G node in Fig. 1C), and each Go cell accumulates evidence for the j^{th} action
188 associated to the j^{th} order (see Frank, 2006 and Ratcliff and Frank, 2012 for related computational
189 models of evidence accumulation in these units, and Doi et al., 2020 for empirical data). Striatal Go cells,
190 via the basal ganglia direct pathway machinery (Alexander and Crutcher, 1990; Mink, 1996), facilitate

191 response execution by projecting towards the corresponding motor thalamus neurons, from here on
192 termed Action nodes for simplicity (blue arrow 2 from Go to Action nodes in Fig. 1C).

193 Action nodes possess a negative bias, which acts as a decision threshold, i.e., the net input needs to
194 exceed this bias in order for action to be executed. This feature again summarizes the computational
195 role of the output of the BG, which serves to inhibit action execution until sufficient evidence reaches
196 the threshold for action gating (Frank, 2006; Wiecki and Frank, 2013; see also Lo and Wang, 2006).
197 Therefore, the weight values between Go and Action nodes control the speed of action execution: the
198 BG encode the rhythm. Action execution can be expressed either as a transient or persistent response
199 (see simulations; Pereira and Brunel, 2020).

200 In turn, Action nodes project excitatory connections to three distinct parts of the network
201 simultaneously. First, Action nodes project to the cluster of excitatory neurons in the RNN representing
202 the $i+1^{th}$ order in the sequences (blue arrow 3a in Fig. 1). Second, Action nodes project to the inhibitory
203 shared neuron (blue arrow 3b to orange node in Fig. 1), that in turn globally inhibits all the clusters in
204 the RNN. In this manner, thalamic Action nodes can update the cortical representation by separately
205 projecting to both inhibitory and excitatory neurons (Schmitt et al., 2017; Rikhye et al., 2018), enabling
206 the RNN to transition from the current state to the next. That is, the activation of action nodes perturbs
207 the E-I RNN balance in a way that allows the i^{th} cluster to shut down and the $i+1^{th}$ cluster to be expressed
208 (see Appendix A for further details). Third, Action nodes project excitatory connections back to their
209 corresponding No Go cells (blue arrow 3c from j^{th} Action node in the thalamus to j^{th} No Go node in the
210 BG, see Fig. 1C). In turn, No Go cells strongly inhibit their corresponding Go cells (Taverna et al., 2008;
211 Wiecki and Frank, 2013; Dobbs et al., 2016), thereby shutting down evidence in favor of the j^{th} action,
212 and hence stopping the execution of the j^{th} action. This loop is then reproduced with the j^{th+1} RNN cluster
213 and j^{th+1} G-A-N triplet in the BG-thalamus unit, and so forth until the action sequence is performed in its
214 entirety.



215

216 **Figure 1. Left panel: Simplified ACDC model architecture.** An input context layer indicates which
 217 sequence needs to be learned or executed. The premotor cortex (PMC) is subtended by a RNN that
 218 learns (via Hebbian learning) to form clusters of excitatory neurons encoding order in the sequence, and
 219 which are regulated by an inhibitory pool. In turn, each cluster learns to trigger action plans,
 220 topographically represented in the BG. Specific actions are executed in the thalamus at specific times
 221 based on learned connections from BG to thalamus. Motor activity is then fed back to the RNN, closing
 222 the cortico-basal ganglia loop. The unfolding of several iterations of this loop is responsible for the
 223 execution of precisely timed action sequences. **Right panel: ACDC full model architecture. A. Input**
 224 **layer:** codes for contexts indicating the sequence to be learned/produced in a N length binary vector. **B.**
 225 **RNN:** represents recurrently interconnected neurons of the PMC, composed of a subset of
 226 interconnected neurons (i.e., clusters) that can give rise to sequential activation states after learning via
 227 cortico-basal ganglia loops. All excitatory nodes in the RNN project to a shared inhibitory neuron (orange
 228 node), which in turn inhibits all excitatory neurons (purple nodes; shown for just one cluster for visual
 229 simplicity). **C. The BG:** composed of two neuron types G (Go cells) and N (No Go cells). Go nodes
 230 accumulate evidence over time and excite Action (A) nodes in the BG output /thalamus layer. Once
 231 activity in the Go nodes reaches a specific threshold, the corresponding action is executed. Once
 232 executed, Action nodes reciprocally activate No Go nodes which in turn suppress Go nodes, shutting
 233 down action execution. **The thalamus:** is composed of Action nodes whose activity represents action
 234 execution. The j^{th} Action node selectively projects excitatory connections to the i^{th+1} cluster in the RNN,
 235 the shared inhibitory neuron and the j^{th} No Go node in the BG. Light blue arrows represent the i^{th}
 236 cortico-basal ganglia loop instance.

237 Several features of the model should be highlighted. First, each cluster activation within the RNN acts as
 238 an attractor state representing the i^{th} order in the sequence. Interestingly, cells in the monkey PMC code
 239 for the position in sequence, regardless of the actual movement produced during that position (Clower
 240 and Alexander, 1998; Shima and Tanji, 2000; Isoda and Tanji, 2003, 2004; Averbeck et al., 2006;
 241 Berdyeva and Olson, 2009; Salinas, 2009). We therefore assume that the neurons forming each cluster

242 represent rank-order-selective neurons whose activation unfolds sequentially: the RNN encodes order
243 information.

244 Second, the speed at which each action is executed is driven by how quickly the evidence in the Go
245 nodes of the BG can cross the decision threshold in the Action nodes: the BG encode time information.
246 Indeed, several studies suggest that temporal processing is subtended by the BG in the (non)human
247 primates and rodent brain (Jin et al., 2009; Schwartz et al., 2011; Gershman et al., 2014; Jones and
248 Jahanshahi, 2014; Mello et al., 2015; Thura and Cisek, 2017; Paton and Buonomano, 2018). Note that
249 there are multiple routes by which timing can be altered within Go nodes in our model: (i) the learned
250 weight value between Go and Action nodes; (ii) a bias input to Go nodes (in addition to that coming
251 from the RNN cluster); and (iii) a multiplicative gain on Go unit activity (see model simulations). As
252 shown below, these separate routes will become important for providing timing and rhythm flexibility.

253 Third, as in many cortico-BG models (e.g., Gurney et al., 2001; Frank, 2006), and motivated by
254 anatomical data (Alexander et al., 1986) our model is characterized by topographical organization of
255 actions across the BG circuit and its outputs (i.e., indexed in our model by the subscript j associated in
256 the G-A-N triplet projections). Recent evidence further confirms topographical action representations in
257 BG-thalamocortical loops (Oh et al., 2014; Hintiryan et al., 2016; Hunnicutt et al., 2016), whereby causal
258 activation of specific subregions is related to specific output behaviors (Peters et al., 2021), and is also
259 supported by human neuroimaging (Gerardin et al., 2003) and monkey/rodent neurophysiology studies
260 (McHaffie et al., 2005; Jin et al., 2009; Znamenskiy and Zador, 2013; Friedman et al., 2015; Gremel et al.,
261 2016; Hooks et al., 2018; Lee et al., 2020). However, in contrast to previous models in which BG gating
262 affords action selection of the corresponding cortical action, in the ACDC model BG gating triggers a
263 cortical dynamical state that initiates the evolution of the *subsequent* item in the sequence.

264 Fourth, we clarify how the ACDC model combines properties of associative chain and cluster-based
265 models. While the ACDC model does initiate a *chain* via sequential propagation across cortico-BG loops,
266 the timing of such transitions is controlled by learning the weights within the BG-thalamus unit, and
267 moreover, what is learned are transitions between clusters of excitatory RNN neurons representing
268 order in the sequence (Maes et al., 2020). Hence, the ACDC model makes use of two distinct
269 conceptualizations of sequence learning, to achieve greater computational flexibility (as demonstrated
270 in the result section).

271 ***Learning in the ACDC model: Hebbian learning for order and Delta rule for time***

272 Learning in the ACDC model takes place in three distinct loci of the network, comprising Hebbian
273 learning for sequence transitions and error-driven learning for precise timing.

274 First, as previously mentioned, order is coded via persistent activation within clusters of the RNN.
275 However, in contrast to pure associative chain models, the ACDC does not assume any feedforward
276 hard-wired structure, but rather learns it. Selective time-dependent inputs to the RNN (i.e., from the
277 input layer and thalamic Action nodes) activate a subset of neurons within the RNN, which get clustered
278 together through dynamic synaptic weights:

$$279 \quad \frac{dW_{ij}}{dt} = -\alpha_1((1-x_i)\bar{x}_j) + \alpha_2(x_i\bar{x}_j(W_{\max} - W_{ij})) \quad \text{equation 1}$$

280 where \bar{x}_j is presynaptic activity low-pass filtered over a time scale τ_w ; x_i is postsynaptic activity; α_1 and
281 α_2 are learning rate parameters. When \bar{x}_j and x_i are both simultaneously > 0 , W_{ij} goes to W_{\max} ;
282 otherwise W_{ij} goes to 0. Note that $\bar{x}_j(t)$ will be non-zero if unit j is active within the time window from t
283 $-\tau_w \rightarrow t$ (as in Murray and Escola, 2017).

284 Second, Equation 1 is also used to learn connections between the RNN and the Go nodes of the BG
285 module; here, pre- and postsynaptic activity refer respectively to RNN excitatory unit activity and Go
286 nodes activity (weight values between RNN units and Go nodes are randomly initialized from a Gaussian
287 distribution with mean = $0.5 / N$ and s.d. = $0.1 / N$, where N is the number of RNN excitatory units).

288 Third, action specific execution time is coded in the weights connecting Go and Action nodes. Here, we
289 describe time learning as a delta rule, whereby an agent receives a supervisory signal explicitly indicating
290 whether a specific action has been produced before (positively signed signal to increase weights) or
291 after (negatively signed signal to decrease weights) the appropriate time, as described in equation 2:

$$292 \quad \Delta W = \eta(t_{\text{observed}} - t_{\text{desired}}) \quad \text{equation 2}$$

293 where the change in weight (ΔW) between the j^{th} Go and Action nodes is driven by the learning rate η ,
294 and the error computed as the difference between the observed and desired response time (t) for each
295 action. Weight values between Go and Action nodes are randomly initialized and drawn from a random
296 Gaussian distribution (mean = 2, s.d. = 0.2). Learning of precisely timed sequences is shaped sequentially
297 (i.e., in chunks): the model first learns to produce the first action at the appropriate time (i.e. until the
298 error $< \phi$ and ϕ is a low value, see table 1 in Appendix A), then the second, and so forth. Note that

299 learning by chunking improves motor execution (Wymbs et al., 2012; Boutin et al., 2013; Lungu et al.,
300 2014; Graybiel and Grafton, 2015; Doyon et al., 2017), and chunk-based representation is at the base of
301 several theoretical models of motor sequence learning (Abrahamse et al., 2013; Verwey et al., 2014;
302 Diedrichsen and Kornysheva, 2015).

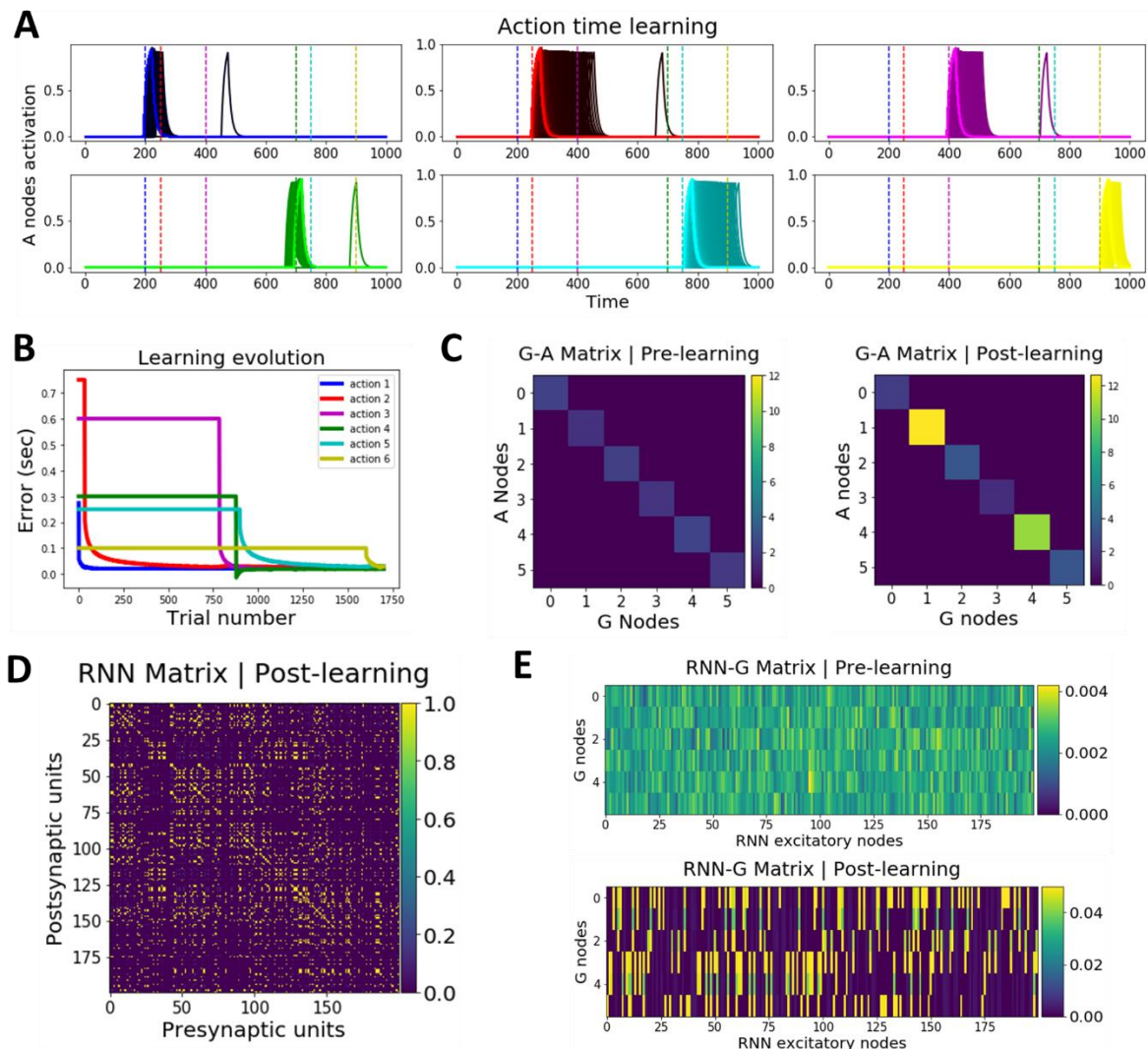
303 Below we describe all the simulations emerging from the ACDC model; parameter values for all
304 simulations are reported in table 1 of Appendix A, and simulation code is available from
305 <https://github.com/CristianBucCalderon/ACDC>. We start by describing how the model can learn to
306 produce precise spatiotemporal sequences. We then simulate all the temporal properties of the model:
307 reproduction of an action sequence with temporal asynchrony, temporal shifting, rescaling, and
308 compositionality, and sustained motor activation. Finally, we describe and simulate empirical and
309 neurophysiological observations.

310 **Results**

311 ***Learning precise spatiotemporal sequences***

312 Figure 2 shows the result of the first simulation, where the ACDC model learns to produce a precisely
313 timed, temporally asynchronous, action sequence. For the purpose of clarity, we limit the sequence to 6
314 actions. The goal of the model in this simulation is to produce each action sequentially at the
315 appropriate time, i.e. action 1 through 6 have to be executed respectively at times 200, 250, 400, 700,
316 750 and 900 ms (within a 1 second window). Note that this is an arbitrarily chosen timing sequence; the
317 model can (learn to) produce any timed, synchronous (see Thunderstruck simulation below) or
318 asynchronous, sequence. Figure 2A shows how the activity of each Action node goes down the gradient
319 and progressively reaches the optimal time (depicted by color coded vertical dashed lines), reflected in a
320 decrease in the action timing error (Fig. 2B) and in the weight changes between Go and Action nodes
321 (Fig. 2C).

322 Figure 2D depicts the RNN connectivity matrix after learning (weights are zero before learning).
323 Excitatory projections to the RNN from the input and motor layer are pseudo-random, with the
324 restriction that two different projections never excite the same RNN neuron. These pseudo-random
325 projections make it hard to visually identify the presence of clusters in fig2D; importantly however, this
326 connectivity matrix does induce clustered dynamics (see Video 2 and Fig. 7A below). Finally, figure 2E
327 shows how the i^{th} cluster in the RNN learns to be (almost) selectively wired with the j^{th} Go node.



328

329 **Figure 2. ACDC's learning dynamics. A. Learning a precisely timed action sequence.** Each action
 330 execution (A node activation) is progressively shifted towards the optimal action time (depicted by the
 331 color coded vertical dashed line; x-axis represents time). Learning progresses from darker to brightest
 332 colors. **B. Learning evolution.** Color coded traces represent the evolution of the error as a function of
 333 trial number for each action in the sequence. Learning unfolds sequentially, whereby timing errors are t
 334 minimized for the first action before the second action starts learning. Therefore, each action (except
 335 action 1) starts off with a plateaued error level until the preceding action reaches the optimal time.
 336 Some action timings are learned faster than others because their optimal time weight value is closer to
 337 their initial value. The error is computed by subtracting the observed from the desired response time
 338 and plotted in seconds. **C. BG weights encode time.** Action timing is learned by changing the weights
 339 from BG Go nodes to thalamus Action nodes. The left and right panel show respectively the weights
 340 values before and after learning. For instance, the second action (red trace in B) starts off being
 341 produced too slowly. Hence, weights increase until they produce the optimal action time for action 2.
 342 Color bars indicate weight values. **D. RNN connectivity matrix after learning.** The RNN connectivity
 343 matrix is initialized as a blank slate (all values are set to 0). After learning, the RNN connectivity matrix

344 displays the appearance of clusters, whereby groups of 20 neurons are fully interconnected with each
345 other and not connected with other neurons in the RNN (please refer to Video 2 and Fig. 7A for better
346 visualization of clusters and their transitions as the sequence unfolds). Color bar represents weight
347 values. **E. RNN i^{th} cluster learns to project to j^{th} Go node.** The top panel shows the randomly initialized
348 weight values between the RNN excitatory units (before learning). The bottom panel shows how each
349 cluster (represented by a subset of RNN neurons) is connected to a specific Go node after learning. Color
350 bars represent weight values.
351

352 ***Temporal flexibility properties of the ACDC model***

353 Having established learned clusters within the RNN and learned sequences in the ACDC model, we now
354 focus on the flexibility properties of the model after learning, without having to overwrite learned
355 weights. First, we show that a previously learnt action sequence with *temporal asynchrony* can be
356 flexibly reproduced. Second, we show that this sequence can be initiated earlier or later in time; we call
357 this property *temporal shifting*. Third, we demonstrate how action sequences can be compressed or
358 dilated, i.e., *temporal rescaling*. Fourth, we show how a given ordered sequence can be produced with a
359 completely different tempo, a property that we refer to as *temporal compositionality*. Fifth, we describe
360 how the model can also output sustained action execution. Finally, we show how the ACDC model can
361 learn (a part of) the Thunderstruck song, which is then flexibly played on a bossa nova tempo; thereby
362 recapitulating the temporal flexibility properties.

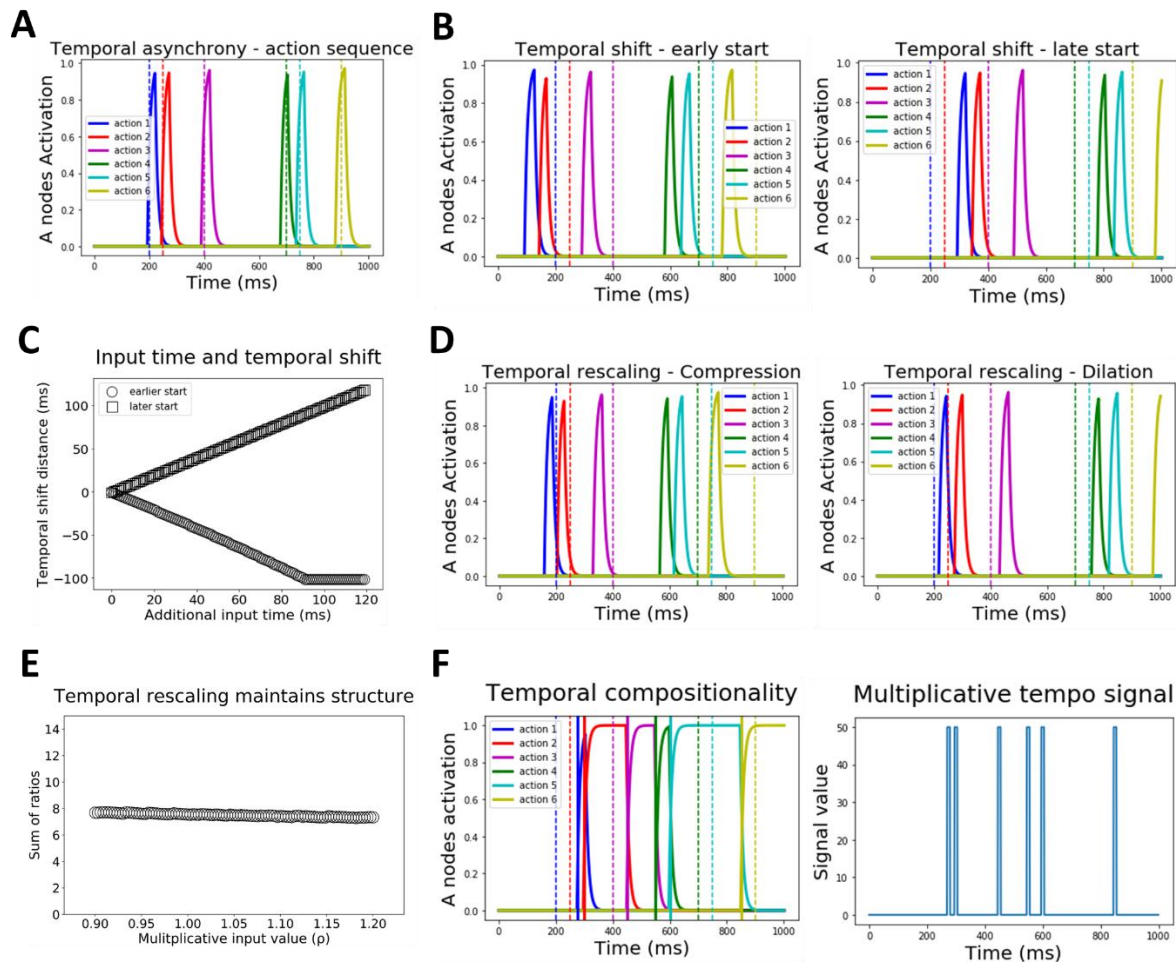
363 **Simulation 2: Reproduction of previously learnt action sequence displaying temporal asynchrony.** In
364 simulation 1, we demonstrated that the ACDC model can learn precisely timed, temporally
365 asynchronous, action sequences. In simulation 2, we now clamp learning (i.e., freeze the weights). We
366 provide the network with the same input and observe that the network can reproduce the sequence
367 maintaining its precision in action timing. Figure 3A shows that, given the clamped set of learned
368 weights, each Action node (color coded for order) within the thalamus layer gets activated at the
369 previously learnt precise timing, in a phasic/transient fashion.

370 **Simulation 3: temporal shifting.** The previous action sequence can be shifted in time, i.e., started earlier
371 or later. Importantly, this shift can occur without changing the timing between actions (i.e., sequence
372 timing is preserved). The ACDC model achieves flexible temporal shifting by either adding an additional
373 positive (to start the sequence earlier) or negative (to start it later) input to the first Go node of the
374 sequence, analogous to the top-down input from pre-SMA to striatum thought to bias starting points for
375 evidence accumulation (Forstmann et al., 2008; although similar effects could be implemented by

376 dopaminergic modulation; see Discussion). In simulation 3, we inject an additional input of +1 or -1 to
377 the first Go node during the first 100 ms of the 1 second time window. Figure 3B shows how the
378 sequence is shifted earlier in time for the positive input (left panel) and later in time for the negative
379 input (right panel). Moreover, figure 3C shows that as this additional input lasts longer, the distance (in
380 time) between the first action of the shifted sequence and that of the original sequence increases
381 linearly. Naturally, intrinsic temporal constraints of the model limit this distance for earlier shifts, as
382 shown by the negative plateau in figure 3C (black circles).

383 **Simulation 4: Temporal rescaling.** Musicians possess the ability to learn a rhythm, i.e., a precisely timed
384 action sequence, and instantly temporally rescale (compress or dilate) that rhythm without additional
385 learning. In our model, flexible rescaling is achieved by sending a multiplicative input (ρ) to all Go nodes
386 simultaneously; if $\rho > 1$ or $0 < \rho < 1$ the sequence is respectively compressed or dilated. Figure 3D shows
387 temporal rescaling for ρ values of 1.2 (compression, left panel in Fig. 3D) and 0.9 (dilation, right panel in
388 Fig. 3D). Importantly, temporally rescaling the sequence does not affect the temporal structure of action
389 sequences. For 100 values of ρ , ranging from 0.9 to 1.2, we computed the relative ratio between a
390 sequence of 3 actions. The ratio was computed by subtracting the time of action 1 from that of action 2
391 (subtraction 1), then the time of action 2 from that of action 3 (subtraction 2), and dividing subtraction 2
392 / subtraction 1. We performed this computation for the action triplets 1-2-3, 2-3-4, 3-4-5 and 4-5-6, and
393 summed the ratios. Figure 3E shows that this sum of ratios stays constant (mean = 7.5, s.d. = 0.12),
394 thereby indicating that temporal structure is maintained albeit rescaled.

395 **Simulation 5: Temporal compositionality.** Musicians must also be capable of temporal compositionality;
396 that is, apply a different tempo to an action sequence that was learned in a different tempo (e.g., apply
397 a bossa nova tempo to a rock song; see below). In simulation 5, we assume that the brain can extract
398 and store a tempo, which then can be used as a dynamical multiplicative signal to all Go nodes as in
399 simulation 4. In simulation 5, we apply a dynamical multiplicative signal (Fig. 3F right panel) to the Go
400 nodes. The result is to produce the learned sequence (described in Fig. 3A) to the tempo described by
401 the multiplicative signal. Figure 3F (left panel) shows how the time of each action in the sequence does
402 not fall on the previous tempo (color coded vertical dashed lines), but now rather is produced at the
403 novel timing (vertical solid lines).



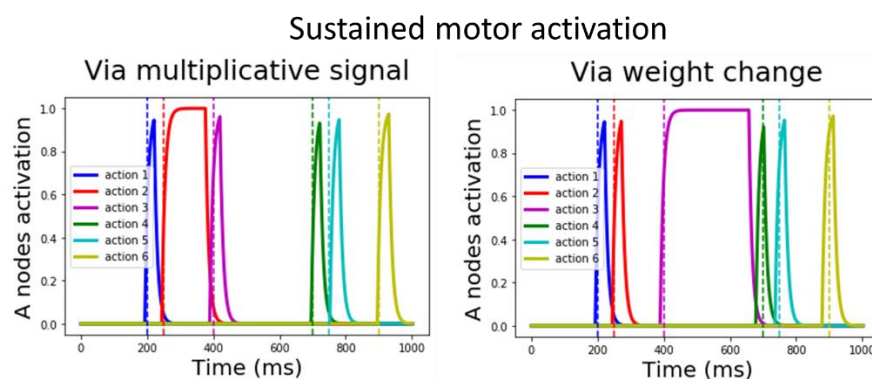
404

405 **Figure 3. Temporal properties of the ACDC model. A. Simulation 2: Reproduction of action sequence**
 406 **with temporal asynchrony.** Each action (i.e. A node activation, color coded) is produced at the precise
 407 desired time indicated by the vertical dashed line (also color coded), within a 1 second time window.
 408 Inter-action interval varies as the sequence unfolds. **B. Simulation 3: Temporal shifting.** A precisely
 409 timed action sequence can be started earlier (left panel) or later (right panel) by respectively injecting an
 410 additional positive or negative input to the first G node (i.e. associated to accumulating evidence in
 411 favor of the first action). Importantly, the temporal structure of the action sequence is not altered. **C.**
 412 **Simulation 3: Temporal shifting varies linearly with additional input time.** Applying longer input times
 413 leads to increasingly earlier or later shifts in sequence initiation times, depending on whether additional
 414 input is positive (circles) or negative (squares). **D. Simulation 4: Temporal rescaling.** Action sequences
 415 can be compressed (left panel) or dilated (right panel) by adding a multiplicative input to all G nodes
 416 simultaneously. **E. Simulation 4: Temporal rescaling preserves action sequence structure.** Importantly,
 417 when temporal rescaling is applied to the action sequence, the relative timing between each action (i.e.
 418 the structure) is preserved. Here, we plot the sum of ratios (y-axis, see main text) as a function of the
 419 multiplicative input ρ (x-axis). The sum of ratios value (black circles) stays constant as a function of ρ ,
 420 indicating a preserved temporal structure even though the sequence is rescaled. **F. Simulation 5:**
 421 **Temporal compositionality.** The left panel shows how A nodes activity are activated on the tempo
 422 described by the multiplicative signal (left panel). Vertical dashed and solid lines on the left panel

423 indicate the timing of each action for the previous and novel tempo respectively. As shown, the
424 respective A nodes become active on the novel tempo.

425

426 **Simulation 6: Sustained motor activation.** The ACDC model is also capable of producing sustained
427 motor activation for any element within the sequence, for instance sustained notes in a musical scale.
428 Our model can achieve sustained motor activation via two mechanisms. First, via a flexible mechanism
429 similar to that of rhythm compositionality, a multiplicative signal ($\rho = 0.1$) is sent to the Go node during
430 the period in which sustained motor activation is needed. On the left panel of figure 4, we show the
431 results of applying such a signal during the period between the start of the second action and the
432 beginning of the third one. The motor activation of the second Action node (red trace) is sustained until
433 the third action is executed (purple trace). Second, via a learning mechanism, the weight value between
434 a specific Action-No Go nodes pair can be decreased to induce sustained activation of the Action node.
435 On the right panel of figure 4, we decreased the weight value connecting the third Action-No go nodes
436 pair. Such a weight change produced a similar result to that of implementing a multiplicative input to
437 the No Go node, i.e. sustained activation of the corresponding Action node.

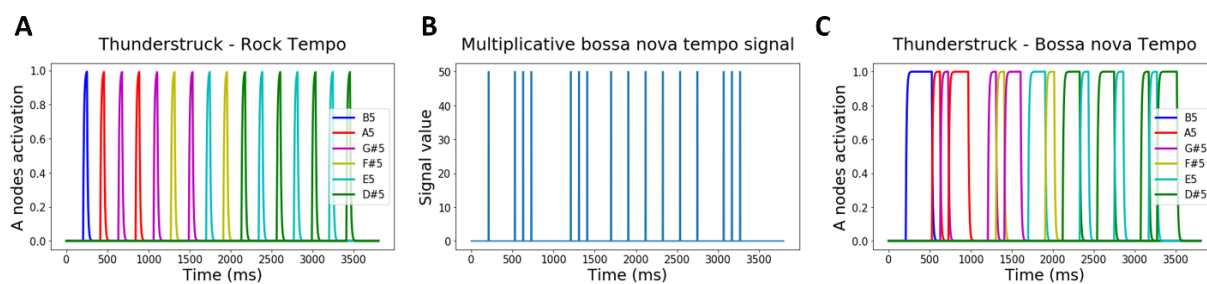


438

439 **Figure 4. Simulation 6: Sustained motor activation.** Both panels demonstrate that the ACDC model is
440 able to output sustained motor activation as desired within a sequence. The left panel shows the results
441 of applying a multiplicative signal ($\rho = 0.1$) to the second No Go node, inducing a sustained activation of
442 the second action (red trace). The right panel shows a similar effect this time by decreasing the value of
443 the Action-No Go connection of the third action, in turn inducing sustained activation of the third Action
444 node (purple trace).

445 **Simulation 7: The ACDC model in action and sound.** Here, the ACDC model learns to produce the
446 second guitar riff of ACDC's (the rock group) Thunderstruck song. This riff is composed of 16 actions
447 hitting six different notes (B5, A5, G#5, F#5, E5, D#5) following an isosynchronous rock tempo (Fig. 5A).
448 By allowing the model to record each note corresponding to each sequential action (following Fig. 5A),

449 the ACDC model was able to musically reproduce the riff ([Audio file 1](#)). Notably, [video 1](#) shows that the
450 RNN dynamics (during the song) represent sequential attractor states, encoding order and leading to the
451 production of each action (and sound) in the sequence (for a slowed down demonstration of similar
452 dynamics with a less complex action sequence see Video 2 below). Next, we imposed the ACDC model to
453 play the riff but now based on a bossa nova tempo without further training (Fig. 5B). The ACDC model
454 was able to flexibly reproduce the riff following the bossa nova tempo (Fig. 5C and Audio file 2). The
455 model thereby displays the ability to produce complex temporal compositionality. Further note that,
456 altogether, this simulation encapsulates distinct temporal flexibility properties. First, flexibly
457 reproducing the Thunderstruck song following a bossa nova tempo requires the ability to generate an
458 action sequence with temporal asynchrony. Second, temporal rescaling is applied to parts of the song as
459 the sequential execution of consecutive notes need to be sped up or slowed down. Third, the model
460 displays its ability to produce sustained motor activation (see [Audio file 2](#)).



461

462 **Figure 5. Simulation 7: the ACDC produces the Thunderstruck song. A. Second guitar riff from ACDC's**
463 **(the group) Thunderstruck song.** The riff is composed of 16 sequential actions creating a isosynchronous
464 rock rhythm over a window of 3500 ms (given a 140 bpm tempo). Each action is associated to a color
465 coded note). **B. Generic bossa nova tempo.** We imposed the model to replay the thunderstruck rock
466 tempo song following a bossa nova rhythm whose tempo is described by the blue trace multiplicative
467 signal. **C. Flexible generation of the Thunderstruck song following a bossa Nova tempo.** When the
468 multiplicative input (Fig. 3B) is given to the Go nodes of the BG, the ACDC model flexibly reproduces the
469 Thunderstruck song but now following the bossa nova tempo.

470

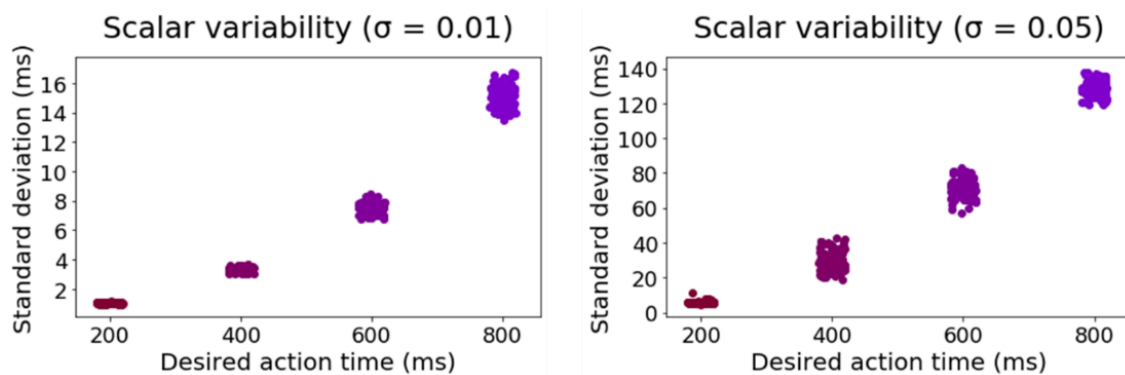
471 --Insert Video 1--

472 **Video 1. Simulation 7: Dynamical visualization of RNN and Action nodes activity coupled with**
473 **simulation-based Thunderstruck song sound.** The top left panel shows how RNN sequential and
474 persistent activity unfolds as a function of time. The bottom left panel is a visualization of RNN dynamics
475 as a neural trajectory in PC space. The neural trajectory displays a pattern of sequential attractor states.
476 The right panel displays how activity in each Action node (and hence Thunderstruck song note) is
477 executed at the learned action time.

478

479 **Behavioral and neurophysiological simulations**

480 **Simulation 8: Behavioral simulation.** In the motor timing literature, a ubiquitous finding is *scalar*
481 *variability*: when asked to produce an action after a specific time interval, the variability in action
482 execution timing increases with the length of interval timing (Ivry and Hazeltine, 1995; Rakitin et al.,
483 1998; Jazayeri and Shadlen, 2010; Acerbi et al., 2012). In simulation 8, our model learns to produce a
484 single action at distinct interval timings (i.e. 200, 400, 600 and 800 ms). For each timing, the model
485 produces 500 reaction times (RTs), from which we extract the standard deviation (SD), and reproduce
486 this process for 100 simulations and two noise values (i.e. gaussian random noise with zero mean and SD
487 of 0.01 or 0.05 is added to the model equations 3-5 and 7). As predicted by empirical work, figure 6
488 shows how the SD of RTs increases as a function of interval timing for both noise values, and thereby
489 demonstrates that the ACDC model displays scalar variability (see also Egger et al., 2020). Furthermore,
490 the SD value range also increases with noise values. This effect is explained in our model by having a
491 fixed negative bias on the Action nodes in the motor layer. Such a feature reduces to having an
492 accumulation-to-bound process for action execution. Hence, given a specific amount of noise, longer
493 RTs are associated to wider RT distributions (i.e. larger SD, Ratcliff and Rouder, 1998). The underlying
494 reason is that the effect of noise on evidence accumulation is amplified as time elapses.

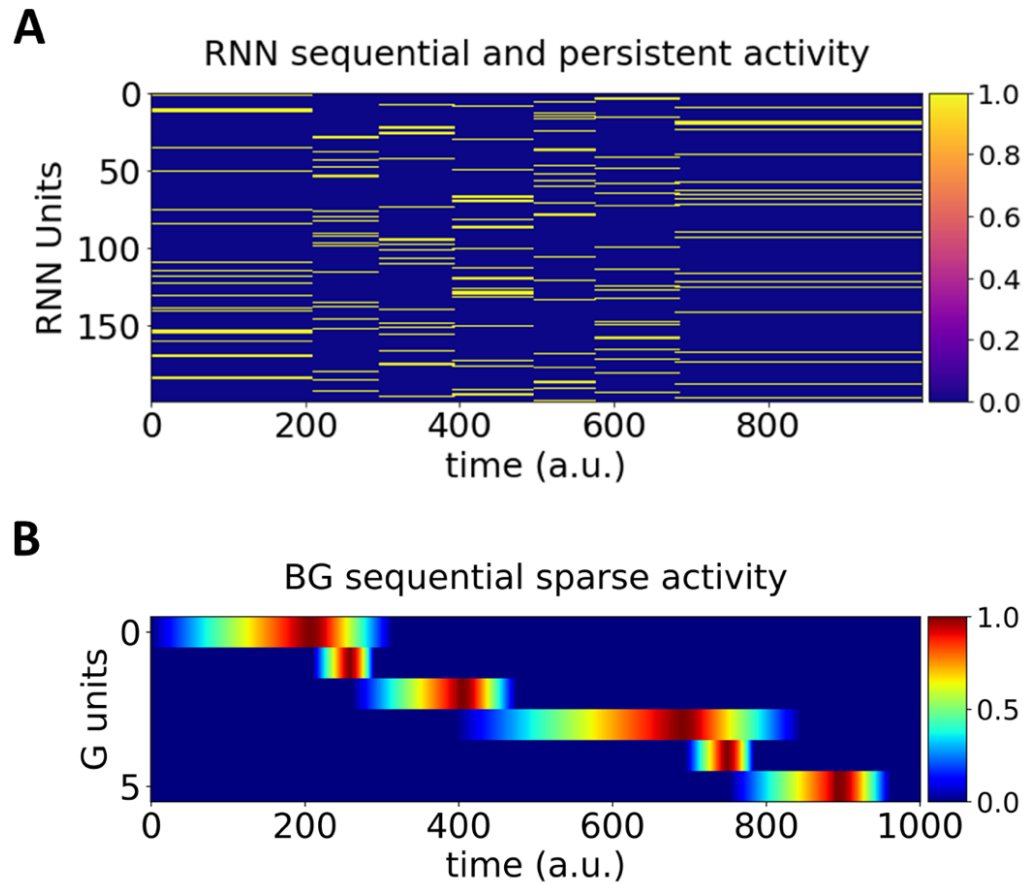


496 **Figure 6. Simulation 8: The ACDC model displays scalar variability.** Left (low noise value = 0.01) and
497 right (high noise value = 0.05) panels show that the standard deviation of RTs increases as a function of
498 the desired action time (i.e. interval timing). Moreover, higher noise values increase the range of
499 standard deviation. Each dot is the result of 1 out of 100 simulations for each interval timing.

500 **Simulation 9: Neurophysiological simulations.** Two other ubiquitous findings are persistent and
501 sequential neural activity. First, several studies have observed persistent neuronal firing rates in
502 temporal (Miyashita and Chang, 1988; Nakamura and Kubota, 1995; Erickson and Desimone, 1999),
503 parietal (Koch and Fuster, 1989; Chafee and Goldman-Rakic, 1998; Gail and Andersen, 2006; Klaes et al.,

504 2011), premotor (Cisek and Kalaska, 2005) and prefrontal (Funahashi et al., 1989, 1990; Miller et al.,
505 1996) cortices whenever an agent has to hold in working memory task-relevant stimulus features (e.g.,
506 spatial location). Theoretical work suggests that persistent activation patterns emerge from recurrently
507 connected networks that settle in one of multiple potential attractor state (Durstewitz et al., 2000;
508 Wang, 2001; Brunel, 2003). Second, as motivated in the introduction, sequential activity has also been
509 observed in distinct sequential behaviors such as spatial navigation (Eichenbaum, 2014) and bird song
510 (Hahnloser et al., 2002; Kozhevnikov and Fee, 2007; Amador et al., 2013; Okubo et al., 2015).

511 Interestingly, recent work suggests that sequential switches in attractor states (and hence persistent
512 neural activity), are associated to the timing of behavioral switches in action sequences (Recanatesi et
513 al., 2020). Therefore, persistent and sequential activity may emerge from the same mechanism. In our
514 model, the RNN activation dynamics display such switches from one attractor to another as the action
515 sequence unfolds. Each attractor state is associated to the persistent activity of neurons forming a
516 cluster in the PMC (RNN). When the action associated to that attractor state (i.e. the j^{th} action
517 associated to the j^{th} order) is executed, this triggers a switch in attractor state in the RNN (via cortico-
518 cortical projections from M1 to PMC), as empirically observed by Recanatesi et. al (2020). In simulation
519 1, the ACDC learns to produce an arbitrary sequence of 6 actions, each with their own desired execution
520 time within a window of 1 sec (i.e. at 200, 250, 400, 700, 750, 900 ms). Figure 7A shows the RNN
521 dynamics after learning. Each cluster of activation displays persistent neural activity until the action is
522 executed, which triggers the following cluster of persistent neural activity. Hence, activity in the RNN is
523 both persistent and sequential in nature.



524

525 **Figure 7. Simulation 9: A. Sequential and persistent activation of clustered neural populations within**
526 **the RNN.** The y-axis represents each RNN unit, the x-axis represents time. The first cluster is activated by
527 the input layer, and maintains persistent activity until the first action is executed. At that moment, via
528 excitatory projections from the Action nodes (Fig. 1C) to the following ($i+1^{th}$) cluster in the RNN (Fig. 1B)
529 gets activated, and thus displays persistent activation, and so forth via the cortico-basal ganglia loops
530 (light blue arrows in Fig. 1). Color bar represents firing rate. **B. Sequential and sparse activation in the**
531 **BG.** The y-axis represents the G unit activity over time (x-axis). Each G unit responds in a sequential and
532 transient manner, as has been shown in neurophysiological single-cell recordings of the BG (e.g., Gouvêa
533 et al., 2015). Color bar represents normalized firing rate.

534 To gain better visual intuition on the RNN dynamics, we performed dimension reduction on the row
535 space of the unit (i.e., neuron) by time matrix displayed in figure 7A. We then dynamically plotted the
536 first 3 principal components (PCs) as a function of time. [Video 2](#) shows that each cluster of persistent
537 neural activity acts as an attractor state (within the highly dimensional space of the RNN), and the
538 dynamics in the RNN switch from one attractor to the other when an action is executed, again displaying
539 both persistent and sequential neural dynamics.

540

--Insert Video 2--

541 **Video 2. Simulation 9: Dynamical visualization of RNN and Action nodes activity.** The left panel shows
542 how activity in each Action node is executed at the learned action time, each color represents the
543 activation of a specific A node in the thalamus. Given the structure and mechanism described in figure 1,
544 the right panel displays the neural RNN trajectory showing that each action execution triggers a switch
545 from the i^{th} to the i^{th+1} attractor state.

546 The qualitative pattern of the RNN sequential and persistent dynamics (Fig. 7A) is different than
547 observed in rodent (Harvey et al., 2012; Eichenbaum, 2014; Mello et al., 2015) or monkey (Jin et al.,
548 2009) neurophysiological recordings, which reveal sequential sparse activation (individual neurons
549 display quick and transient activation as behavior unfolds). Notably however, the Go nodes in the BG
550 module of our model display qualitatively similar sequential and sparse activation patterns as that seen
551 empirically in the BG (Fig. 7B; see figures 2A, 3B, 1E, 8C, respectively of Mello et al., 2015; Rueda-Orozco
552 and Robbe, 2015; Bakhurin et al., 2017; Dhawale et al., 2017)

553 Discussion

554 We have presented a neurocomputational model combining the strength of associative chains (e.g.,
555 Pereira and Brunel, 2020) and cluster-dependent (e.g., Maes et al., 2020) models, while also providing a
556 model of how the BG contribute to recurrent cortical dynamics in sequential behaviors. Our model
557 factorizes action order, identity, and time, which are represented in distinct loci of the cortico-basal
558 ganglia neural network. Crucially, factorizing these features provides the network with the ability to
559 independently manipulate the building blocks of precisely timed action sequences, thereby increasing
560 the computational power of our model. This increased power is illustrated through several interesting
561 emergent properties. First, we demonstrated that the ACDC model can learn and reproduce precise
562 spatiotemporal action sequences with temporal synchrony or asynchrony (i.e., constant or varying inter
563 action intervals). Second, our model displays several flexibility properties: temporal shifting, rescaling
564 and compositionality, and sustained motor activation; culminating in our model's ability to reproduce
565 the Thunderstruck song and change it to a bossa nova tempo. Third, the model can account for
566 behavioral and neurophysiological empirical observations.

567 *Encoding order as attractor state switches in the RNN*

568 Recent work suggests that dynamic representations can be understood as switches in activity of neural
569 networks (Ju and Bassett, 2020). Within this framework, one can envision action sequences as neural
570 states unfolding over time. By analyzing the neural activity of secondary motor cortex in rodents,
571 Recanaseti et al. (2020) showed that sequential behavior was subtended by the sequential unfolding of

572 attractor states, whereby each action in the sequence was subtended by a particular attractor state.
573 Furthermore, these authors were able to model variability in action timing by adding correlated noise to
574 the dynamics of a RNN. This addition allowed their RNN to display dynamics that jump from one
575 attractor state to another, but at random times (hence explaining the variability in action timing). Our
576 model is based on a similar mechanism. The dynamical activity of the RNN reflects switches in attractor
577 states induced by excitatory projections to the RNN that transiently modify its E-I balance. However, via
578 BG learning and modulation, our model precisely controls the switch to another cortical attractor state,
579 thereby influencing output timing. Within our conceptualization, we suggest that persistent activity
580 within a cluster indicates the latent state that the system is in (e.g., Collins and Frank, 2013), which in
581 this case reflects the ordinal position in the sequence. Moreover, in contrast to previous models, the
582 clusters themselves were not assumed to be anatomically hard-wired but emerged within the RNN via
583 learning.

584 Alternative models have proposed different mechanisms for encoding ordinal position. Some models
585 possess a temporal context layer whose state is modified dynamically as time passes. The nature of this
586 activity can take the form of a cyclical signal (Hartley and Houghton, 1996), decaying start signal (Page
587 and Norris, 1998), or a sequence of overlapping states (Burgess and Hitch, 1999, 2006). Other models
588 assume that the network input (used to learn the sequence) is itself sequential in nature (Murray and
589 Escola, 2017; Maes et al., 2020), and learning the spatiotemporal signal depends on the sequential
590 nature of the input. Our model is free of this assumption; the network input is a single pulse of
591 activation, but can nevertheless reproduce a precisely timed spatiotemporal signal. This ability emerges
592 from the feedback loop from thalamic Action nodes to the cortical RNN, triggering transitions to a
593 subsequent cortical attractor. One can therefore consider motor output as part of the teaching input
594 signal to the RNN; because motor activation unfolds sequentially in our model, the sequential nature of
595 the teaching signal emerges from our network architecture.

596 Interestingly, the idea that the motor cortex (presumably via motor thalamus neurons) acts as a
597 teaching signal to other brain areas has received strong support from rodent lesion studies. For instance,
598 rats are unable to learn a precisely-timed lever press when their M1 cortex is lesioned (Kawai et al.,
599 2015), and transiently inactivated or disturbed via optogenetic manipulation (Otchy et al., 2015). More
600 generally, the notion that motor output can influence cognitive representations and transitions is
601 consistent with the emerging literature on how cognitive functions scaffold on top of motor functions in
602 cortico-basal ganglia circuits (Koziol and Budding, 2009; Collins and Frank, 2016a).

603 ***Motor sequence flexibility as inputs to the basal ganglia***

604 Humans can adapt their motor output almost instantaneously given external or internal stimuli. For
605 instance, musicians can modify the tempo of a song upon signaling of the conductor. Such flexibility
606 necessarily needs to stem from fast reconfiguration of neural dynamics, rather than emerge from
607 changes in networks weights (Remington et al., 2018). Murray and Escola (2017) proposed a model of
608 interconnected medium spiny neurons in the striatum that can apply such dynamic reconfiguration. In
609 particular, their model could perform temporal rescaling of sparse sequential activity. Yet, flexibility in
610 this model is constrained to isosynchronous sequences (see also Egger et al., 2020; Kozachkov and
611 Michmizos, 2020)(see also Egger et al., 2020). However, a recent model making use of eligibility traces
612 (Florian, 2007; Izhikevich, 2007; Frémaux et al., 2010; Soltoggio and Steil, 2013; Bellec et al., 2019),
613 manages to learn precise non-isosynchronous spatiotemporal sequence learning (Cone and Shouval,
614 2021). Still, it is unclear how such a model can rescale non-isosynchronous sequences, and neither of
615 these models is capable of exhibiting temporal compositionality. Crucially, the ACDC model can perform
616 temporal rescaling for both isocynchronous and non-isosynchronous sequencing, and it can also flexibly
617 switch the tempo altogether through a multiplicative signal to the BG.

618 The temporal properties of our model discussed in the previous paragraph emerge from additional
619 inputs to the BG. What is the nature of this input? One possibility could be dopaminergic. Indeed,
620 midbrain dopaminergic nuclei massively broadcast to the striatum (Watabe-Uchida et al., 2017), and
621 several studies have implicated dopamine in controlling movement vigor (Beierholm et al., 2013; Hamid
622 et al., 2015, 2021; Panigrahi et al., 2015; Zénon et al., 2016; Berke, 2018; Gaidica et al., 2018; Sedaghat-
623 Nejad et al., 2019; Augustin et al., 2020). Dopamine has also been extensively implicated in impulsive
624 (i.e. pathologically speeded) behavior (van Gaalen et al., 2006; Frank et al., 2007; Pattij and
625 Vanderschuren, 2008; Buckholtz et al., 2010; Pine et al., 2010; Dalley and Roiser, 2012; Economidou et
626 al., 2012). Furthermore, administration of amphetamine and haloperidol to human participants,
627 respectively increasing and decreasing tonic dopamine levels, has been associated to faster and slower
628 response times during a simple reaction time task (Lake and Meck, 2013).

629 If dopamine can flexibly modulate (i.e. speed up or slow down) action execution timing, the question
630 remains upon which psychological process this neuromodulatory effect takes place. Within the
631 accumulation-to-bound framework (Ratcliff, 1978; Ratcliff and Rouder, 1998), this effect could
632 potentially alter two distinct processes. First, dopamine could play a role on the speed (or rate) of
633 evidence accumulation. In line with this hypothesis, several studies have highlighted a clear effect of

634 dopamine on the drift rate of evidence accumulation in perceptual (Yousif et al., 2016; Beste et al.,
635 2018) or reward-based (Westbrook et al., 2020) decision-making tasks. Our model implements this
636 possibility. Indeed, inputs to Go nodes modify (i.e. increase or decrease) the drift rate of evidence
637 accumulation. Yet, the speed at which an action is produced also depends on the response threshold,
638 with lower thresholds increasing speed at the expense of accuracy (Heitz, 2014). Therefore, a second
639 alternative is that dopamine or other BG modulations may modify the threshold of action execution
640 (Wiecki and Frank, 2013; Lloyd and Dayan, 2015). Interestingly, Parkinson's disease patients on
641 subthalamic deep brain stimulation tend to behave impulsively (Frank et al., 2007), due to modulation of
642 the decision threshold (Frank, 2006; Cavanagh et al., 2011; Herz et al., 2016). Naturally, both hypotheses
643 are not mutually exclusive; further research should investigate the effects of dopaminergic and
644 subthalamic modulations regarding motor sequence flexibility.

645 ***Biological basis and learning***

646 In line with recent models (Murray and Escola, 2017; Maes et al., 2020; Cone and Shouval, 2021), the
647 ACDC model implements a certain level of biological plausibility, and still is able to capture a plethora of
648 data both at the neurophysiological and behavioral level. For instance, we demonstrate that the model
649 can replicate sequential sparse activation observed within the basal ganglia (Gouvêa et al., 2015).
650 Another model making use of RNNs set at a near chaotic regime (Rajan et al., 2016) has been able to
651 replicate sparse sequential activations as recorded in mice parietal cortex during spatial navigation
652 (Harvey et al., 2012). Yet, training in these networks is based on highly supervised mechanisms that are
653 not biologically plausible (Sussillo and Abbott, 2009; Laje and Buonomano, 2013; Hardy et al., 2018).
654 Therefore, future research should analyze whether more biological plausible RNNs (see Miconi, 2017)
655 can reproduce such action patterns. Note that some of the implementation details of our model have
656 still to be worked out (see limitations section below).

657 ***Encoding and executing multiple sequences***

658 One important advantage of cluster-based models is the potential to encode multiple sequences within
659 the same network of interconnected neurons. Within our model, this would tantamount to having
660 several sequential attractor state neural trajectories, each of which subtends the execution of one
661 specific action sequence. Therefore, action sequence selection is seen as targeting a specific cluster of
662 units within the RNN, leading to the execution of the corresponding sequence. Moreover, the ability to
663 produce various sequences simultaneously would reduce to simply activating more than one cluster
664 (Murray and Escola, 2017).

665 Another important question focuses on investigating how sequences are chained one after the other in
666 order to produce adaptive behavior. Neurophysiological recordings in mice have revealed the existence
667 of specific neural codes during sequences that signal the beginning and end (or boundaries) of a
668 sequence (Jin et al., 2014; Jin and Costa, 2015). These go and stop signal may be used to signal the
669 system to transition from one neural trajectory to the other, thereby allowing action sequences to be
670 chained (Logiaco et al., 2019).

671 ***Limitations and future directions***

672 As previously noted, some of the implementation details of our model have still to be worked out. For
673 instance, reinforcement learning of action timing is conceptually thought to take the form of a tri-factor
674 hebbian learning rule (Montague et al., 1995; Bailey et al., 2000; Izhikevich, 2007; Hoerzer et al., 2014;
675 Miconi, 2017), where neurons subtending a rewarding behavior (and hence forming a specific cortical
676 activity patterns) increase their connectivity to D1 receptor striatal populations (also known as Go cells)
677 via dopaminergic activity bursts stemming from midbrain nuclei (Frank, 2005; Collins and Frank, 2013).
678 Our implementation is slightly different. Reinforcement is shifted later in the information processing
679 pipeline, and action time learning takes place between Go nodes (which we also consider as D1 receptor
680 cells in the striatum) and thalamic motor neurons. We applied a delta rule within the BG-thalamus
681 module. Much evidence indicates that the BG learn via reinforcement learning (e.g., McClure et al.,
682 2003; O'Doherty et al., 2004; Badre and Frank, 2012), but the brain also makes use of signed errors for
683 precise timing e.g., in the cerebellum. Our learning rule in the BG-thalamus thus summarizes the
684 contributions of these systems in conjunction. In contrast, classical Hebbian learning rule was applied
685 within the RNN and between the RNN and BG. Indeed, these projections simply carry “chaining”
686 information (i.e. they allow for the sequential structure of the chain to emerge during learning), and
687 therefore do not need to be fine-tuned to a specific value for the emergence of precise action timing.
688 Future work should consider both biological learning constraints and implement a more detailed
689 architecture of the basal ganglia networks.

690 Moreover, future work on the ACDC should focus on investigating the limits with which RNNs can
691 encode multiple sequences without creating interference. This entails exploring two aspects of the
692 model. First, our model forces orthogonalization of the inputs to the RNN in order to make sure that no
693 clusters are interconnected. Novel versions of the ACDC need to investigate how this orthogonalization
694 may emerge from specific learning rules. Second, cluster size matter, as bigger cluster size may be more
695 robust to noise. Therefore, given an initial number of RNN excitatory neurons, only but a limited amount

696 of sequences can be encoded. Therefore, the interaction between learning, cluster size and sequence
697 interference should also be investigated more systematically.

698 Our model simulates action sequences such as those needed to play the guitar or the piano. Within this
699 context, each action is represented as a discrete entity. However, many daily life action sequences are
700 subtended by more continuous actions, as for instance when playing violin with a bow. The ACDC could
701 be expanded by having more continuous representations of action plans and execution in our BG-
702 thalamus module. Based on dynamic field theory, one potential approach would be to represent actions
703 as dynamic neural fields (Erlhagen and Schöner, 2002; Cisek, 2006; Klaes et al., 2012), which have been
704 shown to successfully model more continuous reaching actions (Christopoulos et al., 2015). Moreover,
705 these continuous action representations in the BG may require additional inputs from the cerebellum
706 for movement coordination (Thach et al., 1992) or sequence prediction for motor control (Bastian,
707 2006).

708 Finally, recent research focused on how humans extract abstract knowledge, and generalize this
709 knowledge to other situations (Collins and Frank, 2016b) (Collins and Frank, 2013; Whittington et al.,
710 2020; Baram et al., 2021). Indeed, abstracting the action sequence structure of the Thunderstruck song
711 may be useful for future learning. Transferring the abstract structure of the Thunderstruck song when
712 learning a novel song that shares a similar structure should improve learning (Lehnert et al., 2020).

713

Conclusion

714 Separating time and order information in two distinct loci of a biologically inspired model of action
715 sequences allowed us to increase computational power, and to capture a significant amount of data at
716 the neurophysiological and behavioral levels. Although some specific aspect of the implementation of
717 this functional specialization need still to be resolved, we demonstrate that such an architecture
718 increased motor flexibility. We propose a concrete and tractable mechanism of this flexibility, and thus
719 suggest a model of how humans and animals can learn and effortlessly manipulate precise
720 spatiotemporal signals at the basis of complex behavior.

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1050

1051 **Appendix A**

1052 Our ACDC model contains three main modules (Fig. 1): an input layer (Fig. 1A), an RNN (representing
1053 premotor cortex; Fig. 1B), a BG-thalamus unit (Fig. 1C).

1054 The input layer reflects a vector of $N=200$ neurons of which a subset (20) is activated and each neuron
1055 excites only one neuron in the RNN.

1056 The dynamics within the ACDC model represent the sequential unfolding of RNN-BG-thalamus-RNN (i.e.,
1057 cortico-basal ganglia) loops, depicted by the light blue arrows in figure 1. The loop starts with the
1058 activation of a cluster of excitatory RNN neurons, and the dynamics of the RNN excitatory neurons are
1059 governed by equation 3:

$$1060 \quad \tau_{mn} \frac{dx_i}{dt} = -x_i + \Theta \left(\sum_{j=1}^N W_{ij} x_j - J^{EI} x_I + J^{EA} (x_A \gamma_E) + x_i^{in} \right) \quad \text{equation 3}$$

1061 where x_i and x_j represent post and pre-synaptic RNN unit activity (purple nodes in Fig. 1B) and W_{ij} is the
1062 recurrent weight matrix. J^{EI} and J^{EA} represent respectively the weights from the shared inhibitory neuron
1063 (orange node in Fig. 1B) and from the motor thalamus neurons (from here on termed Action nodes for
1064 simplicity) to the excitatory RNN units. x_I , x_A and x_i^{in} represent respectively the activity of the shared
1065 inhibitory neuron, Action nodes (see below), and the input to the excitatory RNN units. γ_E is the gain on
1066 Action nodes activation projected to the excitatory RNN neurons (see below for the functional property
1067 of this parameter). Θ , the non-linear transformation function, is governed by $\Theta(x) = (2 / (1 + e^{-\lambda x})) - 1$

1068 (where λ is the gain parameter and with additional non-linearity at zero, i.e. $\Theta(x) = 0$ if $\Theta(x) < 0$); and τ_{rnn}
 1069 is the encoding constant. Note that input projections and all Action nodes to RNN projections are
 1070 orthogonal (i.e. some RNN excitatory neurons receive inputs from the input layer, whereas others
 1071 receive from inputs from Action nodes; each projection excites 20 RNN units). The shared inhibitory x_I
 1072 activation is described by equation 4:

$$1073 \quad \tau_{rnn} \frac{dx_I}{dt} = -x_I + J^{IE} x_i + J^{IA} (x_A \gamma_I) \quad \text{equation 4}$$

1074 where J^{IE} , J^{IA} and γ_I respectively represent the weights from the excitatory RNN neurons to their shared
 1075 inhibitory neuron, the weights from the Action nodes to the shared inhibitory neuron, and the gain on
 1076 Action nodes activation for the projections towards the inhibitory neuron in the RNN.

1077 In turn, each excitatory RNN cluster projects to its corresponding “Go” cell in the BG (blue arrow 1 from
 1078 Fig. 1B to Go node in Fig. 1C), and each Go cell accumulates evidence for the j^{th} action associated to the
 1079 j^{th} order, following equation 5:

$$1080 \quad \tau_g \frac{dg_j}{dt} = -g_j + \sum_i^N W_{ij} x_i - J^{GN} n_j \quad \text{equation 5}$$

1081 where g_j is the activation of the j^{th} Go units, W_{ij} is the weight matrix representing connectivity between
 1082 RNN and Go units, x_i is the activity of the RNN excitatory units, J^{GN} is the inhibitory weight between the
 1083 j^{th} No Go and Go nodes, n_j is the activation of the j^{th} No Go node, and τ_g is the encoding constant (with τ_g
 1084 $\gg 0$, thereby simulating evidence accumulation-like dynamics).

1085 Striatal Go cells facilitate response execution by projecting towards the corresponding Action nodes
 1086 (blue arrow 2 from the Go to Action nodes in Fig. 1C), whose dynamics are governed by equation 6:

$$1087 \quad \tau_a \frac{da_j}{dt} = -a_j + \Theta(J^{AG} g_j - b) \quad \text{equation 6}$$

1088 where a_j is the activation of the j^{th} action, g_j is the activation of the j^{th} Go unit, b is the negative bias (i.e.
 1089 threshold), Θ is a nonlinear function as in equation 3, and τ_a is the encoding constant. J^{AG} is the weight
 1090 from the j^{th} Go unit to the j^{th} Action unit, and was randomly drawn from a Gaussian distribution with
 1091 mean = 2 and s.d. = 0.2. In turn, Action nodes project excitatory connections to three distinct parts of

1092 the network simultaneously. First, Action nodes project to the cluster of excitatory neurons in the RNN
 1093 representing the $i+1^{th}$ order in the sequences (blue arrow 3a in Fig. 1). Second, Action nodes project to
 1094 the inhibitory shared neuron (blue arrow 3b to orange node in Fig. 1), that in turn globally inhibits all the
 1095 clusters in the RNN. Note that the gain parameter values on Action nodes activity are larger for
 1096 projections to the excitatory clusters vs inhibitory neuron of the RNN (i.e. $\gamma_E > \gamma_I$). This allows the
 1097 activation of Action nodes to perturb the E-I RNN balance in a way that allows the j^{th} cluster to shut
 1098 down and the $i+1^{th}$ cluster to be expressed. Third, Action nodes project excitatory connections back to
 1099 their corresponding No Go cells (blue arrow 3c from j^{th} Action node in the thalamus to j^{th} No Go node in
 1100 the BG, see Fig. 1C). The dynamics of No Go cells are in turn dictated by equation 7:

$$1101 \quad \tau_n \frac{dn_j}{dt} = -n_j + J^{NA} a_j \quad \text{equation 7}$$

1102 where n_j is the activation of the j^{th} No Go node, J^{NA} is the weight from the j^{th} Action unit to the j^{th} No Go
 1103 unit, a_j is the activation of the j^{th} Action node, and τ_n is the encoding constant.

1104 In Table 1 we report the parameter values used for all eight simulations described in the main text.

1105 Table 1. Parameter values for all simulations

Parameters	Values
α_1 (RNN) / α_1 (RNN-Go)	0.01 / 0.00002
α_2 (RNN) / α_2 (RNN-Go)	0.1 / 0.4
W_{\max} (RNN) / W_{\max} (RNN-Go)	1 / 0.05
τ_w, b	0.5
η	0.4
ϕ	0.01
τ_a, τ_n, J^E	0.1
γ_E / γ_I	21.4 / 21
$\tau_{rnn}, \chi^{in}, J^{EI}, J^{EA}, J^{IA}, J^{GN}, J^{NA}$	1
$\lambda_{rnn} / \lambda_a$	10 / 10000
τ_g	0.001

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