The basal ganglia and cerebellum communicate with the cerebellum

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The basal ganglia and cerebellum are major subcortical structures that influence not only movement, but putatively also cognition and affect. Both structures receive input from and send output to the cerebral cortex. Thus, the basal ganglia and cerebellum form multisinaptic loops with the cerebral cortex. Basal ganglia and cerebellar loops have been assumed to be anatomically separate and to perform distinct functional operations. We investigated whether there is any direct route for basal ganglia output to influence cerebellar function that is independent of the cerebral cortex. We injected rabies virus (RV) into selected regions of the cerebellar cortex in cebus monkeys and used retrograde transneuronal transport of the virus to determine the origin of multisinaptic inputs to the injection sites. We found that the subthalamic nucleus of the basal ganglia has a substantial disynaptic projection to the cerebellar cortex. This pathway provides a means for both normal and abnormal signals from the basal ganglia to influence cerebellar function. We previously showed that the dentate nucleus of the cerebellum has a disynaptic projection to an input stage of basal ganglia processing, the striatum. Taken together these results provide the anatomical substrate for substantial two-way communication between the basal ganglia and cerebellum. Thus, the two subcortical structures may be linked together to form an integrated functional network.

The basal ganglia and cerebellum are major subcortical structures that influence not only movement, but putatively also cognition and affect (1, 2). Both structures receive input from and send output to the cerebral cortex. Thus, the basal ganglia and cerebellum form multisinaptic loops with the cerebral cortex. The major interactions between these loops were thought to occur largely at the cortical level (3). Recently, we showed that one of the output nuclei of the cerebellum, the dentate nucleus, has a disynaptic projection to an input stage of basal ganglia processing, the striatum (4). This pathway enables cerebellar output to influence basal ganglia function. Here, we investigated whether a comparable pathway allows basal ganglia output to influence cerebellar function. We injected rabies virus (RV) into regions of the cerebellar cortex in cebus monkeys and used retrograde transneuronal transport of the virus to determine the origin of multisinaptic inputs to the injection sites. Our results indicate that the subthalamic nucleus (STN) of the basal ganglia has substantial disynaptic projections to the cerebellar cortex.

Results

We injected the N2c strain of RV into selected sites within the cerebellar cortex of three cebus monkeys (Fig. 1 and Table S1). RV is transported transneuronally in the retrograde direction in a time-dependent fashion in nonhuman primates (4–8). We set the survival time at 42 h to allow two stages of transport: retrograde transport of RV to first-order neurons that project to the injection site and then, retrograde transneuronal transport of the virus to second-order neurons that make synaptic connections with the first-order neurons. The suitability of the survival time was confirmed by the presence of second-order neurons labeled in cortical layer V, the site of corticopontine neurons (9), and by the absence of third-order neurons labeled in layer III.

Our injections targeted Crus Iip (n = 2) and the hemispheric expansion of lobule VIIb (HVIIB) (n = 1) (Fig. 2A). In all cases we mixed RV with a conventional tracer, the β subunit of cholera toxin (CTb, 0.02%). We used this mixture to facilitate identification of the RV injection site and to label neurons that project directly to it (first-order neurons). After RV-CTb injections into Crus Iip and HVIIB, we found first-order neurons labeled with CTb and RV in regions of the pontine nuclei (Fig. S1) and the inferior olive that are known to project to the injected regions of the cerebellar cortex (10–12). We found second-order neurons labeled with RV in cortical areas and in regions of the parvo-cellular portion of the red nucleus (Fig. 3A) that are known to project to the first-order neurons in the pontine nuclei and the inferior olive (9, 13).

Surprisingly, we also found substantial numbers of second-order neurons labeled with RV in the STN predominantly on the side contralateral to the injection site (Figs. 1, 2B, and 3A). We counted labeled neurons on every other section through the STN of the two animals illustrated in the figures and found 1,160 second-order neurons in the STN after the Crus Iip injection (AB2) and 923 second-order neurons after the HVIIB injection (AB3). In all animals, second-order neurons labeled with RV were present throughout the entire rostrocaudal extent of the STN (Figs. 2B and 4A and B). However, larger numbers of second-order neurons were located in the rostral half of the STN following the Crus Iip injections (AB1 and AB2) (Fig. 4A red bars), whereas larger numbers of second-order neurons were located in the caudal half of the STN following the HVIIB injections (AB3) (Fig. 4A blue bars). The second-order neurons labeled from Crus Iip and HVIIB injections also differed in their dorsoventral and mediolateral distribution. Second-order neurons projecting to Crus Iip were located more ventromedially in the STN than those projecting to HVIIB (Figs. 2B and 4B). These observations indicate that there is a disynaptic connection between the STN and the cerebellar cortex and that this connection is topographically organized.

The STN has been subdivided into three functional territories: sensorimotor, associative, and limbic (Fig. 4C). These subdivisions are based on STN interconnections with regions of the globus pallidus and the ventral pallidum (14–16). The pattern of inputs from the cerebral cortex to the STN also imposes a functional topography on the STN (Fig. 4D). A comparison of our data with these functional subdivisions indicates that most of the STN neurons that project to Crus Iip are located in its associative territory, which receives input from the frontal eye fields and regions of prefrontal cortex. In contrast, most of the STN neurons that project to HVIIB are located in its sensorimotor territory, which receives input from the primary motor cortex and several of the premotor areas in the frontal lobe. Although we have examined only a relatively small...
The nuclei, which mediate the disynaptic connection between the STN and the cerebellum, remain to be determined. However, the STN is known to project to the nucleus reticularis tegmenti pontis (NRTP) and several basal pontine nuclei (17). As noted above, we observed first-order neurons labeled with CTb and RV in the NRTP and multiple basal pontine nuclei after our tracer injections (Fig. S1). Thus, we view the pontine nuclei as the most likely candidates for mediating the disynaptic connection, but this proposal remains to be tested in future experiments.

Discussion

The STN has been described as the “driving force of the basal ganglia” (18). Our results indicate that this driving force extends well beyond the nuclei of the basal ganglia to the cerebellum. As a consequence, the anatomical substrate exists for both normal and abnormal signals from the STN to influence cerebellar processing. The topographic organization of this disynaptic pathway suggests that STN output could have an impact on cerebellar function during motor and nonmotor behavior. In the following paragraphs we will briefly describe some of the potential implications of this pathway for cerebellar involvement in (i) prototypical basal ganglia disorders and (ii) reward processing.

Parkinson’s disease (PD) and dystonia are traditionally considered to be “basal ganglia disorders.” PD is associated with degeneration of a specific set of dopaminergic neurons in the pars compacta of the substantia nigra. Acquired (secondary) dystonia also is often associated with lesions of the basal ganglia (19). Although no overt neurodegeneration has been identified in idiopathic (primary) dystonia, there is evidence for alterations in the basal ganglia in this form of the disorder as well (20). Despite these results, a number of observations have suggested that alterations in cerebellar activity may contribute to the motor symptoms of both PD and dystonia. For example, imaging studies report abnormal increases in cerebellar activity in PD patients and in subjects with idiopathic dystonia (20–22). In PD, deep brain stimulation of the STN improves the motor signs and normalizes cerebellar activation (22, 23). In addition, one of the cardinal symptoms of Parkinson’s disease (PD) and dystonia is traditionally considered to be “basal ganglia disorders.” PD is associated with degeneration of a specific set of dopaminergic neurons in the pars compacta of the substantia nigra. Acquired (secondary) dystonia also is often associated with lesions of the basal ganglia (19). Although no overt neurodegeneration has been identified in idiopathic (primary) dystonia, there is evidence for alterations in the basal ganglia in this form of the disorder as well (20). Despite these results, a number of observations have suggested that alterations in cerebellar activity may contribute to the motor symptoms of both PD and dystonia. For example, imaging studies report abnormal increases in cerebellar activity in PD patients and in subjects with idiopathic dystonia (20–22). In PD, deep brain stimulation of the STN improves the motor signs and normalizes cerebellar activation (22, 23). In addition, one of the cardinal symptoms of Parkinson’s disease (PD) and dystonia is traditionally considered to be “basal ganglia disorders.” PD is associated with degeneration of a specific set of dopaminergic neurons in the pars compacta of the substantia nigra. Acquired (secondary) dystonia also is often associated with lesions of the basal ganglia (19). Although no overt neurodegeneration has been identified in idiopathic (primary) dystonia, there is evidence for alterations in the basal ganglia in this form of the disorder as well (20). Despite these results, a number of observations have suggested that alterations in cerebellar activity may contribute to the motor symptoms of both PD and dystonia. For example, imaging studies report abnormal increases in cerebellar activity in PD patients and in subjects with idiopathic dystonia (20–22). In PD, deep brain stimulation of the STN improves the motor signs and normalizes cerebellar activation (22, 23).
of PD, tremor at rest, is abolished by stimulating or lesioning the ventral intermediate nucleus of the thalamus, which is a target of cerebellar efferents (24). Similarly, in a mouse model of dystonia, pharmacological stimulation of the cerebellar vermis elicited dystonic postures of the trunk and limbs (25).

The discovery of a disynaptic connection between the basal ganglia and cerebellum provides a unique framework for interpreting these results. It is notable that in both PD and idiopathic dystonia, neural activity in the STN is higher than normal and is characterized by abnormal bursting and oscillatory activity (26). Abnormal signals from the STN to the cerebellar cortex could evoke the increased cerebellar activation that is present in both disorders and alter cerebellothalamocortical input to the cerebral cortex. Further attempts to ameliorate the symptoms of PD might benefit by focusing specifically on normalizing activity in the disynaptic pathway from STN to the cerebellum. In fact, part of the effectiveness of deep brain stimulation of the STN might be achieved through this mechanism.

Our findings also provide a potential explanation for the presence of cerebellar activation in imaging studies that were explicitly designed to study the normal functions of the basal ganglia. For example, several imaging studies have examined whether regions of the basal ganglia and related cortical areas display functional activation consistent with their involvement in “temporal difference” models of reward-related learning (27, 28). It is noteworthy that robust cerebellar activation was present in these experiments along with activation in the dorsal and ventral striatum. The disynaptic connection between the STN and the cerebellum provides an anatomical substrate for reward-related signals in the basal ganglia to influence cerebellar function during learning.

From a computational perspective, the basal ganglia and cerebellum have been viewed as segregated modules that implement different learning algorithms—reinforcement learning in the case...
of the basal ganglia and supervised learning in the case of the cerebellum (29, 30). A previous study from our lab demonstrated that an output stage of cerebellar processing, the dentate nucleus, has a disynaptic connection with the input stage of basal ganglia processing, the striatum (4) (Fig. 1). The current report provides evidence for the reciprocal connection. Taken together these results provide the neural basis for substantial two-way communication between the basal ganglia and cerebellum. Thus, the two subcortical structures may be linked together to form an integrated functional network. One might then ask what new computational operations emerge by interconnecting a reinforcement learning module with a supervised learning module.

Methods

Subjects. This report is based on observations from three cebus monkeys (Cebus apella, 1.9–2.6 kg, 2 males and 1 female). In each monkey, a mixture of the N2c strain of the RV and a conventional tracer \( \text{CTb} \) was injected into the cortex of the cerebellar hemisphere. The protocol was approved by the institutional animal care and use committee and the biosafety committee. Biosafety practices conformed to the biosafety level 2 regulations outlined in Biosafety in Microbiological and Biomedical Laboratories (Department of Health and Human Services publication no. 93–8395). Details of the procedures for handling virus and virus-infected animals have been published previously (5).

Experimental Procedures. All surgical procedures were performed under aseptic conditions. The night before surgery, the monkeys were administered dexamethasone (0.5 mg/kg, i.m.). Monkeys were sedated with ketamine (20 mg/kg, i.m.), intubated, and maintained on gas anesthesia (enflurane; 1.5–2.5%). Dexamethasone (0.5 mg/kg, i.m.), glycopyrrolate (0.01 mg/kg, i.m.), and an antibiotic (ceftriaxone; 75 mg/kg, i.m.) were administered at the time of surgery. Respiratory rate, blood oxygen level, body temperature, and sensitivity to noxious stimuli were monitored at regular intervals during the procedure. Each monkey had its head restrained in a Kopf stereotaxic frame (Kopf Instruments). A craniotomy was performed to expose the ventral portions of the occipital cortex and the lateral portion of the posterior cerebellum. With the aid of a surgical microscope, we used a Hamilton syringe (30-gauge needle) to place multiple injection tracks into the cerebellar hemisphere (Crispus lucifer in animals AB1 and AB2, HVIIB in AB3). We injected small amounts (0.2 μL) of a mixture of RV (4.5 × 10^9 pfu/mL; provided by M. Schnell) and CTb (0.02%; List Biological Laboratories) at every 0.5 mm along the depth of each injection track (Table S1). The depths of these injections were based on prior structural magnetic resonance images of each cerebellum. When all injections were completed, the cerebellum was covered with artificial dura and the incision was sutured. The monkeys were placed in an isolation chamber and administered an analgesic (proporphine; 0.01 mg/kg) and dexamethasone (0.25 mg/kg) every 12 h.

Prior studies have demonstrated that RV is transported exclusively in the retrograde direction in a time-dependent fashion (4–8). The available evidence suggests that the spread of RV is exclusively transynaptic and that the virus is neither taken up by fibers of passage nor transported between neurons and glia (5). The time to infect first-, second-, and third-order neurons depends on the strain of RV and its concentration. The N2c strain used in the present experiments is transported at a higher transfer rate than other strains used for tracing (e.g., CV-S-11). In the current experiments, we set a survival time following the cerebellar injections to 42 h (Fig. 1 and Table S1). This survival time was based on a series of experiments that examined a range of survival times following central injections of the N2c strain (4). A 42-h time period is long enough to allow transport of the virus only to second-order neurons.

At the end of the survival time, the monkeys were deeply anesthetized using ketamine (25 mg/kg, i.p.) followed by pentobarbital sodium (40 mg/kg, i.p.). They were perfused transcardially with 0.1 M phosphate buffer (pH 7.4), followed by 10% buffered formalin, and finally a mixture of 10% buffered formalin and 10% glycerol at 4 °C. The brain and spinal cord were removed from the skull and stored overnight in 10% buffered formalin and 10% glycerol at 4 °C and then placed in 10% buffered formalin and 20% glycerol at 4 °C for 2 weeks. Blocks of tissue (cerebral cortex, brainstem, and cerebellum) were individually frozen and sectioned at 50 μm. Every 10th section was stained with cresyl violet for cytoarchitecture analysis. Brain sections were immunohistochemically reacted according to the avidin-biotin peroxidase method (Vectastain; Vector Laboratories). Alternating sections were reacted with mouse anti-M957 (supplied by A. Wandel, 1:300) and goat anti-choleragenoid (List Biological Laboratories, 1:10,000) to detect rabies virus or CTb, respectively. Reacted tissue sections were mounted on gelatin-coated glass slides, air dried, and coverslipped.

Data Analysis. Brain sections through the cerebral cortex, brainstem, and the cerebellum were examined for immunostaining using bright field and polarized illumination. Images of selected anatomical structures (Fig. 3 B and C) were obtained using a digital camera (RT3 monochrome camera, Diagnostic Instruments) coupled to a personal computer. The images were adjusted for contrast, brightness, and intensity using Corel Photopaint. Data were plotted using a computerized plotting system (MD2; Accustage). This system uses optical encoders to measure the X–Y movements of the microscope stage and stores the coordinates of section outlines and labeled neurons.

Injection Sites. We used the CTb labeling to identify and reconstruct the injection sites (8). The plotted sections with outlines of the injection site were used to create a flattened map of the cerebellum (following a procedure adapted from ref. 7). Flattened maps of the cerebellar map and corresponding injection sites were created for each animal, using custom laboratory software. The injection sites were then outlined on a representative map of a cebus monkey cerebellar cortex (adapted from ref. 7) (Fig. 2A).

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Supporting Information

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Fig. S1. Pontine projections to regions of cerebellar cortex. Charts display the location of first-order neurons (small dots) in the pontine nuclei labeled after retrograde transport of cholera toxin subunit β (CTb) from injections into the cerebellar cortex. The results from injections into Crus Iip (AB2) are on the Left and those from injections into HVIIB (AB3) are on the Right. To generate these charts we resized and overlapped data from each animal onto outlines of the pontine nuclei adapted from Schmahmann and Pandya (1). Roman numerals (IV and VIII) represent the rostrocaudal levels according to Schmahmann and Pandya (1). Although there was some overlap, the distribution of first-order neurons displayed some marked differences in AB2 and AB3. At rostral levels, first-order neurons in AB2 were most concentrated in ventral and medial aspects of the pons, whereas in AB3 they were most concentrated in dorsal and lateral aspects. At caudal levels, more first-order neurons were present in AB3 than in AB2.


Table S1. Experimental animals

<table>
<thead>
<tr>
<th>Exp</th>
<th>Age (yr)</th>
<th>Sex</th>
<th>Weight (kg)</th>
<th>Injection site</th>
<th>Number of injections</th>
<th>Volume injected (μL)</th>
<th>Survival time (h)</th>
<th>Virus batch</th>
<th>Virus concentration (PFU)</th>
</tr>
</thead>
<tbody>
<tr>
<td>AB1</td>
<td>14</td>
<td>F</td>
<td>2.6</td>
<td>Crus Iip</td>
<td>59</td>
<td>11.8</td>
<td>42</td>
<td>12/06</td>
<td>4.5 × 10⁹</td>
</tr>
<tr>
<td>AB2</td>
<td>6</td>
<td>M</td>
<td>3.4</td>
<td>Crus Iip</td>
<td>60</td>
<td>12</td>
<td>42</td>
<td>12/06</td>
<td>4.5 × 10⁹</td>
</tr>
<tr>
<td>AB3</td>
<td>2.5</td>
<td>M</td>
<td>1.9</td>
<td>HVIIB</td>
<td>46</td>
<td>9.2</td>
<td>42</td>
<td>12/06</td>
<td>4.5 × 10⁹</td>
</tr>
</tbody>
</table>

The table lists information about the three animals included in this report. We counted labeled neurons on every other section through the STN of AB2 and AB3. We found 1,160 second-order neurons in the STN after the Crus Iip injection (AB2) and 923 second-order neurons after the HVIIB injection (AB3). If one assumes that we sampled half of the relevant sections, then we labeled ~194–200 neurons for each microliter of virus injected. Exp, experimental animal.