the surface of CST-treated cells may have been more extensively folded and therefore competent for transport (Fig. 3A). Alternatively, they may have been incompletely folded molecules not retained in the ER by a CST-compromised quality control system. Calnexin has been shown to be part of the retention apparatus present in the ER (19, 20). The observation that incorporation of G protein from the cell surface into virus particles drops by 90% in the presence of CST suggests that much of it is, in fact, defective (21). The apparent transport of defective G protein to the cell surface suggests that calnexin serves not only as a folding factor for G protein but also as a retention factor.

Our results reveal that calnexin is a true chaperone in the sense that it associates transiently with G protein and promotes its folding. Because our data were obtained in living cells, they are likely to reflect a physiologically relevant activity. Our observations, moreover, provide evidence for sequential BiP and calnexin action during G protein folding in the ER. Indications of ordered chaperone binding have been found in mitochondria in vivo and for protein refolding in the presence of bacterial chaperones in vitro (22, 23). In all these cases, including that of G protein, the initial interaction involves a member of the Hsp70 family of chaperones. In the mitochondrial and bacterial systems, subsequent interaction occurs with a member of the Hsp60 family. In the ER, which seems to be devoid of Hsp60 homologs, calnexin may have the role of secondary chaperone.

Although a large number of proteins have been shown to associate transiently with calnexin, including about 20 different proteins in CHO cells (Fig. 2) (8), only some are likely to be strictly dependent on it for folding. In fact, G protein belongs to a subfraction of glycoproteins whose folding is inhibited by CST and by other glucosidase inhibitors (13, 24). Among the proteins that can fold independently is HA. Although it normally forms a complex with calnexin, HA folding is not inhibited by CST (6). Apparently calnexin constitutes a link in the folding, assembly, and retention machinery of the ER that is used by many glycoproteins but is essential for only some.

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 Before pulse labeling, cells were washed twice with
- b) Derote pulse labeling, cells were washed twice with phosphate-buffered saline (PBS) and incubated for 15 min in methionine-deficient media. Pulse media contained 0.4 mCi/ml of [³⁵S]methionine (L-[³⁵S] in vitro cell labeling mix; >1000 Ci/mM; Amersham). After labeling, we initiated incubations by adding media containing 4.5 mM methionine and 500 μM cycloheximide [I. Braakman, H. Hoover-Litty, K. Wagner, A. Helenius, J. Cell Biol. **114**, 401 (1991)].
- 27. Antibodies to calnexin and VSVG were as previously described (17). Cells were lysed in a buffer containing 2% CHAPS, 200 mM NaCl, 50 mM Hepes (pH 7.6), and chymostatin, leupeptin, antipain, and pepstatin (10 μ g/ml each). Nuclei were pelleted by centrifugation for 5 min at 15,000g, and lysates were rotated at 4°C with antibody and Protein A sepharose beads (Sigma) for 3 hours.
- Triton X-100–SDS wash buffer contained 0.1% SDS, 0.05% Triton X-100, 10 mM tris (pH 8.0), and 300 mM NaCl. CHAPS wash buffer contained 0.5% CHAPS, 50 mM Hepes (pH 7.6), and 200 mM NaCl.
- 29. Loss of G protein signal in Fig. 1A was seen when cells were solubilized in CHAPS, which was necessary for coprecipitation. Signal loss was not seen when cells were solubilized in 1% SDS.
- The authors thank J. Saraste, D. Bole, and K. Simons for providing antibodies to p58, BiP, and VSV G, respectively. We also thank H. Tan for help with photography. Supported by grants from NIH (RO1 GM38346 and PO1 CA46128).

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Anatomical Evidence for Cerebellar and Basal Ganglia Involvement in Higher Cognitive Function

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The possibility that neurons in the basal ganglia and cerebellum innervate areas of the cerebral cortex that are involved in cognitive function has been a controversial subject. Here, retrograde transneuronal transport of herpes simplex virus type 1 (HSV1) was used to identify subcortical neurons that project via the thalamus to area 46 of the primate prefrontal cortex. This cortical area is known to be involved in spatial working memory. Many neurons in restricted regions of the dentate nucleus of the cerebellum and in the internal segment of the globus pallidus were labeled by transneuronal transport of virus from area 46. The location of these neurons was different from those labeled after HSV1 transport from motor areas of the cerebral cortex. These observations define an anatomical substrate for the involvement of basal ganglia and cerebellar output in higher cognitive function.

The basal ganglia and cerebellum have long been regarded as contributing to the planning and execution of movement; however, there have been suggestions that these

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two structures are also involved in nonmotor or cognitive function. For example, Alexander, DeLong, and Strick (1) proposed that the basal ganglia participate in five separate loops with motor and nonmotor areas of the cerebral cortex. According to their scheme, the nonmotor output of the basal ganglia targets three cortical areas via the thalamus: dorsolateral prefrontal cortex, lateral orbitofrontal cortex, and anterior cingulate cortex. As a result of these connections, the output of the basal ganglia is

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thought to influence the higher order functions subserved by each of these cortical areas.

Similarly, Leiner, Leiner, and Dow (2) have suggested that cerebellar output is directed to prefrontal as well as motor areas of the cerebral cortex. They noted that, in the course of hominid evolution, the lateral output nucleus of the cerebellum (the dentate) undergoes a marked expansion that parallels the expansion of cerebral cortex in the frontal lobe. They argued that the increase in the size of the dentate is accompanied by an increase in the extent of the cortical areas in the frontal lobe that are influenced by dentate output. As a consequence, they proposed that cerebellar function in humans has expanded to include involvement in certain language and cognitive tasks. The absence of experimental anatomical support for a cerebellar projection to the prefrontal cortex via the thalamus has led to considerable controversy concerning the participation of the cerebellum in cognitive processing (2, 3).

In this study we examined whether the dorsolateral prefrontal cortex (dIPFC) is the target of output from the basal ganglia and the cerebellum. We chose to study the dIPFC, Walker's area 46 (4), because it is one of the best-characterized nonmotor regions of the frontal lobe (5). There is considerable evidence that the dIPFC is involved in "spatial working memory" and guides behavior on the basis of transiently stored information rather than immediate external cues (6). The dIPFC also appears to be involved in planning the order and timing of future behavior (5).

Many output neurons in the basal ganglia and cerebellum project to thalamic neurons which, in turn, innervate regions of the cerebral cortex. To examine whether some of these basal ganglia and cerebellar neurons connect with thalamic neurons that innervate the dIPFC, we injected the McIntyre-B strain of herpes simplex virus type 1 (HSV1) into area 46 of cebus monkevs (Cebus apella, n = 3) (Fig. 1) (7). The McIntyre-B strain of HSV1 is transported transneuronally in the retrograde direction by neurons in cerebello-thalamocortical and pallido-thalamocortical pathways of primates (8, 9). All injection sites were confined to the region of area 46 in the principal sulcus (PS) and did not spread to adjacent cortical regions, such as the frontal eve field (FEF) in area 8 (Fig. 1) (10).

The cortical injections of HSV1 labeled many neurons in portions of three thalamic nuclei known to innervate the dlPFC: ventralis anterior pars parvocellularis (VApc), medialis dorsalis (MD), and ventralis lateralis pars caudalis (VLc) (11). In addition, retrograde transneuronal transport of HSV1 from the dlPFC labeled many neurons in

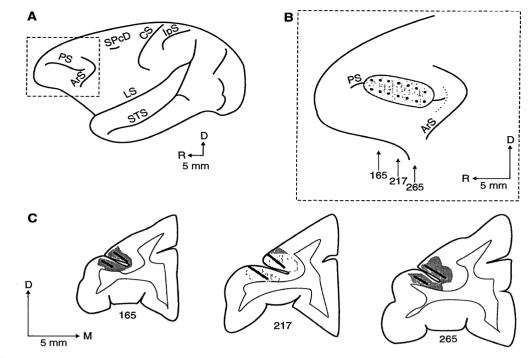
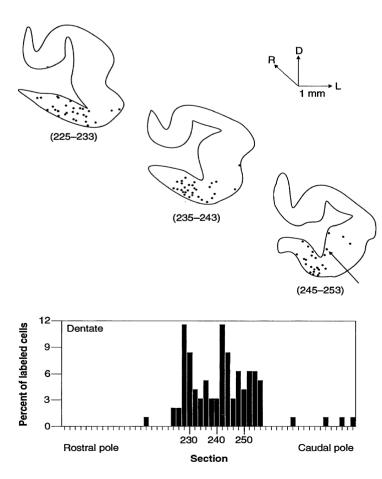


Fig. 1. HSV1 injection sites along the principal sulcus in a cebus monkey (F1). (**A**) Lateral view of the cebus brain. (**B**) Enlargement of the area enclosed by the dashed line in (A). Solid circles represent needle entry points, and the shaded area indicates the spread of HSV1 from the injection sites. The dotted line defines the boundary between Walker's areas 46 and 8 in F1. The numbered arrows indicate the location of sections in (C). (**C**) Coronal sections through the injection site. Heavy lines indicate needle tracks, and the shaded areas indicate the spread of HSV1. ArS, superior limb of the arcuate sulcus; CS, central sulcus; D, dorsal; IpS, intraparietal sulcus; LS, lateral sulcus; M, medial; PS, principal sulcus; R, rostral; SPcD, superior precentral dimple; STS, superior temporal sulcus.



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Fig. 2. Labeled neurons in the dentate nucleus. (Top) Coronal sections at three representative levels through the dentate. The solid dots indicate the positions of neurons labeled by retrograde transneuronal transport observed in five sections spaced 100 to 150 µm apart (section numbers are at the bottom in parentheses). The arrow indicates the location of the labeled neuron shown in Fig. 3A. (Bottom) Plot of the rostro-caudal distribution of labeled cells in the dentate. The location of some sections is shown along the abscissa. L, lateral.

the dentate nucleus of the cerebellum (mean = 122, range = 91-181) and in the internal segment of the globus pallidus (GPi) (mean = 332, range = 318-346) (12). Most of the labeled neurons in the dentate were found contralateral to the cortical injection site. These neurons were confined to the most ventral portion of the nucleus and were concentrated rostro-caudally in the middle third of the dentate (Fig. 2). This region of the dentate clearly differs from the more dorsal regions of this nucleus, which were labeled by retrograde transneuronal transport from the primary motor cortex (M1) or ventral premotor area (PMv) (13), and the more caudal region of the dentate, which was labeled by retrograde transneuronal transport from the FEF (14). The dorsal part of the dentate is where neurons with marked changes in activity during single joint or reaching movements have been found (15).

The labeled neurons in the dentate had round cell bodies, with multiple dendrites originating from the soma (Fig. 3A), features typical of dentate neurons that project to the thalamus (13, 14). Prior studies have shown that some of these dentate neurons terminate in the MD and VLc (16, 17). Thus, our results provide evidence that the dlPFC is a cortical target of a cerebello-thalamocortical pathway from the dentate, and that this pathway is distinct from those innervating motor areas of the cerebral cortex.

In the GPi, labeled neurons were found largely ipsilateral to the cortical injection site in the dlPFC. These neurons were located in the most dorsomedial region of the inner and outer portions of the internal segment and were concentrated rostro-caudally in the middle third of the nucleus (Fig. 4). This region of the GPi clearly differs from the more ventral and lateral regions of the GPi which were labeled by retrograde

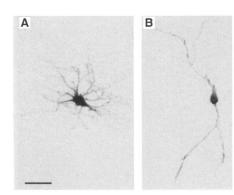


Fig. 3. Neurons labeled by retrograde transneuronal transport of HSV1 from area 46. (A) Dentate neuron. The location of the cell is indicated by the arrow in Fig. 2. (B) GPi neuron. The location of the cell is indicated by the arrow in Fig. 4. Scale bar, 50 μ m.

the supplementary motor area (SMA) (13). These more ventral and lateral regions of the GPi contain neurons that display marked changes in activity during single joint or reaching movements (18). Labeled GPi neurons had elliptical cell

bodies, with one or more dendrites radiating from each pole of the soma (Fig. 3B), features typical of GPi neurons that project to the thalamus (19). It is generally agreed that some of these GPi neurons terminate in the VApc (20). Thus, our results provide evidence that the dlPFC is a cortical target of a distinct pallido-thalamocortical pathway from the GPi.

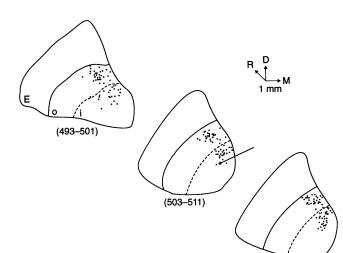
transneuronal transport from M1, PMv, or

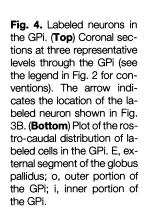
These observations have important implications for theories about the functional organization of basal ganglia and cerebellar loops with the cerebral cortex. According to one view, these loops provide a means for linking widespread regions of the cerebral cortex, such as prefrontal and posterior parietal cortex, with motor output at the level of the primary motor cortex. Such loops would serve to "funnel" information into the motor system to generate commands for movement (21). Our results support an alternative view (1, 2), one in which part of the output of the basal ganglia and cerebellum is directed back to regions of the prefrontal cortex that are known to project to these subcortical structures

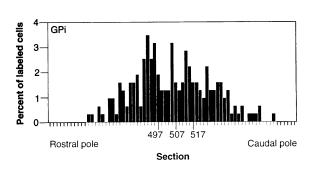
(22). This creates the potential for closed loops between the prefrontal cortex and both the basal ganglia and cerebellum. These loops would operate in parallel with those serving motor areas of the cerebral cortex but would have a "cognitive" rather than a "motor" function.

Considerable evidence indicates that the basal ganglia participate in aspects of cognitive function. Individuals affected by Parkinson's disease and Huntington's disease, two well-known basal ganglia disorders, show cognitive deficits as well as motor symptoms (23). Patients with focal lesions of the GPi have deficits on tests of working memory and rule-based learning (for example, Wisconsin Card Sorting Test) (24). These deficits are considered indices of frontal lobe dysfunction.

The concept that the cerebellum is involved in cognitive function is a relatively recent one which has lacked extensive experimental support (2, 3). There are a number of reports that patients with cerebellar pathology have some cognitive deficits (25). In addition, a positron emission tomography (PET) study found that an inferior and lateral part of the right cerebellar hemisphere was activated during a task that required rule-based generation of words (26, 27). This activation was spatially separate from that found during motor tasks, including speech (26). Finally, support for the







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involvement of cerebellar output in cognitive function has come from a study of functional activation in the human dentate nucleus with magnetic resonance imaging (28). In this study the dentate displayed a large bilateral activation when subjects attempted to solve a pegboard puzzle. The extent of this activation was three to four times greater than that seen during visually guided movements of the pegs.

In conclusion, our results demonstrate that cerebellar and basal ganglia outputs gain access to the prefrontal cortex. These connections provide part of the anatomical substrate for the involvement of these subcortical nuclei in cognitive processing. Thus, we believe that the cerebellum and basal ganglia should no longer be considered as purely motor structures. Instead, concepts about their function should be broadened to include involvement in cognitive processes such as working memory, rule-based learning, and planning future behavior.

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HSV1 (0.05 to 0.10 $\mu l;$ 3.2 to 8.2 \times 10 8 plaque forming units/ml) were injected at sites 1.5 to 7.2 mm below the cortical surface. Between 42 to 71 sites were injected in each animal. Animals were eutha-nized 5 days after virus inoculation, and the brain of each animal was processed to demonstrate the location of virus-specific antigen [for complete details, see P. L. Strick and J. P. Card, in Experimental Neuroanatomy: A Practical Approach, J. P. Bolam, Ed. (Oxford Univ. Press, Oxford, 1993), pp. 81-101]. All procedures were in accordance with institutional . quidelines for the care and use of animals.

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