

Regional cerebral metabolic patterns demonstrate the role of anterior forebrain mesocircuit dysfunction in the severely injured brain

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Although disorders of consciousness (DOCs) demonstrate widely varying clinical presentations and patterns of structural injury, global down-regulation and bilateral reductions in metabolism of the thalamus and frontoparietal network are consistent findings. We test the hypothesis that global reductions of background synaptic activity in DOCs will associate with changes in the pattern of metabolic activity in the central thalamus and globus pallidus. We compared 32 [¹⁸F]fluorodeoxyglucose PETs obtained from severely brain-injured patients (BIs) and 10 normal volunteers (NVs). We defined components of the anterior forebrain mesocircuit on high-resolution T1-MRI (ventral, associative, and sensorimotor striatum; globus pallidus; central thalamus and noncentral thalamus). Metabolic profiles for BI and NV demonstrated distinct changes in the pattern of uptake: ventral and association striatum (but not sensorimotor) were significantly reduced relative to global mean uptake after BI; a relative increase in globus pallidus metabolism was evident in BI subjects who also showed a relative reduction of metabolism in the central thalamus. The reversal of globus pallidus and central thalamus profiles across BIs and NVs supports the mesocircuit hypothesis that broad functional (or anatomic) deafferentation may combine to reduce central thalamus activity and release globus pallidus activity in DOCs. In addition, BI subjects showed broad frontoparietal metabolic down-regulation consistent with prior studies supporting the link between central thalamic/pallidal metabolism and down-regulation of the frontoparietal network. Recovery of left hemisphere frontoparietal metabolic activity was further associated with command following.

vegetative state | minimally conscious state | thalamocortical loops | fronto-striato-thalamic circuit

Disorders of consciousness (DOCs) following severe brain injuries arise in the setting of both cellular and circuit-level dysfunction secondary to deafferentation, neuronal death, and a wide range of changes in cellular function that remain poorly understood. Many studies demonstrate that patients with DOCs may continue to improve slowly over long time intervals (1, 2). In this context, measurements of cerebral metabolism can provide important insight into underlying mechanisms of brain function after severe brain injury (BI).

Historically, [¹⁸F]fluorodeoxyglucose-PET (FDG-PET) measurements provided the first demonstrations that the vegetative state (VS) was characterized by sharp, global reductions in cerebral metabolism to levels consistent with those measured in coma induced by pharmacologic anesthesia (3–5). FDG-PET measurements in the minimally conscious state (MCS) demonstrate similar levels of metabolic depression despite evidence of greater cerebral integrative function in this state (6–8). Assessments of regional changes in metabolism across the brain after severe BI indicate a key role for the posterior medial complex (6, 8), the cortical areas demonstrating the highest resting metabolism in the normal brain (9). Graded increases in metabolism in

the posterior medial complex index recovery from VS to normal function (6) and provide a proxy, along with global metabolic rates for overall changes in background synaptic activity in BI based on known physiological correlation of glucose metabolism with neuronal firing rates (10).

Recent anatomical studies point to a key role of structural connectivity of the thalamus and the posterior medial complex as indexing anatomical substrates correlated with graded outcomes across VS and levels of function within the category of MCS (11). These observations provide an important link to measurements of restoration of central thalamic metabolism across recovery of consciousness (12) and direct restoration of behavioral responsiveness in MCS with electrical stimulation of the central thalamus (c-TH) (13). Theoretical considerations based on known anatomical connectivity, physiological principles, and pathophysiological observations have combined into a proposed model for graded variations in metabolic activity within the anterior forebrain “mesocircuit” across DOCs (14). The mesocircuit model predicts that several related phenomena should arise at the circuit level following BI and also suggests important links to the metabolic variations observed in the posterior medial complex in DOC (8, 15). Most importantly, the model predicts that changes in overall metabolism observed in BI and generally associated with frontoparietal metabolic suppression (4, 8) should covary with a reversal of metabolic activity in the globus pallidus (GP) and c-TH.

Significance

Establishing consciousness in the human brain requires an ensemble activity of multiple cortical and subcortical structures. Following severe brain injuries, widespread deafferentation and neuronal death alter function across the corticothalamic system and its projections to the basal ganglia. In this paper, we test a theoretical model of a mesocircuit mechanism likely to arise in all severe brain injuries. Specifically, reduction of resting glucose metabolism within the central thalamus, striatum, and frontoparietal cortices and concomitant increases in metabolism within the globus pallidus as a result of disinhibition are proposed to arise across etiologies of injury. Here we verify this prediction and demonstrate a reversal of metabolic profile of central thalamus and globus pallidus that characterizes and indexes the patients' behavioral level.

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Here we test these predictions of the mesocircuit model using FDG-PET and behavior measurements in a cohort of severely BI subjects and a group of normal volunteers (NVs). We tested the related hypotheses that the metabolism of the c-TH and the striatum is reduced in severe BI compared with NV subjects and that GP metabolism is increased as a consequence of decreased c-TH outflow to both frontal cortex and striatum. Finally we test the hypothesis that in severe BI the metabolism would be most reduced over cortical regions critical to higher-order cognitive function.

Results

Comparison of Patterns Resting Metabolism in Subcortical Structures of Brain-Injured and Healthy Controls. We first sought to assess whether patterns of resting metabolism differed across subcortical structures of BI and NV groups (Fig. 1 *A–D*). As a first step, we compared the groups using a factorial ANOVA that demonstrated significant effects of mean normalized uptake values (mn-UV) for the factors group ($F = 5.7, P = 0.017$) and basal ganglia and thalamic region of interest (ROI) ($F = 59.3, P < 0.0001$), as well as in their interaction ($F = 21.9, P < 0.0001$); no lateralization effect or interaction of lateralization with group or basal ganglia and thalamic ROI appeared. Therefore, we combined results from subcortical structures of both hemispheres. One-way between-group comparison of the regional mn-UV of the functional subdivisions of the striatum and thalamus demonstrated the following differences between patients and controls: glucose metabolism of the ventral striatum (VST) and the associative striatum (AST) were significantly reduced after severe BI (mn-UV \pm SEM for VST: NV = 1.19 ± 0.11 , BI = 0.48 ± 0.08 ; $F = 21.8, P < 0.0001$; AST: NV = 1.20 ± 0.09 ,

BI = 0.54 ± 0.09 ; $F = 14.5, P = 0.003$), but metabolism of sensorimotor striatum (SMST) did not differ [NV = 1.06 ± 0.13 , BI = 1.30 ± 0.10 ; $F = 1.5, P =$ not significant (NS)]. A significant increase of mn-UV for the GP is present for the BI group (NV = -0.49 ± 0.10 , BI = 0.52 ± 0.07 ; $F = 48.3, P < 0.0001$), whereas the c-TH showed a significant reduction of glucose metabolism for the BI group (NV = 0.43 ± 0.07 , BI = -0.34 ± 0.06 ; $F = 45.2, P < 0.0001$), and the noncentral thalamus (non-c-TH) remained unaffected (NV = -0.25 ± 0.08 ; BI = -0.23 ± 0.05 ; $F = 0.8, P =$ NS; Fig. 2*A*). The relationship of these significant changes in GP and c-TH metabolic rates of the BI subjects importantly represents an exact reversal of the pattern demonstrated by the healthy controls (*Discussion*).

As a second step of statistical verification, a post hoc Bonferroni-corrected analysis of the two-way ANOVA (within group and ROI) demonstrated (*i*) a significant reduction of the metabolism in VST and AST compared with SMST for the BI group only; (*ii*) an increase in GP metabolism for the BI group (showing an equivalent metabolism compared with AST and VST) compared with NV group (where GP metabolism is significantly lower than in the VST, AST, and SMST); (*iii*) a significant difference between c-TH and GP for both BI and NV groups with a clear pattern reversal (see above); and (*iv*) an equivalence of c-TH and non-c-TH metabolism for the BI group in contrast with a significantly higher metabolism observed in c-TH compared with non-c-TH for NV (Fig. 2*B*).

We next carried out a multiple regression analysis using only the initial FDG-PET for the six patients longitudinally tested to further assess the relationship between c-TH glucose metabolism against VST, AST, SMST, and non-c-TH ($R^2_{adj} = 0.72, F = 35.9, P < 0.0001$). In this omnibus comparison, a significant inverse correlation of c-TH and GP glucose metabolism was identified ($F = 23.3, P < 0.0001$; Durbin-Watson = 1.97, serial autocorrelation = -0.014): higher GP glucose metabolic rates covaried with lower c-TH glucose metabolic rates (Fig. 2*C*).

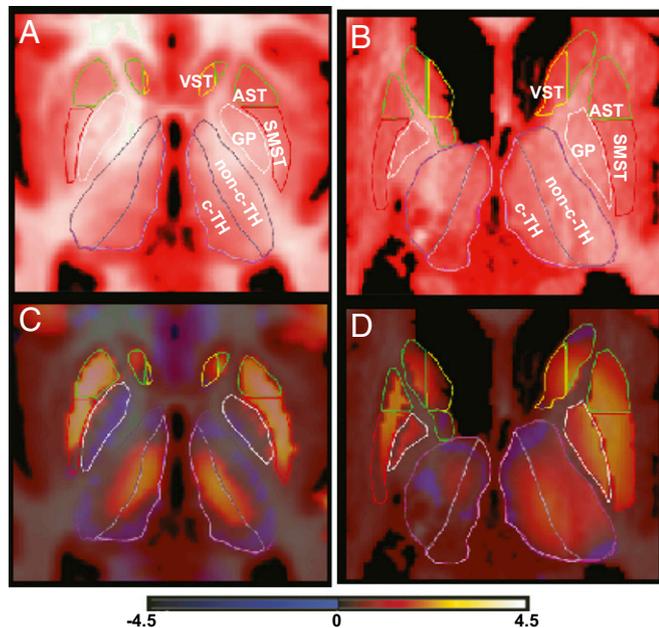


Fig. 1. (*A* and *B*) Axial anatomical high-resolution MRI in representative NV and BI subjects, respectively, showing locations and manual delimitation for ventral, associative, and sensorimotor striatum (VST, AST, and SMST, respectively), globus pallidus (GP), central and noncentral thalamus (c-TH and non-c-TH, respectively), and color scheme for regions of interest (VST, yellow; AST, green; SMST, red; GP, white; c-TH, magenta; non-c-TH, violet). (*C* and *D*) [^{18}F]FDG-PET, T1-MRI fusion in representative NV and BI subject, respectively. NV demonstrates symmetric pattern of relatively increased c-TH metabolism compared with non-c-TH. A marked asymmetry of thalamic metabolism with loss of contrast in c-TH compared with non-c-TH metabolism is evident in BI subject.

Cortical Glucose Metabolism After Severe BI. We compared the regional glucose metabolism across cortical ROIs selected for analysis. A three-way ANOVA demonstrated significant effects of factors group ($F = 244.9, P < 0.0001$), ROI ($F = 33.2, P < 0.0001$), and side ($F = 8.5, P = 0.0035$), as well their interactions ($F = 2.4, P = 0.003$). Accordingly, for comparisons of cortical metabolism, left and right results are reported separately (Fig. 3 and *Table S1*). One-way ANOVA showed a significant reduction of glucose metabolism in the BI group bilaterally across frontal cortical regions [i.e., orbitofrontal cortex (OFC), ventromedial prefrontal cortex (vmPFC), dorsolateral prefrontal cortex (dlPFC), and premotor cortex (PMC)], with the exception of the primary sensorimotor cortex (SM1) (Fig. 3 *A* and *B*). Deep-seated mesial cortical structures including the anterior cingulate cortex (aCGc) did not show group differences; however, the medial cingulate cortex (mCGc), the precuneus, and the left posterior cingulate cortex (pCGc) show a marked symmetric reduction of glucose metabolism in BI. Finally, the BI group showed significant reductions of metabolism of posterior cortical structures on the lateral surface [i.e., bilateral superior posterior parietal cortex (sPPc), right inferior posterior parietal cortex (iPPc), and primary visual cortex (V1)]. No alteration of glucose metabolism was observed on the posterior one-third of the superior temporal sulcus (post-1/3 STS).

Multiple regression analysis of all cortical ROIs only showed a moderate significant linear relationship ($R^2_{adj} = 0.56, F = 4.3; P < 0.005$) between the right c-TH/aCGc ($\beta = -0.58; P < 0.01$) and the c-TH/post-1/3 STS ($\beta = -0.53; P < 0.01$) and on the left side ($R^2_{adj} = 0.50, F = 3.5; P < 0.01$) between the c-TH/post-1/3 STS ($\beta = -0.55; P < 0.01$). Because activity in the precuneus has been specifically associated with recovery of consciousness (11), we further tested in a simple regression model its metabolic interactions with c-TH. Results show a (+) linear relationship

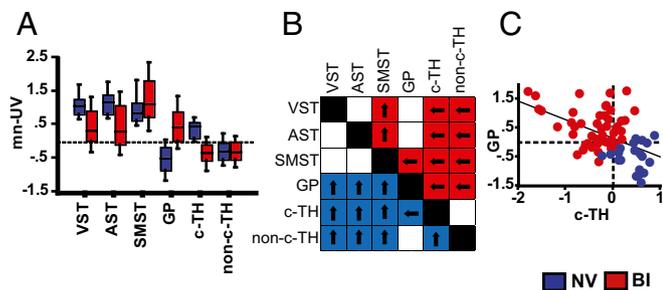


Fig. 2. Group data displaying mn-UV of glucose metabolism in deep brain structures measured in NV and BI subjects. (A) Box plot. A significant reduction in relative glucose metabolism of the ventral striatum (VST), associative striatum (AST), and central thalamus (c-TH) in BI subjects is seen compared with NV. No difference in sensorimotor striatum (SMST) mn-UV is present between NV and BI subjects. A significant increase in globus pallidus (GP) metabolism is present in the group of BI subjects. (B) Contrast showing the results of the post hoc Bonferroni corrected analysis of a two-way ANOVA (group and ROI). Significant results are shown in blue for NV and red for BI patients, whereas white boxes denote no significant differences; arrows indicate the direction of the significance (i.e., pointing toward the higher mn-UV values). (C) Bivariate scattergram demonstrates an inverse linear correlation between glucose metabolic rate of the c-TH (x axis) and the GP (y axis), $P < 0.001$ (between groups).

between metabolism in precuneus and c-TH ($F = 18.449$; $P < 0.0001$; Durbin-Watson = 2.23; serial autocorrelation = -0.12).

Separation of BI Subjects with Evidence for Command Following: Subcortical Structures. As a separate comparison, we segregated the BI group into subsets of subjects capable of demonstrating command following from those unable to show behavioral evidence of command following at the bedside (Fig. 4). This analysis followed an initial evaluation of the Coma Recovery Scale Revised (CRS-R) total scores that did not significantly covary with the brain glucose metabolic rates across the patient subject pool. Membership assignments within these two patient groups were based on the cutoff of the auditory function scale of the CRS-R: command following and no command following. The no command following included 3 patients in VS and 10 subjects in the MCS(-) classification, whereas all command following patients fulfill the MCS(+) category (16) (we excluded patients that emerged from MCS and were in locked-in syndrome who were not tracked on this measure). One-way comparisons between the subgroup of BI subjects able to follow commands and those unable to follow commands demonstrated a difference in glucose metabolism of the c-TH; c-TH demonstrated significantly increased glucose metabolism for the command following subgroup (command following = -0.13 ± 0.06 vs. no command following = -0.45 ± 0.09 ; $F = 8.3$, $P = 0.005$). An opposite finding was present for the VST (command following = 0.26 ± 0.12 vs. no command following = 0.61 ± 0.09 ; $F = 5.6$, $P = 0.02$; Fig. 4A). To assess the linkage of the expressed pattern of c-TH/GP metabolic reversal to behavioral profiles, we created a ratio for c-TH and GP glucose metabolism using the individual raw standardized uptake value (SUV) (not normalized to the global mean). One-way ANOVA comparisons of c-TH/GP ratio between NVs and the BI command following and no command following groups showed that those BI subjects who followed commands demonstrated a significantly higher c-TH/GP ratio even though they also significantly differed from NVs [command following = 0.89 ± 0.031 vs. no command following = 0.75 ± 0.034 ($P = 0.0019$) and vs. NV = 1.26 ± 0.030 ($P \leq 0.0001$); $F = 59.842$, $P < 0.0001$; Fig. 4B].

Separation of BI Subjects with Evidence for Command Following: Cortical Regions. A one-way ANOVA comparison of mn-UV of cortical regions between the command following and no

command following groups showed a significant increment of the regional glucose metabolism in three left-sided cortical regions for the command following group: the SM1 (command following = 0.73 ± 0.09 vs. no command following = 0.30 ± 0.10 ; $F = 7.4$, $P = 0.007$), the iPPc (command following = 1.07 ± 0.08 vs. no command following = 0.32 ± 0.13 ; $F = 23.0$, $P < 0.0001$), and the sPPc (command following = 0.04 ± 0.09 vs. no command following = -0.55 ± 0.21 ; $F = 6.5$, $P = 0.017$). A significant difference between command following and no command following was also observed for the right OFC, which showed a reduction of glucose metabolism in the command following group (command following = -0.50 ± 0.10 vs. no command following = 0.02 ± 0.09 ; $F = 15.2$, $P = 0.0001$; Fig. 4C).

Discussion

The main finding of the present study is that specific changes in the pattern of resting metabolic activity measured by FDG-PET across the individual substructures of the striatal-pallidal-thalamocortical circuit demonstrates clear differences in severe BI patients with DOCs in comparison with NVs.

Reversal of Resting Metabolic Profile of Central Thalamus and Globus Pallidus Characterizes the Severely Injured Brain and Supports the Key Role of the Anterior Forebrain Mesocircuit. A recently proposed model predicts that global reductions of background synaptic activity in DOC subjects will associate with a specific pattern of metabolic down-regulation in the c-TH and GP (14).

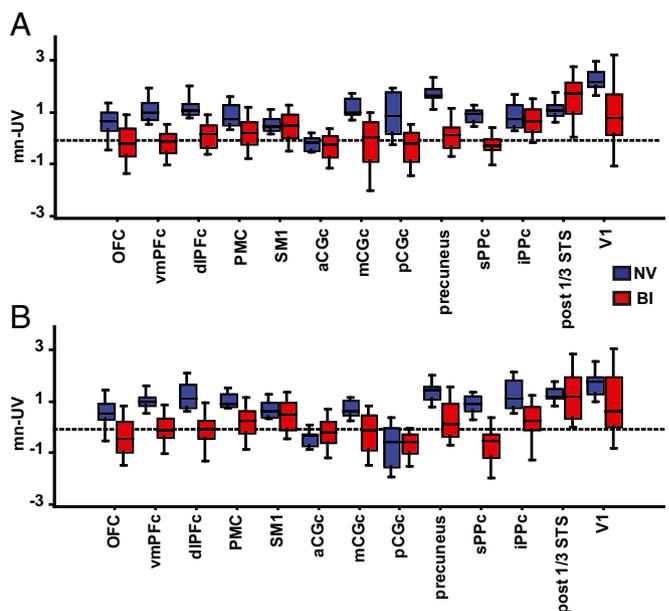


Fig. 3. Group data displaying mn-UV of regional cortical glucose metabolism in NV and BI subjects. (A) Box plot for the left hemisphere cortical regions showing a significant reduction of glucose metabolism across anterior frontal cortices, medial cingulate cortex (mCGc)/precuneus, and posterior parietal cortices in BI subjects. Note a distinct pattern of rostrocaudal progression of increasing mn-UV glucose metabolic values in BI patients as arranged [orbitofrontal, ventromedial, and dorsolateral prefrontal cortices, premotor cortices and primary sensorimotor (OFC/vmPFC/dIPFC/PMC/SM1, respectively)] that converges with NV mn-UV at SM1. BI subjects demonstrate significant decreases in superior posterior parietal cortex (sPPc), posterior cingulate cortex (pCGc), precuneus, and V1. (B) Box plot for the right hemisphere showing similar pattern of rostrocaudal progression of increasing mn-UV glucose metabolic values in BI patients as arranged (OFC/vmPFC/dIPFC/PMC/SM1). Note that right pCGc does not show differences in observed for left hemisphere in posterior structures (for significance, see Table S1).

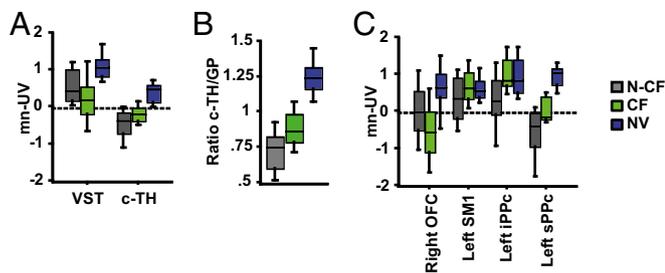


Fig. 4. Group data displaying mn-UV of glucose metabolism for deep brain and cortical structures comparing subsets of severe BI patients demonstrating command following (CF; green), those in the noncommand following (N-CF) group (gray), and NV (blue). (A) Box plot of two deep brain structures showing a significant difference in glucose metabolism. Increases of mn-UV are seen in c-TH for the CF group compared with N-CF, whereas decreases in VST are present. (B) Box plot of the ratio c-TH/GP of raw standard uptake values body weight normalized (i_{c-TH}/GP); index separates the CF and N-CF groups with increased ratio in CF group that remains lower than NV. (C) Box plot of four cortical structures showing a significant difference in glucose metabolism between the two groups. A significant increase of glucose metabolism in the CF group is present for SM1, iPPc, and sPPc in the left hemisphere. An opposite metabolic pattern is seen for the right OFC, which shows a reduction of metabolism for CF group. This finding is consistent with that observed in the VST (A).

Specifically, as a result of the high-threshold UP states of the medium spiny neurons in the striatum (17), it is probable that in the setting of broad deafferentation of corticostriatal and thalamostriatal inputs that the striatal inhibition of the GP will weaken and allow increased pallidal activity and consequent inhibition of c-TH (SI Text and Fig. S1). Our findings of reduced c-TH metabolism and increased GP metabolism in DOC subjects compared with the NVs supports this mesocircuit hypothesis.

The c-TH is known to have dopamine (DA) expression with levels approximating those found in the substantia nigra (SN) (18, 19) and a preponderant distribution of D_2 -like receptors (i.e., D_2 and D_3) (20). Indeed, c-TH DA may constitute a fourth dopaminergic pathway (21) sometimes referred as mesothalamic (22, 23). Therefore, we designed an indirect method to segment the thalamus into two main subregions, the c-TH and the non-c-TH, using a dopamine D_2 -like receptor ligand template (SI Text). We were able to separate the most representative map of D_2 -like thalamic receptors in NVs with [^{11}C]raclopride to show then that if this map is applied to analyze the glucose metabolism in NVs, a clear higher metabolism exists in the c-TH when contrasted to the noncentral aspect of the structure. More importantly, when this same thalamic subdivision was implemented in severe BI patients and results were contrasted to those obtained from NVs, there was a metabolic reduction strictly confined to the c-TH. Moreover, when the group of patients was divided based on their ability to follow commands, c-TH metabolism was higher in the group of responders. Hemodynamic thalamic dysfunction in DOCs has long been shown (24) and is further supported by metabolic studies using [^{18}F]-FDG (5, 25). However, this is to our knowledge, the first evidence of a sustained deficit in glucose metabolism after severe BI at a substructural thalamic level (here, the c-TH).

Here, the striatum was functionally divided based on its cortical loops (SI Text). We find that severe BI patients demonstrate a disproportionate striatal metabolic deficit compromising selectively the VST and AST but sparing the SMST. This distinction in metabolic profiles can be related to our findings of relatively preserved cortical metabolism in SM1, which shows no difference across BI and NV groups. A relative integrity of SM1 neurons and their activity likely accounts for the maintained

mn-UV in SMST. Conversely, AST and VST compartments of the striatum are linked to frontal association cortices known to be functionally engaged during highly elaborated cognitive processing (26–28), all of which show clear metabolic reduction in BI; loss of corticostriatal and thalamostriatal input from c-TH is proposed to account for these measured differences.

There are clear distinctions in the functional relationship of thalamic nuclei and the striatum. Thalamostriatal networks are divided into two segregated subsystems: the centromedian/parafascicular complex striatal projection and the non-centromedian/parafascicular striatal projection (29). Although the thalamic segmentation implemented in this work has mostly likely aggregated the c-TH to include the centromedian/parafascicular complex along with additional anterior midline structures, our results in severe BI patients are supportive of a major selective failure of both the main thalamostriatal and corticostriatal projections to the associative and limbic brain with a preservation of the sensorimotor system. Specifically, this inference is supported by (i) the low metabolism of the c-TH, VST, and AST; (ii) the relatively equal metabolic profiles of the SMST and SM1 in the NV and BI groups; (iii) the lack of alteration of the non-c-TH metabolic profile; and (iv) the observed metabolic increases in GP that show inverse correlation with the c-TH metabolic deficit evident in severe BI patients, suggesting a greater vulnerability of the parafascicular over the centromedian within the centromedian/parafascicular complex and its striatal projection (30).

Fronto-Parietal Cortical Metabolic Down-Regulation in the Severely Injured Brain: Evidence for Lateralization and Mechanistic Link to Subcortical Metabolism.

Similar to prior FDG-PET findings in patients with DOCs, we found here a marked frontoparietal metabolic down-regulation, most prominent in the precuneus (Fig. 3). The present findings extend these observations and provide a demonstration of hemispheric lateralization in frontoparietal metabolic rates distinguishing those patients able to follow commands from those who cannot. Increased metabolism in cortical regions correlated with this distinction are primarily in the left posterior parietal cortex and its connections with the SM1, with the exception of a relative suppression of metabolic activity in the right OFC. A similar dissociation in the metabolic rates of subcortical structures is also evident across the two patient groups, with increased c-TH activity linked to the command following group along with relatively suppressed VST metabolic activity. Collectively, the findings indicate that the patient subjects in the no command following group have both a down-regulation of the left posterior parietal cortex and posterior medial complex (6, 8), as well as an excess of disinhibition of limbic frontal cortices (Fig. 4C). In addition, a more global down-regulation of the anterior forebrain indexed by the c-TH/GP ratio in the no command following group is also present (Fig. 4B). Both of these findings may be related to the metabolic suppression of the c-TH. Anatomical and physiological studies have established the close link between the c-TH and the posterior medial cortical regions; such studies include track tracings in nonhuman primates (31) and demonstrations that the c-TH and precuneus show covarying decreases in regional blood flow during anesthetic coma compared with wakeful baselines that reverse together when consciousness is briefly restored (32). Importantly, our findings here that a linear relationship exists between metabolism in precuneus and c-TH supports this physiological linkage of the central thalamus to the key cortical structure demonstrating highest resting metabolism and correlation with level of recovery of consciousness (6, 9, 33). Thus, the present findings provide further detail of the known graded increases in cerebral metabolic rate in the posterior medial complex indexing recovery from VS to normal function (6) and suggest that regional patterns of metabolic rate within the

frontoparietal network may be informative of mechanisms underlying differences in clinical presentation of patients with DOC. Another detail evident in the differences in frontoparietal metabolism across the patients is a metabolic gradient with apparently more pronounced metabolic differences across a rostrocaudal distribution from prefrontal cortex to SM1 (Fig. 3); this finding is also consistent with the mesocircuit model and comparable to a tendency for the more anterior prefrontal cortical regions to demonstrate reduced blood flow under anesthesia, sleep, and even early awakening sleep inertia (34, 35). Collectively, these findings support a mechanistic link between the key role of the posterior medial complex in outcomes of DOCs (36), restoration of normal resting metabolic activity (33), and restoration of anterior forebrain mesocircuit function via activity of the c-TH (13, 14).

Therapeutic Implications of Metabolic Profiles of the Severely Injured Brain: Toward Molecular Neuroimaging of the Mesocircuit Model of DOCs. Our findings suggest implications for therapeutics at three different sites of action, c-TH, striatum, and GP, and point to testable predictions for future studies that use receptor binding ligands, PET, and pharmacologic intervention.

Dopaminergic drugs may directly increase the dopaminergic tone within the c-TH, enhancing its output (37, 38). It is also likely that dopaminergic drugs at the striatum may counterbalance the observed deficits in the direct and indirect pathway (via D_1 and D_2 , respectively) (39), and the net gain would be the reduction of the inhibitory input to the c-TH. Last, dopaminergic drugs can directly stimulate D_1 -postsynaptic receptors at the GP and consequently disinhibit the c-TH or stimulate D_2 -postsynaptic receptors to indirectly inhibit GABAergic pallidothalamic neurons (40). In addition, it is possible to directly target GABAergic- α -1 receptors distributed at the GP to suppress their activity (14, 34) as reported in some severely BI patients with zolpidem (41, 42). Finally, the findings provide further evidence of the mechanistic basis for arousal regulation effects of activation of the c-TH by electrical stimulation (13, 43) or indirectly via regularization of the globus pallidus interna (44–46), which has also been shown to be effective in increasing arousal states.

Limitations of Our Study. The heterogeneity in our group of patients in terms of lesion source, location, extension, and clinical expression may have hidden some relevant changes occurring at some important levels of the mesocircuit (i.e., AST), principally when comparisons were done between the two groups of patients. In addition, the preserved relative metabolism of SMST in the BI subjects may raise concerns about a bias of increased measured GP metabolism arising from the SMST due to a partial volume effect. Two observations mitigate this concern: first, we find that SMST was equally active in the pool of severe BI patients and NVs. Thus, a similar spill-in to the GP of NVs would have been expected as well but did not occur, with the exception that structural demarcations are clearer in the NV group. Second, after correction for partial volume effect (*SI Text* and Fig. S2) demonstrated that the significant difference in GP between patients and controls remained intact with no evidence of difference of SMST glucose metabolism. Limitations in our results may also arise from quantitative differences in acquisition due to the use of different PET cameras in NV and BI subjects, although prior studies indicate such effects are very small, if present at all (47, 48). We believe that the almost equal glucose metabolic pattern observed between our severe BI patients and controls over SMST and SM1, the reversal of the pattern of c-TH/GP, and the opposing patterns observed with respect to associative frontoparietal cortices and VST-AST make it very unlikely that a systematic source of variation could account for our findings on a nonphysiological basis. Finally, our results may be biased by the introduction of a manual determination of the deep brain

structure target ROIs. However, manual anatomical determinations of these areas were carried out before the MRI-PET coregistration and were not reoriented thereafter based on the metabolic display on FDG-PET.

Materials and Methods

Participants. A prospective sample of 24 severely BI patients with varying etiologies, who fulfilled criteria for VS, MCS, or emerged from MCS, were assessed at Weill Cornell Medical College (Table 1). A total set of 32 PET studies individually treated were included in the final group analysis, and clinical examination was performed using CRS-R (49) (Table S2). A group of 10 NVs (4 females/6 males; 45.2 ± 14.3 y) recruited at the Faculty of Medicine of the University of Liège acted as the age-matched control group. All patients' informed consent was obtained from their legally authorized representative and the medical team. All NVs gave informed consent under the University of Liège Ethics Committee. Ethical approval for the study was provided by the Weill Cornell Medical College Institutional Review Board.

MRI Data Acquisition. A structural anatomical image was acquired on each severe BI patient on a 3-T GE MRI using a T1 3D fast spoiled gradient echo sequence (matrix size, $256 \times 256 \times 120$; voxel size, $0.93 \times 0.93 \times 1.2$ mm; note: in the first four studies, parameters were $256 \times 256 \times 156$; voxel size, $0.93 \times 0.93 \times 1.0$ mm). NV data were acquired on a 3-T Siemens MRI using a T1 3D magnetization prepared rapid gradient echo sequence (matrix size, $256 \times 256 \times 176$; voxel size, $0.9 \times 0.9 \times 0.9$ mm).

PET Data Acquisition. Each BI patient underwent injection of [^{18}F]-labeled FDG (~ 370 MBq). All PET studies but four were acquired on a GE PET-CT LS Discovery unit under a standard resting condition in 2D dynamic high-sensitivity emission mode, matrix size = $128 \times 128 \times 35$, axial field of view = 25 cm, 4.25-mm slice thickness (studies 29–32 on a Siemens Biograph mCT scanner matrix size = $512 \times 512 \times 83$). NV data were acquired after i.v. injection of ~ 370 MBq FDG on a Siemens CTI 951 R16/31 scanner as described in ref. 50.

Using PMOD v.3.309 (PMOD Technologies), SUV was computed, normalized by body weight, and scaled to the global mean signal (mn-UV) across the region defined by their skull-stripped T1-MRI (51), similarly to the first step involved during spatial covariance analysis of resting state metabolic images (52). Coregistration to each subject's MRI was done using a rigid matching method using the PET to determine the transform (dissimilarity function: normalized mutual information with a trilinear interpolation method). Last, smoothing (6-mm full width at half maximum) was applied to the PET data.

Overview of the Approach to [^{18}F]-FDG-PET ROI Definition. To assess the hypotheses tested in this study, we required a detailed evaluation of metabolism within the subcortical structures of the basal ganglia and thalamus. As standard atlas approaches are inadequate to capture the marked anatomical variation present across patients with severe BI (i.e., striatal and/or thalamic templates overlaid the lateral ventricles), we carried out several initial methodological steps using structural and metabolic data obtained from our NVs and from reanalysis of another previously published dataset of normal subjects from a study of D_2 -like receptor expression using [^{11}C]raclopride [8 NVs; 26.0 ± 5.5 y; 8 females (53)]. For all normal subjects and patient analyses, striatal and thalamic ROIs were hand drawn onto 3D MRI (Fig. 1 and *SI Text*).

The thalamic ROIs in NV and BI subjects were first defined manually and then divided into a central and noncentral component (c-TH and non-c-TH,

Table 1. Demographics

Variable	Result
Total number of PET studies	32
Total number of patients	24
PET studies per patients: 1/2/3 studies	18/4/2
Age (y)	40.3 ± 12.8
Sex (female/male)	13/19
Dx (VS/MCS/EMCS)	3/2/7/2
Etiology (TBI/AE/miscellaneous)	19/6/7
Months postinjury	77 ± 84.9
CRS-R	12.7 ± 1.0
MCS command following (yes/no)*	13/14

*Not included: two emerged from MCS; one locked-in syndrome, and two cases with data not recorded.

respectively) based on the use of standard structural automatic segmentation results (SI Text and Fig. S3) and the pattern of D₂-like receptors expression obtained using [¹¹C]raclopride to identify the boundaries of the c-TH.

To obtain cortical ROIs, previously defined cortical regions were extracted using a standard nonrigid normalization to the MNI space (54) and then applied, using an inverse transformation, to the individual NV or BI subjects' natural space (SI Text and Fig. S4).

Statistical Analysis. For the final analysis, the individual ROI mn-UV was used. All data were assessed for normality [Shapiro-Wilk test, $W = 0.99 >$ critical value of $W = 0.947$ (5% significance level)]. For statistical analysis, factorial

ANOVA design with group (NV/BI), side, and ROI was used. A one-way ANOVA was then computed to compare differences between groups (combined for those without lateralization differences). Multiple regression analysis was performed using the c-TH as the dependent variable. Results were considered significant at $P < 0.05$. Bonferroni correction was applied for within-group multiple comparisons. Last, we created a post hoc ratio between the raw SUV (i.e., not normalized to the global mean) in the c-TH and GP and contrasted between groups using a one-way ANOVA.

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