Glucagon-like peptide-1 receptor stimulation increases blood pressure and heart rate and activates autonomic regulatory neurons

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Glucagon-like peptide-1 (GLP-1) released from the gut functions as an incretin that stimulates insulin secretion. GLP-1 is also a brain neuropeptide that controls feeding and drinking behavior and gastrointestinal emptying and elicits neuroendocrine responses including development of conditioned taste aversion. Although GLP-1 receptor (GLP-1R) agonists are under development for the treatment of diabetes, GLP-1 administration may increase blood pressure and heart rate in vivo. We report here that centrally and peripherally administered GLP-1R agonists dose-dependently increased blood pressure and heart rate. GLP-1R activation induced c-fos expression in the adrenal medulla and neurons in autonomic control sites in the rat brain, including medullary catecholamine neurons providing input to sympathetic preganglionic neurons. Furthermore, GLP-1R agonists rapidly activated tyrosine hydroxylase transcription in brainstem catecholamine neurons. These findings suggest that the central GLP-1 system represents a regulator of sympathetic outflow leading to downstream activation of cardiovascular responses in vivo.

satiety or acts as a mediator of aversive stimuli remains uncertain, as several experiments demonstrate development of taste aversion (12, 13) and activation of signaling pathways mediating interoceptive stress responses (14–16) following GLP-1 administration. Consistent with the involvement of GLP-1 signaling in stress responses, central administration of GLP-1 activates neurons containing corticotropin-releasing hormone (CRH) in the PVH and elevates plasma corticosterone (17). Furthermore, both GLP-1 and EXN-4 produce chronotropic and pressor responses in anesthetized animals (18–21); however, whether these effects are direct, due to activation of cardiac GLP-1Rs, or indirect remains unclear. Despite the accumulating data linking GLP-1R signaling to autonomic and neuroendocrine responses, the neural pathways underlying these actions of central GLP-1 are not fully understood. To determine the mechanisms and potential physiological significance of central GLP-1 pathways impinging on sympathetic outflow systems, we investigated the distribution of GLP-1–activated neurons that innervate sympathetic preganglionic neurons and assessed GLP-1 effects on cardiovascular parameters.

Methods

**Animals and surgical procedures.** Adult male pathogen-free Sprague-Dawley rats (250–350 g; Taconic, Germantown, New York, USA) were housed in a light-controlled (12 hours on, 12 hours off) and temperature-controlled (21.5–22.5°C) environment. The animals and procedures used were approved by the Harvard Medical School and Beth Israel Deaconess Medical Center institutional animal care and use committees.

Rats were anesthetized with isoflurane (induction 5%, maintenance 2%) and were implanted in the abdominal aorta with a catheter coupled with a sensor and transmitter (TA11PA-C40; Data Sciences International, St. Paul, Minnesota, USA) (22). Telemetry signals were received by an antenna below the rat’s cage that relays the signals to a signal processor (DataQuest A.R.T. Gold version 2.1 system; Data Sciences International) connected to a Compaq Deskpro PC (Compaq Computer Corp., Houston, Texas). These transmitters allow remote monitoring of blood pressure (BP) and heart rate (HR) in unanesthetized and unrestrained animals. In the same surgical session, rats were implanted intravenously (i.v.) with a silastic catheter into the femoral vein (23) or i.c.v. with a guide cannula into the lateral ventricle (24). In a subset of the animals, the retrograde tracer Fluorogold (FG) (2.5% in pyrogen-free saline) (Fluorochrome, Denver, Colorado, USA) was injected with a glass micropipette and air pressure injection system (25, 26). The injections included the intermediolateral cell column (IML) and consisted of two injections of 50 nl in the rostral (T1–T4) or caudal (T6–T10) thoracic spinal cord.

Ten to fourteen days after surgery, base-line mean arterial blood pressure (MAP) and HR were recorded for the preceding day and night, and rats were injected with EXN-4 (30, 300, or 3000 ng per rat i.c.v.; n = 4–6 for each group) or pyrogen-free saline (PFS) (0.25 ml i.v. or 5 µl i.c.v., total volume per rat; n = 4–6 for each group). In our initial experiments, we observed that both i.v. and i.c.v. injections of PFS induced a transient increase in MAP and HR. Thus, we used a long line to avoid a handling stress during injection in a subsequent experiment. The long line was connected 2 hours before i.c.v. injection of GLP-1(7-36) (300 ng per rat; n = 3) or PFS (5 µl total volume per rat; n = 3). The action of [des-His, Glu]exendin-4 of the GLP-1R antagonist (13) was investigated by i.c.v. administration of 1 µg of the peptide 30 minutes before i.c.v. administration of 30 ng of EXN-4. All drugs were given to unrestrained rats between 2 and 5 hours after lights on. The cardiovascular data were collected and stored in 1-minute bins, and, before further analysis, the data were averaged into 10-minute bins. Results are expressed as means ± SEM from groups of four to six rats. Statistical comparisons between animals were performed at each time. ANOVA for repeated measurement was used to estimate dose-dependent effects within each group, and the differences between groups in each time were assessed by Fisher’s protected least significant difference (PLSD).

**Animal perfusion and histology.** Two hours after administration of peptides or PFS, rats were deeply anesthetized with isoflurane and perfused transcardially with diethyl pyrocarbonate–treated 0.9% saline followed by 10% neutral formalin (Sigma-Aldrich, St. Louis, Missouri, USA). For tyrosine hydroxylase (TH) heteronuclear RNA (hnRNA) in situ hybridization studies, rats were perfused 0, 15, or 30 minutes or 1, 2, 6, or 24 hours after administration of EXN-4 or PFS (n = 3 for each group). Before perfusion, blood was collected from the heart, and blood glucose was measured using a glucose oxidase–based method (LifeScan OneTouch; Johnson & Johnson, Milpitas, California, USA). Tissues were removed and postfixed in the same fixative for 4 hours, submerged in 20% sucrose, and cut at 30 µm (1.5 series).

**Immunohistochemistry.** The procedures for immunohistochemistry were performed as reported previously (23, 26, 27). For single-label immunohistochemistry, sections were pretreated with 0.3% hydrogen peroxide in PBS for 30 minutes, and then in 3% normal donkey serum (Jackson ImmunoResearch Laboratories Inc., West Grove, Pennsylvania, USA) with 0.25% Triton X-100 in PBS (PDT), followed by overnight incubation in Fos rabbit primary antisera (Ab5; Oncogene Research Products, Cambridge, Massachusetts, USA; 1:100,000 dilution in PDT) at room temperature. After washing in PBS, sections were incubated in biotinylated donkey anti-rabbit IgG (Jackson ImmunoResearch Laboratories Inc.; 1:1,000) and then incubated with avidin-biotin complex (Vector Elite ABC Kit; Vector Laboratories Inc., Burlingame, California, USA; 1:500 in PBS). A combination of 0.04% diaminobenzidine tetrahydrochloride (DAB; Sigma-Aldrich), 0.01% nickel ammonium sulfate (Fisher Scientific Co., Pittsburgh,
Pennsylvania, USA), 0.01% cobalt chloride (Fisher Scientific Co.), and 0.01% hydrogen peroxide dissolved in PBS was used for the chromogen reaction with two successive rinses in PBS. The tissue sections were mounted onto subbed slides, air-dried, dehydrated in alcohol, cleared in xylene, and then coverslipped with Permaslip (Alban Scientific, St. Louis, Missouri, USA).

For double-label immunohistochemistry, the sections stained with Fos were blocked and then incubated overnight in FG rabbit primary antisera (Chemicon International, Temecula, California, USA; 1:20,000 in PDT) or mouse anti-TH (Chemicon International; 1:20,000 in PDT). The sections were processed as described above using DAB as a chromogen.

For triple-label immunohistochemistry, the tissue sections were incubated in Fos antisera and were processed as described above but incubated in the DAB solution without nickel sulfate and cobalt chloride, resulting in a brown nuclear reaction product. Sections were then incubated in FG rabbit primary antisera and mouse anti-TH overnight at room temperature. After washing in PBS, sections were incubated in FITC-conjugated donkey anti-rabbit IgG (Jackson ImmunoResearch Laboratories Inc.; 1:200 in PDT) and Cy3-conjugated donkey anti-mouse IgG (Jackson ImmunoResearch Laboratories Inc.; 1:200 in PDT). The Fos-IR was observed with brightfield optics, the TH-immunoreactivity (TH-IR) with a rhodamine filter system, and the retrograde label with a FITC filter system.

In situ hybridization histochemistry. A fragment of the rat TH gene (28) including intron 2 was amplified by PCR with rat genomic DNA (BD Biosciences Clontech, Palo Alto, California, USA) as a template. The following primers were used for amplification: primer 1, 5′-TGTCTCGGGCTGTAAAAGTATTTGA-3′; and primer 2, 5′-AGTGGTGAAATTTGGCTTCAAATGT-3′. The insert was cloned into the pGEM-T Easy Vector (Promega Corp., Madison, Wisconsin, USA). The PCR product was cloned and the identity of the insert as TH intron was confirmed by sequence analysis and comparison to the GenBank database (accession number AF014956). For generation of 550 bp of antisense 35S-labeled cRNA, the plasmid was linearized by digestion with NcoI and subjected to in vitro transcription with SP6 polymerase. The protocol for in situ hybridization histochemistry was a modification of that previously reported by our laboratory (29) and several other laboratories (30). Control procedures to confirm the specificity of our in situ hybridization histochemistry procedures included hybridization with antisense probes after tissue pretreatment with RNase A (200 µg/ml).

Estimates of cell counts. In all procedures, the estimates of cell counts (Fos, TH, and FG) were done with a grid reticule and a 10× objective by an observer blinded to the treatment groups (23, 26). The data were not corrected for double counting and a stereological technique was not used, because the objects we were counting (nuclei and retrogradely labeled cells) did not change in size, and section thickness did not vary between groups. Hence, as all double-label studies are inherently qualitative, our results are meant to provide relative data but are not meant to be accurate estimates of absolute cell counts. Data were analyzed by ANOVA and differences between groups by Fischer’s PLSD.

Production of photomicrographs. Photomicrographs were captured with a digital camera (Spot3.0; Diagnostic Instruments, Sterling Heights, Michigan, USA) mounted directly on the microscope (Axioskop 2; Carl Zeiss Inc., Thornwood, New York, USA) and an Apple Macintosh G3 (Apple Computer Inc., Cupertino, California, USA) computer. Image editing software (Adobe Photoshop; Adobe Systems Inc., Mountain View, California, USA) was used to combine photomicrographs into plates. Only the sharpness, contrast, and brightness were adjusted. All figures were printed on a dye-sublimation printer (Kodak 8670; Eastman Kodak Co. Scientific Imaging Systems, New Haven, Connecticut, USA). For drawings (see Figure 4), cytoarchitectonic details were added using a camera lucida.

Results

GLP-1 agonists increase BP and HR. To ascertain the effects of GLP-1R activation on cardiovascular responses in non-anes-
In anesthetized rats, we assessed BP and HR in freely moving rats using telemetric monitoring. MAP and HR transiently increased after handling and after i.v. or i.c.v. administration of PFS. Both parameters returned to basal levels within 40–50 minutes (Figure 1, a–d). In contrast, i.v. administration of EXN-4 produced a dose-dependent increase in MAP and HR (Figure 1, a and b). Similarly, i.c.v. administration of EXN-4 dose-dependently increased MAP and HR (Figure 1, c and d). MAP and HR remained significantly increased for up to 2 hours after both i.v. (300 and 3000 ng) and i.c.v. (30 and 300 ng) administration of EXN-4.

As EXN-4 is a protease-resistant long-acting lizard GLP-1R agonist, we also examined MAP and HR in freely moving rats using telemetric monitoring. MAP and HR transiently increased after handling and after i.v. or i.c.v. administration of PFS. Both parameters returned to basal levels within 40–50 minutes (Figure 1, a–d). In contrast, i.v. administration of EXN-4 produced a dose-dependent increase in MAP and HR (Figure 1, a and b). Similarly, i.c.v. administration of EXN-4 dose-dependently increased MAP and HR (Figure 1, c and d). MAP and HR remained significantly increased for up to 2 hours after both i.v. (300 and 3000 ng) and i.c.v. (30 and 300 ng) administration of EXN-4.

As peripheral GLP-1 increases insulin secretion in a glucose-dependent fashion, we assessed whether the observed cardiovascular responses were secondary to GLP-1 agonist–induced hypoglycemia. Blood glucose levels increased modestly after i.c.v. administrations of both EXN-4 (basal, 96 ± 1.6; 15 minutes, 105 ± 3.9; 30 minutes, 109 ± 2.4; 60 minutes, 110 ± 3.0; 120 minutes, 110 ± 1.4; mean ± SEM) and PFS (basal, 98 ± 2.0; 15 minutes, 106 ± 4.9; 30 minutes, 104 ± 2.4; 60 minutes, 108 ± 5.6; 120 minutes, 106 ± 4.3; mean ± SEM). Importantly, i.c.v. administration of EXN-4 did not lower blood glucose at any time points after the injection.

Distribution of EXN-4–induced Fos-IR. To determine potential downstream mediators of GLP-1R activation leading to an increase in MAP and HR, we investigated the distribution of Fos-IR 2 hours after i.v. or i.c.v. EXN-4 administration at doses that increase MAP and HR (i.v., 300 ng; i.c.v., 30 ng). Intravenous administration induced Fos-IR in several autonomic control sites (31)
that showed little or no Fos-IR after PFS injection (Figure 2, a–t). These regions included the PVH, the lateral hypothalamic area (LHA), the arcuate nucleus (Arc) and the retrochiasmatic area (RCA), the external lateral parabrachial nucleus (PBel), the locus coeruleus (LC), the A5 catecholamine cell group (A5), the area postrema (AP), the NTS, the rostral ventrolateral medulla (RVLM), the caudal ventrolateral medulla (CVLM), and the IML of the spinal cord. The distribution of Fos-immunoreactive neurons induced by i.c.v. EXN-4 was similar to that induced by i.v. EXN-4 (Figure 3, a–t).

**Distribution of EXN-4–activated neurons projecting to the spinal cord.** To ascertain whether the observed increase in MAP and HR was due to activation of GLP-1–sensitive cell groups that also innervate sympathetic pre-ganglionic neurons, we injected the retrograde tracer FG) into the spinal thoracic cord centered on the IML (26, 27), with or without concomitant i.c.v. EXN-4 administration. The distribution of retrogradely labeled (FG-immunoreactive) neurons in the hypothalamus and the brainstem was similar to that in previous reports (26, 27, 32, 33). Following administration of EXN-4, we observed many double-labeled cells (Fos + FG) in the hindbrain, including the LC, the A5, and the RVLM (see Figure 5, n and p; Table 1). In the NTS, despite the existence of separate groups of neurons expressing either Fos-IR or FG-IR, we did not observe any double-labeled neurons. Many double-labeled neurons were observed in the hypothalamus, including the PVH, the Arc/RCA, and the LHA (Figure 4; Figure 5, a–c; Table 1).

GLP-IR agonists activate catecholamine neurons. As the topographical distribution of neurons displaying EXN-4–induced Fos-IR in the brainstem resembles the distribution of catecholamine neurons that are known to play essential roles in regulating the sympathetic nervous system (34–36), we assessed whether medullary catecholamine neurons were activated by doses of EXN-4 that increase MAP and HR. We found double-labeled (TH + Fos) neurons following i.c.v. EXN-4 administration in the LC, the A5, the AP, the NTS, the RVLM, and the CVLM (Figure 5, g–i, m, and o).

Using triple labeling, we observed that approximately 25% of TH-immunoreactive neurons in the RVLM contain Fos-IR following i.c.v. EXN-4 and also project to the IML (Figure 5, m and n). Similarly, approximately 30% of the TH-immunoreactive neurons in the A5 group contained both Fos and FG (Figure 5, o and...

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**Figure 3**

Distribution of i.c.v. EXN-4–induced Fos-IR in the brain and the adrenal gland. A series of photomicrographs demonstrates Fos-IR in neurons 2 hours after i.c.v. administration of EXN-4 (a–t, first and third columns; v–x) or PFS (a–t, second and fourth columns; u) in several brain regions and the adrenal gland. The brain regions include (a and b) the PVH; (c and d) the LHA; (e and f) the Arc/RCA; (g and h) the PBel; (i and j) the LC; (k and l) the A5; (m and n) the AP and the NTS; (o and p) the RVML; (q and r) the CVLM; and (s and t) the IML in the spinal cord. In the adrenal gland, after administration of PFS, little Fos-IR is visible (u). After i.c.v. EXN-4 at subthreshold dose (v; 3 µg), a little Fos-IR is visible. In contrast, i.c.v. EXN-4 at threshold dose (w; 30 µg) or a higher dose (x; 300 µg) induces Fos-IR in the adrenal medulla. Note that at the higher dose (x), Fos-IR is also visible in the cortex. Scale bar = 500 µm in a–d, g, and h; 200 µm in e, f, and i–t; and 50 µm in u–x.
In contrast, although we observed many double-labeled neurons (Fos + TH) in the NTS, we did not observe any triple-labeled neurons in this region. In the LC, nearly all retrogradely labeled neurons activated by EXN-4 contained TH.

Previous studies have demonstrated that stimuli that induce sympathoadrenal response also display Fos-IR in the adrenal gland (37). To determine whether a dose of i.c.v. EXN-4 that increases MAP and HR is associated with activation of both central and peripheral catecholaminergic pathways, we investigated the distribution of Fos-IR in the adrenal gland 2 hours after EXN-4 administration. Intracerebroventricular administration of EXN-4 at doses that increase MAP and HR also markedly induced Fos-IR in the adrenal medulla (Figure 3, w and x). PFS injection or EXN-4 injection at subthreshold doses (3 ng) induced little or no Fos (Figure 3, u and v). The higher dose of i.c.v. EXN-4 (300 ng) also induced Fos-IR in the adrenal cortex (Figure 3x).

GLP-1R agonists activate TH gene transcription. TH catalyzes the rate-limiting step in catecholamine biosynthesis, and the concentration and activity of TH correlate with neuronal activity (38). Both pharmacological stimulation and electrical stimulation increase TH transcription (36, 39). However, basal expression of TH mRNA in the brainstem is relatively high, making quantitative assessments of mRNA levels difficult. To combat this problem, we constructed an intron-specific (hnRNA) probe using PCR.

hnRNA probes have been widely used as indices of in vivo transcriptional activity (40, 41). hnRNA probes hybridize to intron-specific RNA sequences prior to processing to mature mRNA. The short half-life of hnRNA makes them useful tools to study recent transcriptional activation in individual neurons following specific stimuli (40, 41). We found that the relative levels of TH hnRNA were comparatively high in the LC (Figure 6b) even after PFS administration. In addition, a few neurons in the A5, the NTS, the RVLM, and the CVLM displayed a nuclear hybridization signal (Figure 6, c, e, g, and i). In these regions, neither i.v. nor i.c.v. administration of PFS further induced TH hnRNA expression at any time points examined. In contrast, both i.v. and i.c.v. administration of EXN-4 (i.v., 300 ng; i.c.v., 30 ng) provoked a robust and rapid increase in TH hnRNA expression in the A5, the NTS, the RVLM, and the CVLM (Figure 6, c, e, g, and i). Specifically, TH hnRNA expression was induced 15 minutes to 6 hours after i.c.v. injection, and the TH hnRNA expression returned to baseline by 24 hours.

Discussion

Using telemetric methods, we found that central GLP-1R agonists dose-dependently increased MAP and HR in unrestrained, conscious rats. In addition, we found that doses of EXN-4 that increase MAP and HR also induced Fos-IR in several autonomic control sites in the brain.

Table 1

Intracerebroventricular EXN-4 activates hypothalamic and brainstem neurons projecting to the spinal cord

<table>
<thead>
<tr>
<th>Region (atlas level)</th>
<th>FG-IR (i.c.v. PFS)</th>
<th>FG-IR (i.c.v. EXN-4)</th>
<th>Double-labeled: Fos-IR + FG-IR (i.c.v. PFS)</th>
<th>Double-labeled: Fos-IR + FG-IR (i.c.v. EXN-4)</th>
<th>% Double-labeled: Fos-IR + FG-IR / Total FG-IR (i.c.v. PFS)</th>
<th>% Double-labeled: Fos-IR + FG-IR / Total FG-IR (i.c.v. EXN-4)</th>
</tr>
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<tbody>
<tr>
<td>PVH (26/26)</td>
<td>60 ± 7.9</td>
<td>95 ± 16.1</td>
<td>5 ± 0.7</td>
<td>19 ± 2.6^</td>
<td>8 ± 1.2</td>
<td>21 ± 1.0^</td>
</tr>
<tr>
<td>RCA (27/27)</td>
<td>12 ± 2.2</td>
<td>14 ± 1.1</td>
<td>2 ± 0.0</td>
<td>6 ± 0.3^</td>
<td>19 ± 3.2</td>
<td>41 ± 1.2^</td>
</tr>
<tr>
<td>LHA (32/31)</td>
<td>86 ± 3.7</td>
<td>108 ± 19.7</td>
<td>5 ± 0.7</td>
<td>13 ± 1.8^</td>
<td>5 ± 0.7</td>
<td>13 ± 1.1^</td>
</tr>
<tr>
<td>LC (60/52)</td>
<td>12 ± 2.2</td>
<td>12 ± 2.6</td>
<td>1 ± 0.3</td>
<td>8 ± 1.2^</td>
<td>12 ± 2.6</td>
<td>50 ± 2.7^</td>
</tr>
<tr>
<td>A5 (60/52)</td>
<td>13 ± 1.7</td>
<td>14 ± 2.3</td>
<td>1 ± 0.6</td>
<td>4 ± 0.8^</td>
<td>3 ± 3.6</td>
<td>29 ± 2.4^</td>
</tr>
<tr>
<td>RVLM (70/64)</td>
<td>35 ± 2.2</td>
<td>18 ± 0.7</td>
<td>1 ± 0.6</td>
<td>9 ± 0.9^</td>
<td>5 ± 3.5</td>
<td>44 ± 3.4^</td>
</tr>
</tbody>
</table>

Values represent estimates of mean counts of cells ± SEM. Data were analyzed by ANOVA and differences between groups by Fischer’s PLSD. The atlas level designations correspond to those described by Paxinos and Watson (53) and Swanson (54), respectively. ^P < 0.01 compared with PFS-treated rats; n = 4 for PFS group, and n = 4 for EXN-4-treated rats.
regions and in the adrenal medulla. In the hypothalamus and the brainstem, we also observed Fos-IR in neurons (including catecholamine neurons) that innervate sympathetic preganglionic neurons. Furthermore, EXN-4 increased TH transcription in medullary catecholamine neurons. These findings suggest that CNS pathways known to be critical regulators of sympathetic outflow are engaged by central GLP-1 neurons to regulate physiological processes including BP and HR responses.

GLP-1 neurons are expressed predominantly in the caudal medulla. However, the GLP-1R is widely expressed in the CNS. Available data suggest that central GLP-1 action produces taste aversion (13) and that GLP-1R signaling systems are activated following exposure to agents that serve visceral illness such as lithium chloride (LiCl) or LPS (12, 15). These models are internal stressors and induced responses defined as interoceptive stress (16). Interestingly, GLP-1R antagonists attenuate the increase in Fos-IR induced by the administration of LiCl or LPS. Notably, several interoceptive stress models also activate sympathetic outflow (27, 42). Taken together, these data suggest that the autonomic alterations characteristic of interoceptive stress may be mediated by central GLP-1 systems.

Our current findings extend previous reports linking GLP-1 or EXN-4 to increases in MAP and HR in anesthetized rats (18–21) and provide a mechanism invoking catecholaminergic neurons as downstream mediators of GLP-1 action. In anesthetized animals, bilateral vagotomy blocks the stimulatory effects of i.c.v. GLP-1 on BP and HR (20), suggesting that the cardiovascular effects of central GLP-1 systems may be due to mechanisms including inhibition of vagal tone to the cardiovascular system. However, the inherent confounding effects of anesthesia on cardiovascular physiology must be considered (43, 44). Difference in potential mechanisms of GLP-1 action on the cardiovascular system between various studies may be due in part to experimental differences in the preparation and handling of the animals and the doses of the agonists used in each study. In our system, the threshold dose of i.c.v. EXN-4 to increase MAP is approximately one-tenth to one-hundredth the threshold dose of i.v. EXN-4, and i.c.v. administration of GLP-1R antagonist blocked the effect of i.c.v. EXN-4 on MAP and HR. As GLP-1 is well known to be a potent stimulator of glucose-dependent insulin secretion, we excluded the possibility that changes in blood glucose following GLP-1 administration might be responsible for induction of Fos activation and counter-regulatory responses leading to sympathetic nervous system activation (3). The demonstration that GLP-1 agonists activate Fos, TH gene transcription, and cardiovascular responses at doses that do not lower blood glucose strongly suggests that central GLP-1 systems are capable of regulating BP and HR independent of the established peripheral actions of GLP-1 on glucoregulation.

Based on our findings, we propose that two possible effector pathways downstream of central GLP-1 neurons regulate sympathetic outflow. These two

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**Figure 5**

EXN-4 activates medullary catecholamine neurons projecting to the spinal cord. (a–c) Double-label immunohistochemistry demonstrates that neurons that innervate sympathetic preganglionic neurons (brown cytoplasm) also contain i.c.v. EXN-4–induced Fos-IR (black nuclei) in (a) the PVH, (b) the Arc/RCA, and (c) the LHA of the hypothalamus. (d–f) Higher-magnification view of the boxed areas in a–c, respectively. (g–i) Double-label immunohistochemistry demonstrates that TH-immunoreactive neurons (brown cytoplasm) also contain i.c.v. EXN-4–induced Fos-IR (black nuclei) in (g) the NTS, (h) the LC, and (i) the CVLM. (j–l) Higher-magnification view of the boxed areas in g–i, respectively. (m–p) Triple-label immunohistochemistry reveals that many medullary catecholamine neurons (red cytoplasm in m and o) that project to the IML in the spinal cord (green neurons in n and p) also contain i.c.v. EXN-4–induced Fos-IR (black nuclei) in the RVLM (m and n) and the A5 (o and p). Scale bar = 100 μm in a–c and g–i, 25 μm in d–f and j–l, and 50 μm in m–p.
pathways include monosynaptic hypothalamo-spinal and bulbo-spinal projections to sympathetic preganglionic neurons (32, 45). Several studies have demonstrated that neurons in the RVLM and A5 provide descending input to sympathetic preganglionic neurons. Additionally, the RVLM provides tonic excitatory input to the cardiovascular system (34, 46–48). Therefore, we suggest that central GLP-1 systems are involved in regulating cardiovascular function in part by engaging catecholamine neurons that project to the IML to activate the adrenal medulla. The PVH is composed of functionally distinct subnuclei that regulate endocrine and autonomic responses (31). Within the PVH, EXN-4 induced Fos-IR in the medial parvicellular neurons and in the ventral, dorsal, and lateral subdivisions, suggesting that central GLP-1 systems may regulate the hypothalamo-pituitary-adrenocortical (HPA) axis and autonomic outflow. Indeed, neurons in the caudal NTS that project to the PVH contain GLP-1 (17). GLP-1 also activates the HPA axis, presumably through stimulation of CRH neurons (17). With respect to autonomic responses, the transneuronal tracing has identified the PVH as one of a few brain sites that regulate the entire sympathetic outflow (49). Our current study suggests that GLP-1 acts in the PVH to mediate neuroendocrine responses and to modulate sympathetic outflow.

In our current study, both peripheral and central administration of GLP-1 agonists rapidly increased BP and HR and induced similar patterns of Fos-IR throughout the brain, including several key autonomic regulatory sites. The induction of Fos-IR in these sites is consistent with the concept that the increased activity of these neurons contributes to the increases in BP and HR. However, we cannot rule out the possibility that the expression of Fos in some of these sites may simply be secondary to the increased BP and HR and may not represent activation of neurons inducing the observed responses. Still, the Fos patterns induced by GLP-1 agonists in the brainstem are distinct from patterns induced by phenylephrine-induced hypertension, which does not induce Fos-IR in catecholamine neurons (50). Indeed, the patterns of GLP-1–induced Fos are more similar to those of stimuli that increase HR and BP, increase Fos-IR in TH neurons, and activate TH mRNA (36, 50). Thus,
we suggest that Fos induction in central autonomic control sites by GLP-1 agonists is not likely to be a secondary consequence of increased BP and HR.

One interesting difference in the patterns of i.v. and i.c.v. injections was that in the AP. Specifically, i.v. EXN-4 induced much more Fos-IR than did i.c.v. EXN-4 in the AP. GLP-1Rs are prominently expressed in the AP, a circumventricular organ that lacks a blood-brain barrier. Therefore, it is likely that neurons in the AP have access to circulating GLP-1. Indeed, recent studies using 125I-GLP-1 have demonstrated that GLP-1Rs are accessible to peripheral GLP-1 in the subfornical organ and the AP in the rat (51). Furthermore, in our preliminary study, we found that i.v. EXN-4 induced Fos-IR in GLP-1 neurons, suggesting that peripheral EXN-4 activates the central GLP-1 system, which may contribute to the effects we observed. Consistent with this hypothesis, Barragan et al. (20) showed that i.c.v. administration of a GLP-1 antagonist blocked cardiovascular responses to systemically injected GLP-1. Therefore, it is plausible that circulating GLP-1 activates GLP-1-responsive neurons in the AP and also activates GLP-1 neurons in the underlying NTS. This model would predict that GLP-1 neurons in turn activate neurons expressing GLP-1Rs in the hypothalamus and the brainstem, leading to coordinated endocrine and autonomic effects (Figure 7).

The data reported here suggest that the central GLP-1 system can regulate sympathetic outflow. In normal humans and rodents, circulating levels of GLP-1 are low in the fasting state, rise following nutrient ingestion, and fall within minutes due to rapid enzymatic inactivation by DPP-IV. Indeed, continuous infusion of GLP-1 does not increase BP and HR in human subjects with type 2 diabetes (52). In contrast, more potent degradation-resistant long-acting GLP-1 analogues, including EXN-4, are being considered as therapeutic agents for the treatment of type 2 diabetes. Our current results demonstrate a dose-dependent increase in BP and HR following both central and peripheral administration of EXN-4. However, the pathophysiological significance of our findings in rodents must be assessed in further studies that carefully examine dose response relationships in both acute and chronic paradigms, especially in diabetic subjects. Nonetheless, our results suggest that the central GLP-1 system is a regulator of sympathetic outflow and raise questions about the potential consequences of sustained GLP-1R activation for the treatment of patients with type 2 diabetes.

Acknowledgments

The authors thank Quan Ha, Toshiro Kishi, Im Jihee, Brian J. Choi, Hiromichi Kashihara, Takuma Kishimoto, and Hiroshi Muraki for technical assistance. This work was supported by NIH grants DK-59751 and DK-56116 and Juvenile Diabetes Research Foundation grant 2000-559.


