

# NIH Public Access

Author Manuscript

*Neuroimage*. Author manuscript; available in PMC 2014 September 03.

Published in final edited form as:

Neuroimage. 2012 March ; 60(1): 592-600. doi:10.1016/j.neuroimage.2011.12.023.

# Postprandial Plasma PYY Concentrations are Associated with Increased Regional Gray Matter Volume and rCBF Declines in Caudate Nuclei – a combined MRI and H<sub>2</sub><sup>15</sup>O PET study

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# Abstract

The anorexigenic gastrointestinal hormone Peptide YY plays an important role in the communication between the gastrointestinal tract and the central nervous system. PYY has been shown to modulate brain activity in regions implicated in reward and food related behavior. Its effects on brain structure however, remain unknown. Voxel-based morphometry was used to investigate the relationship between fasting and postprandial plasma PYY concentrations and regional gray matter volume (GMV). For this analysis twenty adult, non diabetic Caucasians were included (18F/2M, age  $31\pm9$  y, percentage of body fat [PFAT]  $32\pm8\%$ ) who had volumetric brain magnetic resonance images and underwent  $H_2^{15}O$  positron emission tomographic (PET) measurements of regional cerebral blood flow (rCBF), a marker of local neuronal activity, and measurements of plasma total PYY, prior to (fasting) and following a satiating liquid meal. Voxelwise analysis revealed a regional positive association between postprandial PYY and gray matter volume bilaterally in the caudate nuclei. These associations remained significant (p < 0.05) after small volume correction for multiple comparisons. Based on these findings we investigated whether postprandial PYY is associated with PET measured rCBF of the caudate nucleus. We found a significant negative association between average postprandial caudate rCBF and postprandial plasma PYY concentrations (r=-0.60, p<0.02, age, sex and PFAT adjusted). Average postprandial caudate rCBF was also negatively associated with rCBF in the right medial orbitofrontal cortex and the right hippocampal formation (p<0.05, corrected for multiple comparison). Total PYY is positively associated with gray matter but negatively with postprandial activity in the caudate nuclei while caudate activity is negatively associated with rCBF in prefrontal and paralimbic regions implicated in reward behavior. Thus, PYY may act centrally to modulate eating behavior via striatal networks.

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#### Keywords

PYY; caudate nucleus; striatum; gray matter; VBM; MRI; PET; rCBF

# 1. Introduction

In the past decades, obesity has become a medical and socioeconomic problem of pandemic proportions in industrialized countries. A vast number of illnesses have been associated with excessive overweight including the major causes of death in western countries, cardiovascular disease, certain types of cancer, and stroke (Guh et al., 2009). In this respect, understanding the physiological events underlying feeding behavior and the development of obesity has become a research question of major importance. Gastrointestinal hormones play a crucial role in the communication between the gastrointestinal tract and the brain, mediating signals of both hunger and satiety. The gastrointestinal hormone Peptide YY (PYY) is a member of the pancreatic peptide fold family along with neuropeptide Y (NPY) and pancreatic polypeptide (PP) and is known to decrease appetite and food intake in lean and healthy humans (Batterham et al., 2003, 2007; Degen et al., 2005; Sloth et al., 2007), inhibit stomach emptying (Witte et al., 2009) and increase gastrointestinal water and electrolyte absorption (Cox, 2007). Synthesized by L-type Endocrine cells in the distal gastrointestinal tract, PYY is secreted into the circulation in response to a meal and postprandial plasma concentrations remain elevated for approximately 6 hours (Adrian et al., 1985). Two endogenous and physiologically active forms have been identified, PYY1-36 and PYY3–36. PYY3–36 is formed by the ubiquitously expressed enzyme dipeptidyl peptidase IV (DP-IV) via cleavage of the first three n-terminal amino acids of PYY1-36 (Mentlein et al., 1993). Effects of PYY are mediated by Y-receptors, a G-protein coupled receptor family (Cabrele and Beck-Sickinger, 2000), widely distributed throughout the gastrointestinal tract and the central nervous system (Widdowson, 1993; Parker and Herzog, 1999).

So far four subtypes have been identified in humans (i.e. Y1, Y2, Y4, Y5) (Gehlert, 1998; Michel et al., 1998; Berglund et al., 2003). The anorectic effects of PYY have been mainly attributed to PYY3-36 and the Y2-receptor. Animal and human studies have shown a reduction in appetite and food intake after systemic administration of PYY 3-36 (Batterham et al., 2003, 2007; Degen et al., 2005; Sloth et al., 2007). In rodents, this effect is absent in Y2R-knockouts (Batterham et al., 2002) or after administration of a Y2R antagonists (Abbott et al., 2005). While PYY3-36 exhibits a highly selective binding profile with a strong affinity for Y2-receptors, PYY1–36 has affinity for Y1, Y2 and Y5-receptors. (Dumont et al., 1995; Batterham and Bloom, 2003). Although the primary effect of PYY on appetite and eating behavior is believed to be mediated via the Y2 receptor of the arcuate nucleus within the hypothalamus, a significant decrease in high-fat food seeking in response to systemic PYY that was independent of arcuate nucleus Y2R signaling was reported in rodents (Ghitza et al., 2007). There is evidence indicating that postprandial rises of PYY are lower in obese compared to lean individuals leading to reduced satiety and relatively higher food intake, yet ratios of PYY1-36 and PYY3-36 do not seem to change with adiposity (le Roux et al., 2006). Importantly, obese individuals do not appear to develop a resistance to

the anorectic effects of PYY as occurs with the adipokine leptin (Batterham et al., 2003). PYY may also play an important role in weight regain following gastric bypass surgery as attenuated postprandial PYY profiles have been found in individuals with poor weight loss (Meguid et al., 2008). Intravenous administration of PYY3–36 modulates brain regions implicated in both homeostatic (e.g. hypothalamus) and reward related feeding behavior (e.g. prefrontal cortical regions, ventral tegmental area, putamen, globus pallidus) (Batterham et al., 2007).

Peripheral PYY clearly has important effects on brain function and has sustained postprandial elevations. Furthermore, other peripherally circulating hormones have been shown to influence brain morphology (Starkman et al., 1999; Bourdeau et al., 2002; Matochik et al., 2005; Pannacciulli et al., 2007). Thus, we sought to investigate the effects of total PYY concentrations in the fasting state and in response to a satiating liquid meal on gray and white matter volume using voxel-based morphometry (VBM). After demonstrating associations with bilateral caudate gray matter, we investigated associations with Oxygen-15 water positron emission tomographic (PET) measurements of regional cerebral blood flow in a caudate nucleus region-of-interest (ROI). Associations between caudate nucleus rCBF and other brain regions were furthermore investigated in a voxel-based analysis, hypothesizing a modulation of prefrontal and limbic/paralimbic neuronal activity.

# 2. Subject and Methods

#### 2.1. Subjects

All subjects in this study had previously participated in brain imaging studies of hunger, satiation, and the predisposition to obesity. Twenty adult, non-diabetic, right-handed Caucasians with a wide range of adiposity  $(18F/2M; age 31\pm9y; percentage of body fat$ [PFAT] 32.2±8.4%; BMI 31.2±9.6) were included who had available MRI and PET scans and measurements of PYY. All subjects were recruited from the Phoenix, AZ metropolitan area by newspaper advertisements. All subjects were free of medical disorders, not taking any medications, as determined by medical history, physical examination, and screening laboratory tests. Female subjects were studied while in the follicular phase of the menstrual cycle. Subjects with a history of substance or alcohol abuse or addiction; endocrine disorders (including abnormal thyroid function and type 2 diabetes); hypertension, pulmonary, cardiovascular, gastrointestinal, hepatic, renal or central nervous system disorders were excluded from the study at screening. The Structured Clinical Interview for DSM-III-R (Spitzer et al., 1990) was used to screen for behavioral and psychiatric conditions (claustrophobia, major depression, the presence of psychotic symptoms, anorexia nervosa, or bulimia nervosa) incompatible with safe and successful participation in the study. All subjects were admitted for one week to the metabolic unit of the Obesity and Diabetes Clinical Research Section of the National Institute of Diabetes and Digestive and Kidney Diseases (NIDDK) in Phoenix, AZ. Subjects were restricted to the research ward and were limited to sedentary activity for the duration of the study. The protocol was approved by the Institutional Review Boards of the NIDDK and the Banner Good Samaritan Regional Medical Center. All subjects provided a written informed consent prior to

#### 2.2 Experimental protocol

The experimental procedures have been described previously (Tataranni et al., 1999). In brief, upon admission, subjects received a weight maintaining diet (50% of calories from carbohydrate, 30% fat and 20% protein). Body composition was assessed by dual energy xray absorptiomery (DPX-1; Lunar Corp, Madison, WI); resting energy expenditure (REE) was measured for 45 minutes using a ventilated-hood system (DeltaTrac, SensorMedics, Yorba Linda, CA). Prior to the brain imaging session, subjects fasted for 36 h. Study subjects had free access to water and non-caloric, non-caffeinated beverages during the fast.

#### 2.3 Metabolite analysis

Plasma concentrations of fasting and postprandial total PYY (including PYY1–36 and PYY3–36) were measured by a commercially available radioimmunoassay kit (Phoenix Pharmaceuticals, Belmont, California, USA) with a 100% crossreactivity between PYY1–36 and PYY3–36. Plasma glucose concentrations were measured by the glucose oxidase method (Beckman Instruments, Fullerton, CA). Plasma insulin concentrations were determined by an automated radioimmunoassay (Concept 4; ICN, Costa Mesa, CA).

#### 2.4 Imaging procedures

Magnetic Resonance Imaging (MRI) and Positron Emission Tomography (PET) procedures were carried out at the Banner Good Samaritan Regional Medical Center (Phoenix, AZ). MRI scans were performed on a 1.5 Tesla Signa system (General Electric, Milwaukee, WI, USA). A set of high-resolution T1-weighted images was acquired with a fast spoiled gradient echo (FSPGR) 3d sequence (repetition time [TR]/ echo time [TE] = 12/5.2; inversion time (TI) = 300ms, number of excitations (NEX) = 1; field-of-view [FOV] =  $24 \times$ 24 cm; 256 × 256 matrix); the whole-brain data were acquired in an axial plane yielding 120 contiguous slices with slice thickness of 1 mm. Each subject's MRI scans were evaluated by an experienced neuroradiologist in order to exclude individuals with anatomic abnormalities.

O-15 water positron emission tomographic measurement of regional cerebral blood flow (counts/voxel/min) was performed on an ECAT-951/31 scanner (Siemens, Knoxville, TN). To adjust for attenuation of  $\gamma$ -radiation by the brain and skull, a 10 min transmission scan was performed using a retractable external ring source of 68Ga/68Ge. For each 1-min PET scan, subjects remained motionless in the supine position and were requested to keep their eyes closed and pointing forward. Subjects received a 50-mCi intravenous bolus of <sup>15</sup>O-water during each scan. Each individual underwent two scans at baseline (fasting, premeal condition) and two after oral administration of a satiating amount of a liquid formula meal (Ensure Plus, 1.5 kcal/ml, Ross-Abbott Laboratories, Columbus, OH) providing 50% of the subject's measured REE, with intervals of 10 min between scans. Formula flavor (strawberry, vanilla or chocolate) was chosen by the subject and the meal was administered continuously over 25 min via a plastic tube placed into the subject's mouth using a peristaltic pump. To avoid swallowing during the scan, subjects received 2 ml of water at room temperature via a plastic tube, administered by a syringe 30 s before each scan and

were asked to retain and swallow. Immediately following each scan, blood samples were drawn for hormonal and metabolic measurements. Using 100-mm visual analogue scales (Raben et al., 1995) subjective ratings of hunger and fullness were obtained after each scan. All subjects had been fully acquainted with the experimental procedures, thus minimizing the risk of learning-related artifacts and anticipated receiving the satiating liquid meal.

#### 2.5 Image data analysis

Voxel-based morphometry (Ashburner and Friston, 2000) was performed using the VBM8 toolkit in Statistical Parametric Mapping package (SPM8, Wellcome Department of Imaging Neuroscience, London, UK; www.fil.ion.ucl.ac.uk/spm). In brief, native MRI scans were normalized to the standard Montreal Neurological Institute (MNI) T1 MRI template (voxelsize:1.5mm×1.5mm×1.5mm) and segmented into grey matter (GM), white matter (WM), and cerebrospinal fluid (CSF) using automated algorithms implemented in VBM8 and the default settings. Finally, segmented images of GM and WM were smoothed with a 12mm full-width at half-maximum isotropic Gaussian kernel. Total intracranial volume (TIV) was calculated based on segmented GM, WM and CSF images over the individual participant's native brain space and was included as a confounding covariate in an analysis of covariance (ANCOVA). In SPM8 voxel-wise multiple variable regression analysis of GM and WM images was performed, entering plasma PYY concentrations (fasting, postprandial, response), age, sex and percentage body fat (PFAT) as covariates with a p-threshold of <0.001 and an extent threshold of 100 continuous voxels (voxelsize:

1.5mm×1.5mm×1.5mm). Small volume correction (SVC) for regional multiple comparisons was performed on regions (GM or adjacent WM) associated with PYY, eating and reward behavior respectively (i.e. caudate nucleus, globus pallidus, thalamus, prefrontal regions, anterior cingulate gyrus and the cerebellum (Batterham et al., 2007; Neary and Batterham, 2010)) by using anatomical masks or a sphere with a 10mm radius when masks were not applicable. Small volume corrected results are reported as significant at p<0.05 (family-wise error correction on the voxel-level). Masks for data extraction and SVC were created by using the WFU pickatlas tool (Maldjian et al., 2003) and the integrated automatic anatomic labeling (AAL) tool (Tzourio-Mazoyer et al., 2002).

PET scans were aligned, spatially normalized to the standard Montreal Neurological Institute (MNI) stereotactic space and smoothed using a 15- mm full-width-at-halfmaximum Gaussian filter (Friston et al., 1995). SAS Software (SAS Institute Inc, version 9.2, Cary, NC) was used for statistical analyses of non-imaging data and extracted data. The hypothesis of brain regions modulated by caudate activity was tested by performing a multiple voxel-wise regression analysis of PET scans (fasting and postprandial) entering average bilateral caudate activity into a multiple regression model including age, sex and PFAT as covariates. Statistical parametric maps were thresholded at p<0.005 and an extent threshold of 50 continuous voxels (voxelsize: 2mm×2mm×2mm). Results are reported as significant at p<0.05 cluster-level corrected (FWE). ROI and whole brain analyses were adjusted for age, sex and PFAT. Pearson correlation coefficients were Fisher Z-transformed to test for differences between the pre- and postmeal condition. Anatomical regions of both VBM and PET analyses were defined by using the BioImage Suite MNI to Talaraich Coordinate Converter (www.bioimagesuite.org) and the Talaraich Client v2.4.2 (Lancaster

et al., 2000) with a 2mm search range. For PET data a search range of 5mm was used to define closest Brodmann Areas.

# 3. Results

### 3.1 Subject characteristics

Table 1 summarizes subject characteristics, hormonal-metabolic measures and subjective appetite ratings in the fasting and postprandial state in our study group. As anticipated, plasma concentrations of glucose, insulin and total PYY significantly increased in response to the meal (all p<0.001), while hunger and fullness scores were respectively significantly decreased and increased (all p<0.001).

#### 3.2 VBM analysis

Fasting PYY concentrations were positively associated with regional GM volume in the bilateral cerebellum, reaching significance after SVC ( $p_{voxel}=0.047$ ;  $k_E=484$ ; MNI<sub>*xyz*</sub> [-6 -64 -20]) (Table 2). Postprandial PYY levels were positively associated with GM volume bilaterally in caudate nuclei (i.e. caudate body) (Fig.1A, B, D). This association remained significant after SVC for the sublobar brain volume and comprised two significant peaks within the left and right caudate body (left:  $p_{voxel}=0.02$ ; MNI<sub>*xyz*</sub> [-3 2 9]; right:  $p_{voxel}=0.049$ ; MNI<sub>*xyz*</sub> [6 4 9];  $k_E = 679$ ).

PYY response (postprandial minus fasting PYY concentrations) was positively associated with GM volume in the left fusiform gyrus, left globus pallidus (extending to the left caudate), right caudate nucleus (i.e. head), left and right middle temporal gyrus and right anterior cingulate gyrus (ACC) (Fig. 1C), uncorrected for multiple comparisons.

After SVC for the sublobar brain volume, subcortical peaks partially reached significance (left globus pallidus  $p_{voxel}=0.038$ ,  $k_E=222$ ,  $MNI_{xyz}[-8\ 2\ 1]$ ; right caudate,  $p_{voxel}=0.093$ ,  $k_E=362$ ,  $MNI_{xyz}[10\ 4\ 1]$ . However, SVC for the bilateral caudate volume revealed two significant peaks in the left ( $p_{voxel}=0.019$ ;  $k_E=130$ ;  $MNI_{xyz}[-6\ 6\ 3]$ ) and right caudate nucleus ( $p_{voxel}=0.016$ ;  $k_E=18$ ;  $MNI_{xyz}[9\ 9\ 3]$ ). SVC for the right ACC volume showed a significant peak ( $p_{voxel}=0.023$ ;  $k_E=121$ ;  $MNI_{xyz}[15\ 23\ 27]$ ) within the right ACC.

No negative associations were observed for fasting PYY, postprandial PYY or PYY response.

VBM analysis of the WM revealed a positive association between fasting PYY and the right inferior frontal gyrus (IFG) and a negative association between PYY response and extranuclear WM in the vicinity of the left thalamus, both significant after SVC correction using a 10mm sphere (IFG:  $p_{voxel}=0.001$ ;  $k_E = 145$ ; MNI<sub>xyz</sub>[44 8 25]; left Thalamus:  $p_{voxel}=0.01$ ;  $k_E = 145$ ; MNI<sub>xyz</sub>[-3 -24 12]). No negative associations were found for postprandial PYY concentrations.

#### 3.3 ROI PET analysis

Caudate rCBF values (fasting and postprandial) were extracted from those with available PET measurements of rCBF in the fasting and postprandial state (n=19). Left and right

caudate rCBF strongly correlated with each other(fasting: r=0.89, p<0.001; postprandial: r=0.94, p<0.001). Caudate rCBF significantly decreased in response to the liquid meal (p=0.03) and postprandial caudate rCBF was significantly negatively associated with postprandial PYY concentrations (bilateral mean: r=-0.60, p<0.02, left: r=-0.57, p=0.02, right: r=-0.61, p<0.01; Pearson partial correlation, adjusted for age, sex, PFAT). Further segmentation into the anatomical subdivision of the caudate nucleus revealed the strongest association in the caudate body and tail (Table 3, Fig.2A, B). Regarding possible correlations between rCBF (fasting rCBF- postprandial rCBF) of the caudate and PYY response (postprandial - fasting) we only observed a positive correlation of rCBF of the right caudate body with PYY response (r=0.56, p<0.03; Pearson partial correlation with age sex and PFAT as partial covariates). Total caudate rCBF and all other anatomical subdivisions showed no significant correlations with PYY response (data not shown). All of the reported associations between caudate rCBF and postprandial PYY did not change after further adjustment for insulin concentrations (data not shown).

Pearson partial correlation (age, PFAT and sex as partial covariates) showed a significant negative association between TIV adjusted GM values of the left and right peak voxel and postprandial rCBF values of the left and right caudate (left: r=-0.64, p=0.008; right: r=-0.71p=0.002). These associations were no longer significant after adjusting for postprandial PYY concentrations (left: p=0.19, r=-0.36; right: p=0.09, r=-0.46).

#### 3.4 Voxel-based PET analysis

Caudate activity appeared to modulate several other brain regions, as determined using our voxel-based analysis. Results are summarized in Table 4, S1–2. In the pre-meal condition, areas positively associated with mean caudate rCBF included the right thalamus (extending to the left thalamus), right anterior cingulate gyrus and the left cuneus (supplementary material, table S1). In the postmeal condition we observed positive associations in the right anterior cingulate gyrus, right putamen extending to the right thalamus, left inferior parietal lobule and the right superior and left occipital gyrus (supplementary material, table S2). Significant negative associations in the fasting state were only observed in the right cerebellum. Notably, in the postprandial state however negative associations were found in the right medial orbitofrontal cortex and the right hippocampal formation (Table 4, Fig. 2C).

Fisher Z transformed Pearson moment-activity correlation coefficient of the right mOFC and right hippocampal formation did not show statistically significant differences between the pre-and postmeal condition (data not shown).

# 4. Discussion

We explored the associations of fasting and postprandial concentrations of the gastrointestinal hormone PYY using a voxel-based analysis of brain morphology and a subsequent ROI analysis of rCBF in a structurally affected brain region. We demonstrated that postprandial plasma concentrations of total PYY are positively associated with GM volume of the bilateral caudate nucleus. PYY response was positively associated with GM of the right anterior cingulate gyrus, temporal regions, the left globus pallidus (inclusive of the left caudate) as well as the right caudate nucleus. In addition, using a region of interest

approach to examine rCBF, we found a significant decrease in caudate rCBF between fasting and post prandial states and a negative association between postpandrial PYY and caudate rCBF. Caudate activity was furthermore negatively associated with rCBF of the right orbitofrontal cortex and the right hippocampus. In humans, systemic application of PYY3-36 has been shown to modulate activity in several brain regions associated with feeding-behavior including prefrontal cortical regions, ventral tegmental area, putamen, globus pallidus and the hypothalamus (Batterham et al., 2007). As postprandial PYY concentrations remain elevated for approximately 6 hours after a meal (Adrian et al., 1985), variability in postprandial PYY profiles could affect human brain morphology. Other peripheral circulating hormones have been associated with effects on human brain tissue composition. Replacement therapy of the adipokine leptin in genetically leptin deficient humans increased cerebellar, inferior parietal lobule and anterior cingulate GM volume (Matochik et al., 2005). In addition, a VBM analysis of healthy humans demonstrated associations of fasting leptin concentrations and GM volume of several brain regions, independent of percent body fat (Pannacciulli et al., 2007). Furthermore changes in brain morphology, specifically in the hippocampus have been found following resolution of hypercortisolemia in individuals with Cushing's syndrome (Starkman et al., 1999; Bourdeau et al., 2002).

We further investigated rCBF changes in the caudate using an ROI approach and demonstrated that caudate activity decreases between fasting and postprandial states and is negatively associated with postprandial PYY. Batterham et al. showed a modulation of striatal BOLD levels in response to PYY3-36 infusion. However, this effect was restricted to the right putamen and the left globus pallidus and did not include the caudate. Furthermore, PYY3–36 infusion showed a positive association with striatal activity (Batterham et al., 2007). Several reasons might explain our divergent findings. First, the paradigms used in the study by Batterham et al. and ours differ substantially and therefore do not allow direct comparisons. Batterham et al. analyzed fMRI measured BOLD levels in response to systemic application of PYY3-36, while we analyzed metabolic/hormonal responses and PET measured rCBF in response to an actual satiating meal. Also, we measured total PYY, which includes both endogenous forms of PYY. These two subtypes, PYY1-36 and PYY3-36 have different receptor selectivity profiles (Y1,Y2,Y5 vs. Y2>>Y1) (Dumont et al., 1995; Batterham and Bloom, 2003) possibly leading to different neuronal responses as the central distribution of Y-receptor subtypes varies substantially between different brain regions (Widdowson, 1993; Parker and Herzog, 1999). The caudate has the highest Y1 receptor density based on a recent study using a specific PET radiotracer (Hostetler et al., 2011). Considering the strong affinity of PYY1–36 and the comparatively low affinity of PYY3–36 for the Y1 receptor (Keire et al., 2000) allows speculation that PYY1-36 is more responsible for the reported findings.

The reason for the positive association between postprandial PYY and caudate GMV is not clear. One possible explanation is that PYY mediates changes in the caudate neurochemical milieu, specifically via its effect on dopamine. Dopamine is known to have trophic effects on neuronal and synaptic plasticity (Nieoullon, 2002). Genetic variants of the dopamine D2 receptor (DRD2) and the dopamine transporter (DAT) that are associated with reduced dopaminergic neurotransmission (Bertolino et al., 2009) have been shown to modulate

caudate GMV in healthy individuals. Rodent studies showed that orexigenic effects of centrally administered PYY are dependent on DA signaling in the caudate nucleus (Hnasko et al., 2004). In vitro experiments have furthermore revealed a modulation of striatal DA synthesis via Y-receptor activation. Selective activation of striatal Y2 receptors with the synthetic Y2 agonist PYY13–36 increased DA synthesis. However, the administration of PYY1–36 and PYY3–36 provoked a significant attenuation of striatal DA synthesis (PYY1–36>PYY3–36) (Adewale et al., 2005). In contrast to these in vitro studies, central administration of PYY in rodents, led to a significant increase of arcuate nucleus tyrosine hydroxylase (TH) mRNA expression, the rate-limiting enzyme for DA synthesis (Hong et al., 1995). Differentiating central versus peripheral effects of PYY is not straightforward as central administration of PYY stimulates appetite and feeding, while peripherally administered PYY has anorectic effects. In addition, central Y-receptor subtypes distribution is different between humans and rats so effects in rat brains may not apply to humans (Widdowson, 1993).

Regarding the observed negative association between postprandial PYY and postprandial caudate rCBF, it has to be noted that the relationship between striatal rCBF and dopaminergic activity remains controversial as animal and human studies using pharmacological dopaminergic challenges have shown differing results. D-amphetamine, a dopamine-releasing agent, administered in four different doses followed by measurements of local DA (via microdialysis) and regional cerebral blood volume (rCBV) by MRI demonstrated increased rCBV in response to higher doses of D-amphetamine which also produced dose related higher concentrations of extracellular DA but an actual decrease in rCBV with lower doses. This finding was interpreted as a switch in dopamine receptor subtype stimulation (D2/D3 vs. D1/D5) associated with dose escalation (Ren et al., 2009) and indicates that different effects on brain structure and function may be seen in pharmacologic versus physiologic situations. L-DOPA, the direct precursor of dopamine, increased striatal rCBF in awake and but decreased rCBF in anaesthesized healthy, nonhuman primates (Hershey et al., 2000). In awake humans, L-DOPA increased striatal rCBF (i.e. bilateral putamen) (Hershey et al., 1998). However, pharmacologic challenge studies using dopamine receptor blocking agents have shown increases in striatal rCBF after systemic administration of haloperidol in both healthy humans and schizophrenic patients(Bartlett et al., 1994; Lahti et al., 2003, 2005).

Historically, the striatum has been most commonly associated with motor control. However, the caudate nucleus has complex interconnections within the striato-thalamo-cortical networks (Alexander et al., 1990) and recent studies indicate a more diverse role for the caudate in human behavior and cognition, including working memory, decision-making, reward and reward based learning (Haruno et al., 2004; Balleine et al., 2007; Ding and Gold, 2010) and food-related behavior (Stice et al., 2008; Neary and Batterham, 2010; Green et al., 2011). We showed significant positive associations between relative mean caudate rCBF and rCBF in the right cingulate gyrus and subcortical nuclei (i.e. thalamic nuclei, putamen, globus pallidus) in both fasting and postprandial state and significant positive associations with parietal and occipital region rCBF in the postprandial state only. Most interestingly, in the postprandial but not the fasting state we observed significant negative associations with right mOFC and right hippocampus rCBF, both significant on the cluster-level. The OFC is

a polyfunctional region that has been linked to reward processing, decision-making and inhibitory control (i.e. the capacity to control a pre-potent reaction in response to a stimulus) (Kringelbach and Rolls, 2004; Perry et al., 2011). Alterations of these cognitive processes and their underlying neuronal systems (i.e. frontostriatal networks) have been associated with several disorders, including obesity, eating disorders (ED) and addictive behavior in general (Volkow, Wang, Telang, Fowler, Goldstein, et al., 2008; Volkow, Wang, Telang, Fowler, Thanos, et al., 2008; Brogan et al., 2010, 2011; Koob and Volkow, 2010; Oberndorfer et al., 2011). Interestingly, the caudate nucleus also seems to play an important role in inhibitory control, as disturbed caudate function and low inhibitory control are characteristics of obsessive-compulsive disorder (OCD) and attention deficit hyperactivity disorder (ADHD) (Tripp and Wickens, 2009; Fineberg et al., 2010). The hippocampus is primarily implicated in memory but also appears to be part of the regulation of reward related behavior as demonstrated in human neuroimaging studies (Liu et al., 2011). The dorsal striatum together with the hippocampus, prefrontal cortical regions and the amygdala may form a network relevant in reward processing and reward based learning in particular (Haber et al., 2006).

Several other limitations must be acknowledged. This is a post-hoc analysis of a study designed to assess neuronal correlates of hunger and satiety. Although we adjusted for sex, results of this study might not entirely apply to the male gender due to the unbalanced sex distribution of our study population, however the results were consistent when women only were analyzed. Imaging procedures and hormonal/metabolic measurements were done after a prolonged fast of 36h. Furthermore, we measured only total PYY concentrations even though the anorectic effects of PYY are majorly attributed to PYY3–36. However, ratios of PYY1–36 and PYY3–36 appear to be stable in both healthy and obese individuals (le Roux et al., 2006) and both endogenous forms (PYY1–36 and PYY3–36) are physiologically active (Pfluger et al., 2007).

As this study is of correlational nature only, we cannot rule out the possibility of other physiological events that occur in response to a meal contributing to the reported associations, as for example changes in plasma concentrations of other hormones (e.g. GLP1, CCK, Pancreatic Polypeptide) or changes in vagal tone. However, adjusting for insulin concentrations did not show an effect on our results. MRI scans were performed on a 1.5T scanner and the VBM analysis presented here is of an exploratory nature in a limited number of subjects. Although the segmentation, especially of subcortical structures, has been improved in VBM8 by including the Maximum A Posterior (MAP) technique and Partial Volume Estimation (PVE) (Gaser, 2009) these results need to be repeated in a larger cohort using different volumetric methodologies. We did not observe differences in functional connectivity between the fasting and the postprandial state of the caudate nucleus and the hippocampus and right mOFC respectively. Nevertheless this may be due to our relatively small sample size. We also acknowledge that no attempt was made in this current study to correct the effects of combined partial volume effects and existence of variation of regional gray matter volume. Thus, additional studies are needed to address this issue and to confirm the significance of our findings independently of such effects.

# 5. Conclusions

We demonstrate that postprandial PYY concentrations are positively associated with gray matter volume of the bilateral caudate nucleus and negatively correlated with caudate rCBF. Furthermore, caudate activity is negatively associated with prefrontal and limbic regions implicated in reward and food related behavior, indicating that peripheral PYY may modulate eating behavior via central effects on striatal networks.

# Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Weise et al.



#### Figure 1.

Glass brain (A) and t-score map (B) of regional gray matter volume (GMV) positively associated with postprandial PYY concentrations. (C) t-score map of positive associations between regional GMV and PYY response.

Maps are thresholded at p<0.001 and a cluster size of 100 continuous voxels (glass brain only), uncorrected for multiple comparison; rACC right anterior cingulate, IGP left globus pallidus, ICN left caudate nucleus, rCN right caudate nucleus, IMTG left medial temporal gyrus, IFG left fusiform gyrus, rMTG right medial temporal gyrus;

(D) Correlation between gray matter (GM) values (adjusted for TIV) of the left and right caudate peak voxel with postprandial PYY values (adjusted for age, sex, PFAT).

Weise et al.



#### Figure 2.

Pearson correlation of postprandial PYY (adjusted for age, sex, PFAT) and mean rCBF of the bilateral caudate nucleus (A) and the bilateral caudate body (B). T-score map (C) of prefrontal and limbic/paralimbic brain regions (rCBF) showing significant (p<0.05 cluster-level corrected) negative associations with mean caudate nuclei rCBF. Maps are thresholded at p<0.005 and a cluster size of 50 continuous voxels, uncorrected. T-score is indicated by color bar.

#### Table 1

# Characteristics of the study population

Age	31.3±8.7		
Sex female/male*	18/2		
PFAT (%)	32.2±8.4		
BMI	31.2±9.6		
Hormonal / metabolic	measures	Fasting	Postprandial
PYY (pg/ml)		25.0±10.6	42.5±18.7
Glucose (mmol/l)		4.5±0.5	5.4±0.5
Insulin (pmol/l)		155.6±31.3	485.5±319.5
Appetite ratings			
Hunger		68.9±27.7	18.9±23.2
Desire		69.9±25.5	21.5±25.5
Fullness		16.6±16.3	75.1±24.7
Prospective foodintake		64.3±24.7	20.0±18.3

All results apart from \*are presented as mean  $\pm$  SD; PFAT percentage body fat;

Table 2

Associations of gray and white matter regions with  $\mathrm{PYY}^*$ 

Region, Brodmann Area <sup>a</sup>	<u>INM</u>	, coordi	nates	L	Cluster size
	x	y	2		
GM regions positively associated with PYY					
Fasting PYY					
L. cerebellum, posterior lobe	9-	-64	-20	5.30	509
R. cerebellum, posterior lobe	-10	-58	-21	3.19	
Postprandial PYY					
L. caudate nucleus, body	ε	7	6	6.18	069
R. caudate nucleus, body	9	4	6	5.48	
PYY response					
L. fusiform gyrus, BA37	-39	-40	-15	6.71	168
L. globus pallidus	<b>%</b>	7	1	5.72	224
R. temporal lobe, BA37	50	-42	-12	5.53	335
R. caudate nucleus, head	10	4	1	5.01	373
L. middle temporal gyrus, BA37	-45	-58	-2	4.75	170
R. cingulate gyrus, BA32	15	33	28	4.37	139
R. anterior cingulate gyrus, BA32	4	33	24	4.25	
GM regions negatively associated with PYY					
none					
WM regions positively associated with PYY					
Fasting PYY					
R. inferior frontal gyrus, BA9	4	×	25	6.04	145
Postprandial PYY					
none					
PYY response					
none					
WM regions negatively associated with PYY					
Fasting PYY					
none					

Region, Brodmann Area <sup>d</sup>	<u>qINW</u>	coordin	lates	Ŧ	Cluster size
	x	у	ч		
Postprandial PYY					
none					
PYY response					
Extranuclear WM, L. Thalamus <sup>d</sup>	-3	-24	12	4.49	123
* Results significant at a threshold of p<0.001 and i	cluster	size of	100 coi	atinuous	s voxels, uncorrec
$^{a}$ Indicates nearest gray matter.					
$^{b}$ MNI: Montreal Neurological Institute.					

#### Table 3

Correlations between caudate rCBF and PYY concentrations in the postprandial state\*

Region	R	Р
Caudate nuclei, bilateral	-0.60	<0.02
Left	-0.57	0.02
Right	-0.61	0.01
Caudate body, bilateral	-0.65	<0.01
Left	-0.62	< 0.01
Right	-0.64	< 0.01
Caudate head, bilateral	-0.12	0.67
Left	-0.25	0.36
Right	0.07	0.81
Caudate tail, bilateral	-0.69	<0.01
Left	-0.81	< 0.001
Right	-0.46	0.07

\* Pearson partial correlation adjusted for age, sex and PFAT; rCBF regional cerebral blood flow, PFAT percentage body fat;

Table 4

Brain regions negatively associated with caudate nucleus  $\mathrm{rCBF}^*$ 

Revion. Brodmann Area (BA) <sup>d</sup>	MNI	coord	inates	F	Cluster size
	×	v	й		
Fasting state					
R. middle frontal gyrus, BA8	26	22	42	6.56	70
R. rectal gyrus, BA11	7	28	-24	6.52	85
R. rectal gyrus, BA11	9	38	-24	4.25	
R. cerebllum, posterior lobe	4	-60	-20	6.02	76
<b>R</b> . cerebellum, anterior lobe $^{\mathcal{C}}$	32	-40	-28	5.12	216
R. fusiform gyrus, BA37	56	-46	-28	4.48	
R. fusiform gyrus, BA36	48	-36	-28	4.40	
R. inferior occipital gyrus, BA18	30	-88	-14	4.82	52
R. inferior occipital gyrus, BA18	42	-86	-14	4.11	
R. lingual gyrus, BA19	32	-76	-16	3.14	
R. medial frontal gyrus, BA10	æ	50	ŝ	4.17	50
R. medial frontal gyrus, BA11	7	60	-12	3.32	
L. cerebellum, posterior lobe	-38	-60	-18	4.00	51
L. cerebellum, posterior lobe	-30	-62	-16	3.39	
L. fusiform gyrus, BA37	-42	-54	-24	3.16	
Postprandial state					
R. medial frontal gyrus, $BA10^{\mathcal{C}}$	9	56	ŝ	7.18	172
R. medial frontal gyrus, BA11	9	48	-14	3.40	
L. middle temporal gyrus, BA21	-44	10	-36	689	55
R. hippocampus $^{c}$	34	-10	-18	6.72	205
R. hippocampus	26	-10	-22	4.87	
R. hippocampus	32	-9	-28	4.61	
R. cerebellum, anterior lobe	18	-34	-30	6.49	69
R. cerebellum, anterior lobe	32	-44	-28	4.98	
R. cerebellum, anterior lobe	40	-40	-30	4.79	

Region, Brodmann Area (BA) <sup>a</sup>	<b>MNI</b> <sup>b</sup>	coordi	nates	L	Cluster size
	x	y	2		
R. cerebellum, posterior lobe $^{c}$	16	-66	-20	5.43	403
R. middle occipital gyrus, BA19	36	-80	-14	5.19	
R. lingual gyrus, BA18	20	-84	-12	5.16	
L. cerebellum, anterior lobe	-24	-42	-26	5.25	133
L. cerebellum, posterior lobe	-36	-54	-20	5.09	
L. fusiform gyrus, BA20	-42	-40	-24	4.67	
R. cerebellum, posterior lobe	56	-58	-24	4.91	82
R. cerebellum, posterior lobe	38	-62	-22	4.20	
R. Occipital lobe, subgyral WM, BA37	44	-52	-18	3.95	
*					

Results significant at a threshold of p<0.005 and a cluster size of 50 continuos voxels, uncorrected for multiple comparisons.

a indicates nearest gray matter region

b\_\_\_\_\_\_\_MNNI: Montreal Neurological Institute.

 $c_{p<0.05}$  cluster-level corrected