IN VIVO QUANTIFICATION OF GLUTAMATERGIC ABNORMALITIES IN BEHAVIORAL VARIANT FRONTOTEMPORAL DEMENTIA

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ABSTRACT

Introduction: Although the pathogenesis underlying behavioural variant frontotemporal dementia (bvFTD) has yet to be fully understood, glutamatergic abnormalities have been hypothesized to play an important role. The aim of the present study was to determine the availability of the metabotropic glutamate receptor type 5 (mGluR5) using a novel positron emission tomography (PET) radiopharmaceutical with high selectivity for mGluR5 ([¹¹C]ABP688) in a sample of bvFTD patients. In addition, we sought to determine the overlap between availability of mGluR5 and hypometabolism, as measured using [¹⁸F]FDG PET and voxel based morphometry (VBM). Methods: Availability of mGluR5 and glucose metabolism ([¹⁸F]FDG) were measured in bvFTD (n=5) and cognitively normal (CN) subjects (n=10). ^{[11}C]ABP688 binding potential maps (BP_{ND}) were calculated using the cerebellum as a reference region, with $[^{18}F]FDG$ standardized uptake ratio maps (SUV_R) normalized to the pons. Grev matter (GM) concentrations were determined using VBM. Voxel-based group differences were obtained using RMINC. Results: BvFTD patients showed widespread decrements in ^{[11}C]ABP688 BP_{ND} throughout frontal, temporal and subcortical areas. These areas were likewise characterized by significant hypometabolism and GM loss, with overlap between reduced [¹¹C]ABP688 BP_{ND} and hypometabolism superior to that for GM atrophy. Several regions were characterized only by decreased binding of [¹¹C]ABP688. **Conclusion:** The present findings represent the first in vivo report of decreased availability of mGluR5 in bvFTD. This study suggests that glutamate excitotoxicity may play a role in the pathogenesis of bvFTD and that [¹¹C]ABP688 may prove a suitable marker of glutamatergic neurotransmission *in vivo*.

RÉSUMÉ

Introduction: Bien que la pathogenèse sous-jacente à la variante comportementale de la démence fronto-temporale (vcDFT) n'ait pas encore été bien comprise, les anomalies glutamatergique ont été retenues comme ayant un rôle important dans cette équation. Le but de la présente étude était de déterminer la disponibilité du récepteur métabotropique du glutamate de type 5 (mGluR5) à l'aide d'une nouvelle tomographie par émission de positons (TEP) radiopharmaceutique possédant une haute sélectivité pour les mGluR5 ([¹¹C]ABP688) testé sur un échantillon de patients vcFDT. De plus, nous avons cherché à déterminer le chevauchement entre la disponibilité des mGluR5 et la neurodégénérescence, telle que mesurée en utilisant la base de [¹⁸F]FDG et voxel based morphométrie (VBM). Méthodes: Disponibilité des mGluR5 et le métabolisme du glucose ont été mesurés dans vcDFT (n=5) et sujets normaux (SN; n=10). $[^{11}C]ABP688$ binding potential maps (BP_{ND}) ont été calculées en utilisant le cervelet comme région de référence, avec [¹⁸F]FDG standardized uptake ratio maps (SUV_R) normalisée au pons. Les concentrations de la matière grise (MG) ont été déterminées à l'aide de VBM. Les différences identifiées entre les groupes assujettis à voxel ont été obtenues par le biais de RMINC. Résultats: Les patients vcDFT ont montré des décroissances généralisées dans [¹¹C]ABP688 BP_{ND} dans les zones frontales, temporales et sous-corticales. Ces zones étaient également caractérisées par un hypométabolisme significatif et la perte de MG, avec un chevauchement entre la réduction $[^{11}C]ABP688 BP_{ND}$ et un hypométabolisme supérieur à celle de l'atrophie MG. De plus, plusieurs régions ont été caractérisées uniquement par la diminution de la liaison de [¹¹C]ABP688. Conclusion: Les résultats actuels représentent une première dans le rapport in vivo de diminution de la disponibilité de mGluR5 dans vcDFT. Cette étude suggère que l'excitotoxicité

du glutamate peut jouer un rôle dans la pathogenèse de vcDFT et que [¹¹C]ABP688 peut s'avérer un marqueur approprié de neurotransmission glutamatergique *in vivo*.

CHAPTER 1: INTRODUCTION AND BACKGROUND

INTRODUCTION

Frontotemporal dementia (FTD) is the clinical diagnostic term used to describe patients presenting with a range of dementia syndromes secondary to focal neurodegeneration of the frontal and anterior temporal lobes. The spectrum of pathologies underlying these focal changes are regrouped under the heading of frontotemporal lobar degeneration (FTLD), and include the microtubule associated protein tau (MAPT), the TAR DNA binding protein of 43 kDa (TDP), and the tumor associated protein fused in sarcoma (FUS) (Mackenzie et al. 2010). In addition to the diagnostic subgroups of progressive non-fluent aphasia (PNFA) and semantic dementia (SD), FTD encompasses a behavioral variant (bvFTD), characterized by executive impairments and progressive deterioration of personality, and social function (Neary et al. 1998). Epidemiological studies suggest that FTD is the second most common cause of early onset (before 65 years of age) dementia after Alzheimer's disease (Ratnavalli et al. 2002; Rosso et al. 2003), with bvFTD representing an estimated 60% of cases (Josephs et al. 2006).

While the mechanisms of FTLD underlying bvFTD have yet to be solved, altered glutamatergic transmission is hypothesized to play a role given the selective loss of glutamatergic projection neurons in this patient population (Ferrer 1999). As the main excitatory neurotransmitter in the central nervous system, glutamate exerts its effects via ionotropic and metabotropic receptor families. In contrast to ionotropic receptors such as N-methyl-D-aspartate (NMDA)—which are coupled to ion fluxes through membrane ion channels (Bowie 2008; Lodge 2009)—metabotropic receptors are coupled via G proteins to phospholipase C activation (Masu et al. 1991). In the case of bvFTD, postmortem studies have shown declines in NMDA receptor populations within frontotemporal areas (Procter et al. 1999), suggesting that glutamate

excitotoxicity (Dodd 2002)—a detrimental scenario involving loss of cortical and subcortical neurons secondary to enhanced glutamatergic calcium signaling—may be at play, possibly in a manner similar to that seen in other neurodegenerative diseases (Rothstein 1995; Shaw 1994).

RATIONALE

Despite an evidence base suggesting glutamatergic dysfunction in bvFTD, these abnormalities have yet to be systematically characterized in *vivo* due the absence of suitable probes for use with molecular neuroimaging. Quantification of the metabotropic glutamate receptor type (mGluR5) however, has become possible *in vivo* using [¹¹C]ABP688, a positron emission tomography (PET) radiopharmaceutical with high selectivity for mGluR5 (Ametamey et al. 2006; Ametamey et al. 2007). Recent work in the area of depression has shown [¹¹C]ABP688 to be capable of identifying reductions in mGluR5, with decreased binding found in the prefrontal cortex, cingulate, and insula, regions known to be affected in bvFTD (Deschwanden et al. 2011). Given the extensive co-localization of NMDA and mGluR5 receptors (Alagarsamy et al. 2002; Luccini et al. 2007)—as well as their physiological (Attucci et al. 2001; Benquet et al. 2002; Groveman et al. 2012) and physical links (Ango et al. 2002; Thomas 2002)—decreased binding of [¹¹C]ABP688 may reflect elevated levels of synaptic glutamate, a possible sign of early NMDA receptor mediated excitotoxicity.

RESEARCH OBJECTIVES

The overall aim of the study presented herein was to gain new insight into the potential role of mGluR5 in bvFTD. The main objectives of the study conducted were:

1) To quantify the availability of mGluR5 in bvFTD using $[^{11}C]ABP688$.

2) To determine the topographic overlap between [¹¹C]ABP688 findings and neurodegeneration, as measured using [¹⁸F]fluorodeoxyglucose ([¹⁸F]FDG) PET and voxel based morphometry (VBM).

HYPOTHESES

1) Binding of [¹¹C]ABP688 would be reduced in frontal, temporal and subcortical regions.

2) Overlap between [¹¹C]ABP688, [¹⁸F]FDG and VBM would be partial.

BACKGROUND

FTD Diagnostic Criteria —Historical Aspects

In the 1980s, research groups in Lund, Sweden (Brun 1987) and Manchester, UK (Neary et al. 1988) began publishing large clinicopathological case series of patients exhibiting progressive focal degeneration of frontal and anterior temporal lobes. This new neurodegenerative entity was named FTD, with consensus based diagnostic and research criteria—specifying core diagnostic, supportive, and exclusion features—released in 1994 (Brun et al. 1994). Core affective and behavioral symptoms included loss of insight as well as personal and social awareness, disinhibition, impulsivity, stereotypies, hyperorality, and utilization behaviors. Poor performance on measures of executive functioning—in the absence of significant amnesia, aphasia, or perceptuospatial impairment—and a progressive reduction of speech were likewise consistent with a diagnosis of FTD. Though an important first attempt at defining FTD, the Lund-Manchester criteria possessed several limitations, including the absence of operational definitions and no mention of the relative importance of the various diagnostic features (Rascovsky et al. 2007).

In 1998, the Lund and Manchester criteria were further refined by Neary and colleagues (Neary et al. 1998). In addition to renaming the pathological spectrum underlying FTD as FTLD, clinical descriptions were provided for the 3 most common FTLD phenotypes: behavioral variant FTD (bvFTD), and the language variants of progressive nonfluent aphasia (PNFA) (Mesulam 1982) and semantic dementia (SD) (Hodges and Patterson 1996; Hodges et al. 1992; Snowden 1999). Further, a distinction was made between core and supportive diagnostic features, with operational definitions and occasional examples provided for each.

Recognizing that previous criteria for FTLD and associated neurobehavioral syndromes were largely designed for research purposes, McKhann and colleagues (McKhann et al. 2001) proposed a third set of criteria designed to facilitate the identification and timely referral of patients. The clinical spectrum associated with FTLD was renamed FTD, with criteria simplified into those for an insidious and progressive change in behavior and those for gradual and progressive changes in language function, effectively collapsing PNFA and SD into a progressive aphasic category. Though a useful heuristic on clinical grounds, the McKhann criteria lacked the specificity required for research purposes (Rascovsky et al. 2007).

Limitations of the Neary Criteria for bvFTD and the Revised FTDC Criteria

Despite their widespread acceptance and use in both research and clinical practice, several limitations characterize the Neary Criteria. Among them, the large number of features, making routine application in clinical settings problematic, as well as the use of ambiguous descriptors and the lack of a probabilistic framework with respect to diagnostic certainty (Rascovsky et al. 2007). Moreover, a number of studies have shown their relative restrictiveness early on the disease course, an important limitation given that disease-modifying treatments are likely to be most effective in the early stages of bvFTD (Mendez et al. 2007; Piguet et al. 2009; Rascovsky et al. 2007).

In an effort to address the limitations of the 1998 criteria and to incorporate advances in the characterization and diagnosis of bvFTD, the International bvFTD Criteria Consortium (FTDC) was formed, and tasked with the development of revised diagnostic guidelines. In contrast to earlier guidelines, the FTDC criteria are hierarchical, incorporating levels of diagnostic certainty (Rascovsky et al. 2011). A diagnosis of possible bvFTD is made on clinical

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grounds alone, requiring three of six behavioral/cognitive symptoms, including behavioral disinhibition, apathy inertia, loss of empathy, perseverative, stereotyped or or compulsive/ritualistic behavior, hyperorality and a neuropsychological profile showing executive deficits with relative preservation of memory and visuospatial functions. The flexibility of this classification scheme allows for variability of symptom presentation at onset, with the attendant aim of identifying patients at the mildest stages of disease. In addition to meeting criteria for possible bvFTD, significant functional decline and supportive imaging results are required for the diagnosis of probable byFTD. Finally, the classification of byFTD with definite FTLD pathology is limited to patients who, in addition to meeting criteria for possible or probable bvFTD, have histopathological evidence of FTLD at biopsy or post-mortem, or harbor a known pathogenic mutation. In a large multicentric study, these criteria proved more sensitive than the Neary criteria—86% versus 53%—in a sample of pathologically verified bvFTD (Rascovsky et al. 2011), though prospective studies addressing the reliability and specificity of these revised guidelines have yet to be conducted.

Epidemiology

Of the different clinical FTD syndromes, the behavioral variant is the most common, accounting for approximately 60% of cases (Josephs et al. 2006). The small number of studies and the inclusion of language variants, however, complicate the accurate ascertainment of bvFTD prevalence. Estimates from the three studies that focused on bvFTD, however, ranged from 2 to 15.4 per 100 000 persons in the 45 to 64 year age range (Harvey et al. 2003; Ikejima et al. 2009; Rosso et al. 2003). The low estimates of prevalence reported in two of these studies are likely

due the use of the 1994 Lund and Manchester Groups criteria (Brun et al. 1994), which are known to possess relatively poor sensitivity (Rascovsky et al. 2007).

Though several studies have reported disease onset in older age (Johnson et al. 2005; Shinagawa et al. 2008)—including 3% in 85 year olds (Gislason et al. 2003)—onset is typically below age 65 (Boronni et al. 2008). With respect to sex distribution, several large studies have reported a male preponderance (Hodges et al. 2003; Rascovsky et al. 2011), however other studies reported no differences (Rascovsky et al. 2005; Roberson et al. 2005; Ratnavalli et al. 2002). Studies of survival in pathologically confirmed cases of bvFTD report median survival times of 8.7 and 9.9 years from diagnosis (Roberson et al. 2005; Chiu et al. 2010), in line with the average of 7.8 years reported in a recent study using autopsy confirmed cases of bvFTD (Rascovsky et al. 2011). Overall, bvFTD is associated with shorter survival and more rapid cognitive and functional decline relative to both additional FTD subtypes and AD (Rascovsky et al. 2005; Roberson et al. 2005).

Clinical Features

The initial symptoms of bvFTD are typically behavioral and personality based, with patients often exhibiting an admixture of apathy and disinhibition (Swartz et al. 1997). Apathy follows loss of integrity in dorsomedial prefrontal structures, and manifests as passivity, social disengagement, a lack of interest in personal affairs, and neglect of personal hygiene (Rabinovici et al. 2008). Disinhibition follows dysfunction within orbital and ventromedial prefrontal areas, and is characterized by an affective shift toward inappropriate joviality, a range of impulsive behaviors including economic extravagances, sexually and socially inappropriate behaviors, verbal dysdecorum (Neary et al. 1998), psychomotor hyperactivity and, more rarely, sociopathic

acts (Mendez et al. 2005). Infrequently, presenting features can include new onset ludomania and the tetrad of hyper-religiosity, hypergraphia, hyposexuality, and viscosity characteristic of Geshwind syndrome (Manes et al. 2010; Postiglione et al. 2008). While all neuropsychiatric features found in bvFTD can occur in other dementias, their early emergence and predominance throughout the disease course typify bvFTD. Stereotyped behaviors are apparent in a majority of patients, and range from repetitive motor behaviors-such as picking, hand rubbing, clapping, and rocking (Mendez et al. 2005)-to more complex compulsions including checking, cleaning, wandering a fixed route, and idiosyncratic hoarding (Mendez et al. 2008; Perry and Miller, 2001). Compulsive self-injurious behavior has also been reported in the form of trichotillomania, picking at fingers to the point of excoriation, and self-biting (Mendez et al. 1997; Passant et al. 2005). In addition, changes in eating behavior are frequent, encompassing overeating despite endorsing satiety, overfilling of the mouth, and idiosyncratic food fads (Woolley et al. 2007). Hyperorality is also frequent, and may include excessive drinking, gum chewing, alcohol and/or tobacco consumption (Gustafson 1987). In a minority of patients with more advanced bvFTD, hyperorality may manifest as the mouthing of non-food items and consumption of inedible objects (Gustafson et al. 1992; Mendez and Foti 1997b).

Social cognition is severely affected in bvFTD. Invariably, patients lose the ability to adopt the perspectives of others, displaying a lack of concern for the implications of their behavior, alongside emotional shallowness and loss of warmth (Gregory and Hodges, 1996; Neary et al. 1998; Rankin et al. 2005). While loss of insight is common at presentation, its presence is not universal (Evers et al. 2007) with reports of both relative preservation and a wide range of impairment on Frontal Behavioral Inventory (FBI) discrepancy scores tied to symptom presence and severity (Banks and Weintraub 2009). More recent work, moreover, suggest that

the loss of insight frequently seen in bvFTD is perhaps more accurately described as indifference or anosodiaphoria rather than anosognosia (Mendez et al. 2008). Considered a core feature in the 1998 diagnostic criteria—alongside insidious onset, gradual progression, and impairment in regulation of personal conduct—loss of insight has been omitted from the recently proposed International Criteria (Rascovsky et al. 2011) on the grounds that its continued inclusion may result in suboptimal discrimination due the fact that its presence is common across numerous neurodegenerative diseases, including AD (Rascovsky et al. 2007).

Nonprogressive bvFTD: bvFTD 'phenocopies'

Recent studies have identified a subset of patients referred to as bvFTD 'phenocopies' owing to the presence of behavioral features characteristic of bvFTD in the absence of brain atrophy or hypometabolism at baseline (Kipps et al. 2009; Kipps et al. 2007). Further, they do not exhibit progressive volume loss, leading some authors to suggest that the syndrome is not due an underlying neurodegenerative etiology. Owing to the near normal life expectancy of these patients, large pathological series have yet to become available, with FTLD pathology absent among the few cases that have examined postmortem (Kipps et al. 2010). As such, the neuropathological correlates of these patients remain unclear.

While being indistinguishable from progressive bvFTD behaviorally, phenocopy cases have been identified on the basis of differential performance on activities of daily living (ADL) tasks. Though phenocopy and progressive bvFTD patients did not differ on the caregiver-based Disability Assessment of Dementia (DAD) scale, progressive patients scored significantly lower on the performance-based Assessment of Motor and Process Skills (AMPS) (Mioshi et al. 2009). Although the AMPS discriminated well between the two groups, a grounded theory derived qualitative rating developed to evaluated the assessment session resulted in better discrimination, and correlated well with MRI ratings (Mioshi et al. 2009b). In a follow up study, only those with significantly impaired DAD total scores were found to exhibit functional decline within a 12-month period despite similar levels of impairment at baseline (Mioshi and Hodges 2009b).

Using neuropsychological measures of executive functioning, the performance of phenocopies was found to be in the normal range across the majority of tasks (Hornberger et al. 2008), with global cognitive screening measures likewise paralleling those of controls (Kipps et al. 2008). Of note, however, is the finding that up to 25% of patients with progressive bvFTD proved normal in terms of their performance on neuropsychological measures at presentation (Hornberger et al. 2008), suggesting that normal scores on neuropsychological measures alone cannot be deemed a reliable exclusion criteria, and that such measures should be corroborated when possible by neuroimaging and/or ADL measures. Performance in the area of social cognition has likewise been identified as a potential discriminator owing to the well-established impairment of bvFTD patients in such areas as Theory of Mind (Lough et al. 2006; Sturm et al. 2008). At present, however, the value of everyday application of such tests in clinical practice remains untested owing to the absence of normative age-related data.

Though the etiology of phenocopy cases remains unknown, it has been suggested that the changes seen in these patients may not be due an underlying neurodegenerative process. Given the important overlap between many of the features of bvFTD and those seen in neuropsychiatric conditions, some authors have proposed that the majority of phenocopy cases tend to be atypical presentations of psychiatric disorders, including late onset bipolar disorder, atypical depression, alcohol abuse, and chronic attention-deficit hyperactivity disorder (Manes 2012; Manes et al.

2010). Additional suggestions include decompensated personality disorders or autism spectrum disorders such as Asperger's syndrome manifest at a level below that required for formal psychiatric diagnosis (Kipps et al. 2010). An additional hypothesis recently emerged from work by Khan and colleagues who reported two bvFTD cases characterized by very subtle atrophy, a relatively stable clinical course, and *C90RF72* hexanucleotide expansions (Khan et al. 2012). This finding may suggest that some phenocopy patients have a more indolent form of bvFTD with TDP-43 pathology.

Neuropathology

The FTLDs comprise a heterogeneous group of diseases that share the finding of bilateral frontotemporal atrophy, microvacuolation, and, to a variable degree, astrocytic gliosis (Piguet et al. 2011). Each disease, however, differs with respect to biochemical signature, comprising inclusion composition, morphology, and distribution (Cairns et al. 2007). In this respect, the microtubule associated protein tau (MAPT) (Hutton et al. 1998), the transactive response DNA binding protein of 43 kD (TDP-43) (Arai et al. 2006; Neumann et al. 2006), and the tumor associated protein fused in sarcoma (FUS) (Kwiatkowski et al. 2009) have been identified as key molecular players underlying neurodegeneration seen in FTLD. Virtually all cases of FTLD can therefore be classified into FTLD-tau, FTLD-TDP, or TDP-FUS, with further subclassification on the basis of inclusion morphology/ lesion distribution (Cairns et al. 2007; Mackenzie et al. 2010; Mackenzie et al. 2009) and—in the case of FTLD-tau—the relative predominance of three or four microtubule binding repeats (Cairns et al. 2007). A fourth category—FTLD-other—is reserved for cases with immunohistochemistry against proteins of the ubiquitin proteosomal system or cases for which the major protein remains unknown (Mackenzie et al. 2010).

In contrast to the progressive aphasic syndromes, which typically exhibit greater association with one histological variant of FTLD over another (PNDA with FTLD-tau and SD with FTLD-TDP (Piguet et al. 2011), bvFTD can be associated with all histological variants of FTLD (Josephs et al. 2006b; Hodges et al. 2004; Kertesz et al. 2005). Cases, however, are generally split approximately 50:50 between FTLD-tau and TDP (Hodges et al. 2004; Shi et al. 2005; Snowden et al. 2007), with a small proportion of cases accounted for by FTLD-FUS (Seelaar et al. 2010). In addition, on the basis of recent clinicopathological studies, some investigators have proposed more specific links between apparent byFTD sub-types and specific molecular pathologies. For instance, FTLD-FUS has been tightly linked to a form of bvFTD characterized by striatal pathology, youthful onset, prominent stereotypies, hypersexuality, and hyperphagia (Snowden et al. 2011), a finding that links previous reports of marked striatal atrophy in patients with bvFTD and stereotypy (Josephs et al. 2008) and striking striatal atrophy in FTLD-FUS (Josephs et al. 2010). The general absence, however, of strong ties between byFTD and any of type FTLD subtype, has led some to question whether the syndrome of bvFTD needs to be revised (Josephs et al. 2011).

Genetics

Approximately 50% of bvFTD patients have a family history of dementia (Rohrer et al. 2009), with an autosomal dominant pattern of inheritance noted in 10 to 30% of pedigrees (Goldman et al. 2005; Seelaar et al. 2008). To date, mutations in seven genes have been association with bvFTD, including progranulin *(GRN)* (Baker et al. 2006; Cruts et al. 2006), the microtubule associated protein tau *(MPAT)* (Gass et al. 2006; Hutton et al. 1998), valosin-containing protein *(VCP)* (Neumann et al. 2007; Forman et al. 2006; Watts et al. 2004), chromatin-modifying

protein 2B (*CHMP2B*) (Skibinski et al. 2005), transactive response DNA-binding protein (*TARDBP*) (Sreedharan et al. 2008), fused in sarcoma (*FUS*) (Kwiatkowski et al. 2009; Vance et al. 2009), and the C9 opening reading frame 72 (*C9ORF72*) (DeJesus-Hernandez et al. 2012; Renton et al. 2011) (for review, see Sieben et al. 2012). While mutations in *MAPT* and *GRN* genes are thought to account for 10 to 20% of familial cases, *C9ORF72* mutations are now believed to be the most common genetic abnormalities in familial bvFTD (Riedl et al. 2014). On pathological grounds, both *C9ORF72* and *GRN* mutations are associated with deposition of the transactive response DNA binding protein of 43 kDa (*TDP-43*), while mutations in *MAPT* are associated with deposition of hyperphosphorylated tau (Josephs et al. 2011).

Structural Imaging

As a syndromic entity bvFTD has classically been viewed as a result of atrophy within the frontal cortices, typically symmetric, though right-sided asymmetry has been reported (Fukui and Kertesz 2000; Rosen et al. 2002; Seeley et al. 2008). More recent studies, however, highlight involvement of rostral limbic areas, including the anterior cingulate, anterior insula, limbic ventral striatum, amygdalae, periaqueductal gray, (Barnes et al. 2006; Boccardi et al. 2005; Rosen et al. 2002; Whitwell et al. 2005) as well as temporal lobe structures such as the hippocampus and parahippocampal gyri (Barnes et al. 2006; Galton et al. 2001; Whitwell et al. 2009). VBM studies have leant support to the concept of bvFTD as a fronto-striatal disorder (Looi et al. 2012), on the basis of gray matter volume reduction in the anterior medial frontal cortex, and anterior cingulate, extending to the insular cortex, and subcortical striatal regions (Pan et al. 2012). VBM studies have, furthermore, helped delineate four anatomical subtypes, with two of these of these subtypes exhibiting predominant gray

matter volume loss in the temporal lobe (temporal-dominant, temporoparietal dominant), with gray matter volume loss in the remaining two subtypes predominant in the frontal lobes (frontal dominant, frontotemporal dominant). Importantly, subtypes differed on measures of confrontation naming, episodic memory, and executive function, though not in terms of behavioral severity (Whitwell et al. 2009b).

Using deformation-based morphometry, Cardenas et al. confirmed frontal and anterior temporal lobe atrophy in bvFTD, and, in addition, reported significant atrophy in the pons, midbrain, inferior/superior colliculus, and the thalamus (Cardenas et al. 2007). Atrophy has, moreover, been found to vary as a function of behavioral profile (Cummings 1993; Hodges 2001; Josephs et al. 2006c Le Ber et al. 2006; Liu et al. 2004; Massimo et al. 2009; Snowden et al. 2001; Whitwell et al. 2006). Specifically, apathetic patients have been shown to exhibit greater atrophy within dorsolateral and medial frontal cortices, with disinhibited patients exhibiting greater atrophy within medial orbitofrontal cortices and temporal lobe (Le Ber et al. 2006; Massimo et al. 2009). Recent clinicopathological studies have identified an additional subtype characterized by prominent stereotypy associated with marked striatal atrophy (Josephs et al. 2008).

Functional Imaging

Functional imaging in bvFTD using [¹⁸F]FDG has shown significant hypometabolism in superior, middle, and prefrontal gyri, orbitofrontal and medial frontal areas, as well as the cingulate gyri. Hypometabolism was also noted in the anterior/ventral temporal lobe, the left inferior parietal lobule, the insula, the uncus, right cerebellar tonsil, and in subcortical structures including the putamen, globus pallidus, dorsomedial thalamus, hypothalamus, and pulvinar (Jeong et al. 2005).

In line with previous findings (Grimmer et al. 2004; Kamo et al. 1987) metabolic declines in bvFTD, while predominant in frontal, anterior temporal, and subcortical structures, are more widespread than previously recognized (Ishii et al. 1998). Moreover, subtype specific metabolic signatures have been identified, with the apathetic subtype associated with hypometabolism predominantly in the dorsolateral prefrontal cortex, and the disinhibited subtype associated with marked metabolic reductions in limbic structures including the cingulate cortex, hippocampus/amygdala, and nucleus accumbens (Franceschi et al. 2005).

Studies using ^{99m}Tc-ECD SPECT have revealed considerable anterior hypoperfusion, particularly affecting the frontal cortex, anterior cingulate, anterior temporal cortex, and parahippocampi and hippocampi, with relative sparing of the posterior cingulate, as well as of the parietal and occipital cortices (Basely et al. 2013). Importantly, relative to AD, bvFTD patients showed greater declines in regional cerebral blood flow in medial temporal lobe (MTL) structures, though characterized by a more anterior/parahippocampal topography. Both global and MTL specific findings are in line with findings from studies using ^{99m}Tc-HMPAO SPECT (Neary et al. 1987; Talbot et al. 1998; Varma et al. 2002).

Neuroreceptor Imaging

Postmortem studies of FTLD patients point to dysfunction of the serotonergic system, with significant neuronal loss in the nucleus centralis superior and nucleus raphe dorsalis, suggesting involvement of ascending serotonergic projection fibers (Yang and Schmitt 2001). Furthermore, a significant loss of serotonergic receptors was observed in frontal and temporal cortices of postmortem brain tissue sections from FTLD patients (Procter et al. 1999). PET imaging using $[^{11}C]MDL$ revealed a marked reduction of 5HT_{2A} receptors in orbitofrontal, medial frontal and

cingulate cortices, as well as in the mesencephalon, the latter finding supporting the hypothesis of involvement of the ascending serotonergic system (Franceschi et al. 2005).

Extrapyramidal deficits have been observed in bvFTD following neurodegeneration of the basal ganglia, (Brun and Passant 1996) substantia nigra and locus ceruleus (Mann et al. 1993). Using [¹¹C]CFT PET, presynaptic dopaminergic function within the nigrostriatal system has been shown to be reduced in bvFTD, with decreased binding in both the caudate and putamen, a finding related to severity of extrapyramidal symptomatology (Rinne et al. 2002). Similar findings were obtained using ¹²³I-FP-CIT SPECT, with motor UPDRS scores shown to correlate negatively with ¹²³I-FP-CIT uptake (Sedaghat et al. 2007). Moreover, in a family with FTD and parkinsonism linked to chromosome 17 (FTDP-17), presynaptic dopaminergic deficits within the striatum were demonstrated using SPECT and a tropane derivative targeting the dopamine transporter (Sperfeld et al. 1999).

Amyloid PET Imaging

While cerebral amyloidosis is not a characteristic neuropathological feature of bvFTD many patients with pathologically verified FTLD have dementia due to AD as antemortem clinical diagnosis. Moreover, 10-40% of patients diagnosed with FTLD are found to have AD postmortem (Alladi et al. 2007; Varma et al. 1999). Accurate diagnosis can prove challenging given common clinical features (Galton et al. 2000; Graham et al. 2005) and increasingly apparent anatomic overlap of supposedly distinct topographic signatures of atrophy and metabolic decline, as revealed by MRI and FDG-PET (Foster et al. 2007; Rabinovici et al. 2007b; Womack et al. 2011).

In small scale studies conducted to date, molecular imaging of amyloid is nevertheless

apparently useful in differentiating FTLD from AD (Engler et al. 2008; Rabinovici et al. 2007; Rowe et al. 2007), with the majority of FTD patients showing no retention of [¹¹C]PIB or [¹⁸F]AV45 (Vardy et al. 2010). Few in number, reports of amyloid-positive FTLD cases may represent false positives, comorbid FTLD/AD pathology, or the frontal variant of AD (Rabinovici et al. 2007).

Glutamatergic neurotransmission, and the excitotoxicity hypothesis

As the principle excitatory neurotransmitter in the human central nervous system, glutamate is involved in the regulation of virtually all aspects of cognition, perception, and behavior (Schaeffer and Duplantier 2010). Following its synthesis from glutamine in the synaptic nerve terminal, glutamate is translocated into synaptic vesicles via the action of vesicular glutamate transporters (Liguz-Lecznar and Skangiel-Kramska 2007), and released into the synaptic cleft in response to an action potential. Synaptic clearance of glutamate is mediated by a family of excitatory amino acid transporters, expressed predominantly on astrocytes (Bunch et al. 2009). Following astrocytic reuptake, glutamate is converted to glutamine and stored until additional levels of glutamate are required.

Glutamate signaling is mediated through two major receptor families, ionotropic and metabotropic. Directly coupled to membrane ion channels, ionotropic receptors (iGluRs) are predominantly expressed post-synaptically and include NMDA, (3-hydroxy-5-methylisoxazol-4-yl) propionic acid (AMPA) and kainate (KA) (Dingledine et al. 1999). The metabotropic glutamate receptors (mGluRs) comprise a family of eight G-protein coupled receptor subtypes, and can be divided into three groups on the basis of sequence homology, ligand binding, and G-protein coupling specificity. Group I receptors (mGluR 1 and 5) are typically localized post-

synaptically, and activate the phospholipase C pathway via Gq signaling (Karim et al. 2001; Warwick et al. 2005), Group II (mGluR 2 and 3) and III (mGluR 4, 6, 7, and 8) receptors are preferentially located pre-synaptically, and couple to Gi/Go, signaling via inhibition of adenylate cyclase (Pin and Duvoisin 1995; Conn and Pin 1997; Harrison et al. 2008). In contrast to iGluRs—which are involved primarily in fast synaptic transmission—mGluRs generally exert a neuromodulatory role, mediating slower responses to synaptic glutamate, in keeping with their second messenger signaling mechanism (Niswender and Conn 2010).

While a tightly regulated process under normal physiological conditions, various pathophysiological conditions are capable of disrupting glutamatergic signaling, resulting in excessive levels of extracellular glutamate. The concept of neuronal degeneration due the resulting over-stimulation of glutamate receptors-the so-called excitotoxicity hypothesis-was first formulated by Olney (Olney 1978) and is well supported experimentally, both in vitro and in vivo (Doble 1999). In contrast to classical excitotoxicity—where neuronal damage is induced by acute elevation of glutamate, such as in cerebral ischemia and neurotrama—milder, more chronic elevations of glutamate are believed to be at play in the case of neurodegenerative diseases (Doble 1999). In this scenario, despite normal levels of synaptic glutamate, weakened postsynaptic neurons become sensitized to glutamate stimulation, resulting in elevated levels of intracellular levels of Ca^{2+} and resulting neuronal death. In both forms of excitotoxicity, however, the NMDA receptor is believed to be the primary mediator of Ca^{2+} related neuronal injury, with this mechanism of neuronal damage appearing to be involved in a wide range of neurodegenerative diseases (for review, see Kalia et al. 2008). Importantly, mGluR5 has emerged as an attractive therapeutic target given its ability to modulate NMDA receptor currents (Alagarsamy et al. 1999).

Structure, Function, and Regional Expression of MGluR5

MGlu5 receptors possess a large extracellular N-terminal domain and seven transmembrane (7TM) spanning regions interjoined by three intracellular/extracellular loops (Spooren et al. 2003). The N-terminal domain is bi-lobed in structure, with the lobes separated from the transmembrane region of the receptor by a cysteine-rich region, forming a cavity where glutamate binds (Costantino et al. 2001; Costantino and Pellicciari, 1996). Upon binding of glutamate to its orthosteric site, the lobes enclose the crevice, with glutamate thought to provide conformational stability via the generation of bonds with residues of both lobes (Bessis et al. 2000; Kunishima et al. 2000). In addition to this 'Venus flytrap' mechanism of receptor activation, the cysteine-rich region is though to constitute a flexible spacer, allowing mobilization of the glutamate binding site towards the transmembrane domains (Hermans et al. 2001). Importantly, mGlu5 receptors can assume distinct homooligomeric or heteromeric conformations—usually as disulfide-linked dimers—that seems to be very sensitive to glutamate binding to the orthosteric site (Romano et al. 1996; Cabello et al. 2009). More specifically, glutamate binding seems to stabilize and change the orientation of mGlu5 extracellular domain. As a consequence, the C-terminal part of the extracellular domain, moves closer, which seems to trigger the switch to dimeric states. (Kunishima et al. 2000).

MGluR5 expression levels are moderate in the olfactory tubercle, the anterior olfactory nucleus as well as in the main and accessory bulb. Moderate to high receptor densities characterize the limbic and neocortex (layers I-VI), including the subiculum, entorhinal, cingulate, and piriform cortical regions (Romano et al. 1995). Within the amygdala, moderate receptor levels are found, with high expression levels in basal ganglia input regions (nucleus accumbens and striatum) and moderate levels in output regions (globus pallidus, substantia nigra,

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subthalamic nucleus). The hippocampus shows high receptor levels in the cornu ammonis subfields 1 and 3, as well as in the molecular layer and hilus of the dentate gyrus. With the exception of the lateral septal nucleus, the septum and basal forebrain exhibit relatively low expression levels of mGluR5, as do the thalamus, hypothalamus, midbrain, and cerebellum (Romano et al. 1995).

Quantification of mGluR5 using [¹¹C]ABP688 PET

The discovery of non-amino-acid molecules with high selectivity and affinity for the mGluR5 transmembrane allosteric binding site has led to advances in our ability to quantify mGluR5 density. In this respect, the radiopharmaceutical 3-(6-methyl-pyridin-2-ylethynyl)- cyclohex-2-enone-O-11C-methyloxime ([¹¹C]ABP688) has been developed for assessment of mGluR5 availability *in vivo* using PET imaging (Ametamey et al. 2006). A negative allosteric modulator for mGluR5, [¹¹C]ABP688 exhibits high selectivity and favorable kinetics. After intravenous administration, [¹¹C]ABP688 distributes to various organs, in particular the liver, gall bladder and kidneys, following fast liver metabolism (Treyer et al. 2008; Treyer et al. 2007). Due to its optimal lipophilicity (logD=2.4), [¹¹C]ABP688 easily crosses the blood brain barrier (BBB; fist pass extraction fraction > 0.9) and binds with high affinity (K_d=1.7 ± 0.2 nM) to mGLUR₅ (Treyer et al. 2008). Moreover, [¹¹C]ABP688 metabolites are hydrophilic and do not cross the BBB, and therefore do not contribute to brain activity.

Previous *in vivo* studies of [¹¹C]ABP688 PET in healthy subjects showed high binding in known mGluR5 dense regions such as the striatum, the hippocampus, and the anterior cingulate cortex (Ametamey et al. 2007; Treyer et al. 2007). While preliminary kinetic analysis using arterial input function (gold standard method) indicates that [¹¹C]ABP688 binding is better

described by a 2-tissue compartment model (Ametamey et al. 2007b), simplified reference tissue models have been shown to be suitable for *in vivo* quantification using PET owing to negligible binding in the cerebellum (Elmenhorst et al. 2010). The [¹¹C]ABP688 imaging outcome described in chapter 2 is the non-displaceable binding potential (BPND), which simultaneously reflects the availability and affinity of mGLUR₅ binding sites for [¹¹C]ABP688. It is an "equivalent" *in vivo* measure of the autoradiography single dose experiment parameter B_{max}/K_d.

Quantification using PET and the Simplified Reference Tissue Method

In vivo quantification of radioligand-receptor binding using PET requires the estimation of tracer uptake, washout, retention in tissue as a function of time (tissue response function), and tissue delivery (arterial input function). In order to obtain measurements of plasma tracer levels in arterial blood, arterial cannulation is required. For a majority of receptor radioligands, however, total plasma concentration measurements are insufficient, with the determination of the fraction of labelled metabolites over time an additional requirement (Lammertsma and Hume 1996). In addition, though arterial blood sampling has been shown to be safe in research subjects (Everett et al. 2009), the invasiveness and complexity of this method renders it unsuitable for routine use in certain patient populations.

In order to circumvent the complexities of arterial input function, a reference tissue model was developed (Lammertsma et al. 1996b; Hume et al. 1992) in which arterial plasma sampling is replaced by the time-activity curve of a reference regions exhibiting negligible specific uptake of the radiotracer in question. This original simplified model fit four parameters. Though estimation of the parameter of interest (binding potential; BP) proved robust, estimates of the remaining parameters—including the plasma to tissue transfer rate—proved imprecise, with slow

convergence rates (Lammertsma and Hume 1996). As such, a three-parameter model was developed (Lammertsma and Hume 1996), which, in addition to yielding BP values essentially the same as those obtained using the four-parameter model, is characterized by rapid convergence and greater stability for remaining parameters (Lammertsma and Hume 1996). Following this, the simplified reference tissue method has been implemented at the voxel level to produce parametric BP images using a basis functions approach (Gunn et al. 1997).

Voxel Based Morphometry

In contrast to traditional techniques for the assessment of atrophy using MRI—which include expert based visual assessment and manual delineation of regions of interest—automated techniques allow for the analysis of atrophy across large groups of individuals without the need for subjective visual ratings and laborious manual tracings. VBM is one such technique, and has proved quite popular since its introduction (Wright et al. 1995; Ashburner and Friston, 2000), given that it is relatively easy to use and has provided biologically plausible results (Whitwell and Josephs 2007; Whitwell and Jack 2005b). Typically using T1-weighted volumetric MRI images, VBM identifies differences in the relative concentration of gray or white matter, while controlling for large-scale differences in gross anatomy. Regional differences in the volume of gray or white matter can likewise be determined through the use of a further processing step know as "modulation" (Good et al. 2001). Finally, regression analysis can be used at the voxel level to determine neuroanatomical correlates of clinical features of interest (Whitwell 2009).

Following spatial normalization—generally achieved using a two-step approach comprising a 12-parameter affine transformation followed by a nonlinear registration using a mean squared difference matching function (Ashburner and Friston, 2000)—images are

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segmented in gray matter (GM), white matter (WM), and cerebrospinal fluid (CSF). Typically, analysis is performed on GM, though WM can also be analyzed, depending on the question of interest. Resulting images are then smoothed, with the dimensions of the smoothing kernel determining the number of voxels averaged at each point (Karas et al. 2003; Whitwell et al. 2009b). In addition to increasing data conformity with respect to the Gaussian field model, smoothing reduces intersubject variability, increasing the validity of parametric tests (Ashburner and Friston, 2000; Salmond et al. 2002). Smoothing likewise increase the sensitivity to detect group differences, though excessive smoothing decreases spatial resolution.

Once the segmented images have been smoothed, parametric statistical analysis is carried out using the general linear model, with the null hypothesis assuming no between group differences. The theory of Gaussian random fields applied to determine significance (Ashburner and Friston, 2000), followed by correction for multiple comparisons to reduce the risk of falsepositive error, typically using the family-wise error (FWE) correction (Friston et al. 1993), or the more lenient false discovery rate (FDR) correction (Genovese et al. 2002). Though VBM findings often prove difficult to validate, studies addressing correspondence between VBM analyses and traditional measurement techniques—including manual volumetry—have shown relatively good agreement (Good et al. 2002; Giuliani et al. 2005; Whitwell et al. 2005c; Davies et al. 2009, Focke et al. 2014), lending support to the biological validity of VBM.

CHAPTER 2: RESEARCH MANUSCRIPT

In vivo characterization of metabotropic glutamate receptor type five abnormalities in behavioural variant FTD

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Introduction

Behavioural variant frontotemporal dementia (bvFTD) is a progressive neurodegenerative syndrome characterized by change in personality, social cognition impairment, and executive dysfunction (Mendez et al. 2008; Swartz et al. 1997). Approaching Alzheimer's disease (AD) as the leading cause of early-onset (before 65 years of age) dementia (Ratnavalli et al. 2002; Rosso et al. 2003), bvFTD arises from a heterogeneous range of pathologies—referred to collectively as frontotemporal lobar degeneration (FTLD)—resulting in degenerative changes within frontal paralimbic, temporal and subcortical brain regions. In most cases, patients show either deposition of the microtubule associated protein tau (tau) or the TAR DNA binding protein of 43 kDa (TDP) (Mackenzie et al. 2010). A minority, however, show a defect in metabolism of the tumor associated protein fused in sarcoma (FUS). The majority of FTLDs can therefore be classified into FTLD-tau, FTLD-TDP, or FTLD-FUS, with further subclassification based predominantly on inclusion morphology and lesion distribution (Mackenzie et al. 2010).

Although the pathogenic mechanisms underlying bvFTD have yet to be fully elucidated, aberrant glutamatergic neurotransmission has been hypothesized to play a role. The primary excitatory neurotransmitter in the mammalian brain, glutamate acts via ionotropic and metabotropic receptors (Schaeffer and Duplantier 2010). Whereas ionotropic receptors mediate fast excitatory neurotransmission, metabotropic glutamate receptors (mGluRs) play an important role in synaptic modulation via regulation of neuronal excitability, transmitter release, synaptic plasticity and glial function. In the case of bvFTD, FTLD has been found to accumulate preferentially within paralimbic and homotypical heteromodal brain regions, areas rich in excitatory glutamatergic pyramidal cells. Indeed, several autoradiographic and immunohistochemical studies in post-mortem bvFTD tissue have provided evidence supporting this hypothesis (Dalfo et al. 2005; Ferrer 1999; Procter et al. 1999), highlighting reduced expression of the N-methyl-D-aspartate (NMDA) ionotropic glutamate receptor. Importantly, activation of mGluR5 was shown to regulate glutamatergic neurotransmission via modulation of NMDA receptor functionality (Llansola and Felipo 2010; Niswender and Conn 2010; Perroy et al. 2008). Moreover, mGluR5 signaling has been shown to be critically involved in the normal cognitive functioning of various neuronal

populations (Schaeffer and Duplantier 2010), including those within FTLD predilection sites (Ferraguti and Shigemoto 2006).

Despite a strong *in vitro* evidence base, glutamatergic abnormalities in bvFTD have yet to be systematically characterized *in vivo* owing to the lack of suitable molecular probes. Using [¹¹C]ABP688—a novel Positron Emission Tomography (PET) radiopharmaceutical with high selectivity for mGluR5 (Ametamey et al. 2006; Ametamey et al. 2007)—we sought to measure mGluR5 availability and to determine the topographic overlap with neurodegeneration within frontotemporal and subcortical brain regions, as indexed using [¹⁸F]fluorodeoxyglucose ([¹⁸F]FDG)-PET and voxel based morphometry (VBM).

Methods

Subjects

Five patients meeting research criteria for probable bvFTD (Rascovsky et al. 2011) were recruited from the McGill Centre for Studies in Aging (MCSA) Alzheimer's Disease Research Unit. Exclusion criteria were (i) past or present use of memantine; (ii) presence of other neurological diseases; (iii) premorbid psychiatric disease or intellectual disability; (iv) history of head injuries and loss of consciousness following head trauma; (v) current (within 1 month) use of psychoactive substances; (vi) parkinsonism as identified using the United Parkinson's Disease Rating Scale (Goetz et al. 2007)—(vii) the presence of any major structural anomaly or signs of major vascular pathology on MRI evaluation (Roman et al. 1993); or the presence of amyloid pathology—as indexed using carbon-11 Pittsburgh Compound B ([¹¹C]PIB) PET—given that bvFTD and the frontal variant of AD are often difficult to differentiate on clinical grounds alone (Alladi et al. 2007). The diagnosis of bvFTD was determined during a multidisciplinary conference taking into consideration available medical, imaging, and complementary laboratory information.

The bvFTD subjects were matched by age and gender to a group of 10 cognitively normal (CN) controls, recruited via advertisements in a local newspaper. CN subjects were identified as individuals

who (i) were independently functioning community dwellers; (ii) did not have neurological or a personal or first degree relative history of psychiatric disorders; (iii) had no cognitive complaints; (iv) had a normal neurological and psychometric examination; (v) were not taking any psychoactive medications; (vi) had no history of head trauma; (vii) showed no signs of vascular pathology on MRI evaluation (Roman et al. 1993) and (ix) had a Mini-Mental State Examination (MMSE) (Folstein et al. 1975) score ≥ 29 , a Neuropsychiatric Inventory (NPI) (Cummings et al. 1994) score of 0, and a Frontal Behavioral Inventory (FBI) (Kertesz et al. 1997) score of 0.

Demographic and clinical data for all subjects are shown in Table 1, with ratings of lobar atrophy (Kipps et al. 2007) and hypometabolism (Poljansky et al. 2011) for bvFTD patients shown in Table 2. All subjects and their caregivers provided written informed consent. The study protocol, approved by the Research Ethics Board of the Montreal Neurological Institute as well as by the Faculty of Medicine Research Ethics Office, McGill University, was carried out in accordance with the Declaration of Helsinki.

PET acquisition

3-(6-methyl-pyridin-2-ylethynyl)-cyclohex-2-enone- O^{-11} C-methyl-oxime ([¹¹C]ABP688) was synthesized as described previously (Elmenhorst et al. 2010), with a radiochemical purity > 99%. The study was performed using a High-Resolution Research Tomograph (HRRT) PET scanner (CTI/Siemens, Knoxville, Tennessee), a brain dedicated tomograph combining high spatial image resolution with high sensitivity. Prior to radiopharmaceutical administration, a 6-minute transmission scan was acquired for scatter and attenuation correction using a [¹³⁷Cs] rotating point source. A 60-minute dynamic list-mode emission scan was started concomitantly with the venous injection of 370 MBq (mean specific activity >500 Ci/µmol) of [¹¹C]ABP688, with emission data acquired in list-mode format, and binned into 26 time frames. For each and every time frame, sets of fully 3D sinograms were generated from the list-mode data (2209 sinograms, span 9, with 256 radial bins and 288 azimuthal angle samples). A time-series of 26 3D images (frame duration: 6 × 30s, 4 × 60s, 8 × 120 s, 3 × 240 s, 5 × 300 s) were then reconstructed from
these sinograms, each 3D image being composed of $256 \times 256 \times 207$ cubic voxels (voxel side-length of 1.21875 mm), using an expectation maximization image reconstruction algorithm with an ordinary Poisson model of the acquired PET data. The reconstruction included full accounting for the normalization, attenuation, and time-dependent scatter of random events. To reduce the partial volume effect, resolution modeling with point-spread function was implemented in the reconstruction (Comtat 2008). Subject head-motion correction was implemented using a data-driven motion estimation and correction method (Costes et al. 2009).

All patients underwent an [¹⁸F]FDG PET scan using a Siemens ECAT EXACT HR + PET device (CTI/Siemens, Knoxville, TN, USA) as part of their clinical evaluation. In keeping with the ALARA radiation safety principle (Natarajan et al. 2013), data was not recollected on the HRRT. After fasting overnight, patients received a venous bolus injection of 185 MBq of ¹⁸F-fluorodeoxyglucose ([¹⁸F]FDG) in a quiet environment. A dynamic scan was performed in 3-dimenstional mode for 10 min under standard resting-state conditions with eyes open, recording 63 transaxial slices simultaneously with an axial resolution of 5 mm full width at half maximum (FWHM) and an in-plane resolution of 4.6 mm. Each collected slice had a thickness of 2.45 mm and a matrix size of 128×128 voxels. After correction for attenuation, scatter, decay and scanner-specific dead time, the PET data were reconstructed by filtered back-projection using a Hann filter (4.9 mm FWHM).

CN subjects had their acquisition conducted on the HRRT, with acquisition parameters identical to those for [¹¹C]ABP688, as described above. Images were reconstructed taking into consideration data acquired between 45-60 minutes only, with reconstruction matching that used for the HR+ data. In order to compare data from the HRRT and HR+ PET scanners, the resolution of the HRRT was matched to the partial volume effect of the HR+. To do so, an anisotropic Gaussian kernel of 5.7x5.7x6.7mm FWHM was used, which was found to be the best match of scanner resolutions through an internal phantom study (unpublished data). In the case of [¹⁸F]FDG, two CN subjects were excluded owing to movement with one patient unable to return for the scan, reducing the sample size for [¹⁸F]FDG to 4 bvFTD and 8 CN.

Magnetic Resonance Imaging

For anatomical co-registration and identification of the volumes of interest (VOI), all subjects underwent a high-resolution T-1 weighted MRI using a Siemens TRIO 3T scanner (Siemens Medical Solutions, Erlangen, Germany). Images were acquired in 3-D (voxel size=1mm³; FOV=256x256 mm; TR=22 ms; TE=9.2 ms; Flip angle=30°), with the scan performed on either on the same day or less than 2 weeks apart from the PET acquisitions, depending on the availability of the research slots.

Imaging analysis

[¹¹C]ABP688 binding potential, non-displaceable (BP_{ND}) values were obtained using the simplified reference tissue method (SRTM) (Gunn et al. 1997), using the cerebellum as a reference region (Elmenhorst et al. 2009; Minuzzi et al. 2009). [¹⁸F]FDG PET frames were summed and standardized uptake value ratio (SUV_R) maps calculated by normalizing the summed image to mean pontine activity for each subject. In order to correct for Partial volume error (PVE), a modified version (Greve et al. 2014; Rousset et al. 2007) of the Muller-Gartner method (Muller-Gartner et al. 1992; Rousset et al. 1998) was implemented using the PVElab software package (https://nru.dk/downloads/software/pveout/pveout.html) (Quarantelli et al. 2004).

Following correction for field inhomogeneities (Sled JG 1998), native MRI volumes were nonlinearly resampled into standardized stereotaxic space, using the high-resolution ICBM template as reference (Fonov et al. 2009). Subsequently, normalized images were classified into grey matter (GM), white matter (WM) and cerebrospinal fluid (CSF) using an automatic algorithm (INSECT) (Zijdenbos A 1998). Voxel-based morphometry (VBM) (Ashburner and Friston, 2000) was carried out on the structural segmented GM images nonlinearly resampled to the standard stereotaxic space after blurring with an isotropic Gaussian kernel of 10mm FWHM. Finally, classified images were resampled to an anatomical template and automatically labeled using a probabilistic atlas based approach (ANIMAL) (Collins DL et al. 1999; Collins and Evans 1997). VOIs yielded by this procedure were subsequently applied to PET BP_{ND} (cerebellum) and SUV_R maps (pons). Voxel-wise analysis maps of [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and VBM values were estimated using a basis functions approach (Gunn et al. 1997), with PET images convolved using an isotropic Gaussian kernel of 6mm FWHM. Parametric maps created in native space were then normalized into MNI space in order to allow for group comparisons. The resulting t-maps, calculated using RMINC (Lerch 2006), show the areas with a significant difference in BP_{ND}, SUV_R, and relative concentration of GM between groups. Those areas were subsequently adjusted for a statistical cluster-wise threshold of p <0.05, and corrected for multiple comparisons using random field theory (Worsley et al. 1998). [¹¹C]ABP688 BP_{ND} local maxima coordinates were used to extract [¹⁸F]FDG SUV_R, and VBM values, in order to compare the magnitude of decline.

Brain regions where all patients differed significantly from controls on the basis of Z scores ≥ 2 calculated using the formula [(individual patient value) - (control mean)/ (control standard deviation)]were calculated for [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and VBM t-maps. These areas were then used to extract raw [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and VBM values, which, after reconversion to Z scores, were plotted using GraphPad Prism 5 software. Overlap maps—[¹¹C]ABP688 BP_{ND} and [¹⁸F]FDG SUV_R, [¹¹C]ABP688 BP_{ND} and VBM, [¹⁸F]FDG SUV_R and VBM—as well as areas showing only reduced [¹¹C]ABP688 BP_{ND} , [¹⁸F]FDG SUV_R, and VBM—were created using MINC tools (http://www.bic.mni.mcgill.ca/ServicesSoftware/MINC). For overlap maps, binary masks were generated by applying the cluster-corrected t-map thresholds to each individual t-map-[¹¹C]ABP688 BP_{ND}, $[^{18}F]FDG SUV_R$, and VBM—setting voxels less than the given threshold to 0 and voxels greater than the threshold to 1. Binary masks were then summed, with voxels having a value of 2 indicating overlap. In order to show areas exhibiting only reductions (e.g. in availability of mGluR5) binary masks were subtracted (e.g. $[^{11}C]ABP688 BP_{ND} - [^{18}F]FDG SUV_R - VBM$), with the range of values in the resulting volume restricted to lie between 0 and 1, removing negative values generated as a result of the subtraction. Finally, volumes visualized color-coded using software DISPLAY were and the (http://www.bic.mni.mcgill.ca/software/Display/Display.html).

Results

Groups differed significantly in terms of MMSE, FBI and NPI (see Table 1). No differences were observed for age at scan, education, sex, or handedness. While all patients showed moderate hypometabolism within frontotemporal regions, atrophy ranged from very mild to moderate (see Table 2). *Z* score maps showing regions with significantly reduced [¹¹C]ABP688 BP_{ND}, hypometabolism and atrophy common to all patients—along with plots showing *Z* scores relative to controls (see Fig 1). Location and coordinates of local maxima for the contrast [¹¹C]ABP688 BP_{ND} CN > bvFTD] are reported in Table 3, along with values for [¹⁸F]FDG SUV_R and VBM using these local maxima. Despite neurodegeneration being more widespread than declines in mGluR5 availability, hypometabolism and atrophy were found to be inferior to reductions in [¹¹C]ABP688 BP_{ND} in a wide range of FTLD predilection sites (see Table 3). Relative to controls, bvFTD patients showed reductions of 65 percent, 30 percent, and 15 percent—for [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and VBM, respectively—on the basis of values extracted from common *Z* score maps.

Voxel-wise analysis of group differences in [¹¹C]ABP688 BP_{ND} revealed declines in mGluR5 availability (85 152 mm³) in orbital, ventromedial, and dorsomedial prefrontal areas (corrected for multiple comparisons, p < 0.05, df=13) (see Fig 2). Declines were likewise noted in the gyrus rectus, anterior cingulate (L>R), right posterior cingulate, superior frontal gyrus (L>R), paracentral lobule (L>R), caudate (L>R), left putamen, insula (R>L), thalamus (L>R), right lingual gyrus, and right cuneus. Additional declines were found in the right dorsolateral, right ventrolateral, and anterior prefrontal cortex, the right superior and middle temporal gyri, as well as in the temporal poles. No significant increases of [¹¹C]ABP688 binding were observed in bvFTD subjects.

Significant hypometabolism was noted among bvFTD patients (116 742 mm³) in extensive prefrontal areas, including the orbitofrontal (R>L), ventromedial and dorsomedial prefrontal (L>R), and the cingulate gyrus (L>R) (corrected for multiple comparisons, p < 0.05) (see Fig 3). Metabolism was significantly reduced in the superior, middle, and inferior frontal gyri as well as in the precuneus and paracentral lobule (L>R). Hypometabolism was also found in the bilateral insula (R>L), uncus/amygdala,

and parahippocampus (L>R), as well as in subcortical structures, including the head of the caudatum and the left thalamus. There was also hypometabolism in the superior, middle, and inferior temporal gyri, temporal poles, and cerebellar tonsils.

Among bvFTD subjects, grey matter loss (88 845 mm³) was predominantly focused in the striatum—including the putamen (L>R) and head of the caudate nucleus bilaterally (corrected for multiple comparisons, p < 0.05) (see Fig 4). There was significant involvement of the thalamus and insula bilaterally, as well as the amygdala, and parahippocampus. Atrophy was also observed in the anterior cingulate (L>R), precuneus (R>L) gyrus rectus, orbitofrontal gyrus, as well as the right superior/middle temporal gyri, though to a lesser degree (see table 3).

Overlap between [¹¹C]ABP688 BP_{ND} and [¹⁸F]FDG SUV_R t-maps was observed in the gyrus rectus (L>R), anterior cingulate/ventromedial prefrontal cortex (L>R), dorsomedial prefrontal cortex (L>R), thalamus (L>R) insula (R>L) and temporal poles (22 379 mm³; see Fig 5). [¹¹C]ABP688 BP_{ND} and VBM findings overlapped in the anterior cingulate/ventromedial prefrontal cortex (L>R), orbitofrontal cortex (R>L), thalamus (L>R), head of the caudate nucleus (L>R) and the insula (R>L) (13 463 mm³; see Fig 6). Overlap between hypometabolic regions and atrophy was noted in medial and lateral orbitofrontal areas, anterior cingulate/ventromedial prefrontal cortex (L>R), dorsomedial prefrontal cortex (R>L), insula (R>L), thalamus (L>R), left amygdala, right hippocampal formation, and the head of the caudate nucleus (R>L) (14 179 mm³; see Fig 7). Though hypometabolism and atrophy were found in frontal, temporal, and subcortical brain regions, these declines were found to be inferior relative to those for [¹¹C]ABP688 BP_{ND} in a wide range of areas, including the gyrus rectus, medial and lateral orbitofrontal cortex, ventromedial prefrontal cortex, left dorsomedial prefrontal cortex, paracentral lobule, frontal pole, left putamen, left insula, lingual gyrus, cuneus, temporal poles, right superior temporal gyrus, inferior and middle temporal gyri, and the right dorso- and ventrolateral prefrontal cortex.

Subtraction of binarized t-maps ($[^{11}C]ABP688 BP_{ND} - [^{18}F]FDG SUV_R - VBM$) showed that decreased binding of $[^{11}C]ABP688$ was unique to the gyrus rectus (R>L), orbitofrontal cortex (R>L), lateral portion of the right head of the caudate nucleus, left putamen, left superior temporal lobe, inferior

temporal lobes, temporal poles (R>L), right posterior cingulate, right ventral/dorsolateral prefrontal cortex, left paracentral lobule, right occipital cortex, and right lingual gyrus (55 742 mm³; see Fig 8). A similar subtraction yielded a volume of 86 616 mm³ for regions displaying only hypometabolism ([¹⁸F]FDG SUV_R – [¹¹C]ABP688 BP_{ND} – VBM)—including the uncus/amygdalae (L>R), the parahippocampus (L>R), bilateral cuneus, posterior cingulate/precuneus (L>R), bilateral insula, medial prefrontal cortex, left posterior paracentral lobule, left frontal operculum, anterior temporal poles, orbitofrontal gyrus/gyrus rectus (R>L), and bilateral cerebellar cortex (see Fig 9). In addition, the orbitofrontal gyrus, the right middle temporal gyrus, right temporal operculum, left anterior insula, posterior insula bilaterally (R>L), ventral amygdala (L>R), left posterior inferior temporal gyrus, anterior cingulate gyrus (L>R), the putamen (L>R) and head of the caudate nucleus bilaterally, the thalamus as well as the posterior portion of the hippocampal formation, bilaterally, were found to be characterized only by GM reductions (VBM – [¹⁸F]FDG SUV_R – [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R and VBM were found to co-exist within the orbitofrontal cortex and temporale lobe, relative to [¹⁸F]FDG and VBM, findings for [¹¹C]ABP688 were more ventral and lateral, respectively.

Discussion

The present findings represent the first *in vivo* report of decreased availability of mGluR5 in bvFTD. In line with recent studies showing reduced binding of [¹¹C]ABP688 in disorders characterized by glutamate excitotoxicity—such as major depressive disorder and temporal lobe epilepsy (Choi et al. 2014; Deschwanden et al. 2011)—our findings may indicate increased glutamate levels in bvFTD. Further, we reproduced previous [¹⁸F]FDG and VBM findings in terms of both the topography of neurodegeneration and its partially asymmetric distribution (Diehl-Schmid et al. 2007; Hornberger et al. 2012; Jeong et al. 2005; Pan et al. 2012). In addition, we showed that the volume of decreased mGluR5 availability was inferior to that for hypometabolism and GM atrophy, and that the overlap between reduced [¹¹C]ABP688

mGluR5 availability were unique to several isocortical, limbic, and paralimbic areas, possibly representing an early sign of pyramidal cell dysfunction. In this respect, the focality of [¹¹C]ABP688 BP_{ND} reductions in the present study is striking given the widespread distribution of mGluR5. In addition, several frontotemporal areas showed hypometabolism and/or GM loss in the absence of reduced [¹¹C]ABP688 binding. Taken together, these findings suggest a differential neuronal vulnerability to FTLD pathology in bvFTD—similar to that seen in other neurodegenerative diseases (Double et al. 2010)—with reduced availability of mGluR5 possibly preceding neurodegeneration within select frontotemporal brain regions.

While at physiological concentrations glutamate is known to play a pivotal role in synaptic plasticity (Balschun et al. 2006; Huber et al. 2001)—with any given function of a given cortical region likely to depend on glutamatergic neurotransmission at some level (Francis 2009)—at high concentrations it has been shown to act as a neurotoxin, promoting neuronal injury and death in animal models (Rao et al. 2001; Rothstein 1996) and in neurodegenerative diseases, including AD (Francis 2003). In the case of AD, accumulation of β -amyloid is thought to inhibit astroglial glutamate uptake, resulting in increased extracellular levels of glutamate, which, under chronic conditions, lead to cell death via sustained elevations in intracellular calcium (Harkany et al. 2000). This excitotoxic scenario may explain decreased binding of [¹¹C]ABP688 in that continued high levels of glutamate may alter the availability of its transmembrane allosteric binding site (Ametamey et al. 2007) by altering mGluR5 conformational states (Cabello et al. 2009; Canela et al. 2009; Changeux and Edelstein 2005; Romano et al. 1996). Indeed, affinity shifts in receptor-radioligand interactions have previously been described in the context of dopaminergic neurotransmission, where the affinity of a D2 PET radiopharmaceutical was altered following an amphetamine challenge (Narendran et al. 2004; Seneca et al. 2006; Wilson et al. 2005).

Recently, an expanded hexanucleotide repeat in the chromosome 9 open reading frame 72 (C9ORF72) was identified as the most common cause of familial FTD and amyotrophic lateral sclerosis (ALS), with mutations associated with deposition of TDP-43 pathology (DeJesus-Hernandez et al. 2011;

Renton et al. 2011). While the pathogenic mechanism(s) by which this repeat expansion could cause disease remain unknown, induced pluripotent stem cell differentiated neurons from C9ORF72 ALS patients were shown to be highly susceptible to glutamate excitotoxicity (Donnelly et al. 2013). Related work on primary cells from TDP-43 transgenic mice showed an increase vulnerability to the toxic effects of excess glutamate (Swarup et al. 2011). Moreover, a recent study involving transgenic mice expressing the FTDP-17 mutation P301L in the human tau gene—resulting in the accumulation of hyperphosphorylated tau—showed a tau dependent impairment of glutamate metabolism (Nilsen et al. 2013). These studies suggest that the pathogenicity of hyperphosphorylated tau and TDP-43—the molecular pathologies accounting for most cases of bvFTD (Mackenzie et al. 2011)—may involve glutamatergic excitotoxicity.

Certain methodological aspects, however, limit interpretation of the present findings. In addition to this study's cross sectional design and small sample size, the absence of histopathological data precludes conclusions about the homogeneity of the sample from the perspective of underlying molecular pathology. As such we were not able to address the possible interplay between different FTLD subtypes and possibly differing effects on mGluR5 availability. Moreover, potential limitations may accompany the use of VBM when applied to atrophic brains (Good et al. 2002).

Despite these caveats, our findings shed light on the possible role of glutamate excitotoxicity in the pathogenesis of bvFTD and suggest that [¹¹C]ABP688 may prove a suitable non-invasive marker of glutamatergic neurotransmission *in vivo*. Larger prospective studies are required to validate these findings, to establish the trajectory of reduced mGluR5 availability relative to other biomarkers of neurodegeneration, and to address the potential link between the dysregulation of glutamatergic neurotransmission and bvFTD symptomatology.

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Conflict of Interest

The authors declare no conflict of interest.

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Tables and Figures

	BvFTD (n=5)	CN (n=10)	P ^a value
Age at scan, Med (IQR), years	65 (7)	63 (2.75)	0.65
Education, Med (IQR), years	10 (5)	16 (4)	0.06
Sex, M/F	3/2	7/3	1.00
Handedness, R/L	5/0	9/1	0.52
MMSE, Med (IQR), max=30	26 (1)	30 (1)	0.03
FBI, Med (IQR), max= 72	20 (0)	0 (0)	0.001
NPI Total, Med (IQR), max= 144	32 (8)	0 (0)	0.001

Table 1 Demographic and clinical data for all subjects.

Due to the small group sizes, data are represented as Med (IQR)= median (interquartile range).

Abbreviations: M/F= Male/Female; R/L= Right/Left; MMSE= Mini-mental state examination; FBI= Frontal Behavioral Inventory; NPI= Neuropsychiatric Inventory.

^a The t test for continuous variables, Fischer's exact test for categorical variables.

Table 2 Ratings of hypometabolism and lobar atrophy in patients with bvFTD.

[¹⁸ F]FDG-PET									MRI		
Subject	Frontal Lobe	Temporal Lobe	Parietal Lobe	Occipital Lobe	Cerebellum	Basal Ganglia	Thalamus	Frontal Lobe	Anterior Temporal Lobe		
1	2	2	1	0	0	0	1	3	2		
2	2	2	1	0	0	0	0	1	1		
3	2	2	1	0	0	0	0	3	3		
4	2	2	1	0	0	0	0	1	2		
5	2	2	1	0	0	0	1	1	2		

Ratings for [¹⁸F]FDG-PET: 0=absent, 1=mild, 2=moderate, 3=strong.

Ratings for MRI: 1=very mild, 2=mild, 3=moderate.

Brain region	X	У	Z	t [11C]ABP688	Р	t [18F]FDG	Р	t _{VBM}	Р
Gyrus rectus L	-9.0	28.2	-24.7	-7.12	>0.0001	-4.19	>0.0001	-2.28	0.0401
Gyrus rectus R	13.8	26.2	-24.7	-6.52	>0.0001	-3.22	0.0092	-2.38	0.0333
Medial orbitofrontal cortex L	-16.0	23.1	-15.2	-8.49	>0.0001	-2.40	0.0373	-0.527	0.6071
Medial orbitofrontal cortex R	15.0	-15.2	-9.66	-9.73	>0.0001	-1.98	0.0076	-0.002	0.9998
Lateral orbitofrontal cortex L	-32.1	36.0	-15.2	-4.63	0.0005	-4.14	0.0020	-4.03	0.0014
Lateral orbitofrontal cortex R	28.1	47.2	-15.2	-5.21	0.0002	2.59	0.0269	-4.45	0.0007
Ventromedial prefrontal cortex L	-2.9	51.3	4.0	-7.42	>0.0001	-5.55	0.0002	-5.6t9	>0.0001
Ventromedial prefrontal cortex R	13.3	53.2	2.5	-4.42	0.0007	-2.01	0.0722	-3.32	0.0055
Dorsomedial prefrontal cortex L	-3.6	56.9	28.0	-6.14	>0.0001	-4.56	0.0010	-1.43	0.1763
Dorsomedial prefrontal cortex R	3.3	42.0	34.2	-4.11	0.0012	-4.88	0.0006	-4.29	0.0009
Anterior cingulate L	-2.4	37.0	14.7	-4.93	0.0003	-4.03	0.0024	-5.00	0.0002
Anterior cingulate R	5.0	40.2	14.7	-4.40	0.0007	-3.97	0.0026	-5.66	>0.0001
Frontal pole L	-10.1	71.1	3.0	-4.85	0.0003	-2.79	0.0191	-1.53	0.1500
Frontal pole R	14.2	72.0	3.0	-4.81	0.0003	-2.59	0.0269	-1.42	0.1791
Dorsolateral prefrontal cortex L									
Dorsolateral prefrontal cortex R	52.1	25.1	21.8	-8.3	>0.0001	-5.01	0.0005	-1.51	0.1550
Ventrolateral prefrontal cortex L									
Ventrolateral prefrontal cortex R	52.1	38.9	-4.9	-5.87	>0.0001	-5.28	0.0004	-0.96	0.3546
Paracentral lobule L	-4.6	-26.9	64.5	-5.09	0.0002	-1.98	0.0759	-2.34	0.0359
Paracentral lobule R	7.0	-35.9	-55.1	-4.11	0.0002	-1.98	0.0759	-1.52	0.0152

Table 3 Location and Talairach coordinates of local maxima for areas of reduced [^{11}C]ABP688 BP_{ND} in patients with bvFTD, along with t-values for [^{18}F]FDG SUV_R and VBM findings, using [^{11}C]ABP688 BP_{ND} maxima coordinates.

Thalamus L	-17.8	-31.3	1.8	-6.50	>0.0001	-3.11	0.0111	-8.74	>0.0001
Thalamus R	9.0	-15.9	1.8	-4.74	0.0004	-1.93	0.0824	-9.66	>0.0001
Hypothalamus L	-5.1	-5.9	-11.1	-4.18	0.0011	-2.32	0.0428	-8.49	>0.0001
Hypothalamus R	4.1	-5.9	-11.2	-4.26	0.0009	-2.67	0.0235	-8.34	>0.0001
Caudate L	-11.2	14.1	1.8	-5.43	0.0001	-4.65	0.0009	-5.89	>0.0001
Caudate R	9.0	9.1	7.3	-3.96	0.0016	-4.16	0.0019	-1.05	0.3128
Putamen L	-26.2	17.1	1.8	-4.74	0.0004	-3.64	0.0445	-3.81	0.0022
Putamen R									
Insula L	-41.1	13.0	-9.8	-4.55	0.0005	-3.59	0.0049	-4.11	0.0012
Insula R	41.7	14.1	-2.0	-5.76	>0.0001	-4.32	0.0015	-5.78	>0.0001
Temporal pole L	-38.1	22.2	-35.9	-5.59	>0.0001	-4.13	0.0020	-2.02	0.0645
Temporal pole R	31.1	22.2	-39.0	-6.79	>0.0001	-2.75	0.0205	-1.56	0.1428
Superior temporal gyrus L	-58.2	-10.9	2.0	-3.96	0.0016	-4.07	0.0023	-1.74	0.1055
Superior temporal gyrus R	49.4	-2.4	-4.0	-6.61	>0.0001	-3.21	0.0093	-4.78	0.0004
Middle temporal gyrus L	-58.2	-16.0	-11.0	-4.60	0.0005	-4.00	0.0024	-1.29	0.2195
Middle temporal gyrus R	52.7	-12.1	-24.0	-6.99	>0.0001	-4.53	0.0011	-1.53	0.1500
Inferior temporal gyrus L	-58.2	-24.5	-25.4	-7.19	>0.0001	-3.95	0.0027	-0.16	0.8753
Inferior temporal gyrus R	52.7	-45.1	-22.9	-5.16	0.0002	-3.65	0.0045	-0.09	0.9297
Hippocampal formation L	-35.1	-11.8	-19.2	-4.94	0.0003	-1.00	0.3409	-1.62	0.1292
Hippocampal formation R	34.5	-14.6	-18.5	-1.99	0.0680	-2.77	0.0198	-3.26	0.0062
Cuneus L	-11.2	-92.9	15.9	-5.22	0.0002	-3.35	0.0074	-4.81	0.0003
Cuneus R	13.8	-90.8	15.9	-5.27	0.0002	-2.05	0.0611	-4.69	0.0004
Lingual gyrus L	-12.7	-73.7	-1.7	-6.87	>0.0001	-1.27	0.2328	-5.52	>0.0001
Lingual gyrus R	22.8	-79.1	-3.9	-7.17	>0.0001	-0.41	0.6905	-0.26	0.7983

Abbreviations: standardized uptake ratio (SUV_R); voxel based morphometry (VBM); non-displaceable binding potentials (BP_{ND}). Talairach coordinates (x, y, z) of [¹¹C]ABP688 BP_{ND} local maxima (t-values, corrected for multiple comparisons, p < 0.05) were applied to [¹⁸F]FDG SUV_R and VBM t-maps in order to extra t-values. P values were determined using t-values and degrees of freedom ([¹¹C]ABP688 BP_{ND}, =14, [¹⁸F]FDG SUV_R = 13, VBM =14). R= Right hemisphere; L=Left hemisphere. -- Indicates no findings in that region.

Fig. 1 Z score maps for all bvFTD patients were created for [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and VBM. These maps were then combined to show areas with significantly reduced [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and GM common to all bvFTD patients (top left, top right, bottom left, respectively). These common Z maps were then used to extract raw [¹¹C]ABP688 BP_{ND}, [¹⁸F]FDG SUV_R, and VBM values. After conversion to Z scores, values were plotted, relative to CN subjects (bottom right). *** p < 0.001 ** p < 0.01 * p < 0.05



Fig. 2 Voxel-wise t-maps showing areas of decreased [^{11}C]ABP688 BP_{ND} in patients with bvFTD compared with CN subjects (85 152 mm³; corrected for multiple comparisons, p < 0.05). Leftward asymmetry was noted in the anterior cingulate, superior frontal gyrus, paracentral lobule, caudate, putamen, and thalamus. Rightward asymmetry was found in the posterior cingulate, lingual gyrus, cuneus, dorsolateral/ventrolateral prefrontal cortex, superior/middle temporal gyri and the temporal poles. No significant increases of [^{11}C]ABP688 binding were observed in bvFTD subjects.



Fig. 3 Voxel-wise t-maps showing areas of decreased [¹⁸F]FDG SUV_R in patients with bvFTD compared with CN subjects (116 742 mm³; corrected for multiple comparisons, p < 0.05). The dorso/ventromedial prefrontal cortex, cingulate gyrus, frontal gyri, paracentral lobule, precuneus, parahippocampus, thalamus, caudate and temporal lobes were characterized by leftward asymmetry. Hypometabolism was also noted with rightward asymmetry in the insula and orbitofrontal gyrus and in the cerebellar tonsils bilaterally.



Fig. 4 Voxel-wise t-maps showing areas of reduced VBM derived GM concentration in patients with bvFTD compared with CN subjects (88 845 mm³; corrected for multiple comparisons, p < 0.05). Atrophy was predominant in the thalami (L>R), head of caudate, insula, uncus/amygdala, and parahippocampus. GM loss was likewise noted in the putamen (L>R), precuneus (R>L), anterior cingulate (L>R), gyrus rectus, orbitofrontal gyrus and right superior/middle temporal gyri.



Fig. 5 Overlap (yellow; 22 379 mm³) between binarized [11 C]ABP688 BP_{ND} (red) and [18 F]FDG SUV_R (green) t-maps was found with leftward predominance in the in gyrus rectus, anterior cingulate dorso/ventromedial prefrontal cortex, and thalamus. Overlap was also noted with rightward asymmetry in the insula and in the temporal poles bilaterally.



Fig. 6 Overlap (magenta; 13 463mm³) between binarized [11 C]ABP688 BP_{ND} (red) and VBM (purple) t-maps was found with leftward asymmetry in the anterior cingulate, ventromedial prefrontal cortex, thalamus, and head of the caudate nucleus. Rightward asymmetry was noted in the insula and orbitofrontal cortex.



Fig. 7 Overlap (yellow; 14 179mm³) between binarized [18 F]FDG SUV_R (red) and VBM (purple) t-maps was observed with leftward asymmetry in the orbitofrontal cortex, anterior cingulate/ventromedial prefrontal cortex, thalamus and amygdala. Rightward asymmetry was found in the dorsomedial prefrontal cortex, insula, hippocampal formation, and head of the caudate.



Fig. 8 Subtraction of binarized t-maps ($[^{11}C]ABP688 BP_{ND} - [^{18}F]FDG SUV_R - VBM$) showed areas characterized only by declines in $[^{11}C]ABP688 BP_{ND} (55742 \text{ mm}^3)$. Areas characterized by rightward asymmetry included the inferior temporal lobes, temporal poles gyrus rectus, orbitofrontal cortex, head of the caudate nucleus, posterior cingulate, ventral/dorsolateral prefrontal cortex, lingual gyrus, and occipital cortex. Areas characterized by leftward asymmetry included the putamen, superior temporal lobe, and paracentral lobule.



Fig. 9 Subtraction of binarized t-maps ($[^{18}F]$ FDG SUV_R – $[^{11}C]$ ABP688 BP_{ND} – VBM) showed areas characterized only by hypometabolism (86 616 mm³), including the bilateral insula, medial prefrontal cortex, left posterior paracentral lobule, left frontal operculum, anterior temporal poles, and bilateral cerebellar cortex. Leftward asymmetry was noted for the uncus/amygdalae, parahippocampus, and posterior cingulate/precuneus. Rightward asymmetry was noted for the orbitofrontal gyrus/gyrus rectus.



Fig. 10 Subtraction of binarized t-maps (VBM – $[^{18}F]$ FDG SUV_R – $[^{11}C]$ ABP688 BP_{ND}) showed areas characterized only by reductions in GM (67 635 mm³), including the orbitofrontal gyrus, the right middle temporal gyrus, right temporal operculum, left anterior insula, head of the caudate nucleus bilaterally, left posterior inferior temporal gyrus, the thalamus as well as the posterior portion of the hippocampal formation, bilaterally. Leftward asymmetry was noted for the ventral amygdala, anterior cingulate gyrus, and putamina. Rightward asymmetry was noted for the posterior insula.



CHAPTER 3: SUMMARY, CONCLUSIONS, AND FUTURE DIRECTIONS

Reduced binding of [¹¹C]ABP688 was found among bvFTD patients in a wide range of brain regions, including the prefrontal cortex, lateral and anterior temporal cortex, basal ganglia, and thalamus. These findings overlapped partially with those for [¹⁸F]FDG and VBM, with the volume of tissue affected greatest for impaired metabolism, followed by GM concentrations and mGluR5 availability. Relative to control values, however, reductions were greatest for [¹¹C]ABP688, followed by hypometabolism and GM atrophy. In addition, several areas were characterized only by reduced [¹¹C]ABP688 BP_{ND}, including portions of the gyrus rectus, orbitofrontal gyrus, and occipital cortex.

Our results seemingly refute a previous semi-quantitative immunohistochemistry study in which mGluR5 expression levels were found to be increased in the frontal cortex (Brodmann area 8) of 3 cases with Pick's disease (Dalfo et al. 2005). Since the overexpression of mGluR5 was confined to a subset of cells present in the frontal cortex, this cell specific upregulation may have occurred in the context of declines in mGluR5 availability within the brain as a whole. Alternatively, the difference could be attributable to methodology: while the antibody used in immunohistochemistry binds mGluR5 irrespective of its functional state (i.e. even if the receptor has undergone internalization), availability of the transmembrane allosteric binding site for [¹¹C]ABP688 is dependent on mGluR5 being in its high affinity functional state of mGluR5 and not, per se, on its level of expression. A change in the proportion of mGluR5 in one or the other affinity state induced by FTLD could thus differentially affect results obtained using these two approaches.

The present study should be interpreted in light of the following considerations. 1) Our sample size was small. This was due, in part, to strict entry criteria: to be eligible for this study, patients were required to meet criteria for probable bvFTD, including both compatible structural and functional neuroimaging findings. Further, they had to be free of motor symptoms and vascular anomalies, and show no signs of amyloid pathology as assessed using [¹¹C]PIB PET. Moreover, healthy controls were required to have a score of 0 on both the NPI and FBI. 2) Our cross-sectional design could not determine whether mGluR5 reductions accompany, precede, or follow neurodegeneration. 3) Although TDP-43 and tau are the most likely underlying pathologies, the absence of genetic and post-mortem data prevents speculation as to the possibly differing associations between FTLD subtypes and patterns of mGluR5 availability. 4) As discussed above, [¹¹C]ABP688 binding can be interpreted as reductions in receptor density or lower affinity for the allosteric binding site. Future *post-mortem* studies are required to disambiguate these two possibilities.

In conclusion, the present study provides the first *in vivo* evidence that bvFTD is characterized by decreased availability of mGluR5, possibly as a result of an abnormal glutamatergic neurotransmission. Moreover, this study represents a proof of concept that PET [¹¹C]ABP688 BP_{ND} can be successfully employed to assess the role of mGluR5 in this population. In addition, this study could carry implications for novel treatment strategies in bvFTD since positive allosteric modulators of mGluR5 function are available and effective in reducing glutamate- and NMDA-induced neuronal cell death (Doria et al. 2013). Future studies are required to substantiate these findings in a larger series, to investigate whether reductions in mGLuR5 represent adaptations or vulnerability to FTLD pathology, and to explore the possibility of mGluR5 based treatment strategies for bvFTD.

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