

# Cholinergic and perfusion brain networks in Parkinson disease dementia

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## ABSTRACT

**Objective:** To investigate muscarinic M1/M4 cholinergic networks in Parkinson disease dementia (PDD) and their association with changes in Mini-Mental State Examination (MMSE) after 12 weeks of treatment with donepezil.

**Methods:** Forty-nine participants (25 PDD and 24 elderly controls) underwent  $^{123}\text{I}$ -QNB and  $^{99\text{m}}\text{Tc}$ -exametazime SPECT scanning. We implemented voxel principal components (PC) analysis, producing a series of PC images of patterns of interrelated voxels across individuals. Linear regression analyses derived specific M1/M4 and perfusion spatial covariance patterns (SCPs).

**Results:** We found an M1/M4 SCP of relative decreased binding in basal forebrain, temporal, striatum, insula, and anterior cingulate ( $F_{1,47} = 31.9, p < 0.001$ ) in cholinesterase inhibitor-naïve patients with PDD, implicating limbic-paralimbic and salience cholinergic networks. The corresponding regional cerebral blood flow SCP showed relative decreased uptake in temporoparietal and prefrontal areas ( $F_{1,47} = 177.5, p < 0.001$ ) and nodes of the frontoparietal and default mode networks (DMN). The M1/M4 pattern that correlated with an improvement in MMSE ( $r = 0.58, p = 0.005$ ) revealed relatively preserved/increased pre/medial/orbitofrontal, parietal, and posterior cingulate areas coinciding with the DMN and frontoparietal networks.

**Conclusion:** Dysfunctional limbic-paralimbic and salience cholinergic networks were associated with PDD. Established cholinergic maintenance of the DMN and frontoparietal networks may be prerequisite for cognitive remediation following cholinergic treatment in this condition.

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## GLOSSARY

**AIC** = Akaike information criterion; **CAMCOG** = Cambridge Cognitive Examination; **CAMCOG<sub>exec</sub>** = Cambridge Cognitive Examination executive function subscale; **CAMCOG<sub>memory</sub>** = Cambridge Cognitive Examination memory subscale; **ChEI** = cholinesterase inhibitor; **DLB** = dementia with Lewy bodies; **DMN** = default mode network; **MMSE** = Mini-Mental State Examination; **NBM** = nuclear basalis of Meynert; **NPI** = Neuropsychiatric Inventory; **PC** = principal component; **PD** = Parkinson disease; **PDD** = Parkinson disease dementia; **rCBF** = regional cerebral blood flow; **SCP** = spatial covariance pattern; **SN** = salience network; **SSF** = subject scaling factor.

In Parkinson disease (PD), development of dementia (PDD) occurs in up to 80% of people 15–20 years after PD diagnosis,<sup>1</sup> with 50% developing cognitive impairment within 6 years.<sup>2</sup> In PDD, cholinergic dysfunction is strongly implicated in cognitive deficits, fluctuating cognition, and visual hallucinations.<sup>3</sup> Reductions in choline acetyltransferase are marked in PDD compared to Alzheimer disease and PD,<sup>4</sup> while clinically, cholinesterase inhibitors (ChEIs) can ameliorate cognition and visual hallucinations. However, response is variable with some efficacy.<sup>5</sup>

Since the brain is a networked entity, pathologic change in one area may influence other topographically distant regions. Indeed, distributed network dysfunction is now considered a key contributor to symptoms that manifest in neurodegenerative dementias.<sup>6</sup> In PDD, theoretical models of dysfunctional neural networks have been proposed. In particular, several cholinergic networks arising from the nuclear basalis of Meynert (NBM) projecting to

Supplemental data  
at Neurology.org

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specific brain regions are thought to affect major cognitive domains, e.g., attention (NBM → neocortex), visuoperceptual (NBM → parieto-occipital, parahippocampal-fusiform), and memory (NBM → medial temporal).<sup>7</sup> One way to examine functional brain connectivity is by spatial covariance analysis.

We applied spatial covariance to (R, R) <sup>123</sup>I-QNB SPECT data,<sup>8</sup> acquired in ChEI-naive patients with PDD, to investigate disease-associated M1/M4 cholinergic networks. Also, as cholinergic network dysfunction is implicated in cognitive impairment and amelioration of cholinergic function is an important aspect of treatment, we derived an M1/M4 covariance pattern that correlated with a change in Mini-Mental State Examination (MMSE) score, after 12 weeks of treatment with the ChEI (donepezil), to probe the clinical significance of these networks.

**METHODS Standard protocol approvals, registrations, and patient consents.** Study approval was from the UK Department of Health's Administration of Radioactive Substances Advisory Committee and Newcastle, North Tyneside, and Northumberland research ethics committees. All participants or nearest relatives gave written informed consent for the study including treatment.

**Participants.** The study comprised 49 individuals (25 PDD and 24 similarly aged controls). Patients were recruited from outpatient movement disorder clinics in Newcastle-upon-Tyne and Gateshead, while healthy controls were from patient spouses and friends in this and other studies. Participants had physical, neurologic, and neuropsychiatric assessments, including mental state, history, physical examination, and, for patients, blood screen with B<sub>12</sub> and folate levels. The study battery administered included the MMSE,<sup>9</sup> Neuropsychiatric Inventory (NPI),<sup>10</sup> and Cambridge Cognitive Examination (CAMCOG)<sup>11</sup> with memory and executive function subscales (CAMCOG<sub>memory</sub>, CAMCOG<sub>exec</sub>).

Diagnosis was carried out consensually between 2 clinicians using the diagnostic criteria for PDD.<sup>12</sup> Patients with PDD were on levodopa and carbidopa or benserazide combination therapy and were naive to ChEI treatment at the time of QNB imaging. Participants on any of the following medications were excluded from the study: antipsychotics, cholinergics, anticholinergics, and antidepressant medications. Clinicopathologic diagnosis was established for 11 cases (2 controls, 9 PDD).

**Radiochemistry.** Using the technique of Lee et al.,<sup>13</sup> (R, R) <sup>123</sup>I-QNB radiosynthesis was conducted, the specifics of which are described elsewhere.<sup>14</sup>

**Acquisition.** Participants were scanned with a triple-head gamma camera (Picker 3000XP; Philips, Best, the Netherlands) 5 hours postinjection of (R, R) <sup>123</sup>I-QNB using a previously reported imaging protocol.<sup>8</sup> Within 4 weeks of the (R, R) <sup>123</sup>I-QNB scan, individuals underwent <sup>99m</sup>Tc-exametazime regional cerebral blood flow (rCBF) SPECT imaging in accordance with a past scanning procedure.<sup>8</sup>

**Spatial preprocessing.** All SPECT scans were registered to match, where applicable, a <sup>123</sup>I-QNB or <sup>99m</sup>Tc-exametazime SPECT template in standard stereotactic Montreal Neurological Institute space using linear image registration software (<http://fsl.fmrib.ox.ac.uk/fsl/fslwiki/flirt/>). Generation of specific template images has been described.<sup>14,15</sup> The spatially transformed images were then smoothed with a 10-mm full width at half maximum 3D Gaussian filter.

**Multivariate spatial covariance analysis.** Principal component (PC) analysis was applied on a voxel basis to all processed <sup>123</sup>I-QNB SPECT images using covariance analysis software ([http://www.nitrc.org/projects/gcva\\_pca/](http://www.nitrc.org/projects/gcva_pca/)),<sup>16</sup> producing a series of PC images. For each PC image, voxels had either positive or negative weights that represent the sign and strength of covariance between voxels. In this study, voxels with positive and negative weights were viewed as concurrently preserved/increased and decreased M1/M4 binding, respectively. The extent to which an individual expressed the PC image was by way of a subject scaling factor (SSF) for that PC, calculated by superimposing the PC image onto an individual's processed QNB scan by computation of a dot product, which involves image multiplication on a voxel basis followed by summation of the products generating a score. Higher SSF scores for an individual for that PC image represents greater increased binding in voxels with positive weights and greater concurrent decreased binding in voxels with negative weights. To identify the QNB spatial covariance pattern (SCP) that distinguished PDD from controls, each individual SSF was entered into a linear regression model as explanatory variables with group as the dependent parameter. Akaike information criteria (AICs) determined how many PCs should be included to reach optimal bias-variance tradeoff.<sup>17</sup> The set of PCs yielding the lowest AIC value was used to derive the SCP<sub>QNB</sub>. The degree to which each participant expressed the SCP<sub>QNB</sub> was by the SSF<sub>QNB</sub>.

The same approach was applied to the <sup>99m</sup>Tc-exametazime SPECT images. Therefore, positive and negative weights were interpreted as concurrent increased and decreased rCBF, respectively. The analysis produced the SCP<sub>rCBF</sub> that best separated PDD from controls, while each participant expressed the SCP<sub>rCBF</sub> by his or her SSF<sub>rCBF</sub>.

Following their <sup>123</sup>I-QNB scan, the majority of patients (n = 18) were then treated with the ChEI donepezil titrated up to the standard daily clinical dose of 10 mg. After a period of 12 weeks, patients underwent repeated MMSE assessments. We derived a ChEI-naive M1/M4 SCP that correlated with  $\Delta$ MMSE<sub>rel,b</sub>, which described the percentage change in MMSE relative to baseline. This involved conducting a separate analysis, generating a series of PCs expressed by each participant by the SSFs, which in turn were introduced into a regression model as predictor variables with  $\Delta$ MMSE<sub>rel,b</sub> as the response parameter. The resulting linear combination with the smallest AIC value generated the SCP <sub>$\Delta$ MMSE</sub> ( $R^2 = 0.34$ ,  $p = 0.005$ ), where each individual expressed the pattern by the SSF <sub>$\Delta$ MMSE</sub>.

Stability and reliability of the SCPs were assessed by bootstrap resampling (1,000 iterations) to identify areas that contributed to the patterns with high confidence. This transforms the voxel weights of each SCP into Z maps, computed as the ratio of voxel weight and bootstrap SD. The Z-statistic follows roughly a standard normal distribution where a one-tailed  $p \leq 0.05$  infers a threshold of  $|Z| \geq 1.64$ .<sup>18</sup>

**Statistical analyses.** Continuous variables were tested for normality using visual inspection of histograms and Shapiro-Wilk test. Demographic, clinical, and imaging measures were assessed, where applicable, using parametric (analysis of variance) and nonparametric  $\chi^2$  tests. Correlations were performed using Pearson  $r$  coefficients.

**Table 1** Participant characteristics

	Controls	PDD	Statistic, p value
No.	24	25	
M:F	15:9	17:8	$\chi^2 = 0.2, 0.7$
Age, y	74.1 ± 5.1	72.0 ± 5.0	$F_{1,47} = 2.0, 0.2$
MMSE	28.3 ± 1.5	18.8 ± 4.9	$F_{1,47} = 82.3, <0.001$
CAMCOG	95.0 ± 3.9	63.0 ± 14.2	$F_{1,47} = 113.8, <0.001$
CAMCOG <sub>memory</sub>	22.1 ± 1.9	16.1 ± 4.7	$F_{1,47} = 32.6, <0.001$
CAMCOG <sub>exec</sub>	20.8 ± 4.2	8.5 ± 3.1	$F_{1,47} = 135.2, <0.001$
NPI	1.3 ± 2.3	19.7 ± 17.8	$U = 73.0, 0.002$
NPI <sub>hall</sub>	NA	3.5 ± 2.5	
UPDRS III	0.9 ± 1.5	38.5 ± 11.6	$U = 600.0, <0.001$

Abbreviations: CAMCOG = Cambridge Cognitive Examination; CAMCOG<sub>exec</sub> = Cambridge Cognitive Examination executive function subscale; CAMCOG<sub>memory</sub> = Cambridge Cognitive Examination memory subscale; MMSE = Mini-Mental State Examination; NA = not applicable; NPI<sub>hall</sub> = Neuropsychiatric Inventory hallucinations subscale; PDD = Parkinson disease dementia; UPDRS = Unified Parkinson's Disease Rating Scale. Values denote mean ± 1 SD.

Statistical tests were interpreted as significant if  $p \leq 0.05$ . Data analysis used the Statistical Package for Social Sciences (Chicago, IL) software (SPSS version 22.0; <http://www-01.ibm.com/software/analytics/spss/products/statistics/>).

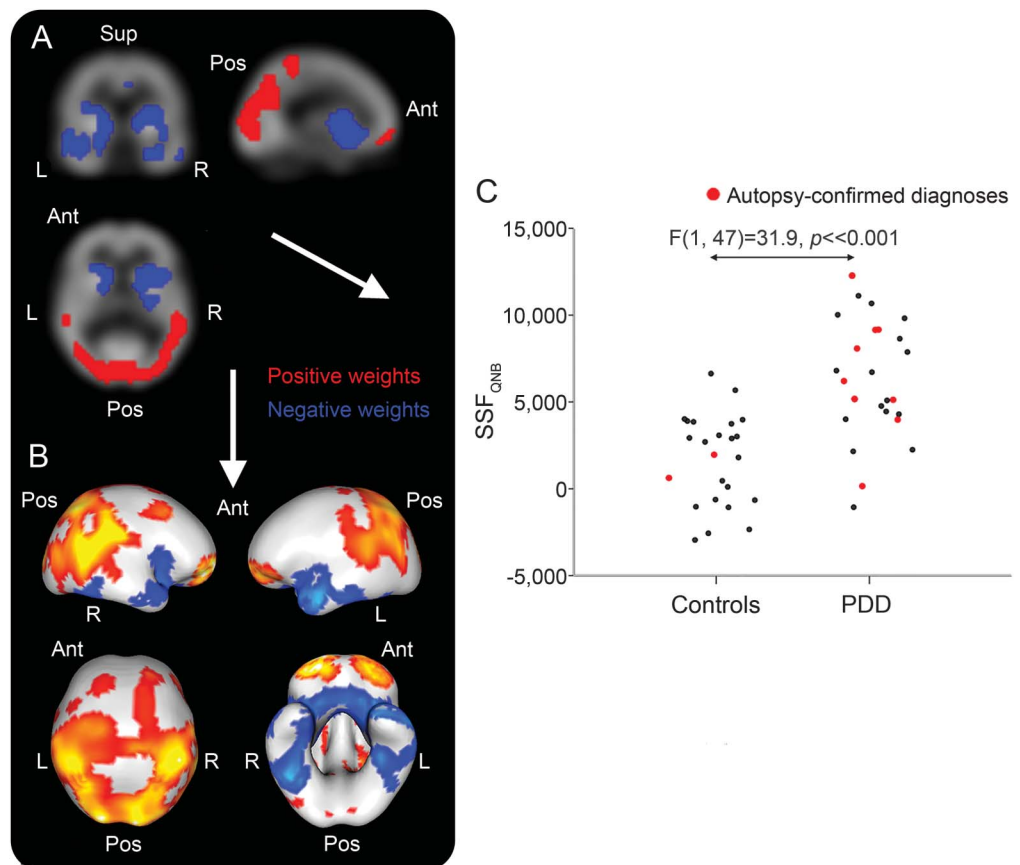
**RESULTS Demographics and clinical characteristics.**

Table 1 shows demographic and clinical characteristics of the study sample. Groups were similar in age and sex, while as expected, all other variables differed ( $p < 0.001$ ).

**Spatial covariance analysis.** The SCP<sub>QNB</sub> that distinguished PDD from controls is shown in figure 1, A and B. SSF<sub>QNB</sub> scores were higher in PDD than controls (mean ± SD; controls = 1.5 ± 2.5, PDD = 6.2 ± 3.3,  $F_{1,47} = 31.9, p < 0.001$ ; figure 1C). The pattern was mainly characterized by concomitant decreases in M1/M4 binding (blue regions) in basal forebrain, temporal, striatal, insula, and anterior cingulate together with concomitant preserved or increases (red regions) in frontal and parieto-occipital areas. Table e-1 on the *Neurology*<sup>®</sup> Web site at Neurology.org presents detailed description of specific regions contributing to the M1/M4 disease-related pattern.

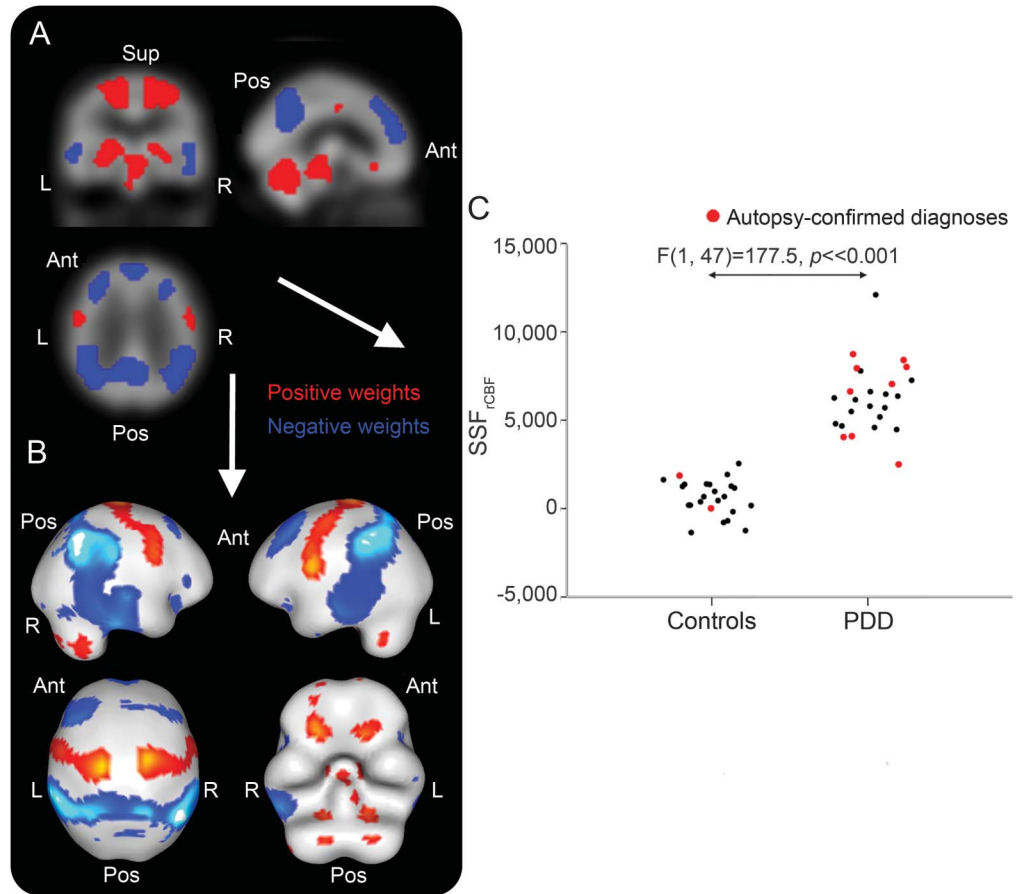
The associated SCP<sub>rCBF</sub> that differentiated PDD from controls is illustrated in figure 2, A and B, where SSF<sub>rCBF</sub> scores differed between groups (controls = 0.4 ± 0.9, PDD = 6.2 ± 1.9,  $F_{1,47} = 177.5, p < 0.001$ ; figure 2C). The pattern mainly comprised relative decreased rCBF (blue) in temporoparietal and

**Figure 1** Muscarinic M1/M4 spatial covariance pattern in Parkinson disease dementia (PDD)



Disease-related M1/M4 spatial covariance pattern in PDD projected onto orthogonal (A) and rendered (B) displays of the QNB template. Distribution of subject scaling factor (SSF<sub>QNB</sub>) scores across groups (C). Ant = anterior; Pos = posterior; Sup = superior.

**Figure 2** Regional cerebral blood flow (rCBF) spatial covariance pattern in Parkinson disease dementia (PDD)



Disease-related rCBF spatial covariance pattern in PDD projected onto orthogonal (A) and rendered (B) displays of the rCBF template. Distribution of subject scaling factor ( $SSF_{rCBF}$ ) scores across groups (C). Ant = anterior; Pos = posterior; Sup = superior.

prefrontal areas with relative increases (red) in cerebellum, brainstem, striathalamic, and motor regions. Detailed description of specific regions participating in the rCBF disease-related pattern is provided (table e-2).

Relationships between SCP expressions and age, MMSE, CAMCOG, CAMCOG<sub>memory</sub>, CAMCOG<sub>exec</sub>, total NPI, NPI hallucinations subscale, and Unified Parkinson's Disease Rating Scale III were investigated in PDD. No correlations were found between  $SSF_{QNB}$  and these measures ( $|r| \leq 0.23$ ,  $p \geq 0.14$ ). For the rCBF pattern expression, total NPI correlated with  $SSF_{rCBF}$  ( $\rho = 0.62$ ,  $p = 0.006$ ), which was not observed for the other variables ( $|r| \leq 0.28$ ,  $p \geq 0.09$ ). An exploratory examination of NPI subscores did not yield any specific relationships with SCP expressions.

Summary data for the donepezil-treated group are shown (table 2). During the observation period, differences in MMSE were identified between baseline and 12-week scores ( $p < 0.001$ ). The resultant  $SCP_{\Delta MMSE}$  is presented in figure 3, A and B, while figure 3C depicts  $SSF_{\Delta MMSE}$  plotted as a function of

$\Delta MMSE_{rel,b}$ . The pattern consists of concurrent decreases in M1/M4 binding (blue) in fusiform, anterior cingulate, lingual gyrus, and precentral areas with concurrent preserved or increases (red) in pre/medial/orbitofrontal, parietal, and posterior cingulate regions.

**Table 2** Summary data of patients with PDD treated with donepezil

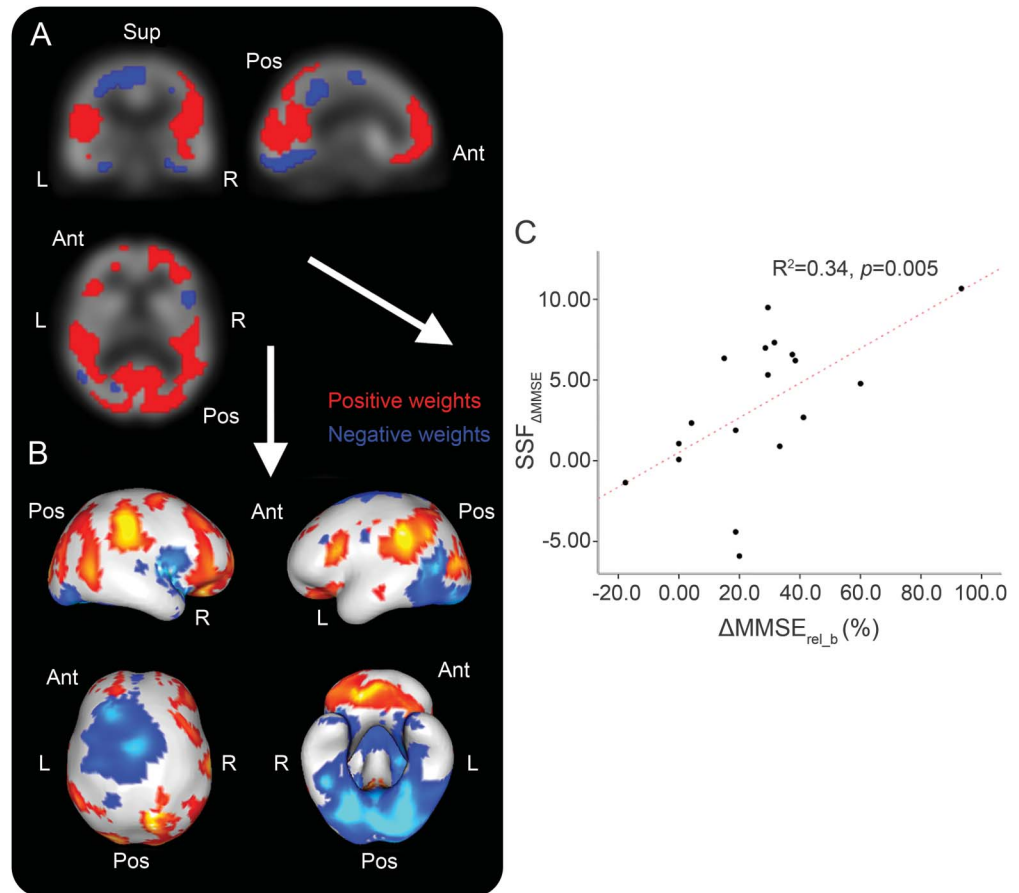
	PDD <sub>donepezil</sub>	Statistic, p value
No.	18	
M:F	11:7	
Age	71.6 ± 4.6	
MMSE <sub>b</sub>	18.1 ± 4.3	
MMSE <sub>12w</sub>	22.4 ± 4.7	
$ MMSE_{12w} - MMSE_b $	4.3 ± 3.8	$t_{17} = 4.8, <0.001$
$\Delta MMSE_{rel,b}$ , %	26.8 ± 24.6	

Abbreviations: 12w = at 12 weeks; b = at baseline; MMSE = Mini-Mental State Examination; PDD = Parkinson disease dementia.

Values denote mean ± 1 SD.

$\Delta MMSE_{rel,b}$  (%) =  $\{(MMSE_{12w} - MMSE_b)/MMSE_b\} \times 100\%$ .

**Figure 3** M1/M4 spatial covariance pattern (SCP)  $\Delta$  Mini-Mental State Examination (MMSE) in Parkinson disease dementia (PDD) (n = 18)



(A-C) M1/M4 SCP in PDD (n = 18) that correlated with  $\Delta$ MMSE<sub>rel\_b</sub>.

Details of specific regions participating in the pattern are supplied (table e-3).

**DISCUSSION** We undertook a multivariate network perspective of (R, R) <sup>123</sup>I-QNB SPECT, a M1/M4 receptor ligand in ChEI-naïve patients with PDD. We derived a disease-related M1/M4 pattern of spatial covariance that appears largely distinct from rCBF, which implies the presence of several dysfunctional cholinergic networks in PDD. We also identified a clear M1/M4 covariance pattern that was associated with an improvement in MMSE; this network had distinctive spatial elements suggesting certain cortical regions and their associated cholinergic innervation may have a more preminent role in cognitive amelioration by cholinergic treatments. Relevant to the present study, this spatial covariance technique has extensively and successfully been utilized in perfusion SPECT and glucose metabolism PET data for the investigation of disease progression and symptomatology in PD.<sup>19–22</sup>

We derived a voxel cholinergic SCP from <sup>123</sup>I-QNB images that differentiated PDD from controls. The disease-related pattern comprised decreased

and preserved/increased M1/M4 uptake in a number of concomitant brain regions or networks. The covariant negative-weighted pattern mainly converged on limbic/paralimbic regions. Notably, this cholinergic receptor network mapped onto previously described resting-state networks, including anterior insula and the anterior cingulate, key nodes of the salience network (SN), which is important for initiation of cognitive control and switching between networks to aid access to working memory and attention resources.<sup>23,24</sup> Networks involving the insula have also been shown to play a role in episodic memory,<sup>25</sup> while hippocampus, parahippocampus, and amygdala are known to be involved in memory storage and retrieval.<sup>26</sup> As such, this pattern would align with a cognitive network deficit implicating the basal forebrain and these structures, i.e., a cholinergic limbic-paralimbic/SN dysfunction. The disease-related pattern also encompassed regions implicated in dorsal (occipital → parietal) and ventral (occipital → temporal → limbic) visual streams,<sup>27</sup> providing indirect evidence for the role of distinct cholinergic networks in visual function in PDD, which would be in keeping with known visuoperceptual deficits and predisposition

to visual hallucinations, symptoms that both show good response to cholinesterase inhibitors.<sup>28,29</sup>

The associated rCBF pattern largely comprised relative decreases in temporoparietal and prefrontal areas along with relative increases in cerebellum, brainstem, striathalamic, and motor regions that implicate a number of functional networks in PDD. Regions that were concomitantly reduced appear to involve hubs of the frontoparietal attention (inferior parietal, dorsolateral prefrontal cortex)<sup>30</sup> and default mode networks (DMNs) (medial prefrontal, posterior cingulate, ventral precuneus, inferior parietal),<sup>31,32</sup> which is of interest since, respectively, attention deficits are one of the most disabling cognitive symptoms in PDD,<sup>33</sup> while network theories have strongly implicated the DMN in contributing to cognitive decline.<sup>34</sup> Our previous studies revealed modulation of the frontoparietal network in PDD that was similar to patients with dementia with Lewy bodies (DLB),<sup>35</sup> while also demonstrating, albeit in DLB, its relationship with severity and frequency of cognitive fluctuations.<sup>36</sup> Other investigations have reported decreases in DMN connectivity in PDD<sup>37</sup> and its association with cognitive dysfunction in PD.<sup>38</sup> These and our rCBF findings appear to provide further evidence that implicate the DMN and frontoparietal networks in the pathogenesis of symptoms in PDD, in particular cognitive. Moreover, perhaps not unexpectedly, the rCBF pattern seemed to represent an extended topography of the PD-related motor and cognitive patterns, which have been previously reported from <sup>18</sup>F-FDG PET studies using similar network approaches,<sup>21,39</sup> thus indirectly validating the analytic methodology used in the present study.

We failed to detect any correlations between the M1/M4 pattern expressions and neuropsychological and neuropsychiatric measures in PDD. For rCBF pattern expression, only total NPI score was found to correlate. Thus, patients with more global severe neuropsychiatric symptoms, a marker of greater disease severity, were more likely to express the perfusion SCP characteristic of PDD. The lack of correlations may be explained by either the notion that each spatial covariance pattern is likely to characterize a number of overlapping and convergent brain networks and thus fails to project on specific cognitive and clinical parameters or that patterns derived from combined (control–dementia) cohorts are less sensitive. Isolating key networks from these patterns could increase sensitivity, but this is methodologically challenging.

We found a clear M1/M4 covariance pattern that correlated with a change in MMSE that could indicate a positive treatment response. This pattern showed relative decreases in fusiform, striatum, anterior cingulate, lingual gyrus, and precentral areas with relative preservation or increase in pre/medial/orbitofrontal,

parietal, and posterior cingulate regions. From a network perspective, there was covariant preservation/upregulation in regions overlapping key nodes of the DMN and frontoparietal networks that could imply that a relative cholinergic maintenance of these networks is prerequisite for ChEI treatment response in PDD, and more generally may point toward the potential relevance of these networks and their cholinergic innervation and its associated cognitive symptoms. Notably, a recent study showed that cholinergic and serotonergic antagonists can impair DMN-like network in mice similarly, suggesting that both neurotransmitter systems are involved in maintaining the integrity of the DMN-like networks.<sup>40</sup> Hence, this pattern appears to provide some evidence that supports the cholinergic DMN maintenance hypothesis, and its potential significance as a predictor of positive treatment response in PDD and perhaps in other neurodegenerative disorders.

Modest sample sizes and uncertainty regarding which receptor subtype is affected (that is, M1 vs M4) are limitations of the study. Another limitation was the use of MMSE rather than Montreal Cognitive Assessment to assess cognitive function in these patients, reflecting the fact that our data were collected before the widespread use of the latter scale. Replication of this study with neuropsychological assessments that align more with the cognitive deficit profile of PDD may provide a more nuanced cholinergic response network pattern. Strengths were scanning and clinically assessing patients with PDD free from cholinergic medications with perfusion and muscarinic SPECT images available for all participants. We also had, in a sizeable minority, autopsy confirmation of diagnoses.

Our findings imply several dysfunctional cholinergic and perfusion networks in PDD. The relevance of these networks may be important in terms of their contribution to cognitive and, in particular, attentional deficits of this condition. The use of ChEIs could improve such deficits, but there is marked heterogeneity in response to these agents and it is not possible to reliably predict on clinical grounds who might respond to these drugs. Although tentative, we observed a SCP that suggests that those with cholinergic maintenance of DMN and frontoparietal networks could experience cognitive improvement with ChEI treatment. These findings provide further neurobiological insights into therapies targeted at improving cholinergic neurotransmission and treatment outcomes in PDD.

#### AUTHOR CONTRIBUTIONS

Dr. Colloby: co-designed the study, conducted all image and data analyses, and wrote the manuscript. Professor McKeith: reviewed the manuscript and secured project funding. Professor Burn: reviewed the manuscript and secured project funding. Professor Wyper: reviewed the manuscript and secured project funding. Professor O'Brien: reviewed

the manuscript and secured project funding. Dr. Taylor: co-designed the study and cowrote the manuscript.

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## DISCLOSURE

S. Colloby reports no disclosures relevant to the manuscript. I. McKeith has been a consultant for GE Healthcare, Bayer Health care, and Nutricia. D. Burn has received an honorarium from Acadia. D. Wyper reports no disclosures relevant to the manuscript. J. O'Brien has been a consultant for GE Healthcare, Lilly, Bayer Healthcare, TauRx, and Nutricia and has received honoraria for talks from GE Healthcare, Lilly, and Novartis. J. Taylor has been a consultant of Lundbeck and received honoraria for talks from GE Healthcare and Flynn Pharmaceuticals. Go to [Neurology.org](http://Neurology.org) for full disclosures.

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