

Distinct brain networks underlie cognitive dysfunction in Parkinson and Alzheimer diseases

Paul J. Mattis, PhD
Martin Niethammer,
MD, PhD
Wataru Sako, MD, PhD
Chris C. Tang, MD, PhD
Amir Nazem, MD, PhD
Marc L. Gordon, MD
Vicky Brandt, BA
Vijay Dhawan, PhD
David Eidelberg, MD

Correspondence to
Dr. Eidelberg:
deidelbe@northwell.edu

ABSTRACT

Objective: To determine whether cognitive impairment in Parkinson disease (PD) and Alzheimer disease (AD) derives from the same network pathology.

Methods: We analyzed ^{18}F -fluorodeoxyglucose PET scans from 40 patients with AD and 40 age-matched healthy controls from the Alzheimer's Disease Neuroimaging Initiative and scanned an additional 10 patients with AD and 10 healthy controls at The Feinstein Institute for Medical Research to derive an AD-related metabolic pattern (ADRP) analogous to our previously established PD cognition-related pattern (PDCP) and PD motor-related pattern (PDRP). We computed individual subject expression values for ADRP and PDCP in 89 patients with PD and correlated summary scores for cognitive functioning with network expression. We also evaluated changes in ADRP and PDCP expression in a separate group of 15 patients with PD scanned serially over a 4-year period.

Results: Analysis revealed a significant AD-related metabolic topography characterized by covarying metabolic reductions in the hippocampus, parahippocampal gyrus, and parietal and temporal association regions. Expression of ADRP, but not PDCP, was elevated in both AD groups and correlated with worse cognitive summary scores. Patients with PD showed slight ADRP expression, due to topographic overlap with the network underlying PD motor-related pattern degeneration, but only their PDCP expression values increased as cognitive function and executive performance declined. Longitudinal data in PD disclosed an analogous dissociation of network expression.

Conclusions: Cognitive dysfunction in PD is associated with a specific brain network that is largely spatially and functionally distinct from that seen in relation to AD. *Neurology*® 2016;87:1925-1933

GLOSSARY

AD = Alzheimer disease; **ADNI** = Alzheimer's Disease Neuroimaging Initiative; **ADRP** = Alzheimer disease-related pattern; **ANOVA** = analysis of variance; **FDG** = ^{18}F -fluorodeoxyglucose; **FIMR** = Feinstein Institute for Medical Research; **MCI** = mild cognitive impairment; **MCI(-)** = cognitively intact; **MCI(m)** = mild cognitive impairment involving multiple cognitive domains; **MCI(s)** = mild cognitive impairment involving a single cognitive domain; **MMSE** = Mini-Mental State Examination; **PD** = Parkinson disease; **PDCP** = Parkinson disease cognition-related pattern; **PDD** = Parkinson disease dementia; **PDRP** = Parkinson disease motor-related pattern; **RMANOVA** = repeated-measures analysis of variance.

Cognitive impairment is commonly observed in patients with Parkinson disease (PD), even early in the clinical course,^{1,2} but its cause remains unclear. The postmortem observation of amyloid- β plaques and tau neurofibrillary tangles, pathologic hallmarks of Alzheimer disease (AD), in individuals with PD and dementia has led to the hypothesis that the cognitive changes in PD are caused by comorbid AD.³⁻⁵ Many patients with PD have substantial cognitive loss without forming plaques and tangles, however, and the severity of neuropsychological deficits in patients with PD with coexisting cortical Lewy body and AD-like pathology correlates only with

Supplemental data at Neurology.org

From the Center for Neurosciences (P.J.M., M.N., W.S., C.C.T., A.N., V.B., V.D., D.E.) and Litwin-Zucker Research Center for the Study of Alzheimer's Disease (M.L.G.), The Feinstein Institute for Medical Research, Manhasset; and Department of Neurology (P.J.M., M.N., M.L.G., D.E.), Northwell Health, Manhasset, NY.

Data used in preparation of this article were obtained from the Alzheimer's Disease Neuroimaging Initiative (ADNI) database (adni.loni.usc.edu). As such, the investigators within the ADNI contributed to the design and implementation of ADNI and/or provided data but did not participate in analysis or writing of this report. A complete listing of ADNI investigators can be found at: http://adni.loni.usc.edu/wp-content/uploads/how_to_apply/ADNI_Acknowledgement_List.pdf. (also available as supplemental material).

Go to Neurology.org for full disclosures. Funding information and disclosures deemed relevant by the authors, if any, are provided at the end of the article.

the former.⁶ Whether AD contributes to the cognitive deficits in PD thus remains an unsettled question.

The fact that neuronal dysfunction in neurodegenerative diseases is propagated along discrete networks makes such questions amenable to neuroimaging approaches. We and others have used ¹⁸F-fluorodeoxyglucose (FDG)-PET and fMRI to identify disease-specific networks in PD, AD, Huntington disease, and other conditions.⁷⁻¹³ We have shown that akinesia-rigidity in PD correlates with the expression of a PD motor-related pattern (PDRP)^{10,11} such that more severe symptoms are reflected in higher PDRP scores; tremor is mediated by a distinct cerebello-thalamo-cortical network¹⁴; and the expression of a distinct PD cognition-related pattern (PDCP), characterized by diminished metabolism in the medial frontal and parietal regions, rises as cognitive function deteriorates.¹⁵⁻¹⁸ Similarly, in AD, neuropsychological test performance correlates with the expression of an AD-related pattern (ADRP) characterized by covarying reductions in hippocampal, superior temporal, and parieto-occipital resting-state activity.⁷ To determine whether patients with PD, with or without cognitive deficits, express ADRP, we compared ADRP and PDCP expression in patients with AD and those with PD. We also analyzed changes in ADRP and PDCP over time in a separate longitudinal cohort of patients with PD.

METHODS **Participants.** *Patients with AD.* To ensure the most rigorous comparison, we used the same spatial covariance analysis approach we previously used to characterize PDRP^{10,19} to identify and validate a reliable ADRP topography for prospective use. We analyzed metabolic images from 40 patients with AD (23 men and 17 women, mean \pm SD age 75.9 \pm 5.6 years, Mini-Mental State Examination [MMSE] score 22.6 \pm 3.3) and 40 age-matched healthy controls (23 men and 17 women, age 76.0 \pm 4.7 years) who had been scanned with FDG-PET as part of the Alzheimer's Disease Neuroimaging Initiative (ADNI; see table e-1 at [Neurology.org](http://www.neurology.org) for demographic and neuropsychological data).

Using a random number generator (<http://www.r-project.org/>), we assigned the patients with AD and controls to 1 of 2 groups. The first group comprised 20 patients with AD (AD1; 13 men and 7 women, age 76.0 \pm 6.2 years, MMSE score 23.6 \pm 2.4) and 20 normal controls (NL1; 10 men and 10 women, age 76.6 \pm 5.2 years). We used scans from these AD1 and NL1 participants to identify a significant ADRP topography and scans from a second group of 20 patients with AD (AD2; 10 men and 10 women, age 75.8 \pm 5.2 years, MMSE score 21.6 \pm 3.8) and 20 normal controls (NL2; 13 men and 7 women, age 75.5 \pm 4.2

years) for testing. A third group constituting an additional validation set, consisting of 10 additional patients with AD (AD3; 6 men and 4 women, age 74.5 \pm 5.3 years, MMSE score 23.9 \pm 4.2) and 10 age-matched healthy volunteers (NL3; 4 men and 6 women, age 73.4 \pm 4.8 years), were scanned with FDG-PET on the GE Advance tomograph (General Electric, Milwaukee, WI) at The Feinstein Institute for Medical Research (FIMR; Manhasset, NY) as described elsewhere.^{19,20} One of the investigators (M.L.G.) made the diagnosis of AD in the AD3 cohort on the basis of published criteria.²¹

Patients with PD. We quantified ADRP expression values and subject scores for the previously characterized PDCP network topography in a cross-sectional sample of 89 patients (61 men and 28 women, age 61.6 \pm 9.0 years, MMSE score 28.0 \pm 2.3) who were recruited at the Movement Disorders Center of Northwell Health (Great Neck, NY) and diagnosed according to the UK Brain Bank criteria for idiopathic PD.²² These patients had pure parkinsonism without a history of known causative factors (such as encephalitis or neuroleptic treatment and without supranuclear gaze abnormalities or ataxia); patients with Lewy body dementia were excluded. Table e-2 summarizes the demographic and neuropsychological data for these participants.

We classified patients with PD as having dementia or not having dementia according to the Dementia Rating Scale²³; 11 had scores <127, classifying them as having PD dementia (PDD); the remaining 78 were classified as not having dementia. Using the neuropsychological battery (see e-Methods), we assigned patients with PD without dementia to 3 previously defined cognitive categories: cognitively intact [PD-MCI(-); n = 18], mild cognitive impairment (MCI) involving a single cognitive domain [PD-MCI(s); n = 30], and MCI involving multiple cognitive domains [PD-MCI(m); n = 30]. The neuropsychological testing criteria for MCI in patients with PD are described elsewhere.^{2,18}

In addition, we compared the progression of ADRP and PDCP in a previously published longitudinal imaging cohort of 15 patients with PD (11 men and 4 women, age 58.0 \pm 10.2 years) who were scanned with FDG-PET at baseline (within 2 years of diagnosis) and again 24 and 48 months later.²⁴ All participants in the cross-sectional and longitudinal PD cohorts underwent FDG-PET at FIMR according to the same protocol used for the AD3 group.

Standard protocol approvals, registrations, and patient consents. For the patients with AD and healthy volunteers participating in ADNI protocols, written informed consent was obtained after approval was granted by the Institutional Review Board of the collaborating institutions. For participants scanned at FIMR, we obtained ethics permission from the Institutional Review Board of Northwell Health and written consent from each participant after a detailed explanation of the procedures.

Image analysis. FDG-PET scans were preprocessed with SPM5 (Wellcome Trust Centre for Neuroimaging, Institute of Neurology, London, UK) running on Matlab 6.0 (Mathworks Inc, Natick, MA). Scans were spatially normalized to a standard PET brain template and smoothed with a 3-dimensional gaussian kernel (full width at half maximum = 10 mm).

Spatial covariance analysis^{10,19,20} identified a distinct ADRP topography in the combined AD1 (n = 20) and NL1 (n = 20) ADNI derivation sample. (The computational procedures used to identify and validate this pattern are detailed in e-Methods.) We computed ADRP expression values (subject scores) for each member of the ADNI (AD2/NL2, n = 40) and FIMR (AD3/NL3, n = 20; PD, n = 89) testing samples using an automated algorithm available at <http://www.feinsteinneuroscience.org>. The resulting measures were z scored with respect to corresponding

measures from the NL1 normal reference sample used in pattern identification. We used analogous procedures to quantify PDCP expression in the same scans in each of the testing cases. To accord with previously published results, PDCP expression values were z scored with respect to values from the group of normal controls (NL4; 8 men and 7 women, age 56.7 ± 12.3 years) used for reference in prior studies.^{18,25,26}

Network correlates of neuropsychological test performance.

We evaluated the relationship between network expression and cognitive performance in the combined ADNI (AD1/AD2) patient cohort and the FIMR PD sample. For the ADNI participants, we used the summary score data for memory and executive functioning provided in the database.²⁷ Because the ADNI database does not provide a summary score for language function, we constructed a language index score based on the measures that were available for each participant, standardized (z scored) with respect to age-corrected normative values, and averaged individual scores to form a language summary index. We used a similar approach to compute summary scores for memory, executive function, and language in members of the cross-sectional PD sample.

Data analysis. We assessed differences in ADRP expression between patients with AD and controls separately for the ADNI (AD2/NL2) and FIMR (AD3/NL3) testing sets and for the PD and age-matched normal reference (NL4) groups scanned at FIMR. Subject scores computed in each of the subgroups within this PD sample, PD-MCI(-), PD-MCI(s), PD-MCI (m), and PDD, were separately compared to corresponding control values. Differences in ADRP expression between patients and controls in each testing sample were assessed with Student t tests, and differences in ADRP expression across the PD subgroups were assessed with 1-way analysis of variance (ANOVA), followed by post-hoc Dunnett tests for multiple comparisons. We used identical statistical procedures to assess group differences in PDCP expression in each testing sample. Potential network \times subgroup interaction effects in the PD data were evaluated with 2-way repeated-measures ANOVA (RMANOVA), with network as the within-subject variable and subgroup as the between-subject variable.

We used a similar approach to evaluate the neuropsychological summary scores for the 2 disease groups. We constructed separate 1-way ANOVA models for each cognitive domain to identify differences across the PD subgroups; p values were adjusted for multiple comparisons with the Dunnett correction. Relationships between individual subject ADRP and PDCP expression values and the neuropsychological summary scores were evaluated separately in the AD and PD groups by calculating the Pearson product-moment correlations.

To compare changes in ADRP and PDCP expression over time in the longitudinal PD cohort, we analyzed the subject scores using 2-way RMANOVA with network and time as within-subject variables. If a significant network \times time interaction effect was evident in the data, we used 1-way RMANOVA to assess changes over time in the expression of each network. All statistical analyses were performed with SAS 9.3 for Windows (SAS Institute, Inc, Cary, NC) and considered significant at $p < 0.05$ (2-tailed).

RESULTS ADRP, but not PDCP, expression was consistently elevated in patients with AD. Spatial covariance analysis of metabolic scans from the combined AD1/NL1 derivation sample revealed a distinct ADRP topography characterized by metabolic reductions in the hippocampus, parahippocampal gyrus,

and temporal and parietal association regions, along with an increase in activity in the sensorimotor cortex and cerebellum (figure 1A, top and table). The voxel weights on the pattern, reflecting local contributions to overall network activity, were found to be stable by bootstrap estimation (absolute value of the inverse coefficient of variation range = $[-2.822$ to $2.814]$, $p < 0.005$, 1,000 iterations). The ADRP was topographically unrelated to the PDCP ($r^2 = 0.013$, voxel weight correlation, figure 1B, top). It is worth noting that the ADRP exhibited some correlation with the PDRP, although the relationship did not reach significance ($r^2 = 0.123$, voxel-weight correlation); there are shared regional metabolic reductions involving the inferior parietal lobule bilaterally, which are more salient for ADRP than PDRP¹¹ (see e-Methods). ADRP subject scores (figure 1A, bottom), which measure pattern expression in individual cases, were consistently elevated in patients with AD relative to their healthy counterparts ($p < 0.0003$, permutation test).

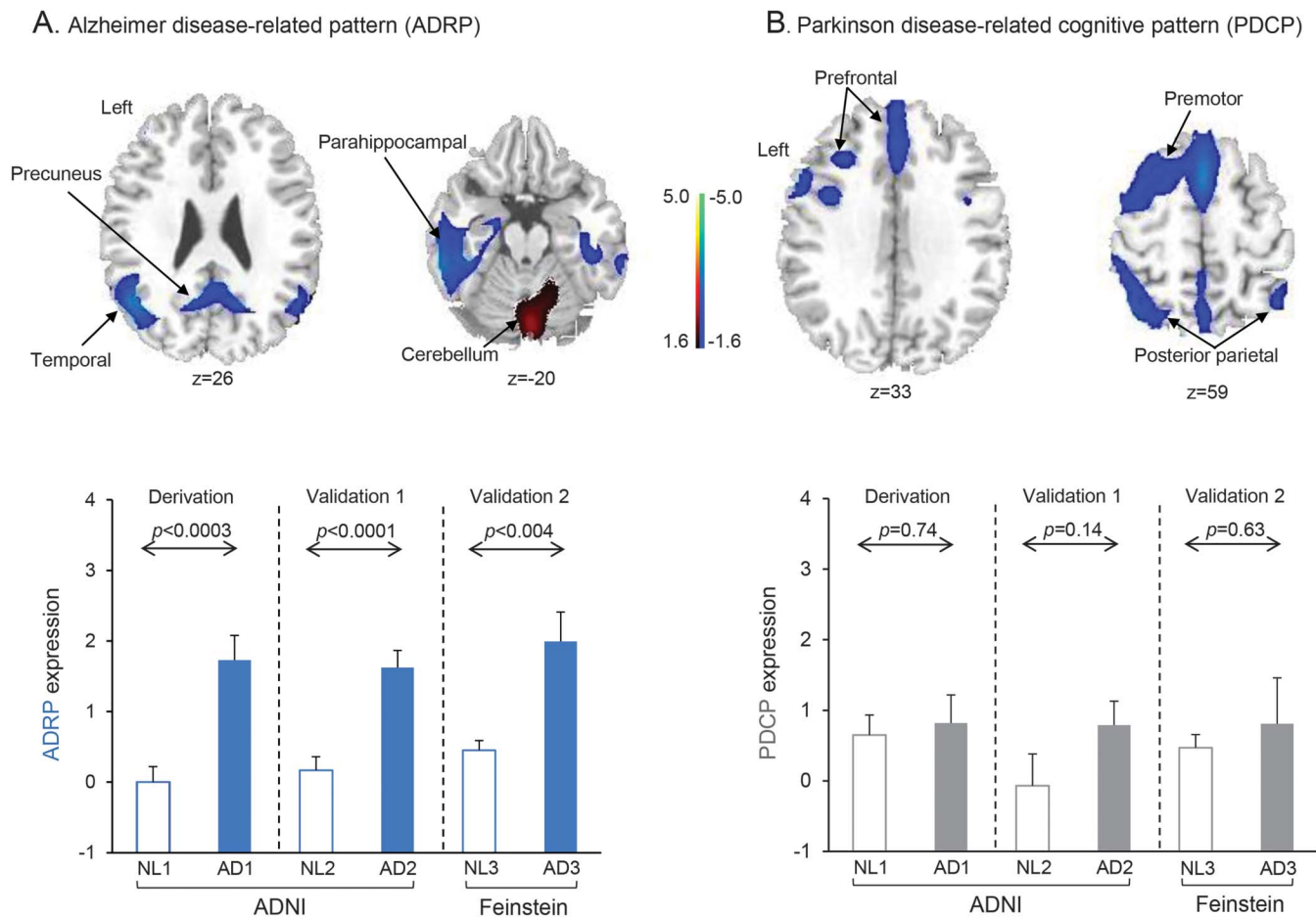
For validation, we used an automated algorithm to quantify ADRP expression in the ADNI (AD2 and NL2) and FIMR (AD3 and NL3) testing sets on a prospective single-scan basis. As in the derivation sample, ADRP subject scores (figure 1A, bottom) were markedly elevated in patients relative to controls in both AD testing sets (AD2, $p < 0.0001$; AD3, $p < 0.004$, Student t tests). In contrast to the ADRP, expression values for the PDCP (figure 1B, bottom) did not differ from normal in the derivation set ($p = 0.74$) or in either of the testing sets (AD2, $p = 0.14$; AD3, $p = 0.63$). There was no discernible correlation between ADRP and PDCP expression values in the AD patient sample ($p > 0.34$) or between ADRP or PDCP expression and subject age ($p > 0.18$).

In patients with AD, neuropsychological scores correlated with ADRP but not PDCP expression levels.

We used the summary score data for memory, language, and executive functioning provided by the ADNI database,²⁷ which included data from a variety of measures, depending on which tests were administered (e-Methods). Summary scores for each of the 3 cognitive domains correlated with ADRP expression in the combined AD cohorts: the worse the summary score, the more elevated the expression of ADRP (memory: $r = -0.32$, $p < 0.05$; language: $r = -0.45$, $p < 0.004$; executive function: $r = -0.34$, $p < 0.03$, Pearson correlations, figure 2, left). There was no such correlation between neuropsychological summary scores and PDCP expression ($p > 0.12$, figure 2, right).

Cognitive impairment in patients with PD correlates with expression of PDCP but not ADRP. As expected,¹⁸ the PD cohort showed PDCP expression ($p < 0.0005$) that increased as cognitive impairment

Figure 1 Alzheimer disease (AD)-related pattern (ADRP) expression, but not Parkinson disease (PD) cognition-related pattern (PDCP) expression, is increased in AD



(A) Top, ADRP, identified by spatial covariance analysis of ^{18}F -fluorodeoxyglucose-PET scans of 20 patients with AD (AD1) and 20 normal controls (NL1) from the Alzheimer's Disease Neuroimaging Initiative (ADNI), was characterized by reduced activity in the hippocampus, parahippocampal gyrus, and parietal and temporal association regions, with relative increases in the cerebellum, sensorimotor cortex, and supplementary motor area (represented by the second principal component, which accounted for 12.8% of the subject \times voxel variation in the data). Voxel weights on the pattern, reflecting local contributions to overall network activity, were found to be stable by bootstrap estimation (absolute value of the inverse coefficient of variation range = $[-2.822$ to $2.814]$, $p < 0.005$, 1,000 iterations). Bottom, In the derivation sample, which comprised 20 patients with AD (AD1) and 20 normal controls (NL1) selected randomly from the ADNI database (see supplemental data), ADRP expression values (subject scores) accurately discriminated patients from healthy controls ($p < 0.0003$, permutation test). Prospectively computed ADRP subject scores achieved comparable group separation ($p < 0.004$, Student t test) in 2 separate testing samples. The first testing set comprised the 20 patients with AD (AD2) and 20 normal ADNI participants (NL2) not used for pattern derivation. The second testing set comprised 10 patients with AD (AD3) and 10 normal controls (NL3) scanned separately at the Feinstein Institute. (B) Top, PDCP was previously identified by spatial covariance analysis of metabolic images from 15 patients with PD with varying levels of cognitive dysfunction.¹⁵ PDCP is characterized by reduced activity in the presupplementary motor area, premotor, and prefrontal regions and in parietal associative cortex, with relative increases in the cerebellar vermis and dentate nuclei. Bottom, PDCP expression did not differ ($p > 0.14$) between patients with AD and controls in the derivation sample or in the ADNI and Feinstein validation samples. (The covariance maps shown on the top in A and B were overlaid on T1-weighted magnetic resonance template images. For each pattern, the display was thresholded at $|Z| = 1.64$ [$p < 0.05$]). Voxels with positive region weights [relative increases] are color-coded red; those with negative region weights [relative decreases] are color-coded blue. Error bars shown on the bottom of A and B represent 1 SEM.)

worsened ($F_{3,85} = 3.42$, $p < 0.03$, 1-way ANOVA, figure 3A). Patients with PD also expressed ADRP ($p < 0.05$, Student t tests), but there was no association with the degree of cognitive impairment ($F_{3,85} = 1.61$, $p = 0.19$, 1-way ANOVA). Indeed, ADRP levels were similarly elevated ($p < 0.02$, Student t test) in all patients with PD (figure 3B), including the cognitively intact PD-MCI(-) group, which showed modest elevations in both PDCP and ADRP expression relative to healthy controls (ADRP,

$p < 0.01$; PDCP, $p < 0.04$, Student t tests); the level of pattern expression in these participants (subject scores ≈ 1.0) was similar for the 2 topographies ($p = 0.76$ for comparison of ADRP and PDCP subject scores, paired Student t test). In the cognitively impaired PD-MCI(m) and PDD subgroups, PDCP expression was greater than ADRP [PD-MCI(m), $p = 0.001$; PDD, $p < 0.04$, paired Student t tests]. The entire PD data set showed a clear network \times subgroup interaction ($F_{3,85} = 3.78$,

Table Brain regions with a significant contribution to Alzheimer disease-related pattern

	Coordinates ^a				
	BA	Z _{max}	x	y	z
Regions with negative weights					
Inferior parietal lobule, left	40	-2.84	-49	-52	37
Precuneus					
Right	7,31	-2.34	13	-59	30
Left		-2.27	-13	-60	27
Superior temporal gyrus, supramarginal gyrus, left	39,40	-2.70	-52	-53	23
Middle temporal gyrus					
Right	37	-2.06	50	-54	-1
Left	21	-3.14	-61	-40	-14
Inferior temporal gyrus, middle temporal gyrus, middle occipital gyrus, left	37	-2.39	-51	-53	-1
Fusiform gyrus, parahippocampal gyrus					
Right	20,36,37	-2.25	39	-35	-13
Left	37	-2.17	-34	-40	-10
Hippocampus, left		-2.00	-30	-17	-14
Caudate, left		-2.08	-9	14	-8
Regions with positive weights					
SMA, right	6	3.42	7	-4	68
Paracentral lobule, postcentral gyrus					
Right	5,6	2.60	3	-28	60
Left		3.02	-5	-44	61
Paracentral lobule, postcentral gyrus					
Right	1,2,3,4	2.15	31	-34	61
Left	2,3,4	1.90	-26	-33	61
Cerebellum, vermis V/VI/VIIA/VIIB, hemisphere lobule V/VI/crus I/crus II^b					
Right		2.08	8	-82	-23
Left		2.01	-7	-84	-23

Abbreviations: BA = Brodmann area; SMA = supplementary motor area.

^aMontreal Neurological Institute (MNI) standard space.

^bAccording to the atlas of Schmahmann et al.³¹

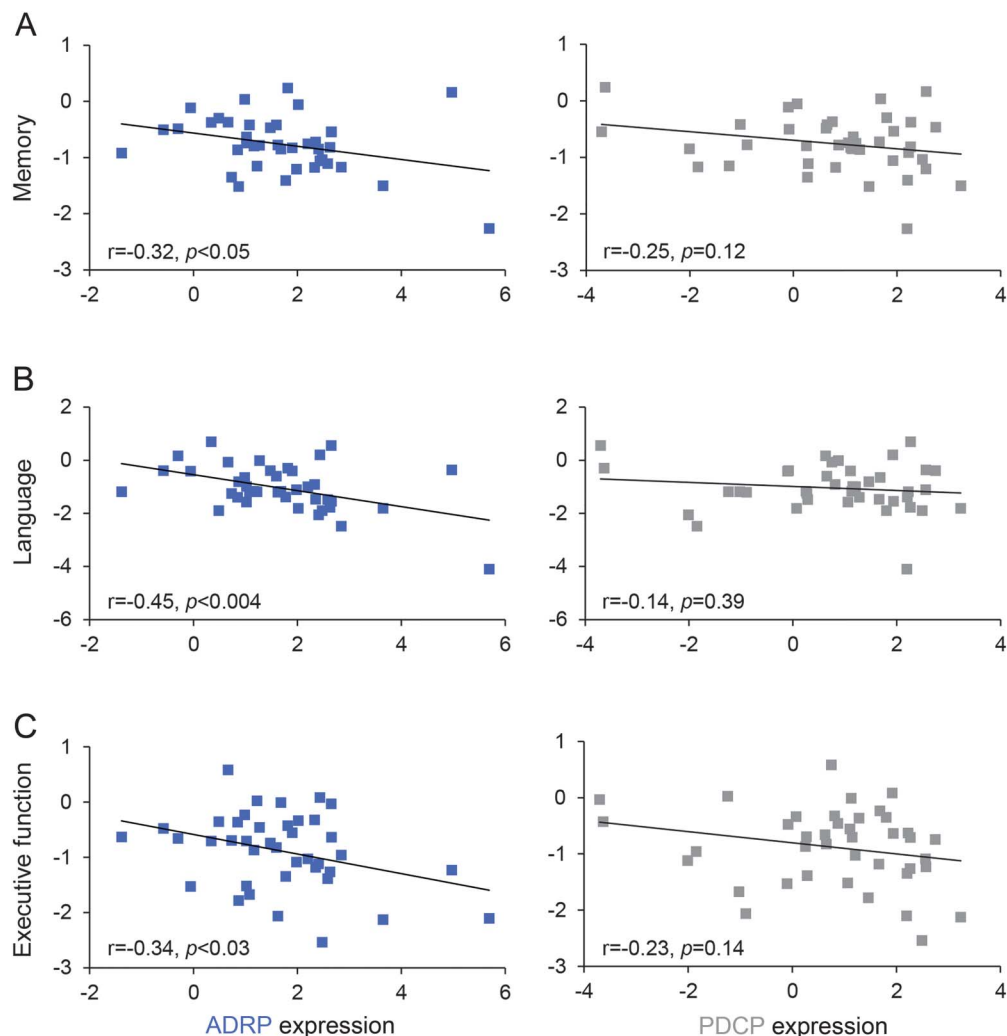
$p = 0.01$, 2×4 RMANOVA) such that stepwise increases in PDCP expression corresponded with more severe cognitive dysfunction but without corresponding differences in ADRP. PDCP and ADRP expression values did not correlate in the PD subgroups with dementia ($p = 0.67$) or without dementia ($p = 0.66$).

If ADRP expression is present in patients with PD but does not correlate with cognitive deficits, what does it represent? Given the aforementioned overlap in the parietal association cortex between ADRP and PDRP topographies, we computed expression values for a modified ADRP topography, called ADRP', which was defined by masking the overlapping regions in each brain before forward application of the native

ADRP pattern (figure e-1A). ADRP' expression (figure 3C) did not differ from normal in any of the PD subgroups ($p > 0.05$, Student t tests) but remained elevated in each of the AD patient samples ($p < 0.01$, figure e-1B). ADRP expression in patients with PD thus largely reflects changes in PDRP due to PD rather than an AD-associated disease process.

PDCP, but not ADRP, expression levels correlate with neuropsychological measures in patients with PD. We chose the cognitive measures for our PD cohorts to match the ADNI composite measures as closely as possible, and we used a similar approach to compute the analogous summary scores (e-Methods). Consistent with MCI classification, neuropsychological summary

Figure 2 Alzheimer disease (AD)-related pattern (ADRP) expression, but not Parkinson disease cognition-related pattern (PDCP) expression, correlates with neuropsychological summary scores in AD



Network correlations with neuropsychological summary scores in the patients with AD from the AD1 and AD2 cohorts (see text). Significant negative correlations ($r < -0.32$, $p < 0.05$) were seen between ADRP expression and summary scores for memory (A, left), language (B, left), and executive function (C, left). By contrast, no cognitive correlations ($p > 0.12$) were observed with PDCP expression values (A-C, right) in the same AD cohort.

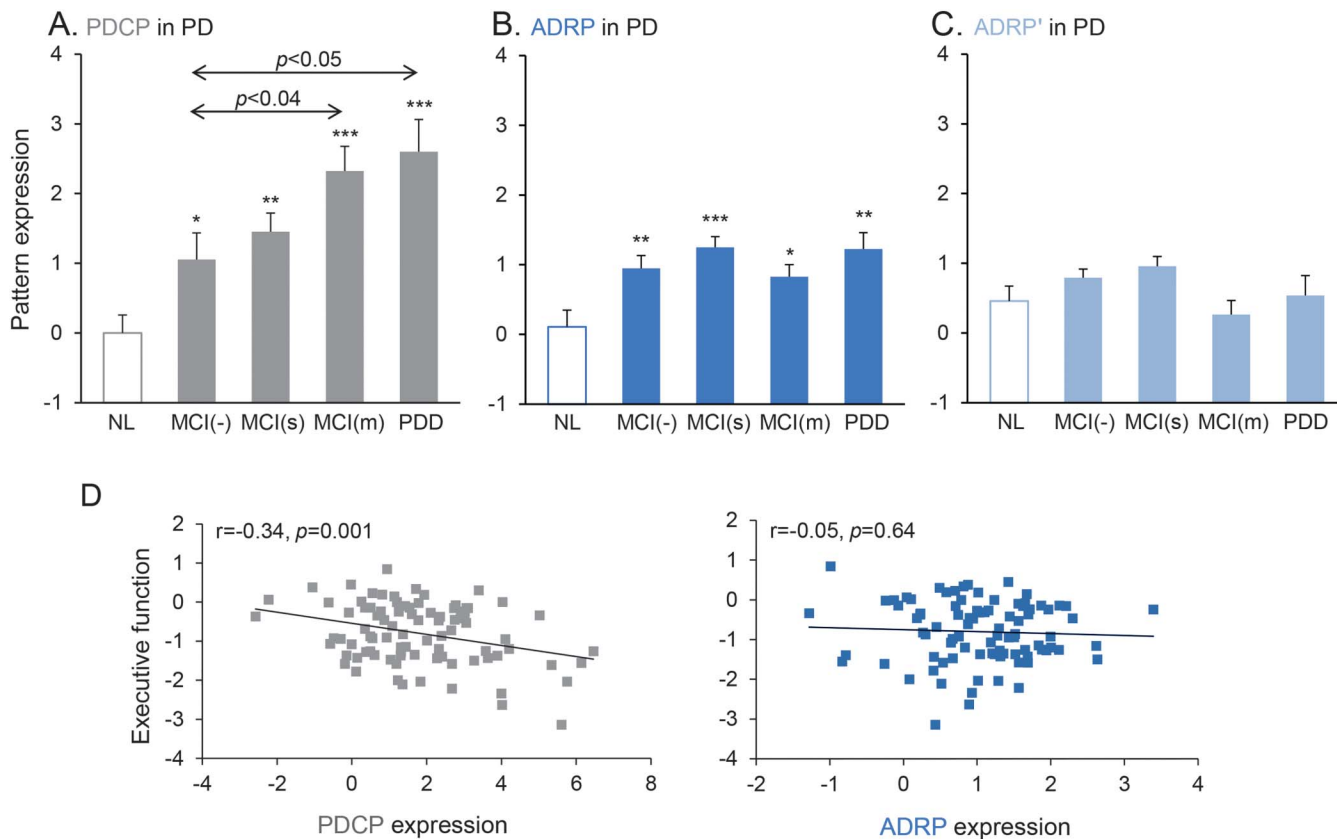
scores (table e-2) declined with worsening cognitive performance in all 3 domains across the PD patient subgroups. Summary scores for memory differed across the PD subgroups ($F_{3,81} = 13.7$, $p < 0.0001$, 1-way ANOVA), with worse performance in the PD-MCI and PDD subgroups [$p < 0.001$ for each subgroup relative to PD-MCI(-); post hoc Dunnett tests]. Executive function ($F_{3,83} = 5.4$, $p < 0.002$) and language ($F_{3,82} = 5.3$, $p < 0.003$) showed similar trends. PD-MCI(m) and patients with PDD exhibited clear deficits in executive performance compared with PD-MCI(-) ($p < 0.01$); language deficits were more pronounced in the PDD group [$p < 0.005$ relative to PD-MCI(-), post hoc Dunnett test].

Higher PDCP expression values reflected worse summary scores for executive performance ($r = -0.34$, $p = 0.001$, figure 3D, left). There

was a weak correlation between PDCP expression and memory impairment ($r = -0.22$, $p < 0.04$) but not with language deficits ($r = -0.09$, $p = 0.42$, data not shown). ADRP values, in contrast, did not correlate with cognitive performance in any of the 3 domains ($p > 0.56$, figure 3D, right). Thus, in patients with AD, individual differences in memory, language, and executive function were reflected in ADRP but not PDCP expression. In patients with PD, in contrast, executive function had a clear relationship with PDCP expression but not with ADRP expression.

PDCP, but not ADRP, expression increases over time in patients with PD. In the longitudinal cohort of early patients with PD (figure 4), a significant network \times time interaction effect ($F_{2,23} = 4.34$, $p < 0.03$, 2-way

Figure 3 Parkinson disease (PD) cognition-related pattern (PDCP) expression, but not Alzheimer disease (AD)-related pattern (ADRP) expression, is associated with cognitive impairment in PD



Mean PDCP (A) and ADRP (B) expression values are displayed for patients with PD with no evidence of mild cognitive impairment [MCI(-); $n = 18$], single-domain MCI [MCI(s); $n = 30$], multiple-domain MCI [MCI(m); $n = 30$], and PD with dementia (PDD; $n = 11$); values for normal controls (NL; $n = 15$) are provided for reference (see text). Arrows indicate post hoc Dunnett test relative to MCI(-) group. (C) ADRP' subject scores, reflecting the expression of this pattern after the exclusion of ADRP/PDRP overlap regions (see text), did not differ significantly from normal in any of the PD subgroups, regardless of cognitive status. (D) PDCP expression in this PD cohort (left) correlated significantly ($r = -0.34, p = 0.001$) with executive function, whereas ADRP did not correlate with summary scores for executive functioning (right) in these patients with PD ($*p < 0.05, **p < 0.01, ***p < 0.001$, Student t tests compared to normal controls).

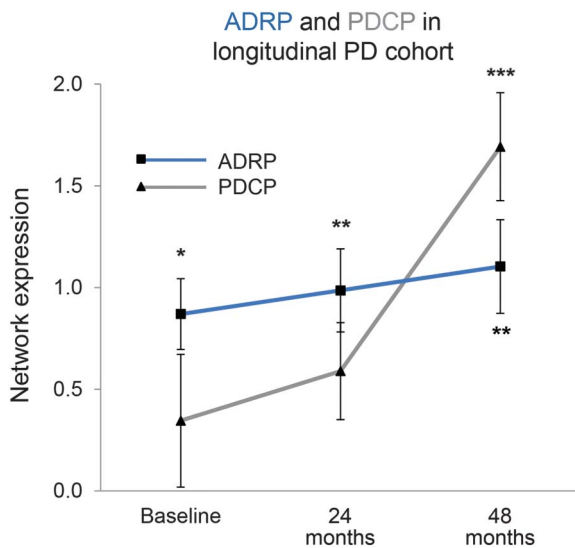
RMANOVA) revealed different progression rates for the 2 networks over the course of disease. PDCP expression in this early PD cohort increased with time ($F_{2,23} = 7.16, p < 0.004$, 1-way RMANOVA), and subject scores for this pattern measured at 4 years were higher than at baseline ($p < 0.003$, post hoc Dunnett test). Expression values for concurrently measured ADRP, however, did not change over the 4 years of follow-up (ADRP: $F_{2,23} = 2.21, p = 0.13$, 1-way RMANOVA), nor did it correlate with age ($p > 0.31$). Moreover, although ADRP expression in these patients was elevated relative to control values ($p < 0.02$, Student t test) at all 3 time points, the expression of ADRP' did not reach significance at any time point ($p > 0.05$, Student t test).

DISCUSSION In this study, we used the distinct network topographies of AD and PD to investigate whether the cognitive deficits that commonly occur

in patients with PD are more likely attributable to PD itself or to comorbidity with AD. Our analyses indicate that, despite overlapping regions of pathology, cognitive dysfunction in PD is largely distinct from that which occurs in AD.

This is not to say, however, that some patients with PD may not also have AD. Although most of the 11 patients with PD with the most severe disease (PDD) had ADRP expression levels tightly clustered around the mean, 2 participants showed relatively high ADRP expression that remained elevated even after masking regions of ADRP/PDRP overlap in the parietal association cortex (ADRP: 2.63 and 2.00, subject score = 0.98 ± 0.76 for the remaining 9 PDD cases; ADRP': 1.40 and 1.85, subject score = 0.29 ± 0.86 for the remaining 9 cases). Even more strikingly, the patient with PDD with the highest ADRP' expression was also exceptional in expressing almost normal PDCP levels; this patient most likely has true comorbid AD. It is worth noting that our

Figure 4 Parkinson disease (PD) cognition-related pattern (PDCP) expression, but not Alzheimer disease-related pattern (ADRP) expression, increases in PD over time



Mean ADRP (squares) and PDCP (triangles) expression in a longitudinal early-stage PD cohort (see text). PDCP expression values (gray line) increased over time in this group ($p < 0.004$, repeated-measures analysis of variance [RMANOVA]), while concurrent ADRP changes (blue line) did not reach significance ($p = 0.13$, RMANOVA). ADRP expression in these patients was elevated relative to control values ($p < 0.02$, Student *t* test) at all time points; PDCP expression, in contrast, reached abnormal levels ($p < 0.001$) only at the final time point. (Error bars represent 1 SE at each time point. * $p < 0.05$, ** $p < 0.01$, *** $p < 0.001$, Student *t* tests compared to normal controls.)

metabolic network findings complement structural imaging data showing AD-like volumetric changes in the cerebral cortex of patients with PD.²⁸ The pattern of volume loss seen with structural MRI in patients with AD²⁹ exhibits some topographic homology with ADRP but not with PDCP; it is possible that Lewy- and Alzheimer-type pathologies affect cognition differently in individuals with PD.⁶

From a clinical standpoint, it is important for physicians and patients to recognize that cognitive dysfunction in PD is likely not caused by comorbid AD. Furthermore, the disease specificity of ADRP and PDCP provides a distinct contrast with resting-state networks found in healthy controls such as the default mode network that show progressive attrition in both disease populations.³⁰ To fully articulate the relationship of the observed imaging changes with the underlying disease process will require multimodal imaging in conjunction with molecular genetic analyses and thorough postmortem assessments. In the meanwhile, having objective imaging biomarkers to differentiate between underlying pathologies will help with prognosis, selection of participants for clinical trials, and choice of treatments when they become available.

AUTHOR CONTRIBUTIONS

Dr. Mattis: study design, data acquisition, statistical analysis, data analysis and interpretation, and drafting and revising of the manuscript for

intellectual content. Dr. Niethammer: data interpretation and revising of the manuscript for intellectual content. Dr. Sako: data analysis and drafting of the manuscript for intellectual content. Dr. Tang: statistical analysis, data analysis and interpretation, and drafting and revising of the manuscript for intellectual content. Dr. Nazem: data analysis and drafting of the manuscript for intellectual content. Dr. Gordon: data acquisition and revising of the manuscript for intellectual content. V. Brandt: revising of the manuscript for intellectual content. Dr. Dhawan: data acquisition, data interpretation, study supervision, and revising of the manuscript for intellectual content. Dr. Eidelberg: study design, statistical analysis, data interpretation, study supervision, obtaining funding, and drafting and revising of the manuscript for intellectual content.

ACKNOWLEDGMENT

The authors wish to thank Ms. Ivana De Lucia for her valuable technical assistance and Ms. Yoon Young Choi and Ms. Toni Fitzpatrick for manuscript preparation/copyediting.

STUDY FUNDING

This work was supported in part by the National Institute of Neurologic Disorders and Stroke Morris K. Udall Center of Excellence for Parkinson's Disease Research at The Feinstein Institute for Medical Research (P50 NS071675 to D.E.) and the Thomas Hartman Foundation for Parkinson's Research. The content is solely the responsibility of the authors and does not necessarily represent the official views of the National Institute of Neurologic Disorders and Stroke or the NIH. The sponsor did not play a role in study design, collection, analysis or interpretation of data, writing the report, or the decision to submit the paper for publication.

DISCLOSURE

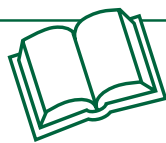
P. Mattis, M. Niethammer, W. Sako, C. Tang, A. Nazem, V. Brandt, and V. Dhawan report no disclosures relevant to the manuscript. M. Gordon received research support, without direct compensation, from Merck, Lundbeck, Genentech, Accera, Baxter, Forest, and Eli Lilly; he has also participated in an advisory board for Lundbeck. D. Eidelberg serves on the scientific advisory board and has received honoraria from The Michael J. Fox Foundation for Parkinson's Research; is listed as coinventor of patents for markers for use in screening patients for nervous system dysfunction and a method and apparatus for using same, without financial gain; and has received research support from the NIH (National Institute of Neurologic Disorders and Stroke, National Institute on Deafness and Other Communication Disorders, National Institute of Allergy and Infectious Diseases) and the Dana Foundation. Go to Neurology.org for full disclosures.

Received March 18, 2016. Accepted in final form July 18, 2016.

REFERENCES

1. Aarsland D, Kurz MW. The epidemiology of dementia associated with Parkinson disease. *J Neurol Sci* 2010; 289:18–22.
2. Janvin CC, Larsen JP, Aarsland D, Hugdahl K. Subtypes of mild cognitive impairment in Parkinson's disease: progression to dementia. *Mov Disord* 2006;21:1343–1349.
3. Irwin DJ, White MT, Toledo JB, et al. Neuropathologic substrates of Parkinson disease dementia. *Ann Neurol* 2012;72:587–598.
4. Irwin DJ, Lee VM, Trojanowski JQ. Parkinson's disease dementia: convergence of alpha-synuclein, tau and amyloid-beta pathologies. *Nat Rev Neurosci* 2013;14:626–636.
5. Jellinger KA, Seppi K, Wenning GK, Poewe W. Impact of coexistent Alzheimer pathology on the natural history of Parkinson's disease. *J Neural Transm* 2002;109:329–339.
6. Compta Y, Parkkinen L, O'Sullivan SS, et al. Lewy- and Alzheimer-type pathologies in Parkinson's disease dementia: which is more important? *Brain* 2011;134:1493–1505.

7. Habeck C, Foster N, Perneczky R, et al. Multivariate and univariate neuroimaging biomarkers of Alzheimer's disease. *Neuroimage* 2008;40:1503–1515.
8. Scarmeas N, Habeck CG, Zarahn E, et al. Covariance PET patterns in early Alzheimer's disease and subjects with cognitive impairment but no dementia: utility in group discrimination and correlations with functional performance. *Neuroimage* 2004;23:35–45.
9. Teune LK, Strijkert F, Renken RJ, et al. The Alzheimer's disease-related glucose metabolic brain pattern. *Curr Alzheimer Res* 2014;11:725–732.
10. Eidelberg D. Metabolic brain networks in neurodegenerative disorders: a functional imaging approach. *Trends Neurosci* 2009;32:548–557.
11. Niethammer M, Eidelberg D. Metabolic brain networks in translational neurology: concepts and applications. *Ann Neurol* 2012;72:635–647.
12. Feigin A, Leenders KL, Moeller JR, et al. Metabolic network abnormalities in early Huntington's disease: an [(18)F]FDG PET study. *J Nucl Med* 2001;42:1591–1595.
13. Tang CC, Feigin A, Ma Y, et al. Metabolic network as a progression biomarker of premanifest Huntington's disease. *J Clin Invest* 2013;123:4076–4088.
14. Mure H, Hirano S, Tang CC, et al. Parkinson's disease tremor-related metabolic network: characterization, progression, and treatment effects. *Neuroimage* 2011;54:1244–1253.
15. Huang C, Mattis P, Tang C, Perrine K, Carbon M, Eidelberg D. Metabolic brain networks associated with cognitive function in Parkinson's disease. *Neuroimage* 2007;34:714–723.
16. Mattis PJ, Tang CC, Ma Y, Dhawan V, Eidelberg D. Network correlates of the cognitive response to levodopa in Parkinson disease. *Neurology* 2011;77:858–865.
17. Meles SK, Tang CC, Teune LK, et al. Abnormal metabolic pattern associated with cognitive impairment in Parkinson's disease: a validation study. *J Cereb Blood Flow Metab* 2015;35:1478–1484.
18. Huang C, Mattis P, Perrine K, Brown N, Dhawan V, Eidelberg D. Metabolic abnormalities associated with mild cognitive impairment in Parkinson disease. *Neurology* 2008;70:1470–1477.
19. Spetsieris P, Ma Y, Peng S, et al. Identification of disease-related spatial covariance patterns using neuroimaging data. *J Vis Exp* 2013;76:e50319.
20. Spetsieris PG, Eidelberg D. Scaled subprofile modeling of resting state imaging data in Parkinson's disease: methodological issues. *Neuroimage* 2011;54:2899–2914.
21. McKhann G, Drachman D, Folstein M, Katzman R, Price D, Stadlan EM. Clinical diagnosis of Alzheimer's disease: report of the NINCDS-ADRDA work group under the auspices of Department of Health and Human Services Task Force on Alzheimer's Disease. *Neurology* 1984;34:939–944.
22. Hughes A, Daniel S, Ben-Shlomo Y, Lees A. The accuracy of diagnosis of parkinsonian syndromes in a specialist movement disorder service. *Brain* 2002;125:861–870.
23. Mattis S. *Dementia Rating Scale: Professional Manual*. Odessa: Psychological Assessment Resources; 1988.
24. Tang C, Poston K, Dhawan V, Eidelberg D. Abnormalities in metabolic network activity precede the onset of motor symptoms in Parkinson's disease. *J Neurosci* 2010;30:1049–1056.
25. Holtbernd F, Ma Y, Peng S, et al. Dopaminergic correlates of metabolic network activity in Parkinson's disease. *Hum Brain Mapp* 2015;36:3575–3585.
26. Niethammer M, Tang CC, Ma Y, et al. Parkinson's disease cognitive network correlates with caudate dopamine. *Neuroimage* 2013;78:204–209.
27. Crane PK, Carle A, Gibbons LE, et al. Development and assessment of a composite score for memory in the Alzheimer's Disease Neuroimaging Initiative (ADNI). *Brain Imaging Behav* 2012;6:502–516.
28. Weintraub D, Dietz N, Duda JE, et al. Alzheimer's disease pattern of brain atrophy predicts cognitive decline in Parkinson's disease. *Brain* 2012;135:170–180.
29. Davatzikos C, Genc A, Xu D, Resnick SM. Voxel-based morphometry using the RAVENS maps: methods and validation using simulated longitudinal atrophy. *Neuroimage* 2001;14:1361–1369.
30. Spetsieris PG, Ko JH, Tang CC, et al. Metabolic resting-state brain networks in health and disease. *Proc Natl Acad Sci U S A* 2015;112:2563–2568.
31. Schmahmann JD, Doyon J, Toga AW, Petrides M, Evans AC. *MRI Atlas of the Human Cerebellum*. San Diego: Academic Press; 2000.



Neurology® Online CME Program

Earn CME while reading *Neurology*. This program is available only to online *Neurology* subscribers. Simply read the articles marked CME, go to Neurology.org, and click on CME. This will provide all of the information necessary to get started. The American Academy of Neurology (AAN) is accredited by the Accreditation Council for Continuing Medical Education (ACCME) to sponsor continuing medical education for physicians. *Neurology* is planned and produced in accordance with the ACCME Essentials. For more information, contact AAN Member Services at 800-879-1960.