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Effects of Alzheimer's and Lewy Body Disease Pathologies on Brain Metabolism

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Abstract

Objective: This study aimed to determine the pattern of ^{18}F -fluorodeoxyglucose positron emission tomography (FDG-PET) related to postmortem Lewy body disease (LBD) pathology in clinical Alzheimer's disease (AD).

Methods: FDG-PET scans were analyzed in 62 autopsy-confirmed patients and 110 controls in the Alzheimer's Disease Neuroimaging Initiative. Based on neuropathologic evaluations on Braak's stage for neurofibrillary tangle, Consortium to Establish a Registry for AD score for neuritic plaque, and Lewy-related pathology, subjects were classified into AD(-)/LBD(-), AD(-)/LBD(+), AD(+)/LBD(-), and AD(+)/LBD(+) groups. The association between postmortem LBD and AD pathologies and antemortem brain metabolism was evaluated.

Results: AD and LBD pathologies had significant interaction effects to decrease metabolism in the cerebellar vermis, bilateral caudate, putamen, basal frontal cortex, and anterior cingulate cortex in addition to the left side of entorhinal cortex and amygdala; and those to increase metabolism in the bilateral parietal and occipital cortices. LBD pathology was associated with hypermetabolism in the cerebellar vermis, bilateral putamen, anterior cingulate cortex, and basal frontal cortex, corresponding to the LB-related hypermetabolic patterns. AD pathology was associated with hypometabolism in the bilateral hippocampus, entorhinal cortex, and posterior cingulate cortex regardless of LBD pathology, while LBD pathology was associated with hypermetabolism in the bilateral putamen and anterior cingulate cortex regardless of AD pathology.

Interpretation: Postmortem LBD and AD pathologies had significant interaction effects on the antemortem brain metabolism in clinical AD patients. Specific metabolic patterns related to AD and LBD pathologies could be elucidated when simultaneously considering the two pathologies.

Keywords: Lewy body disease; Alzheimer's disease; Metabolic pattern;
 ^{18}F -fluorodeoxyglucose PET.

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Introduction

In the two most common forms of neurodegenerative dementia, Alzheimer's disease (AD) and Lewy body disease (LBD), mixed pathology is found in more than 50% of the patients.¹⁻³ In AD, the presence of LBD pathology is associated with younger symptom onset,⁴ parkinsonism,⁵ worse cognition,⁶ and faster progression.⁷ However, even with the evidence of mixed pathologies in cognitively impaired patients, the lack of sensitive *in vivo* biomarkers for LBD pathology prevents precise diagnoses in clinical settings. Therefore, it is not uncommon for patients who were clinically diagnosed with AD to show mixed pathologies of LBD during autopsy.⁸ Failure to evenly distribute patients with mixed LBD may lead to inaccurate evaluation of new therapeutic agents in clinical trials for AD.

Metabolic changes on ¹⁸F-fluorodeoxyglucose (FDG)-positron emission tomography (PET) have clinical implications for the differential diagnosis of neurodegenerative dementia.⁹ Hypometabolism in the temporal, parietal, and medial temporal cortex is the typical metabolic pattern of clinically diagnosed AD patients,^{10,11} and occipital hypometabolism¹² and posterior cingulate island signs¹³ are observed in clinically diagnosed patients with dementia with Lewy bodies (DLB) (Supplementary Figure 1).

Hypermetabolism in the basal ganglia, motor cortex, and cerebellum has been reported in LBD, including idiopathic RBD,¹⁴ Parkinson's disease (PD),^{15,16} and DLB,^{17,18} and these LB-related metabolic patterns (LBRPs) may enhance the diagnostic accuracy of LBD.¹⁹ However, to the best of our knowledge, metabolic changes related to AD and LBD have not been elucidated in autopsy-confirmed patients.

In-vivo detection of mixed pathologies in patients with neurodegenerative dementia would have clinical importance for treatment and predicting prognosis. In this study, we evaluated the metabolic changes of antemortem FDG-PET related to AD and LBD pathologies on postmortem autopsies from the Alzheimer's Disease Neuroimaging Initiative

(ADNI) cohort, which were aimed to include dementia or mild cognitive impairment (MCI) due to AD.²⁰ We hypothesized that AD and LBD pathologies could have significant interaction effects on brain metabolism,²¹⁻²⁵ and pathology-specific metabolic patterns could be identified after considering these interaction effects.

Methods

Study subjects

Our data were obtained from the ADNI database (<http://adni.loni.usc.edu/>). The ADNI is a longitudinal study launched in 2003 as a public-private partnership, to assess biomarkers for the progression of MCI and early AD. The inclusion and exclusion criteria are available at the ADNI protocol (<http://adni.loni.usc.edu/methods/documents/>). Briefly, diagnosis of dementia was based on the National Institute of Neurological and Communicative Disorders and Stroke and the Alzheimer's Disease and Related Disorders Association criteria for probable AD. Individuals with MCI had preserved general cognition and functional performance and were required to have an abnormal memory function, Mini-Mental State Examination (MMSE) between 24 and 30, and a clinical dementia rating (CDR®) score of 0.5. General cognitive status was measured using the MMSE and CDR-sum of boxes (CDR-SOB).

Autopsy consent and neuropathologic evaluation were established by a Neuropathology Core (NPC) of the ADNI database. As of July 2021, autopsy results from 81 subjects were available in the ADNI database. After excluding the subjects who had not received FDG-PET (n=17) and those who maintained normal cognition until autopsy (n=2), a total of 62 subjects were included in the analysis. Among cognitively normal participants, 352 received FDG-PET. We included subjects who were followed up for >2 years, had maintained normal cognition, and had normal CSF p-tau/A β ratios (<0.028).²⁶ After excluding low-quality MRI and PET images (n=4), 110 control subjects were included.

Neuropathologic assessment

Neuropathologic evaluation of the ADNI database²⁷ follows the National Institute on Aging and Alzheimer's Association (NIA-AA) guidelines²⁸ and DLB Consortium classification.²⁹ Details of the neuropathologic protocol of the ADNI-NPC can be found in the ADNI database (<http://adni.loni.usc.edu/>). AD neuropathology was evaluated using the Thal phase for A β plaques, the Braak and Braak stage for neurofibrillary tangles (NFTs) or tau burden, and the Consortium to Establish a Registry for AD (CERAD) score for neuritic plaques.²⁷ A β plaques have little effects on the clinical features in LBD as well as AD,²² and pathologic diagnosis of mixed AD with LBD could be overestimated by including LBD patients with high A β load but low burden of tau or neuritic plaques. Therefore, we defined the presence of AD pathology using the previously reported modified NIA-AA criteria based on NFT Braak stages and CERAD scores.^{3,30} In specific, subjects were classified into: (i) no AD = NFT(0)/CERAD(0), NFT(I-II)/CERAD(0); (ii) low AD = NFT(I-II)/CERAD(1-3), NFT(III-IV)/CERAD(0-1); (iii) intermediate AD = NFT(III-IV)/CERAD(2-3), NFT(V-VI)/CERAD(1); and (iv) high AD = NFT(V-IV)/CERAD(2-3). Subjects with intermediate or high AD pathology were treated as having AD pathology.

LBD neuropathologic changes were evaluated based on the degree and distribution of Lewy-related pathology in six stages: none, brainstem-predominant, limbic (transitional), neocortical (diffuse), amygdala-predominant, or olfactory-only.^{28,29,31} Dichotomization of LBD pathology was based on the presence of brainstem-predominant, limbic, and neocortical Lewy-related pathology, which are mostly related to the core clinical features of LBD.^{23,32} According to dichotomized AD and LBD pathologies, autopsy-confirmed subjects were grouped into four groups: AD(-)/LBD(-), AD(-)/LBD(+), AD(+)/LBD(-), and AD(+)/LBD(+). The control subjects were considered to have AD(-) and LBD(-).

MRI image processing

We used the FMRIB Software Library (FSL, <http://www.fmrib.ox.ac.uk/fsl>) for image processing. Each subject's T1-weighted images were corrected for intensity inhomogeneity, skull-stripped, and registered to the ADNI-Montreal Neurological Institute (MNI) atlas, which is a T1-weighted template for older adults.^{33,34} The brain tissues were classified into white matter, gray matter (GM), or CSF based on the hidden-Markov random field model and the associated expectation-maximization algorithm.³⁵ The GM probability map obtained from this algorithm was non-linearly transformed into the ADNI-MNI template. The striatal regions were segmented using the FMRIB integrated registration and segmentation tool (FIRST) algorithm.³⁶ We included the striatal regions of interest in the GM class. Then, we averaged all of the individual GM probability maps and assigned each voxel into either the foreground or background by binarizing above 30% of the map to generate a study-specific GM mask. For voxel-based MRI statistical analyses, an additional modulation step for the GM probability map was included to keep the total GM amount constant, regardless of local expansion or contraction due to image normalization.

FDG-PET image processing

We linearly registered each subject's FDG-PET image to individual T1-weighted MRI using a rigid body transformation. To generate standardized uptake value ratio (SUVR) maps, we used the cerebellar cortex as a reference region. Then, we spatially normalized the SUVR maps to the ADNI-MNI template using non-linear warping fields acquired in the T1-weighted image processing stage, and then smoothed them using 6-mm full width at a half-maximum Gaussian kernel. To calculate the FDG subject residual profile (FDG-SRP), each data was transformed into logarithmic form, and the data matrix was centered by subtracting each

subject mean and the group mean voxel profile from the data.³⁷ To extract individual LBRP scores, we adopted LBRPs from the Severance Hospital dataset, as described in the previous study.¹⁹ Briefly, to capture the regional covariance patterns associated with disease progression and severity in the LBD spectrum, we computed the principal component coefficients using reduced singular value decomposition from spatially normalized FDG-SRP images of the Severance Hospital dataset (55 DLB patients and 49 healthy controls). The LBRP scores were then computed by applying the first principal component coefficient map that is associated with the LBRPs to individual FDG-SRP images of the ADNI dataset within the GM mask.

Quality assurance

For quality assurance, results from the image processing stages were visually inspected by three researchers (Y. Lee, S. Jeon, and B.S. Ye) who were blinded to the subject information. We excluded seven subjects from the original dataset due to image processing errors in brain masking, tissue classification, and volume registration. Finally, a total of 172 subjects were included in the statistical analysis.

Statistical analysis

Statistical analyses of the demographic and clinical data were performed using IBM SPSS version 26.0 (SPSS Inc., Chicago, IL, USA), and those for voxel-based statistics were performed using the SurfStat toolbox (<http://www.math.mcgill.ca/keith/surfstat>). An analysis of variance, Kruskal-Wallis, and chi-square tests were used to compare the demographic variables between the control group and the four autopsy-confirmed groups. General linear models for LBRP and FDG-SRP were performed using AD and LBD pathologies and their interaction as predictors to evaluate the effect of each pathology on brain metabolism.

Covariates included the age at FDG scan, sex, and education. We also compared the LBRP and FDG-SRPs between the control group and the four disease groups using general linear models after controlling for age, sex, and education. In the voxel-based analyses, false discovery rate (FDR) methods were used to correct for multiple comparisons across the voxels. We displayed voxel-wise statistical outcomes, including effect sizes (r , d), within statistically significant areas (corrected $p < 0.05$, FDR) on the ADNI-MNI stereotaxic space in neurological convention. To interpret the patterns of independent and interaction effects from AD and LBD pathologies, we selected 15 regions of interest (ROIs) where there were significant pathology-related effects or group differences. For the ROI-based analyses, we used the median value of the selected ROIs based on the automated anatomical labeling atlas 3.³⁸

Data availability

All data used in this study can be obtained from the ADNI website (<http://adni.loni.usc.edu/>).

Results

Demographic and clinical characteristics

The demographic and clinical characteristics are shown in Table 1. The control subjects did not differ in age at the time of FDG-PET, sex, and education compared to the four neuropathologic groups. On average, the autopsy was conducted at the age of 84.1 years and 4.9 years after the FDG-PET acquisition. Age at autopsy and interval from FDG-PET to autopsy were comparable between the four neuropathologic groups. Neuropathologic groups had lower MMSE scores and higher CDR-SOB compared to the control group. Among neuropathologic groups, the MMSE score was the lowest in AD(+)/LBD(+) group and the highest in AD(-)/LBD(-) group. Detailed neuropathologic findings of the neuropathologic

groups are described in Supplementary Table 1. Seven of 10 subjects in the AD(-)/LBD(-) group had neuropathologic diagnoses, including frontotemporal lobar degeneration (FTLD), argyrophilic grain disease (AGD), primary age-related tauopathy (PART), and aging-related tau astroglipathy (ARTAG), while three did not have a specific diagnosis (Supplementary Table 2).

Effects of AD and LBD pathologies on brain metabolism

Significant interaction effects of AD and LBD pathologies were observed in the cerebellar vermis, bilateral caudate, putamen, basal frontal cortex, anterior cingulate cortex, parietal cortex, and occipital cortex in addition to the left entorhinal cortex and amygdala (Figure 1). After considering the interaction effects, AD pathology was associated with hypometabolism in the bilateral hippocampus, caudate, entorhinal cortex, posterior cingulate cortex, basal frontal cortex, and parietal cortex, in addition to the left amygdala, while it was associated with hypermetabolism in the cerebellar vermis, bilateral motor cortex, and medial occipital cortex. LBD pathology was associated with hypermetabolism in the cerebellar vermis, bilateral putamen, anterior cingulate cortex, and basal frontal cortex, while it was associated with hypometabolism in the bilateral parietal cortex and posterior cingulate cortex.

Comparison of brain metabolism with the control group

When the brain metabolism of each neuropathologic group was compared to the control group, AD(-)/LBD(+), AD(+)/LBD(-), and AD(+)/LBD(+) groups commonly had hypermetabolism in the cerebellum, pallidum, and motor cortex, while they had hypometabolism in the inferior/lateral temporal and parietal cortices (Figure 2). The AD(+)/LBD(-) and AD(+)/LBD(+) groups had hypometabolism in the hippocampus/entorhinal cortex and hypermetabolism in the medial occipital cortex, while the

AD(-)/LBD(+) group did not. Meanwhile, the AD(-)/LBD(+) and AD(+)/LBD(+) groups had hypermetabolism in the putamen, while the AD(+)/LBD(-) group did not. In the bilateral caudate and left amygdala, the AD(-)/LBD(+) group had hypermetabolism; the AD(+)/LBD(+) group had hypometabolism; and the AD(+)/LBD(-) group did not have significant metabolic changes. Compared to the control group, AD(-)/LBD(-) group had hypometabolism localized in the hippocampus/entorhinal cortex.

ROI-based comparison of brain metabolism

To capture the patterns of independent and interaction effects from AD and LBD, we presented the regional metabolism in the 15 ROIs (Table 2 and Figure 3). Compared to the sum of mean metabolism in the AD(+)/LBD(-) and AD(-)/LBD(+) groups, the AD(+)/LBD(+) group had more decreased metabolism in the bilateral caudate, and left side of amygdala and entorhinal cortex ROIs, but more increased metabolism in the bilateral occipital cortex and parietal cortex ROIs. Hypometabolism in the medial occipital cortex ROI and hypermetabolism in the bilateral basal frontal cortex, caudate, and hippocampus, and left side of amygdala and entorhinal cortex ROIs were observed only in the AD(-)/LBD(+) group, but not in the AD(+)/LBD(-) and AD(+)/LBD(+) groups. AD pathology was associated with hypometabolism in the bilateral hippocampus, posterior cingulate cortex, and left entorhinal cortex ROIs regardless of LBD pathology, while LBD pathology was associated with hypermetabolism in the bilateral putamen and anterior cingulate cortex ROIs regardless of AD pathology.

Lewy body-related metabolic patterns in neuropathologic groups and control group

When the effects of AD and LBD on LBRP were evaluated, both pathologies had significant independent and interaction effects on the LBRP (Table 3). The general linear model for

LBRP showed that the AD(-)/LBD(+), AD(+)/LBD(-), and AD(+)/LBD(+) groups had higher LBRP than did the control and AD(-)/LBD(-) groups (Supplementary Table 3). In the AD(+)/LBD(-) group, GM density in the right insula, bilateral posterior putamen, frontal, temporal, and occipital cortices showed a significant correlation with the LBRPs (Supplementary Figure 2). When the AD(+)/LBD(-) group was further divided into AD patients with amygdala-predominant or olfactory-only Lewy-related pathology (AD(+)/Lewy(+); n = 10) and those without (AD(+)/Lewy(-); n = 19), both groups exhibited increased metabolism in the cerebellum, pallidum, and motor cortex compared to the control group. Although the AD(+)/Lewy(+) group had relatively more hypermetabolism in the cerebellum, pallidum, and putamen and higher LBRPs compared to the AD(+)/Lewy(-) group, statistical significance was not reached (Supplementary Figure 3 and Supplementary Table 4).

Discussion

Herein, we evaluated the antemortem brain metabolic patterns related to the postmortem AD and LBD pathologies. The major findings of our study were as follows. First, AD and LBD pathologies had regionally distinct interaction effects on brain metabolism. Second, after considering the interaction effects, LBD pathology was associated with hypermetabolism in the cerebellar vermis, bilateral putamen, anterior cingulate cortex, and basal frontal cortex. Third, AD pathology was associated with hypometabolism in the bilateral hippocampus, entorhinal cortex, caudate, posterior cingulate cortex, basal frontal cortex, parietal cortex, and left amygdala. Taken together, our results suggest that AD and LBD pathologies are interactively associated with brain metabolic changes, and specific metabolic patterns related to AD and LBD could be elucidated when simultaneously considering the two pathologies.

Our first major finding was that AD and LBD pathologies had regionally distinct

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interaction effects on brain metabolism. First, decremental interacting effects on the cerebellar vermis, bilateral caudate, putamen, basal frontal cortex, anterior cingulate cortex, and left side of entorhinal cortex and amygdala metabolism suggest that two pathologies may aggravate synaptic dysfunction. Cross-fibrillation between tau and α -synuclein could be a possible candidate for these interaction effects to aggravate their spreading and neuronal degeneration.³⁹⁻⁴¹ On the other hand, incremental interacting effects were observed in the bilateral parietal and occipital metabolism. In specific, the metabolism of parietal and occipital cortices in AD(+)/LBD(+) group was greater than the sum of metabolism in AD(+)/LBD(-) and AD(-)/LBD(+) groups. Also, medial occipital hypometabolism was observed only in AD(-)/LBD(+) group, but not in AD(+)/LBD(+). This incremental interaction was interpreted as pathologic rather than compensatory and could be related to previously reported neuronal hyperexcitability in animal models of tau and α -synuclein co-pathologies.^{42,43} Based on the regionally distinct interaction effects between AD and LBD pathologies, it is suggested that metabolic changes due to neurodegenerative diseases cannot be measured by unidirectional hypometabolism.

Our second major finding was that after considering the significant interaction effects of AD and LBD on brain metabolism, LBD pathology was associated with hypermetabolism in the cerebellar vermis, bilateral putamen, anterior cingulate cortex, and basal frontal cortex. This hypermetabolic pattern overlaps with the previously reported PD-^{15,16} or DLB-related metabolic patterns.¹⁷⁻¹⁹ However, to our knowledge, the present study is the first autopsy-based validation of the association between LBD pathology and LBRPs. Furthermore, hypermetabolism in the bilateral putamen and anterior cingulate cortex was evident in both AD(-)/LBD(+) and AD(+)/LBD(+) groups (Figure 3 and Table 2), suggesting that they could be useful biomarkers for the presence of LBD pathology regardless of AD pathology. In contrast, occipital hypometabolism and relative preservation of posterior

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cingulate cortex metabolism were evident only in AD(-)/LBD(+) group, but not in AD(+)/LBD(+) group (Figure 3). Furthermore, independent effect of LBD pathology on occipital metabolism was not significant, in contrast to the negative effect on posterior cingulate cortex (Table 2). This may explain the low diagnostic sensitivity of cingulate island sign and occipital hypometabolism in clinical settings,^{12,13} especially in patients with concomitant AD pathology.

Our third major finding was that after considering the significant interaction effects of AD and LBD on brain metabolism, AD pathology was associated with hypometabolism in the bilateral caudate, hippocampus, entorhinal cortex, posterior cingulate cortex, basal frontal cortex, parietal cortex, and left amygdala. Among these brain regions, hypometabolism in the bilateral hippocampus, posterior cingulate cortex, and left entorhinal cortex was observed in the AD(+)/LBD(-) and AD(+)/LBD(+) groups, suggesting that they could be metabolic hallmarks of AD regardless of the presence of LBD pathology (Figure 3). Although the effect size of AD pathology on brain metabolism was the highest in both left entorhinal cortex and bilateral posterior cingulate cortex (Table 2), as the AD(-)/LBD(+) group also showed hypometabolism in the posterior cingulate cortex compared to the control group (Figure 2), hypometabolism on the left side of entorhinal cortex could be a better biomarker for AD.¹⁷

Notably, AD pathology was associated with hypermetabolism of the cerebellar vermis, bilateral motor cortex, medial occipital cortex, and pallidum (Figure 1). Furthermore, in addition to the AD(-)/LBD(+) and AD(+)/LBD(+) groups, the AD(+)/LBD(-) group also showed hypermetabolism in the cerebellar vermis, bilateral pallidum, and motor cortex (Figure 2). There could be several hypotheses to explain these results. First, the hypermetabolic changes in the AD(+)/LBD(-) group could be attributed to the limited Lewy-related pathology observed in 10 of 29 patients (34.5%) in the AD(+)/LBD(-) group (Supplementary Table 1). In the AD(+)/LBD(-) group, LBRPs were correlated with lower

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GM density in the right insula, bilateral posterior putamen, frontal, temporal, and occipital cortices, where GM changes were observed in LBDs,⁴⁴⁻⁴⁶ with relative sparing of the entorhinal cortex, vermis, and anterior cingulate cortex (Supplementary Figure 2). Second, hypermetabolic brain changes may not be specific to Lewy-related pathology but also be related to AD pathology. To test this hypothesis, we compared the metabolic changes between AD(+)/Lewy(-) and AD(+)/Lewy(+) within the AD(+)/LBD(-) group (n = 29). The AD(+)/Lewy(-) group exhibited increased metabolism in the cerebellum, pallidum, and motor cortex compared to the control group (Supplementary Figure 3) and had comparable LBRPs than AD(+)/Lewy(+) group (Supplementary Table 4). There could be several hypothetical mechanisms for the increased metabolism in these pure AD patients. First, tau-related disruption of inhibitory network could induce relative hypermetabolism.⁴⁷ Second, metabolism could be relatively preserved due to the lack of tau pathology in the cerebellum and motor cortex. Third, synergistic interaction between tau and limited Lewy-related pathology in brain regions where neuropathologic examinations are not routinely conducted could also be a possible candidate. However, since hypermetabolism involving the putamen was observed only in the AD(+)/Lewy(+) group but not in the AD(+)/Lewy(-) group, it could be a biomarker specific for Lewy-related pathology in AD patients. Future autopsy studies with sufficiently large sample sizes with post-mortem global screening of Lewy-related pathology are needed.

Although the underlying pathologies of AD(-)/LBD(-) were heterogeneous, most of the cases (70%) were diagnosed as having tauopathies, including ARTAG and primary tauopathy, such as FTLN, AGD, and PART (Supplementary Table 2). Clinical presentations of these tau-related pathologies are considered to be milder than AD, consistent with the limited hypometabolism of entorhinal cortex observed in the AD(-)/LBD(-) group compared to other neuropathology groups (Figure 2A).⁴⁸⁻⁵⁰ Our results suggest that limited

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entorhinal/hippocampal hypometabolism could be an imaging feature of non-AD tauopathies mimicking clinical AD.

This study had several limitations. First, there were time intervals from FDG-PET scanning to autopsy. We considered that the pathologic burden at the time of FDG-PET scanning would be similar to that in the postmortem autopsy results. Second, due to the rarity of autopsy data, our results were based on small sample size. Also, since we combined the subjects without any Lewy-related pathology and those with limited Lewy-related pathology into a LBD(-) group, we could not analyze the effects of various Lewy-related pathology distributions on metabolic patterns. Future studies with a large sample size are warranted using different categorizations by the distribution of Lewy-related pathology to reveal the effects of LBD staging on metabolism. Despite these limitations, we found that mixed LBD pathologies are common and contribute to brain metabolic changes in clinical AD patients.

In conclusion, we found significant interaction effects of postmortem AD and LBD pathologies on antemortem brain metabolism and demonstrated AD- and LBD-related metabolic changes in patients with the clinical diagnosis of AD. Since our results were based on group-level findings, future studies with a large sample size are needed to investigate the diagnostic implication of AD- and LBD-related metabolic patterns on an individual level.

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Potential conflict of interest: The authors declare that they have no conflict of interest.

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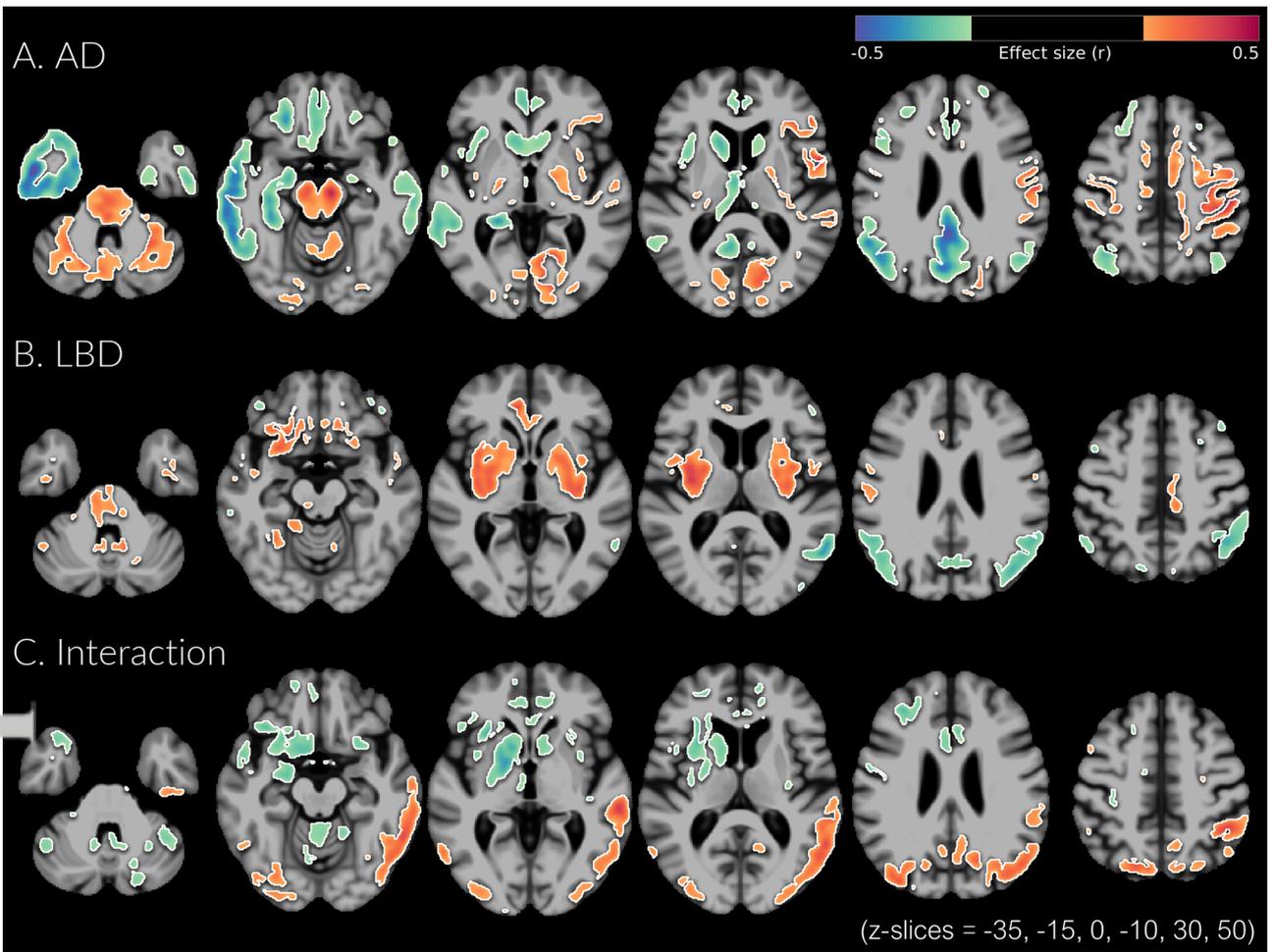
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Figure legends

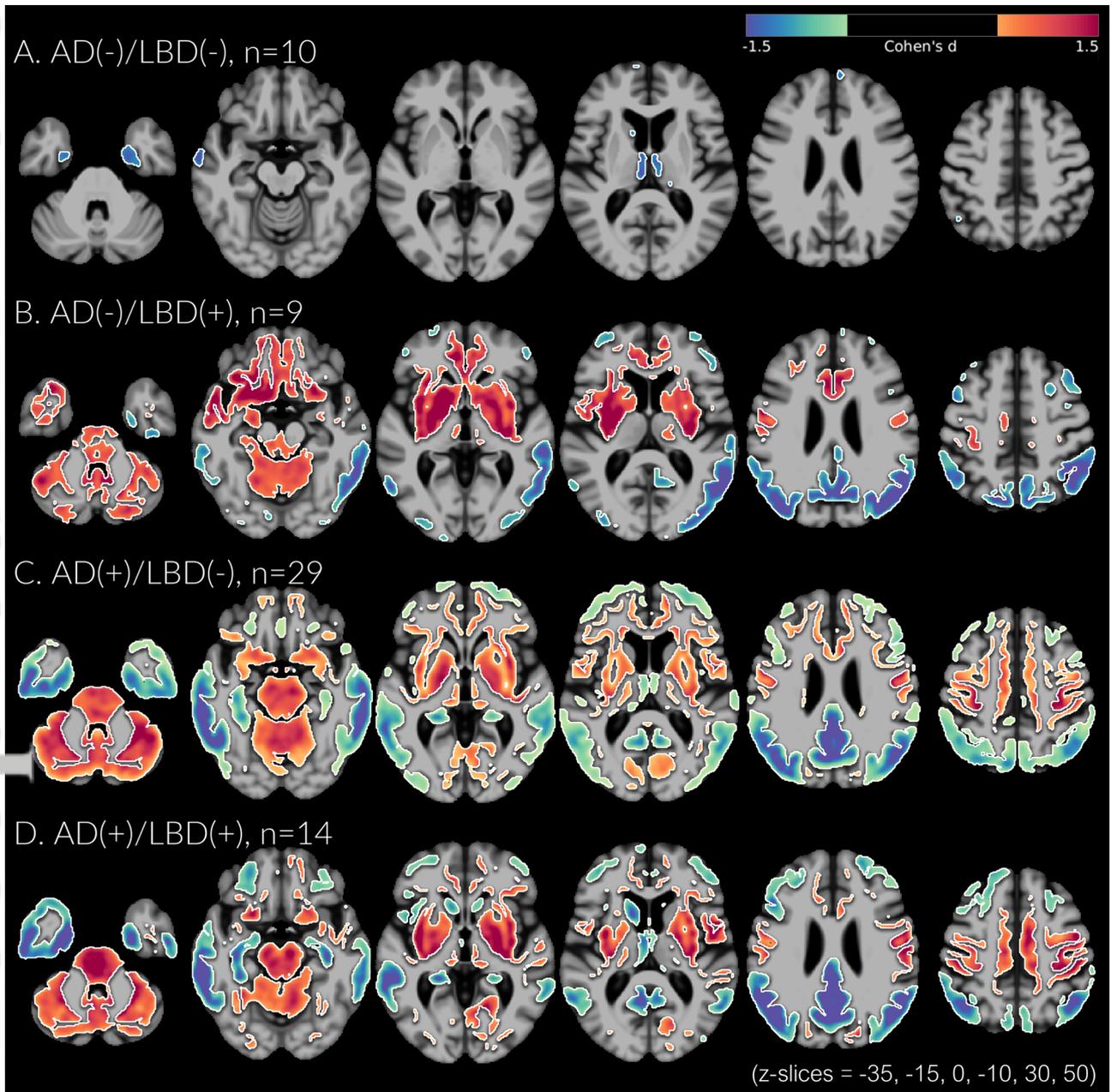
Figure 1. Effects of AD and LBD pathologies on regional brain metabolism. Data are the results of general linear models for regional ^{18}F -FDG metabolism after controlling for age, sex, and education. AD (A), LBD (B), and their interaction term (C) were used as predictors. The color scale indicates effect sizes (r-score) within statistically significant regions after multiple comparisons correction ($p < 0.05$, false discovery rate). The brain images are displayed according to neurological convention. AD, Alzheimer's disease; FDG, fluorodeoxyglucose; LBD, Lewy body disease.

Figure 2. Regional brain metabolism of the neuropathologic groups. Data are the results of group comparison for regional ^{18}F -FDG metabolism after controlling for age, sex, and education. Each neuropathologic group was compared to the control group. The color scale indicates effect sizes (Cohen's d) within statistically significant regions after multiple comparisons correction ($p < 0.05$, false discovery rate). The brain images are displayed according to neurological convention. AD, Alzheimer's disease; FDG, fluorodeoxyglucose; LBD, Lewy body disease.

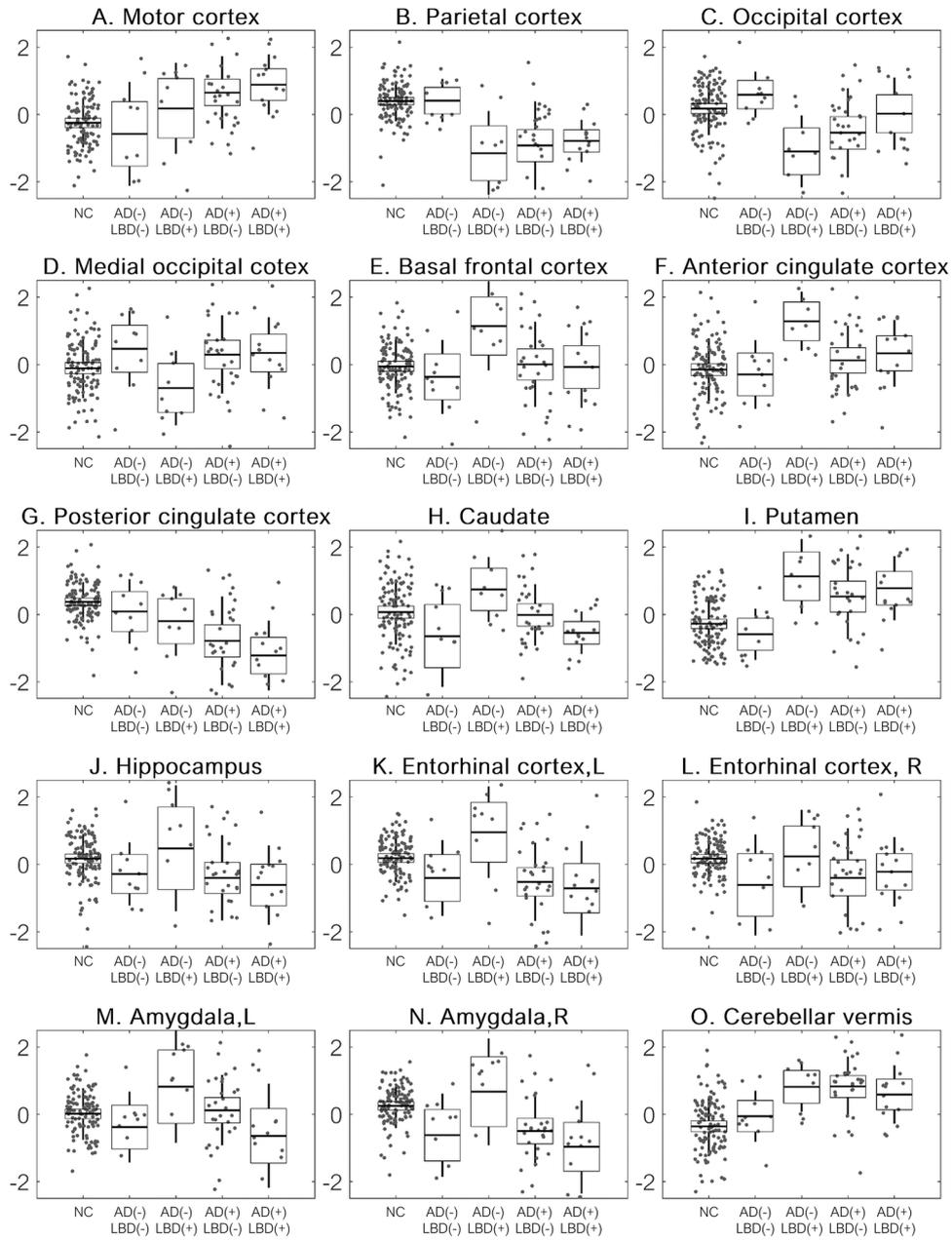
Figure 3. Regional brain metabolism in the neuropathologic groups. Data shows the distribution of the z-transformed residuals in each neuropathologic group from the general linear models for regional ^{18}F -FDG metabolism after controlling for age, sex, and education. Fifteen regions of interests were selected where significant independent effects or interactions effects between AD and LBD pathologies were identified from Figure 1. ACC, anterior cingulate cortex; AD, Alzheimer's disease; FDG, fluorodeoxyglucose; LBD, Lewy body disease; LBRPs, Lewy body-related metabolic patterns; PCC, posterior cingulate cortex.



ANA_26355_Figure1.tif



ANA_26355_Figure2.tif



ANA_26355_Figure3.tif

Table 1. Demographic and clinical characteristics of participants

	Control	AD(-)/LBD(-)	AD(-)/LBD(+)	AD(+)/LBD(-)	AD(+)/LBD(+)	<i>p</i>
Number	110	10	9	29	14	
Age at PET (y)	76.6 (6.0)	80.0 (6.8)	82.0 (8.8)	78.5 (8.1)	78.0 (6.3)	0.093
Age at autopsy (y)	N/A	85.1 (7.5)	86.9 (8.9)	83.6 (8.6)	82.3 (6.6)	0.563
Interval between PET and autopsy	N/A	5.2 (3.8)	4.9 (3.2)	5.1 (2.7)	4.3 (2.8)	0.874
Sex, female	49 (44.5%)	3 (25.0%)	1 (11.1%)	13 (44.8%)	2 (14.3%)	0.053
Education (y)	16.5 (2.7)	15.9 (2.6)	16.4 (3.5)	16.1 (2.4)	16.6 (2.2)	0.934
MMSE	29.0 (1.4)	27.9 (1.6)	24.4 (3.2) ^{a, b, c, d}	20.8 (6.0) ^{a, b, c}	20.4 (5.5) ^{a, b, d}	<0.001
CDR-SOB	0	1.5 (1.4)	4.1 (3.5) ^{a, b, d}	6.2 (4.0) ^{a, b}	6.6 (3.5) ^{a, b, d}	<0.001

Abbreviations: AD, Alzheimer's disease; CDR-SOB, clinical dementia rating sum of boxes; LBD, Lewy body disease; MMSE, Mini-Mental Status Examination; PET, positron emission tomography.

Data are expressed as the mean (standard deviation) or numbers (%). Results are from the analysis of variance or chi-square tests, as appropriate.

^aSignificantly different in comparison with the control group.

^bSignificantly different in comparison with AD(-)/LBD(-) group.

^cSignificantly different in comparison between AD(-)/LBD(+) and AD(+)/LBD(-) groups.

^dSignificantly different in comparison between AD(-)/LBD(+) and AD(+)/LBD(+) groups.

Table 2. Effects of AD and LBD pathologies on regional metabolic activity shown in Figure 1

Regions	AAL3 label	AD		LBD		Interaction	
		r	p^a	r	p^a	r	p^a
Motor cortex	1, 2	0.28	< 0.001	0.14	0.075	-0.05	0.313
Parietal cortex	63 – 70	-0.18	0.013	-0.30	< 0.001	0.34	< 0.001
Occipital cortex	53 – 58	0.07	0.191	-0.14	0.075	0.32	< 0.001
Medial occipital cortex	47, 48	0.22	0.005	-0.10	0.175	0.11	0.102
Basal frontal cortex	17, 18, 21 – 24	-0.19	0.013	0.20	0.018	-0.22	0.005
Anterior cingulate cortex	151, 152 ^b	-0.12	0.080	0.28	< 0.001	-0.22	0.006
Posterior cingulate cortex	39, 40	-0.38	< 0.001	-0.20	0.015	0.03	0.368
Cerebellar vermis	113, 114, 120	0.17	0.024	0.19	0.020	-0.26	< 0.001
Caudate	75, 76	-0.21	0.006	0.03	0.446	-0.21	0.008
Putamen	77, 78	0.08	0.161	0.31	< 0.001	-0.22	0.005
Hippocampus	41, 42	-0.26	< 0.001	0.02	0.451	-0.09	0.151
Entorhinal cortex, L	43 ^c	-0.38	< 0.001	0.00	0.498	-0.16	0.028
Entorhinal cortex, R	44 ^c	-0.16	0.030	0.05	0.401	0.01	0.453
Amygdala, L	45	-0.22	0.005	0.02	0.451	-0.26	< 0.001
Amygdala, R	46	-0.11	0.099	0.02	0.451	-0.13	0.069

Abbreviations: AAL3, Automated Anatomical Labelling atlas 3; AD, Alzheimer's disease; and LBD, Lewy body disease.

Data are the results of general linear models for regional ¹⁸F-FDG metabolism after controlling for age, sex, and education. AD, LBD, and their interaction term (AD×LBD) were used as predictors.

^aCorrected *P* values after multiple comparison correction using false discovery rate method.

^bSubgenual part of ACC on AAL3.

^cThe anterior part of parahippocampal gyrus on AAL3 subdivided in half along the y-axis.

Table 3. Effects of AD and LBD neuropathologies on LBRPs

	AD		LBD		Interaction	
	β (SE)	<i>p</i>	β (SE)	<i>p</i>	β (SE)	<i>p</i>
LBRPs	24.80 (2.69)	<0.001	74.64 (15.74)	<0.001	-25.27 (7.04)	<0.001

Abbreviations: AD, Alzheimer's disease; LBD, Lewy body disease; LBRPs, Lewy body-related metabolic patterns.

Data are the results of general linear models for LBRPs after controlling for age, sex, and education. AD, LBD, and their interaction term (AD×LBD) were used as predictors.