

Original Article





Received: Feb 2, 2025 Revised: Mar 11, 2025 Accepted: Mar 25, 2025 Published online: Apr 9, 2025

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Establishing Regional Aβ Cutoffs and Exploring Subgroup Prevalence Across Cognitive Stages Using BeauBrain Amylo®

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ABSTRACT

Background and Purpose: Amyloid-beta (Aβ) plaques are key in Alzheimer's disease (AD), with Aβ positron emission tomography imaging enabling non-invasive quantification. To address regional Aβ deposition, we developed regional Centiloid scales (rdcCL) and commercialized them through the computed tomography (CT)-based BeauBrain Amylo platform, eliminating the need for three-dimensional T1 magnetic resonance imaging (MRI). **Objective:** We aimed to establish robust regional Aβ cutoffs using the commercialized BeauBrain Amylo platform and to explore the prevalence of subgroups defined by global, regional, and striatal Aβ cutoffs across cognitive stages.

Methods: We included 2,428 individuals recruited from the Korea-Registries to Overcome Dementia and Accelerate Dementia Research project. We calculated regional Aβ cutoffs using Gaussian Mixture Modeling. Participants were classified into subgroups based on global, regional, and striatal Aβ positivity across cognitive stages (cognitively unimpaired [CU], mild cognitive impairment, and dementia of the Alzheimer's type).

Results: MRI-based and CT-based global A β cutoffs were highly comparable and consistent with previously reported Centiloid values. Regional cutoffs revealed both similarities and differences between MRI- and CT-based methods, reflecting modality-specific segmentation processes. Subgroups such as global(-)regional(+) were more frequent in non-dementia stages, while global(+)striatal(-) was primarily observed in CU individuals.

Conclusions: Our study established robust regional Aβ cutoffs using a CT-based rdcCL method and demonstrated its clinical utility in classifying amyloid subgroups across

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Funding

This research was supported by a grant from the Korea Health Technology R&D Project through the Korea Health Industry Development Institute (KHIDI), funded by the Ministry of Health & Welfare and Ministry of Science and ICT, Republic of Korea (grant number: HU20C0111); by a National Research Foundation of Korea (NRF) grant funded by the Korean government (MSIT) (NRF-2019R1A5A2027340); and by funds (2021-ER1006-00) from the Research of Korea Disease Control and Prevention Agency.

Conflict of Interest

The authors have no financial conflicts of interest.

Author Contributions

Conceptualization: Park S, Kim K, Jang H, Na DL, Kim HJ, Moon SH, Kim JP, Seo SW, Kim J, Kwak K; Data curation: Jang H, Na DL, Kim HJ, Moon SH, Kim JP; Investigation: Yoon S, Kim S, Ahn J; Methodology: Lim KY; Supervision: Seo SW, Kwak K; Writing - original draft: Park S; Writing - review & editing: Seo SW, Kim J, Kwak K.

cognitive stages. These findings highlight the importance of regional $A\beta$ quantification in understanding amyloid pathology and its implications for biomarker-guided diagnosis and treatment in AD.

Keywords: Alzheimer Disease; Tomography, X-Ray Computed; Positron-Emission Tomography; Cognition Disorders; Amyloid Plaques

INTRODUCTION

Alzheimer's disease (AD) is characterized by the accumulation of amyloid-beta (A β) plaques in the brain, which begins years before the onset of clinical symptoms. ¹⁻⁴ The advent of A β positron emission tomography (PET) imaging has provided a non-invasive in vivo method to detect amyloid deposition, offering valuable insights into the pathophysiological stages of AD. ⁵⁻⁹ Studies have shown that the prevalence of amyloid positivity increases across cognitive stages, from approximately 20%–40% in cognitively unimpaired (CU) individuals to 50%–60% in those with mild cognitive impairment (MCI) and over 80% in dementia of the Alzheimer's type (DAT). ^{10,11} While visual assessments remain widely used for A β PET interpretation, quantification pipelines offer more objective and reproducible results, emphasizing the need for comparative studies to evaluate these methods.

To standardize amyloid quantification across different PET ligands, the Centiloid (CL) method was introduced, enabling the harmonization of 11C-labeled Pittsburgh compound B and 18F-labeled A β PET tracers through a common scale. This approach has greatly enhanced cross-study comparability and clinical utility in AD research. However, traditional CL methods primarily focus on a global measure of A β burden, potentially overlooking regional variations that can provide critical insights into disease progression.

Recent evidence underscores the need to examine such regional variations in Aβ deposition. For instance, some individuals with globally amyloid-negative scans may still exhibit regional amyloid positivity, which has been associated with subtle cognitive changes¹⁶ and an increased likelihood of conversion to global amyloid positivity.¹⁷ Conversely, for globally amyloid-positive individuals, striatal involvement has been linked to worse clinical outcomes, highlighting the importance of assessing both global and region-specific amyloid patterns.^{16,18,19} Furthermore, CL values are increasingly used in the context of Food and Drug Administration-approved amyloid-targeting therapies, aiding in patient selection, treatment monitoring, and identifying those most likely to benefit from these treatments.²⁰⁻²³

To address the limitations of a purely global approach, our group developed regional Centiloid scales (rdcCL) that harmonize A β uptake across various PET ligands using computed tomography (CT) images acquired during PET-CT scans for anatomical reference. ²⁴ By eliminating the need for additional three-dimensional (3D) T1 magnetic resonance imaging (MRI) data, this method simplifies the analytic process and broadens its applicability in both clinical and research settings. Previous studies have validated the pathological and clinical utility of rdcCL, showing that it effectively detects region-specific A β deposition and can distinguish clinical phenotypes relevant to AD progression. ^{16,19}

Building on these findings, we developed BeauBrain Amylo, a platform designed to standardize and implement rdcCL for broader clinical and research use. A key advantage of



BeauBrain Amylo is its ability to derive regional A β measures from CT-based registration alone, thereby removing the need for separate 3D T1 MRI acquisitions. In this study, we aim to establish robust regional A β cutoffs using this platform and to explore the prevalence of subgroups defined by global, regional, and striatal A β cutoffs across cognitive stages (CU, MCI, and DAT). Through this work, we hope to deepen our understanding of amyloid pathology and its clinical implications, ultimately paving the way for more precise diagnostics and targeted interventions in AD.

METHODS

Participants selection

This study included 2,428 individuals recruited from the Korea-Registries to Overcome Dementia and Accelerate Dementia Research (K-ROAD) project, 25 which is a member of the Worldwide Alzheimer's Disease Neuroimaging Initiative. The K-ROAD project aimed to establish a genotype-phenotype cohort to accelerate the development of innovative diagnostic and therapeutic techniques for neurodegenerative diseases, mainly AD and related dementia syndromes, in collaboration with 25 nationwide university-affiliated hospitals in South Korea between 2016 and 2023. All participants displayed one or more of the following: CU, MCI, and DAT. CU participants (n=581) were selected based on following criteria: (1) absence of medical history that is likely to affect cognitive function based on Christensen's health screening criteria²⁶ and (2) absence of objective cognitive impairment observed on any cognitive domain (above the -1.0 standard deviation [SD] of age- and education-matched norms in memory and above -1.5 SD in other cognitive domains).²⁷ Participants with MCI (n=931) met the specified criteria²⁸: (1) subjective cognitive complaints by the participants or caregiver; (2) objective cognitive impairment in any cognitive domain (below the -1.0 SD of age- and education-matched norms in memory and/or below -1.5 SD in other cognitive domains); (3) no significant impairment in activities of daily living; and (4) no dementia. Participants diagnosed with DAT (n=645) fulfilled the National Institute on Aging and Alzheimer's Association diagnostic criteria.²⁹ The "others" group included a total of 271 participants, comprising 258 with vascular cognitive impairment (VCI), 9 with frontotemporal dementia (FTD), and 4 with dementia with Lewy bodies (DLB). These diagnoses were based on established criteria, including the National Institute of Neurological Disorders and Stroke and the Association Internationale pour la Recherche et l'Enseignement en Neurosciences criteria for VCI, 30 the International Consensus Criteria for FTD,³¹ and the Fourth Consensus Report of the DLB Consortium for DLB.³² This approach ensured consistency with standard diagnostic guidelines in neurodegenerative and vascular conditions.

All participants underwent clinical interviews, neurological examinations, neuropsychological testing, and brain MRI. After these evaluations, clinical diagnoses were determined through agreement among a multidisciplinary team. Furthermore, all participants underwent blood tests, including complete blood counts, blood chemistry tests, vitamin B12/folate measurement, thyroid function test, and syphilis serology, to rule out the possibility of medical conditions causing cognitive decline, and APOE genotyping. Patients were excluded if they had territorial infarctions, brain tumors, or vascular malformations on MRI. Patients with white matter hyperintensities caused by radiation injury, vasculitis, multiple sclerosis, or leukodystrophy were also excluded.



The Institutional Review Board (IRB) of Samsung Medical Center approved the study protocol (IRB approval No. 2021-02-135), and all methods were performed according to the approved guidelines. Written consent was obtained from each participant.

MRI data acquisition

Standardized 3D T1 turbo field echo images were acquired from all participants at Samsung Medical Center using the same 3.0 T MRI scanner (Philips Achieva; Philips Healthcare, Andover, MA, USA). The detailed parameters were as follows: sagittal slice thickness, 1.0 mm, over contiguous slices with 50% overlap; no gap; repetition time of 9.9 ms; echo time of 4.6 ms; flip angle of 8°, and matrix size of 240×240 pixels, reconstructed to 480×480 over a field of view of 240 mm.

Aβ PET-CT data acquisition

Participants underwent ¹⁸F-flutemetamol (FMM) and ¹⁸F-florbetaben (FBB) PET at Samsung Medical Center using a Discovery STe PET/CT scanner (GE Medical Systems, Milwaukee, WI, USA) in 3D scanning mode that examined 47 slices 3.3-mm in thickness spanning the entire brain. ^{33,34} According to the protocols for the ligands proposed by the manufacturers, a 20-minute emission PET scan with dynamic mode (consisting of 4×5 minutes frames) was performed 90 minutes after injection of a mean dose of 185 MBq of FMM or 311.5 MBq of FBB. 3D PET images were reconstructed in a 128×128×47 matrix with a voxel size of 2 mm×2 mm×3.27 mm using the ordered-subsets expectation maximization algorithm (FMM iterations=4 and subset=20; FBB iterations=4 and subset=20).

CT images were acquired using a 16-slice helical CT (140 KeV, 80 mA; 3.75-mm section width) for attenuation correction and were reconstructed in a 512×512 matrix with a voxel size of 0.5 mm×0.5 mm×3.27 mm. The kernel (i.e., smoothing filter) size of 5 mm was used for FMM according to the vendor's recommendation. However, since there was no recommendation of smoothing for FBB, we did not do perform it.

Quantitative analysis of Aβ PET images

We employed BeauBrain Amylo (BeauBrain Healthcare, Inc., Seoul, Korea; beaubrain.bio) to automatically quantify Aβ binding in amyloid PET. This software provides fully automated calculations of standardized uptake value ratio (SUVR) and CL values, ensuring robust quantitative analysis applicable to both PET-CT and PET-MR images. The processing pipeline includes co-registration, spatial normalization to a template, atlas-based segmentation, and calculation of dcSUVR and rdcSUVR. Subsequently, dcCL and rdcCL scales are calculated for the global region as well as six specific regions: frontal, temporal, parietal, posterior cingulate (PC), occipital and striatum. For CT images, and additional Hounsfield unit correction step is included as part of the preprocessing.

Defining subgroups by global, regional, and striatal AB cutoffs

The group was classified into four groups based on global and striatal rdcCL cutoffs. Initially, the participants were classified into 2 subgroup based on global A β rdcCL scales, which reflect the amyloid burden in the global region: global (–) and global (+). The global (–) group was further subdivided into regional (–) and regional (+) groups according to regional cutoffs for at least one region. Similarly, the global (+) group was divided into striatal (–) and striatal (+) groups based on striatal cutoffs. Thus, the cohort was classified into 4 groups: global (–) and regional (–) A β : G(–)R(–), global (–) and regional (+) A β : G(–)R(+), global (+) and striatal (–) A β : G(+)Str(–), and global (+) and striatal (+) A β : G(+)Str(+).



Table 1. Participant demographics and clinical findings

| Variables | Total | CU | MCI | DAT | Others |
|---------------------|-------------------------|---------------------|---------------------|---------------------|--------------------|
| No. of participants | 2,428 | 581 | 931 | 645 | 271 |
| Age (yr) | 71.2±9.5 | 69±10.6 | 72±8.2 | 70±10.2 | 76±7.1 |
| Sex (F) | 1,443 (59.4) | 357 (61.4) | 516 (55.4) | 398 (61.7) | 172 (63.4) |
| Education (yr) | 11.5±4.9 | 11.9±4.8 | 12.0±4.7 | 11.3±4.8 | 9.3±5.2 |
| APOE4 carriers | 931 (38.3) | 150 (25.8) | 394 (42.3) | 321 (49.8) | 66 (24.4) |
| Ligands (FMM/FBB) | 1,378/1,050 (56.8/43.2) | 208/373 (35.8/64.2) | 613/318 (65.8/34.2) | 323/322 (50.1/49.9) | 69/202 (25.5/74.5) |
| MMSE | 24.1±5.2 | 28.0±2.0 | 25.6±3.2 | 19.1±5.1 | 22.0±5.3 |

Values are presented as number (%) or mean ± standard deviation.

CU: cognitively unimpaired, MCI: mild cognitive impairment, DAT: dementia of the Alzheimer's type, F: female, FBB: 18F-florbetaben, FMM: 18F-flutemetamol, MMSE: Mini-Mental State Examination.

Statistical analyses

The cutoff values for amyloid positivity on the dcCL and rdcCL scales for each brain region in the MR and CT datasets were determined using a Gaussian Mixture Model (GMM), a probabilistic machine learning method that models the data distribution as a mixture of multiple-Gaussian distributions. A GMM with 2 components (k=2) was fitted using the 'normalmixEM' function from the 'mixtools' R package (R Studio version 2023.12.1+402). The cutoff was defined as the point where the 2 Gaussian distributions intersect.

RESULTS

Demographics of the participants

Table 1 summarizes the demographic and clinical characteristics of the study participants. A total of 2,428 individuals were included in the analysis, with a mean age of 71.2±9.5 years. Among them, 1,443 (59.4%) were female, and the average years of education was 11.5±4.9. APOE4 carriers comprised 931 participants (38.3%). The PET ligands used in this study included 1,050 participants (43.2%) with FBB and 1,378 participants (56.8%) with FMM. The mean Mini-Mental State Examination (MMSE) score across all participants was 24.1±5.2. The APOE genotype information was unavailable for 53 patients, while education years were not recorded for 25 patients, and MMSE scores were not evaluated for 110 patients.

MRI-based CL cutoffs by brain regions

Table 2 summarizes the CL cutoffs identified for each region. The GMM analysis revealed distinct cutoffs for key regions associated with amyloid deposition. The calculated cutoffs were 28.5 for the global brain, 34.1 for the frontal cortex, 29.5 for the parietal cortex, 28.1 for the temporal cortex, 43.3 for the occipital cortex, 28.4 for the PC cortex, and 25.7 for the striatum (**Fig. 1**).

Table 2. Centiloid cutoffs identified for each brain region

| Brain regions | Centiloid | Centiloid cutoff | | |
|----------------------------|-----------|------------------|--|--|
| | MRI | СТ | | |
| Global | 28.5 | 29.5 | | |
| Frontal cortex | 34.1 | 32.4 | | |
| Parietal cortex | 29.5 | 36.6 | | |
| Temporal cortex | 28.1 | 20.8 | | |
| Occipital cortex | 43.3 | 28.7 | | |
| Posterior cingulate cortex | 28.4 | 39.0 | | |
| Striatum | 25.7 | 38.7 | | |

 $\label{eq:magnetic resonance imaging, CT: computed tomography.} \\$



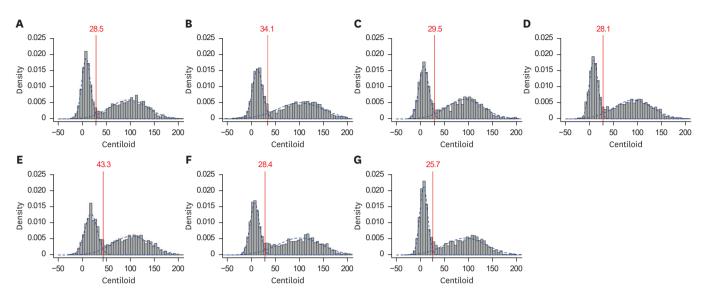


Fig. 1. Centiloid cutoff values derived from Gaussian Mixture Model using magnetic resonance method for each brain region: (A) Global, (B) Frontal, (C) Parietal, (D) Temporal, (E) Occipital, (F) Posterior cingulate, and (G) Striatum.

CT-based CL cutoffs by brain regions

Table 2 summarizes the cutoffs identified for each brain region. The GMM analysis revealed region-specific CL cutoffs, with cutoffs of 29.5 for the global brain, 32.4 for the frontal cortex, 36.6 for the parietal cortex, 20.8 for the temporal cortex, 28.7 for the occipital cortex, 39.0 for the PC cortex, and 38.7 for the striatum (**Fig. 2**).

Defining subgroups by global, regional, and striatal Aβ cutoffs

The cohort was classified into subgroups based on global, regional, and striatal A β cutoffs derived from MRI- and CT-based analyses. No significant differences were observed in A β positivity distributions between the MRI and CT-based methods across the subgroups (CU: χ^2 =1.57, p-value=0.67, MCI: χ^2 =0.40, p-value=0.40, DAT: χ^2 =0.24, p-value=0.97, and others:

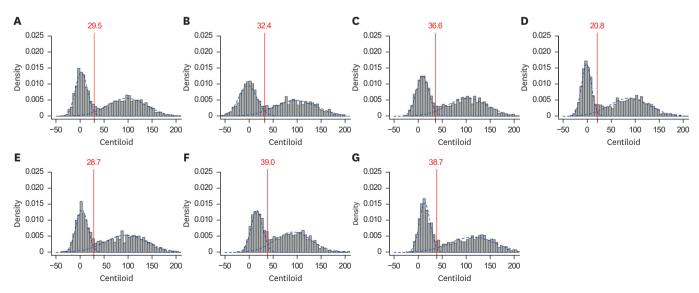


Fig. 2. Centiloid cutoff values derived from Gaussian Mixture Model using computed tomography method for each brain region: (A) Global, (B) Frontal, (C) Parietal, (D) Temporal, (E) Occipital, (F) Posterior cingulate, and (G) Striatum.



 χ^2 =2.65, p-value=0.45). The proportions of each subgroup were further analyzed across the clinical diagnoses of CU, MCI, and DAT. Using MRI-based cutoffs (Fig. 3A), most CU participants were classified as G(-)R(-) (66.8%), followed by G(+)Str(+) (20.5%), G(-)R(+) (7.4%), and G(+)Str(-) (5.3%). Similarly, in the MCI group, the largest proportion was classified as G(+)Str(+) (53.9%), followed by G(-)R(-) (36.2%), G(-)R(+) (7.1%), and G(+) Str(-) (2.8%). Among DAT participants, the majority were G(+)Str(+) (85.9%), with smaller proportions in G(-)R(-) (9.1%), G(+)Str(-) (2.2%), and G(-)R(+) (2.8%). In comparison, using CT-based cutoffs (Fig. 3B), the distribution was slightly shifted. This difference in distribution appears to be due to variations in spatial resolution and tissue contrast between MRI and CT. Among CU participants, G(-)R(-) remained the largest group (67.6%), followed by G(+)Str(+) (18.6%), G(-)R(+) (9.0%), and G(+)Str(-) (4.8%). MCI participants showed a similar trend, with most classified as G(+)Str(+) (53.2%), followed by G(-)R(-) (36.2%), G(-)R(+) (7.8%), and G(+)Str(-) (2.8%). In the DAT group, G(+)Str(+) again dominated (85.6%), with smaller proportions classified as G(-)R(-) (9.0%), G(-)R(+) (3.3%), and G(+)Str(-)(2.2%) (Table 3). The PC cortex had the highest frequency of positivity in both CT-based (56.7%) and MRI-based (58.1%) cutoffs.

Table 3. Distribution of subgroups by clinical diagnosis and imaging modality

| Variables | CU | MCI | DAT | Others |
|------------|------------|------------|------------|------------|
| MRI | | | | |
| G(-)R(-) | 388 (66.8) | 337 (36.2) | 59 (9.1) | 138 (50.9) |
| G(-)R(+) | 43 (7.4) | 66 (7.1) | 18 (2.8) | 22 (8.1) |
| G(+)Str(-) | 31 (5.3) | 26 (2.8) | 14 (2.2) | 10 (3.7) |
| G(+)Str(+) | 119 (20.5) | 502 (53.9) | 554 (85.9) | 101 (37.3) |
| CT | | | | |
| G(-)R(-) | 393 (67.6) | 337 (36.2) | 58 (9.0) | 126 (46.5) |
| G(-)R(+) | 52 (9.0) | 73 (7.8) | 21 (3.3) | 28 (10.3) |
| G(+)Str(-) | 28 (4.8) | 26 (2.8) | 14 (2.2) | 16 (5.9) |
| G(+)Str(+) | 108 (18.6) | 495 (53.2) | 552 (85.6) | 101 (37.3) |

Values are presented as number (%).

MRI: magnetic resonance imaging, CU: cognitively unimpaired, MCI: mild cognitive impairment, DAT: dementia of the Alzheimer's type. G: Global. Str. Striatal. R: Regional. CT: computed tomography.

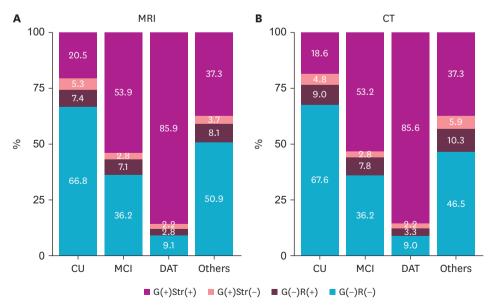


Fig. 3. Distribution of subgroups across clinical diagnoses based on (A) MRI- and (B) CT-derived amyloid-beta cutoffs. MRI: magnetic resonance imaging, CT: computed tomography, G: Global, Str: Striatal, R: Regional, CU: cognitively unimpaired, MCI: mild cognitive impairment, DAT: dementia of the Alzheimer's type.



DISCUSSION

In this study, we developed an algorithm to measure 3D T1 MRI-based and CT-based global and regional CLs using a large-sized dataset, enabling their broader applicability in clinical and research settings. This algorithm was used to calculate MRI-based and CT-based global and regional A β cutoffs through GMM, a statistical method. These cutoffs were then applied to classify participants into subgroups across cognitive stages (CU, MCI, and DAT) by global, regional, and striatal A β positivity, with similar proportions observed between MRI-based and CT-based methods. Taken together, our findings suggest that MRI-based and CT-based algorithms provide comparable classifications of A β uptakes across cognitive stages. These results highlight the potential to optimize resource allocation by reducing the need for additional 3D T1 MRI scans, making these algorithms highly practical for clinical and research applications.

Our study demonstrates that MRI-based and CT-based global Aβ cutoffs were generally similar, with the values closely aligning with previously reported MRI-based global Aβ cutoffs from other studies. ³⁵⁻³⁷ While existing studies vary in their specific cutoff values, a liberal cutoff is often set around 20 and a conservative cutoff around 40, with most studies reporting values within this range. ³⁸⁻⁴² Lower thresholds, typically around 10–20 CLs, are often used to detect early amyloid positivity, ^{38,41,42} while higher thresholds, closer to 40 CLs, are associated with advanced amyloid burden seen in clinical stages of AD. ^{39,40} These ranges are consistent with findings using common tracers such as FBB and FMM, highlighting the variability in cutoffs depending on the population characteristics and imaging methods employed. ⁴³ This consistency highlights the reliability of our findings and supports the broader applicability of these cutoffs in clinical and research contexts.

For regional CLs, our ability to calculate accurate cutoffs was made possible by leveraging a head-to-head dataset incorporating young controls, which was established in our previous studies. 19,44 These efforts led to the development of a commercialized product, BeauBrain Amylo, which enabled us to accurately derive regional CL values. This advantage allowed us to comprehensively compare MRI- and CT-based methods and establish robust regional cutoffs, revealing both similarities and differences between the two modalities. In some regions, such as the frontal cortex, the cutoffs were closely aligned between MRI and CT, whereas in others, like the occipital cortex and PC, notable differences were observed. These variations may reflect differences in the segmentation processes between MRI and CT algorithms, emphasizing the importance of modality-specific considerations in regional amyloid quantification.

These cutoffs were then applied to classify participants into subgroups across cognitive stages (CU, MCI, and DAT) by global, regional, and striatal A β positivity, with similar proportions observed between MRI-based and CT-based methods. The combined amyloid positivity rate [G(+)Str(+) and G(+)Str(-)] in our study was 25.8% and 23.4% for CU, 56.4% and 56.0% for MCI, and 88.1% and 87.8% for DAT in MRI-based and CT-based classifications, respectively. These rates closely align with previously reported amyloid positivity rates of approximately 20%–40% in CU individuals, 50%–60% in MCI, and over 80% in DAT, further validating the accuracy and reliability of our derived cutoffs. Interestingly, the G(-)R(+) subgroup was more frequently observed in non-dementia groups such as CU and MCI, while the G(+)Str(-) subgroup was more prevalent in CU compared to other cognitive stages. These patterns suggest potential transitional stages of amyloid



pathology, with G(-)R(+) possibly representing early amyloid accumulation detectable in regional brain areas, and G(+)Str(-) indicating global amyloid positivity without striatal involvement, often linked to advanced amyloid deposition. 45,46

Although these subgroups [G(-)R(+) and G(+)Str(-)] constitute a relatively small proportion of the cohort, their clinical significance warrants attention. Previous studies, including our own, have shown that individuals in the G(-)R(+) subgroup are more likely to progress to amyloid positivity over time, indicating its relevance as an early marker of amyloid pathology. 16,18,47,48 Furthermore, both G(-)R(+) and G(+)Str(-) groups have been associated with intermediate levels of structural brain changes and cognitive impairments compared to fully amyloidnegative [G(-)R(-)] or fully amyloid-positive [G(+)Str(+)] groups. 49 For example, G(-)R(+) individuals may exhibit subtle cortical thinning, while G(+)Str(-) individuals demonstrate MCIs, consistent with early clinical stages of AD. These findings underscore the need for longitudinal studies to better understand the trajectory and implications of these subgroups, particularly in the context of early intervention and disease prevention.

Our study's primary strength lies in its use of a large, head-to-head dataset incorporating MRI- and CT-based imaging, enabling robust comparisons and accurate determination of both global and regional A β cutoffs. Despite this strength, our study has several limitations. First, the sample size for certain subgroups, such as G(-)R(+) and G(+)Str(-), was relatively small, which may limit the generalizability of findings for these categories. Second, the CT-based segmentation algorithm may have inherent biases compared to MRI-based methods, potentially affecting regional cutoff calculations. Third, while longitudinal data were used to validate some findings, further longitudinal analyses are needed to confirm the clinical trajectories of the defined subgroups. Fourth, the use of a specific PET tracer (e.g., FBB or FMM) may limit the generalizability of our findings to other tracers or cohorts with differing imaging protocols. Nevertheless, our study provides valuable insights into the comparability of MRI- and CT-based A β quantification, highlights the clinical significance of transitional subgroups such as G(-)R(+) and G(+)Str(-), and establishes a framework for integrating multimodal imaging in amyloid research, making it a valuable contribution to the field.

In summary, our study established robust MRI- and CT-based global and regional A β cutoffs, demonstrating their comparability and clinical utility across cognitive stages. These findings highlight the importance of transitional subgroups such as G(-)R(+) and G(+)Str(-) in understanding amyloid pathology and its progression. By providing a framework for multimodal amyloid quantification, our work contributes to advancing precision diagnostics and personalized therapeutic strategies in AD research.

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