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Frequency of LATE neuropathologic change across the spectrum of Alzheimer’s disease neuropathology: combined data from 13 community-based or population-based autopsy cohorts

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Declarations

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Abstract

Limbic-predominant age-related TDP-43 encephalopathy neuropathologic change (LATE-NC) and Alzheimer's disease neuropathologic change (ADNC) are each associated with substantial cognitive impairment in aging populations. However, the prevalence of LATE-NC across the full range of ADNC remains uncertain. To address this knowledge gap, neuropathologic, genetic, and clinical data were compiled from 13 high-quality community- and population-based longitudinal studies. Participants were recruited from United States (8 cohorts, including one focusing on Japanese-American men), United Kingdom (2 cohorts), Brazil, Austria, and Finland. The total number of participants included was 6196, and the average age of death was 88.1 years. Not all data were available on each individual and there were differences between the cohorts in study designs and the amount of missing data. Among those with known cognitive status before death ($n = 5665$), 43.0% were cognitively normal, 14.9% had MCI, and 42.4% had dementia—broadly consistent with epidemiologic data in this age group. Approximately 99% of participants ($n = 6125$) had available CERAD neuritic amyloid plaque score data. In this subsample, 39.4% had autopsy-confirmed LATE-NC of any stage. Among brains with “frequent” neuritic amyloid plaques, 54.9% had comorbid LATE-NC, whereas in brains with no detected neuritic amyloid plaques, 27.0% had LATE-NC. Data on LATE-NC stages were available for 3803 participants, of which 25% had LATE-NC stage > 1 (associated with cognitive impairment). In the subset of individuals with Thal A β phase = 0 (lacking detectable A β plaques), the brains with LATE-NC had relatively more severe primary age-related tauopathy (PART). A total of 3267 participants had available clinical data relevant to frontotemporal dementia (FTD), and none were given the clinical diagnosis of definite FTD nor the pathological diagnosis of frontotemporal lobar degeneration with TDP-43 inclusions (FTLD-TDP). In the 10 cohorts with detailed neurocognitive assessments proximal to death, cognition tended to be worse with LATE-NC across the full spectrum of ADNC severity. This study provided a credible estimate of the current prevalence of LATE-NC

in advanced age. LATE-NC was seen in almost 40% of participants and often, but not always, coexisted with Alzheimer's disease neuropathology.

Keywords

ADRD; Tau; NFT; Nondemented; Oldest-old; Epidemiology; APOE; ROS-MAP; Vantaa 85 +; HAAS; CFAS; CC75C; The 90 + study; ACT; VITA; Nun study; Biobank for aging studies; Mayo clinic study of aging

Introduction

Brain autopsies of persons with documented amnesic dementia often reveal evidence of Alzheimer's disease neuropathologic change (ADNC) [69], limbic predominant age-related TDP-43 encephalopathy neuropathologic change (LATE-NC) [81], or both. However, the independent and joint prevalence of each of these disorders are unclear. There remain uncertainties about optimal classification of LATE-NC and some individual brains are challenging to categorize, as is the case for other subtypes of neurodegenerative disease [8, 29, 43, 54, 65, 82, 97]. Thus, high-quality data, derived from different geographic locations and including autopsy results, are required to shed light on the prevalence and co-existence of these high-morbidity brain pathologies.

The cardinal diagnostic feature of LATE-NC is TDP-43 pathology—aberrant TDP-43 protein deposits visualized with immunohistochemistry [81]. TDP-43 pathology was discovered in 2006 as the primary pathological hallmark of frontotemporal lobar degeneration with TDP-43 inclusions (FTLD-TDP) and amyotrophic lateral sclerosis [84]. However, TDP-43 pathology is now appreciated to occur in many other conditions [19]. Although diagnostic ambiguities still exist in TDP-43 neuropathologic assessments, LATE-NC has distinguishing characteristics including the neuroanatomical distribution of TDP-43 pathology, clinical features, genetic risk factors, and epidemiology [21, 39, 53, 81, 94]. For example, the demographic group most likely to show LATE-NC is persons beyond 85 years of age [81], and, LATE-NC is strongly associated with amnesic dementia, independent of other known brain pathologies [12, 32, 36, 39, 40, 44, 47, 51, 59, 70, 72, 79, 92].

Like LATE-NC, ADNC is prevalent and is associated with amnesic dementia. ADNC and LATE-NC are genetically pleiotropic: the *APOE* ϵ 4 ADNC risk allele is also associated with increased risk for LATE-NC [3, 28, 44, 118]. LATE-NC and ADNC are often present in the same brains [45, 46, 61, 63], and TDP-43 pathology may co-localize with tau-immunoreactive neurofibrillary tangles (NFTs), a hallmark ADNC lesion [44, 103, 111]. The presence of “mixed” pathologies is important because the clinical manifestations vary with different combinations of pathologies [62]. For example, “pure LATE-NC” is, on average, associated with a less severe clinical phenotype than “pure ADNC”, whereas the common combination (ADNC + LATE-NC) is associated with a more aggressive clinical course than either alone [48, 49, 74, 110, 119].

Despite recent progress, questions persist. Investigators have considered whether TDP-43 pathology in aging is best defined as a subtype of ADNC [43, 117]. While there is

heterogeneity in the genetic, pathologic, and clinical features of AD-type dementia [9, 41, 62, 71], there currently are no consensus-based criteria for delineating subtypes of ADNC. Basic related questions include: What is the overall end-of-life frequency of LATE-NC in the brains of older persons? How does the prevalence of LATE-NC vary in different research cohorts? How frequently is LATE-NC seen in brains with no-, low-, intermediate-, or high-severity ADNC, and in those with varying severities of primary age-related tauopathy (PART) [22]?

Addressing questions about the prevalence of different pathologies requires relatively population-representative autopsy cohorts. Dementia clinic- and hospital-based cohorts are invaluable resources for research, but they tend to be substantially enriched for unusual subtypes of dementia [99], early-onset diseases, and genetic risk factors, which limit the generalizability of the findings. While there have been prior reports about LATE-NC from individual research centers, and from various consortia [5, 57, 67], there has not been a prior study bringing together findings from a large number of community-based autopsy cohorts.

In the current study, summary data were gathered related to LATE-NC and ADNC from 13 separate well-established study cohorts with available autopsy information. These cohorts included participants who were mostly recruited without dementia and followed longitudinally to autopsy at research centers in United States (8 cohorts), United Kingdom (2 cohorts), Brazil, Austria, and Finland. Several of the included cohorts can be described as “population-based”, in that the individual donors were recruited from a general population within a geographical boundary in a study design that aimed to recruit from all subgroups within the population (See Supplemental Table 1, online resource). While the cohorts that are not population-based did not use probability-sampling and are not completely generalizable to their target populations, they are likely to be far more representative of the populations from which they were derived than clinic- or hospital-based cohorts. The combined data from multiple research cohorts provided the bases for gaining insights into how commonly LATE-NC is seen at autopsy, with or without comorbid ADNC.

Methods

The main goals of this study were to examine the frequency of LATE-NC at the end of life in community-based research participants and to stratify results by the level of reported ADNC severity. Based on those goals, summary data were requested related to ADNC and LATE-NC from 13 high-quality community-based and population-based cohorts of brain aging and dementia. (The term “community-based” is mostly used from here forward to refer to the present collection of cohorts.) Data were collected from each of the following autopsy cohorts (in alphabetical order): Adult Changes in Thought (ACT) [58]; Brazilian Biobank for Aging Studies (BAS) of the University of Sao Paulo [106]; Cambridge City over-75 s Cohort (CC75C) [16]; Medical Research Council Cognitive Function and Ageing Study (CFAS) [115]; Duke University/University of North Carolina AD Research Center (Duke/UNC-ADRC) [36]; Honolulu Asia-Aging Study (HAAS) [116]; Mayo Clinic Study of Aging (MCSA) [91]; Nun Study [112]; Rush University Religious Orders Study/Memory and Aging Project (ROS-MAP) [10]; University of California Irvine The 90 + Study (The 90 + Study) [50]; University of Kentucky AD Research Center (UKy-ADRC) [98]; Vantaa

85 + Study [52]; and, Vienna Trans-Danube Aging (VITA) study [55]. See Supplemental Table 1, online resource, for more information on each cohort. All study procedures were approved by the respective Institutional Review Boards or Research Ethics Boards. Each participant (or their next of kin if they lacked capacity) provided informed consent for cohort participation. No additional approvals were needed for analysis of the de-identified summary data from each cohort. Many of the research participants were recruited from the community using methods such as local media advertising, health fairs, and presentations to community groups.

The structured data requests sent to a representative of each cohort are shown in Supplemental Table 2, online resource. For the collection of data on ADNC, different pathology-based measures were requested: Braak NFT distribution staging (0–VI scale) [14] performed using anti-phospho-Tau antibodies; CERAD neuritic amyloid plaque density scores (graded as “None”, “Sparse plaques”, “Moderate plaques”, or “Frequent plaques”), which indicate the detected density of neuritic plaques in cerebral cortex [66]; and, Thal A β phases (a 0–5 scale based on anatomic distribution of A β plaques detected with A β immunostaining) [6, 108]. The rationale for incorporating these parameters was that they are all used for determining the presence and severity of ADNC according to current consensus-based criteria [69].

There were differences among the cohorts in the methods of tissue-processing at autopsy, neuropathologic evaluations, and data missingness. See Supplemental Table 3, online resource, for more information about how many participants were included from each cohort. Cohort-specific data format variations were conspicuous in the area of cognitive assessment instruments that were administered to participants. To operationalize global cognitive status, the cohorts used Mini-Mental State Examination (MMSE) [33] scores, except HAAS used the Cognitive Abilities Screening Instrument (CASI) [107], and both the Brazil BAS cohort and MCSA used the Clinical Dementia Rating sum of boxes scores [27]. For the UKy-ADRC, only participants who were recruited while cognitively normal were included and 11 subjects were excluded from the cognitive assessments due to no MMSE scores. For the BAS, participants 50 years or older at death were included and participants were excluded from this cohort with inconsistent clinical information, a post-mortem interval greater than 24 h, or if the brain tissue was incompatible for neuropathological analyses (e.g., cerebrospinal fluid pH < 6.5 or major acute brain lesions including hemorrhages). The Nun Study used MMSE cut points as follows: scores of < 17: dementia; 17–21: mild cognitive impairment (MCI); and, > 21 nondemented. For HAAS, the CASI scores were used at cutoffs ≥ 74 (normal), 60–73.9 (MCI), or < 60 (dementia). ROS/MAP data on clinical status were missing for 1 subject (0.05%). For The 90 + Study, 14 participants were excluded from the MMSE analyses due to missing scores. For the Duke/UNC-ADRC cohort, participants 90 years or over at death were included in the study. Approximately 70% from this cohort were cognitively normal at recruitment, and 29 participants were excluded from the cognitive assessment due to no MMSE score. For the Vantaa 85 + Study, DSM-III-R criteria were used to diagnose dementia and MMSE scores were assessed for most participants in the baseline study and follow-ups. For the MCSA, 37 participants did not have the Clinical Dementia Rating sum of boxes scores within 3 years of death.

Cohorts were also queried as to whether they had clinical evaluations during life and corroborating neuropathologic studies that likely would have captured cases of FTD/FTLD-TDP if they were in the cohort. The specific question posed to each autopsy cohort was: how many clear-cut FTD/FTLD-TDP cases were in the cohort? The symptoms of FTD include behavioral disturbances and language problems [53, 89, 104], but variants of these cognitive signs and symptoms (e.g., disinhibition and aphasia) may also occur in Alzheimer's disease and other dementia disorders, so there was some subjectivity in the clinical diagnosis.

To address whether multiple blinded neuropathologic raters from different institutions would agree with the results of Braak NFT staging, particularly in the context of cases with LATE-NC but lacking substantial ADNC, a multicenter digital pathology study was performed. Brain sections from 10 cognitively impaired individuals were included in this focused study, of which 8 had LATE-NC, 1 had FTLD-TDP, and 1 had severe ADNC. The following slides had been stained for phospho-Tau IHC (PHF-1 antibody [34]): hippocampus at the level of the lateral geniculate nucleus; anterior hippocampus and entorhinal cortex; occipital neocortex (Brodmann Area [BA] 17/18/19); superior and mid-temporal neocortex (BA 21/22); and, middle frontal gyrus (BA 9). Slides were anonymized and then converted to digital format using a Leica/Aperio ScanScope AT2 slide scanner at 40 × resolution. Four separate raters with experience in digital neuropathologic evaluation (coauthors M.D.C., J.D., B.N.D., and J.H.N.) scored the pathologies via internet connection, using either the Aperio ImageScope™ or QuPath open-source software, to derive Braak NFT stages for each case while blinded to other information.

For data analyses, the joint distribution of neuropathologic rating parameters were obtained from each cohort via templated spreadsheets (Supplemental Table 2, online resource). The overall joint distributions were simply summations of each cell in the joint distribution from each cohort. For demographic characteristics (average age at death and sex), a single summary measure was provided by each cohort. To compute the overall summary of age at death and sex distribution, as well as *APOE* ϵ 4 positivity, cohort-specific results were combined by weighting each cohort by its sample size. The association between *APOE* ϵ 4 positivity and LATE-NC rate was evaluated using simple meta-regression that ignored sample weights, did not include the VITA cohort (where *APOE* genotype data were unavailable), and did not factor in *APOE* genotype data missingness. For the comparisons of Braak NFT stages (PART severity [22]) in Thal A β phase = 0 cases (comparing the results with versus without LATE-NC), a Fisher's exact test was applied to determine statistical significance.

Results

Selected demographic, clinical, genetic, and summary neuropathologic data on included participants from each of the 13 community-based cohorts are shown in Table 1. The total number of included participants was 6196. Subset analyses were performed and the included numbers of subjects from each center for each analysis are provided in Supplemental Table 3, online resource. The median number of research participants included per cohort was 321, with a range of 109–1620 participants per cohort. Mean weighted age of death for all

included cohorts was 88.1 years; age ranges for the cohorts was 72.2–97.2 years. Overall, 62.3% of participants were women.

A chart depicting the clinical features of participants at their last clinical evaluation is shown in Fig. 1 ($n = 5665$ participants had those data available). Slightly over 40% were judged to be cognitively normal at their last clinical examination, and approximately the same proportion had documented dementia. In the 12 cohorts reporting the parameter, ~ 15% had MCI (See Supplemental Table 3, online resource).

In terms of FTD/FTLD cases, data were only considered from a cohort if FTD cases (clinically) and/or FTLD-TDP cases (pathologically) would likely have been documented in that cohort. Having applied those criteria, data were provided from 9 different cohorts, comprising $n = 3267$ participants. In this combined subsample, no clinical FTD/FTLD-TDP case was identified (Table 2). Although these participants were evaluated by clinicians, it is conceivable that early FTLD-TDP cases were present but not detected.

APOE $\epsilon 4$ allele genotype data were available from a total of $n = 5157$ included participants (83.2% of the combined cohort). *APOE* allele data missingness by cohort is indicated in Table 1. Of the participants with known *APOE* genotype, 25.5% carried at least one copy of the *APOE* $\epsilon 4$ allele (range: 13.0–33.6%). In the 12 cohorts with available *APOE* genotyping, there was a marginal positive association between *APOE* $\epsilon 4$ allele carrier prevalence and LATE-NC frequency ($r^2 = 0.36$; $p = 0.039$), indicating that cohorts with higher *APOE* $\epsilon 4$ prevalence also had higher LATE-NC frequency (Fig. 2a). By contrast, there was no such statistically significant association between LATE-NC frequency with cohorts' average age, sex (percent female), or percent of included subjects with neocortical Lewy body pathology (Fig. 2b–d).

LATE-NC is classified according to a 0–3 stage system, related to the anatomic distribution of TDP-43 pathology [81] and derived from studies that evaluated brains across a broad spectrum of pathologic severity [45, 73]. Cohort neuropathologists applied different antibodies to detect TDP-43 pathology; most cohorts used antibodies against phosphorylated TDP-43 protein (data not shown). Findings in the various subset analyses, stratified by the subsamples evaluated and the LATE-NC results, are depicted in Table 3.

The full spectrum of ADNC severity was represented in the sample. Among those with known CERAD neuritic plaque scores ($n = 6125$), 31.6% were classified as CERAD “None”, 17.6% “Sparse plaques”, 28.3% “Moderate plaques”, and 22.5% “Frequent plaques” (Table 4, Fig. 3). In participants with known Braak NFT stage ($n = 5985$), 31.5% were Braak NFT stages 0-II, 42.0% III/IV, and 26.5% V-VI (Table 4, Fig. 4). As such, approximately 1/4 of participants had severe ADNC.

In a subset of cases comprising $n = 3803$ participants, data were available including LATE-NC stages (0–3), Braak NFT stages (0–VI), and Thal A β phases (0–5) on each individual subject. The distribution of results stratifying by these parameters is shown in Table 5. Selected findings from those data are presented in chart format in Fig. 5.

Collectively, these data indicated that brains with more severe ADNC were relatively likely to have comorbid LATE-NC. For example, participants with Braak NFT stage 0-II had a 22.4% probability of LATE-NC being diagnosed, whereas those with Braak NFT stage VI had a 54.7% probability of a LATE-NC diagnosis (Table 4, Fig. 3). However, most participants with LATE-NC (61.2%) coincided with Braak NFT stages between 0 and IV (because only ~ 1/4 of participants had severe ADNC). Similar trends were observed for CERAD neuritic amyloid plaque densities (Table 4, Fig. 3), and Thal A β phases (Table 5). Although cohort-to-cohort variation was seen, there was broad agreement in findings, as can be appreciated by the 25th–75th percentile error bars in Figs. 3, 4.

Trends could be identified along the full ranges of ADNC and LATE-NC severities. Note that in the Table 5 data, LATE-NC stage 3 brains comprised only 11% of LATE-NC + cases (168 out of 1469), and LATE-NC stage 3 was associated with a high rate of severe ADNC—approximately the same frequency of severe ADNC as seen in LATE-NC stage 2. Furthermore, in brains lacking A β amyloid deposition (Thal A β phase = 0; $n = 787$), PART pathology was relatively more severe, i.e. higher Braak NFT stages, in persons with comorbid LATE-NC (Fig. 5).

While LATE-NC tended to be more frequent in more severe ADNC cases, LATE-NC was nonetheless present across all ADNC levels and even in those without ADNC. As shown in Table 3, 1935 participants had “None” neuritic amyloid plaques, and of these, 522 (27.0%) had LATE-NC. In the subset of individuals with known Thal A β phase = 0 (i.e. lacking A β plaques), 19.4% had LATE-NC, and 11.6% had LATE-NC Stages > 1, a severity of LATE-NC which has been consistently associated with cognitive impairment [18, 70, 73, 74, 78] (Table 5).

To assess how different neuropathologic raters would diagnose Braak NFT staging of LATE-NC cases that lacked severe ADNC, a convenience sample of phospho-Tau immunostained slides was evaluated by four separate blinded neuropathology diagnosticians, using digital pathology over the internet. As expected [4], there was some variance in Braak NFT staging by the raters, but the median rendered Braak NFT stages were within 1 Braak stage of the initial diagnosis in 8/10 cases and within 1.5 Braak stages in all 10 cases (see Supplemental Table 4, online resource).

Summary information on final cognitive status of included participants was requested from each cohort, with the data stratified by Braak NFT stages (bottom of Supplemental Table 2, online resource). These data were a focal-point because Braak NFT staging is the widely gathered ADNC parameter that correlates most robustly with cognitive impairment [80]. Detailed stratified cognitive testing results were not available from VITA, CC75C, and CFAS cohorts and thus were not included in the clinical-pathological analyses. Among the cohorts with accessible information, the cognitive status data were variable from cohort to cohort. There were different cognitive assessment instruments, different intervals of testing, and different workflows used in administering the tests. The nature of these combined summary data precluded statistical testing. However, a recurrent pattern did emerge across the different study groups, despite the many sources of variance and the smaller sample sizes when using data from single cohorts: there was a tendency for cognitive scores to be lower

in individuals with LATE-NC, across the full spectrum of ADNC severity in terms of Braak NFT stages (Fig. 6). Some of the implications and context of the present study are presented in Fig. 7.

Discussion

Data related to LATE-NC and ADNC were gathered, combined, and analyzed from 13 community-based and population-based longitudinal cohort studies. Overall, almost 40% of autopsied participants had LATE-NC. LATE-NC was relatively common in brains with severe ADNC—approximately half of severe ADNC cases had comorbid LATE-NC. By contrast, approximately one in four brains with no or minimal evidence of ADNC had LATE-NC. PART pathology was relatively more severe in persons with comorbid LATE-NC. There was a tendency for cognitive scores to be worse in persons with LATE-NC, across the full spectrum of ADNC severity. These findings address basic questions about LATE-NC in people with and without comorbid ADNC.

Both the quality and quantity of data were strengths of this study. The community- and population-based study designs of the contributory cohorts included many persons recruited while cognitively normal and followed longitudinally to autopsy. At the last exam before death, clinical features of the combined cohort showed slightly over 40% cognitive normal, and no FTD/FTLD examples were documented. This may underestimate the extent of cognitive impairment experienced, although most of the decedents were assessed in the last year of life. We emphasize that this distribution of clinical findings is in accord with epidemiologic data from human populations of this age group [21, 60, 86, 90]. While no study with autopsies examines all potential subjects, and none is perfectly representative of the variability in human populations across demographic and ethnographic boundaries, community- and population-based autopsy cohorts are the nearest approximation to a generalizable sample. Each cohort included here has provided the basis for published work related to LATE-NC [3, 32, 36–39, 51, 56, 77, 83, 88, 105]. Aggregating these data into a combined cohort comprising > 6000 people provided new insight into the prevalence of LATE-NC in aging, while also highlighting between-cohort variability.

One way to evaluate recruitment bias in a dementia study is to compare the frequency of *APOE* ϵ 4 allele among the reported participants with population-based figures. This is especially relevant because *APOE* ϵ 4 is associated with increased risk for LATE-NC [28, 93, 114]. In most human populations, approximately 25% of individuals carry at least one copy of the *APOE* ϵ 4 allele [20, 101] (the ϵ 4 prevalence tends to be somewhat higher in Scandinavia [30, 101]). It is notable that 25.5% of the genotyped participants in the current study had at least one *APOE* ϵ 4 allele. By contrast, in many dementia research cohorts the *APOE* ϵ 4 prevalence is higher [31]. For example, a recent report on LATE-NC derived from multiple clinic-based cohorts included 495 participants of which 47.4% were *APOE* ϵ 4 + (and 11.7% had FTD clinical syndrome) [49]. Many dementia studies have even higher *APOE* ϵ 4 positivity [23]. These studies may provide important insights (some impossible to achieve in community-based cohorts), but the distribution of pathologic findings in such clinic-based cohorts are unlikely to be representative of a broader population.

The current work has important limitations. Although the community-based cohorts encompassed thousands of research participants from five countries on three continents, human populations other than White Caucasians were underrepresented. Prior studies compared LATE-NC between ethnoracially defined groups [72, 77], but more work is required in this area [31, 85].

There were additional challenges in reconciling the LATE-NC data between cohorts. Neuropathologists used study-specific protocols, including non-identical tissue sampling and different antibodies. Some biologic variance is to be expected given the between-cohort differences in age, cognitive status, geography, and birth cohorts. These factors contribute to the wide variability of frequency of detected LATE-NC across the different included cohorts (range 11–63%). However, this inclusive approach, encompassing a range of diagnostic methods rather than one specific proscribed protocol, reflects the broad range of neuropathologic methods that are applied in everyday practice around the world, as well as true differences in frequency of neuropathologic lesions.

Another consideration is that TDP-43 pathology restricted to the amygdala was included to operationalize the presence of LATE-NC. There were undoubtedly LATE-NC false-negatives because the amygdala was not examined in some cases. LATE-NC stage 1 is hypothesized to be an incipient disease stage, analogous to early pathologic stages of AD and Lewy body diseases [76, 80]. As specific examples, Braak NFT stages I-III, Thal A β phases 1–2, and Braak Parkinson's disease stages 1–2 are all common in persons without documented neurological impairment [35, 42, 109]. Among the 3803 brains in the current study where all the LATE-NC stages were known, LATE-NC stage 1 comprised 36% of the LATE-NC cases and may correlate with limited, if any, cognitive manifestations [24, 73–75, 81]. However, the counterpoint is that 25% of the entire cohort had LATE-NC stage > 1, which is associated robustly with cognitive impairment [12, 32, 36, 39, 40, 44, 47, 51, 59, 70, 72, 79, 92].

Beyond the evaluation of LATE-NC, there are other challenges in reconciling neuropathologic data from different cohorts. The various studies had gathered brain donations over decades, and tissue handling methods have changed over time. One may expect imperfect agreement regarding low-Braak NFT stages as uniform staging requires standard sectioning and staining, and neuroanatomical expertise. (LATE-NC has been associated with NFT anatomical distribution that deviates from conventional Braak NFT staging [103].) Indeed, prior studies reported imperfect agreements in ADNC assessments among neuropathologists [4, 68]. This tendency was also evident in our digital pathological study with four separate raters evaluating the same cases using digital pathology over the internet.

An interpretation of the public health implications of this cross-sectional study should consider that the average age at death for included participants was 88.1 years. The frequency of autopsy-confirmed LATE-NC in this study (slightly under 40%), and other findings, does not represent projected population prevalence, but instead are a readout related to persons dying in that age range and agreeing to research brain donation. The study sample coincides with an age group at relatively high risk for LATE-NC [81]. (The role

of age as a factor in the relative frequencies of neurodegenerative disorders could not be examined thoroughly in the present study.) It may be argued that the included participants were unusually long-lived persons, considering normative data. For example, the average age of death in the United States during 2020 was 80.5 years for women, and 75.1 for men [2]—slightly older in European cohorts. Yet these averaged longevity calculations included many individuals who died at considerably younger ages. US Social Security Administration actuarial data predict that a woman who lives to age 70 years in the United States has a 32% chance to live until age 90 years, and a 70-year-old man a 21% chance to live until age 90 years [1]. Thus, a substantial proportion of adults will probably survive to the ages of participants included in the current study, with high risk for ADNC and LATE-NC.

This study reported summary information from each cohort rather than individual participant-level data, so regression models and other descriptive statistics were not appropriate for evaluating most of the data. In terms of clinical–pathological correlation, only broad trends were described, because robust statistical testing require a more standardized cognitive assessment format. There are many possible sources of data variability, e.g., additional pathologies, and testing variation between cohorts. Importantly, prior studies have established that LATE-NC is independently associated with cognitive impairment in aging when other factors (e.g., pathologic comorbidities) were considered [12, 36, 39, 70, 79, 92]. Thus, the main contribution of the current study is not clinical–pathological correlation, but instead it is a relatively sound estimate of LATE-NC prevalence in community- and population-based elderly autopsy cohorts across the ADNC severity spectrum.

LATE-NC was more common in brains with comorbid ADNC than in those without ADNC. Specifically, there was a 2- to 2.5-fold enrichment for LATE-NC in persons with severe ADNC versus those lacking ADNC. LATE-NC is not the only pathology that tends to be increased in parallel with ADNC. For example, Lewy body pathology subtypes and cerebrovascular pathologies such as arteriolosclerosis are also relatively prevalent in persons with ADNC [11, 17, 88, 95], as are white matter hyperintensities visualized with neuroimaging [7, 102], and other, rarer, phenomena [25, 64, 100]. The tendency for these brain conditions to coexist with ADNC may be due to shared ‘upstream’ risk factors such as the *APOE* ϵ 4 allele which is known to be pleiotropic for multiple diseases (see above), or other causes of brain injury. ‘Downstream’ of genetic and other risk factors, one subtype of pathology may directly promote other deleterious changes in the same cells. In particular, TDP-43 pathology often co-occurs with tau pathology in neurons vulnerable to NFT formation, such as in the entorhinal cortex [44, 111]. Conversely, tau inclusions coexist in cells prone to TDP-43 pathology, such as the hippocampal dentate granule neurons, in LATE-NC [103]. The increased severity of PART pathology in cases with LATE-NC in the present study further underscores the tendency for there to be pathologic synergies between tau and TDP-43 pathologies.

Although often comorbid, LATE-NC and ADNC were also seen in brains that lacked the other pathology. It is notable that ~ 75% of participants overall had some detectable ADNC, as shown previously [15, 109]. Thus, the generalization is true that “most people with LATE-NC have ADNC”, yet most old people’s brains *without* LATE-NC *also* have

ADNC. In this sample with a broad range of pathologies, > 60% of brains with LATE-NC lacked severe ADNC (i.e., had Braak NFT stages 0–IV). Among those with severe ADNC, approximately one-half lacked TDP-43 pathology. These data indicate that LATE-NC is not an integral feature of ADNC. Further support for the idea that LATE-NC and ADNC are distinct disorders come from prior published reports. For example, LATE-NC is an unusual co-pathology (< 10% prevalence) in severe ADNC linked to Down syndrome [113].

There was a substantial subgroup of participants with LATE-NC but with none or very mild ADNC: persons with Braak NFT stages 0–II had a 22.4% probability of LATE-NC whereas persons with “None” neuritic amyloid plaque score had a 26.9% probability of LATE-NC. Remarkably, among 3267 subjects surveyed for the condition, no FTD/FTLD case was identified. Thus, in community dwelling older persons with no or minimal evidence of ADNC, LATE-NC was still common and was not associated with a clinical diagnosis of FTD (in the nine cohorts in which that clinical evaluation was made). It is possible that a handful of FTD/FTLD cases was overlooked. Yet their extreme paucity in such a large combined cohort implies that FTD/FTLD-TDP is very uncommon in community-based cohorts. If the ~ 25% autopsy frequency is considered an estimate, albeit imprecise, of lifetime risk for LATE-NC in persons without ADNC, it can be contrasted with the epidemiologic studies that have found ~ 0.1% lifetime risk for FTLT-DTP [21, 53]. Thus, though there are important intersections between FTLT-DTP and LATE-NC, our results further support the conclusion that LATE-NC should be considered a separate entity from FTD/FTLD.

In summary, the current study found that LATE-NC was a frequent pathology in older brains: ~ 25% of participants overall had LATE-NC stage > 1, which is associated with cognitive impairment. LATE-NC was relatively common in brains with coexisting ADNC, and PART pathology was also relatively more severe in brains with comorbid LATE-NC. However, the presence of LATE-NC or ADNC was neither necessary nor sufficient to predict the presence of the other. Encompassing the full spectrum of ADNC severity, LATE-NC tended to be associated with cognitive impairment. These data are interpreted to indicate that LATE-NC, with or without comorbid ADNC, is highly prevalent and impactful in persons of advanced age.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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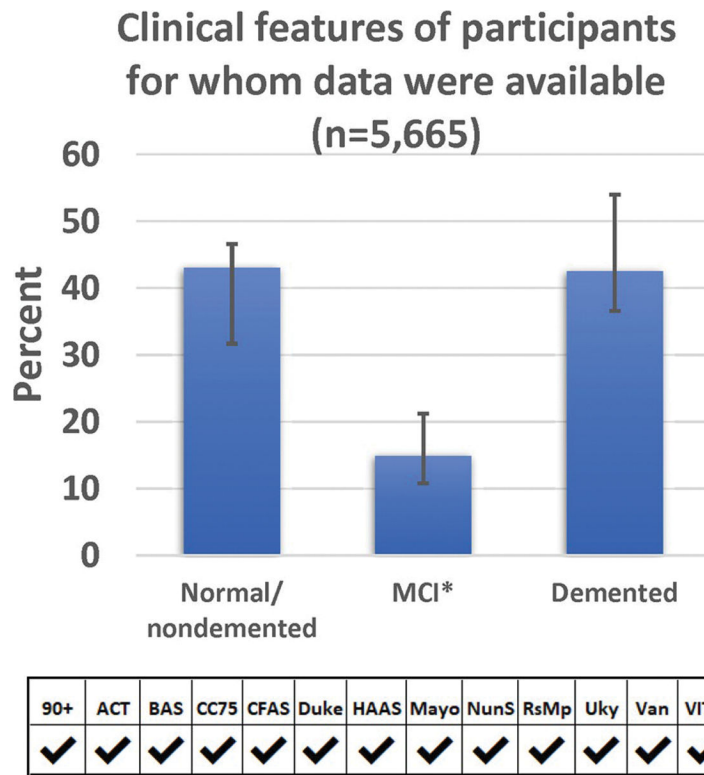


Fig. 1. Frequencies of clinical/cognitive features among the included participants. All cohorts had data about whether participants had normal cognition or dementia prior to death, and most (12 cohorts) had some measure for an intermediate clinical status, usually mild cognitive impairment (MCI). The finding of slightly over 40% cognitive normal prior to death is consistent with epidemiologic data of human populations in this age range [21, 60, 86, 90]. The result of each cohort was weighted equally in order to convey the cohort-to-cohort variance. For numbers of participants included from each cohort, see Table 1. Error bars denote 25th and 75th percentiles. *-MCI data were present for all cohorts except Vantaa 85 +

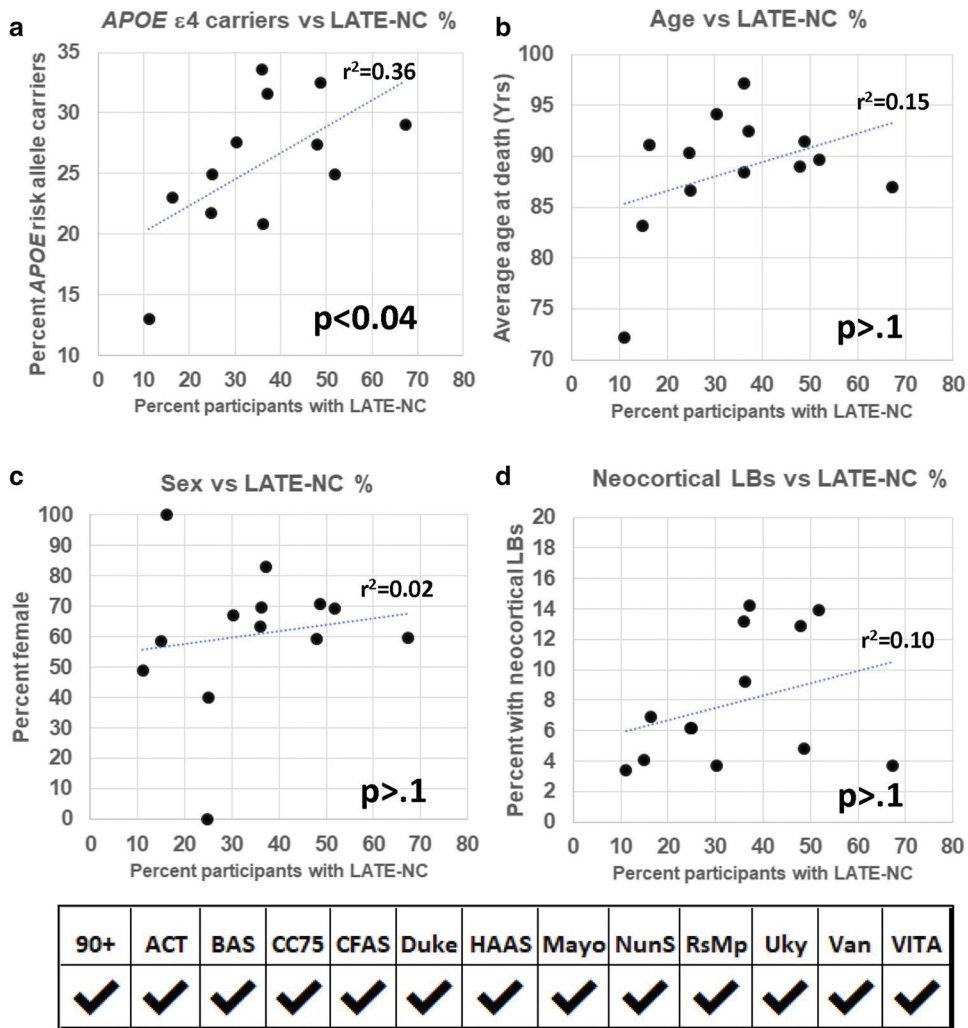


Fig. 2. The association between the percentage of included LATE-NC + participants in each cohort (x-axis) with percentages carrying the *APOE* ε4 allele (a), average age at death (b), sex (percent female (c), and, proportion with neocortical Lewy bodies (LBs), (d) on the y-axes. Each of the autopsy cohorts is indicated by a separate circular marker. The only association that was statistically significant in a simple regression analysis was *APOE* ε4 carrier frequency rate (a). *APOE* data were missing from a single cohort; see Table 1 for the numbers of research participants from each contributory cohort

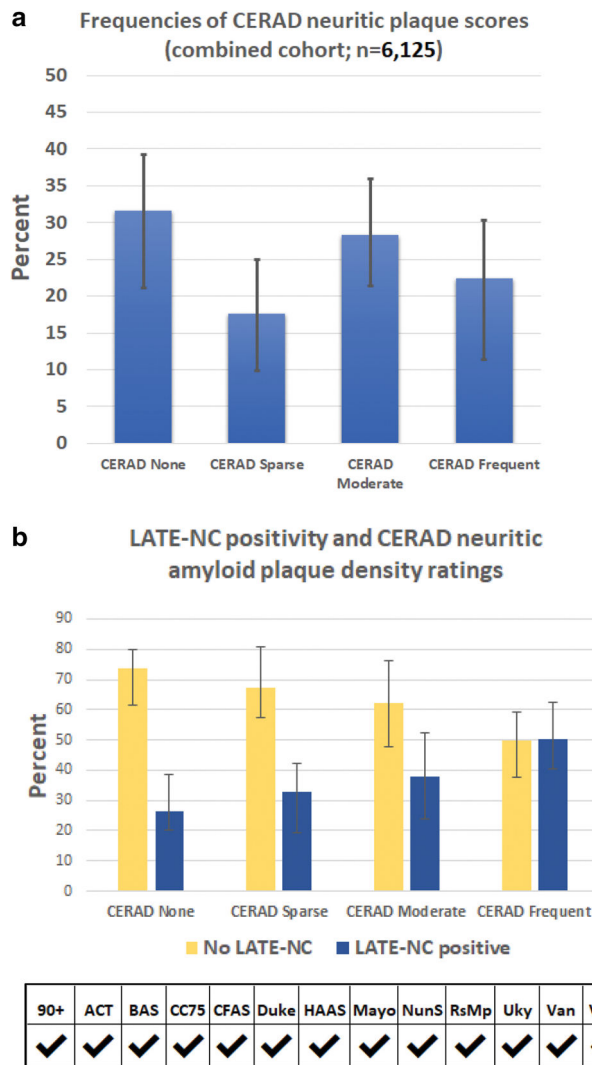


Fig. 3. LATE-NC absence or presence, stratified by CERAD neuritic amyloid plaques scores. All LATE-NC stages were combined and the results from each of the cohorts averaged. The frequency of LATE-NC increased with greater neuritic amyloid plaque densities. The distribution of CERAD plaques by frequencies is shown in (a). Note that subgroups with none or minimal ADNC were the most well represented in this combined meta-cohort (see Table 2). Correlation with LATE-NC status is shown in (b). Given the study design differences between cohorts, the results were generally consistent. For these charts, the results of each cohort were weighted equally in order to convey the cohort-to-cohort variance. For exact numbers of participants included from each cohort, see Supplemental Table 3, online resource. Error bars denote 25th and 75th percentiles

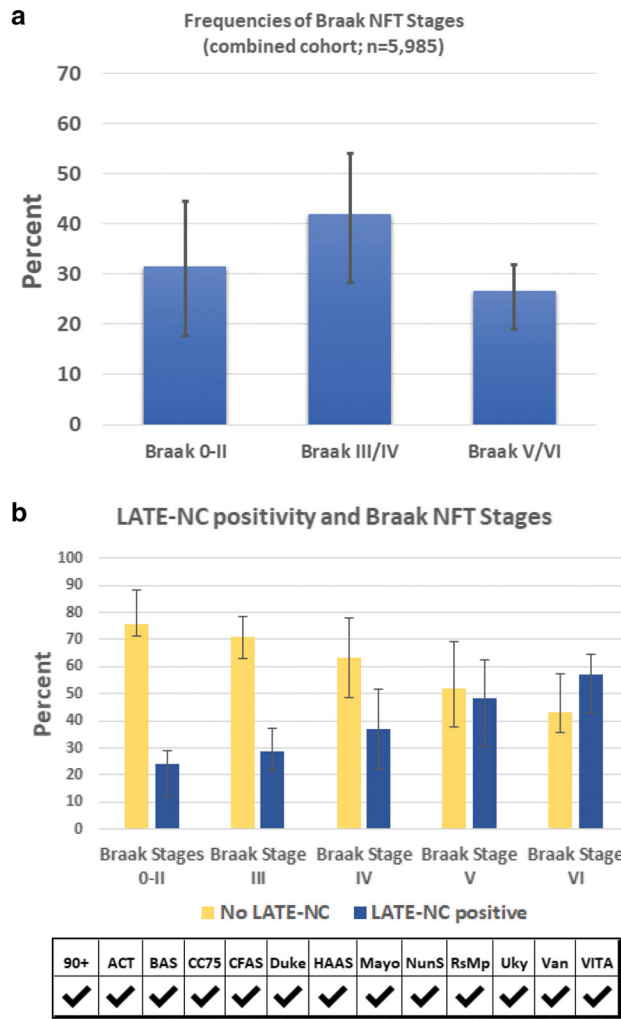


Fig. 4. LATE-NC absence or presence, stratified by Braak NFT stages. Here, all LATE-NC stages were combined and the results from each of the cohorts averaged. The distribution of Braak NFT stage groups by frequencies is shown in (a). Correlation with LATE-NC status is shown in (b). The frequency of LATE-NC increased with higher Braak NFT stages. Given the study design differences between cohorts, the results were generally consistent. For these charts the results of each cohort were weighted equally to convey the cohort-to-cohort variance. For exact numbers of participants included from each cohort, see Supplemental Table 3, online resource. Error bars denote 25th and 75th percentiles

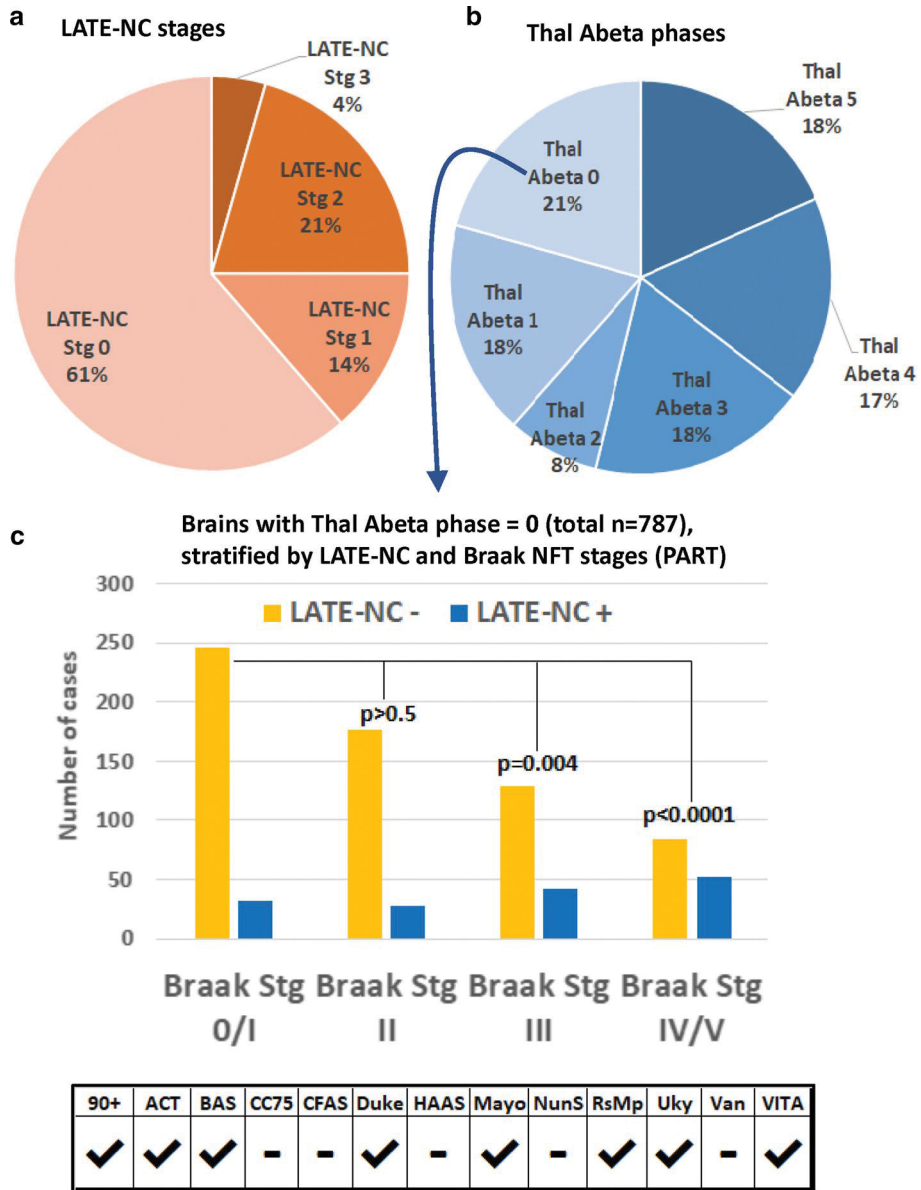


Fig. 5. Findings in the 3803 participants with available LATE-NC stage data (a), Thal Aβ phases (b), and Braak NFT staging, which indicate an association between LATE-NC and PART pathology. A pie chart (a) shows the relative frequencies of the different LATE-NC stages. Note that ~ 25% of participants have LATE-NC stage 2 (21% of participants) or stage 3 (4% of participants). A separate pie chart (b) depicts the relative frequencies of different Thal Aβ phases. The bar chart in panel (c) shows the number of cases with Thal Aβ phase = 0, stratified by Braak NFT stages. In these brains lacking Aβ amyloid pathology, the presence of LATE-NC was associated with higher Braak NFT stages (more severe PART pathology). For exact numbers, see Table 5, and for a breakdown of the numbers of participants included from each cohort, see Supplemental Table 3, online resource

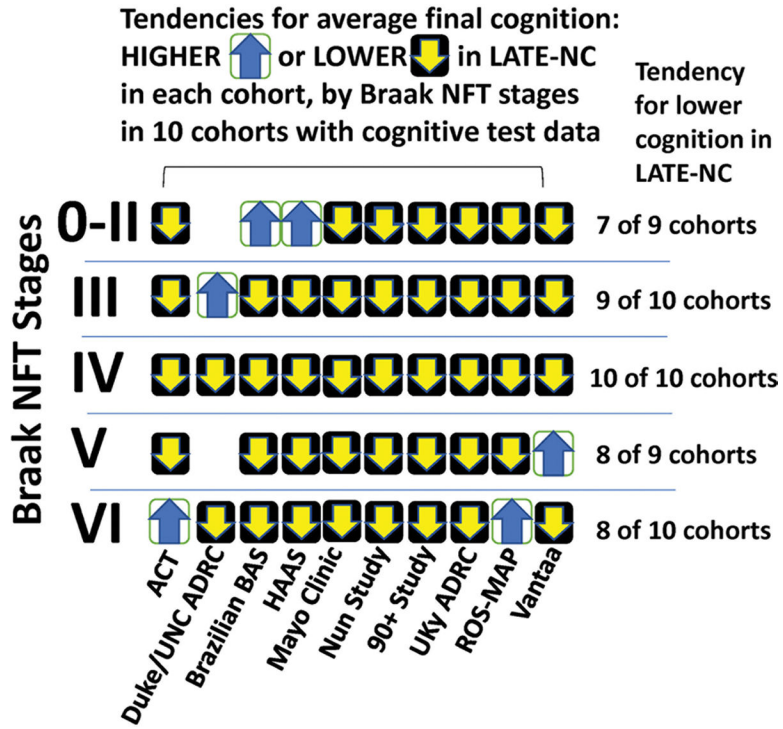


Fig. 6. There is a tendency for LATE-NC to be associated with cognitive impairment, across a broad range of Braak NFT stages, in ten community-based cohorts. Data were gathered on cognitive status, stratifying by LATE-NC status and Braak NFT stages. Trends were evaluated from each cohort as to whether the cognitive status tended to be lower in persons with LATE-NC (down-going black arrow) or higher (up-going white arrow) in given Braak NFT stages. To operationalize global cognitive status, final Mini-Mental State Examination scores [33] were used, except HAAS used the Cognitive Abilities Screening Instrument [107] and the Brazil BAS and MCSA cohorts used the Clinical Dementia Rating sum of boxes scores [27]. There was a tendency for participants with LATE-NC to have lower cognition across the full range of Braak NFT stages

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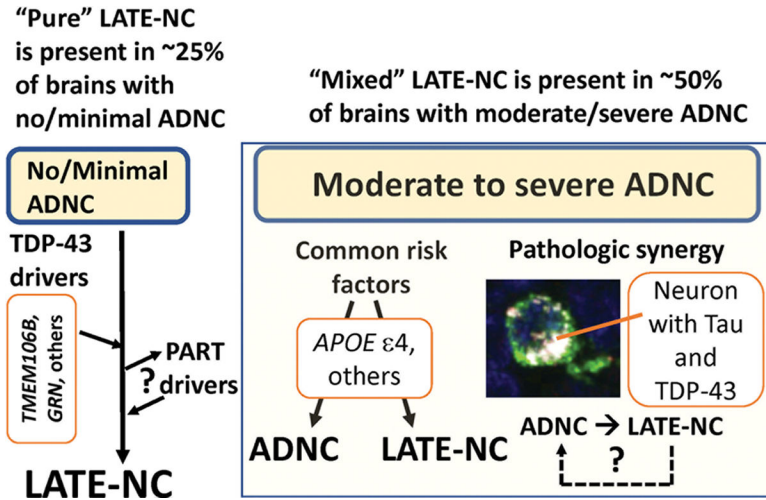


Fig. 7. Selected findings and context of the current study. Data were analyzed from participants in 13 high quality community- and population-based cohorts comprising over 6000 individuals followed longitudinally to autopsy. As such, the findings (with appropriate caveats) have broad implications. In participants that had none or minimal ADNC, a substantial proportion (~ 25%) had LATE-NC. This indicates that there are ADNC-independent TDP-43 pathology-driving mechanisms, which probably include gene variants in *TMEM106B* and *GRN*[26, 87, 96]. LATE-NC also was associated with more severe PART pathology (and vice versa), indicating pathologic synergy between LATE-NC and PART. Approximately 2/3rd of subjects in advanced age showed moderate or severe ADNC at brain autopsy, in concordance with the published literature [15]. In these individuals, there was a relatively high frequency of LATE-NC: approximately 50% of participants with moderate to severe ADNC had LATE-NC. The “mixed” ADNC-LATE-NC may be driven by pleiotropic genetic factors (e.g., *APOE* ε4 allele [114]) and there may also be pathologic synergies downstream from genetics. For example, intracellular tauopathy may promote TDP-43 pathology in the same cell [44, 103, 111]. The neuron shown here is stained with immunofluorescence in the hippocampal dentate gyrus, and is immunolabeled green (tau), and red (phospho-TDP-43) with overlap depicted in white [103]

Table 1
Selected demographic, genetic, clinical, and neuropathologic data on the 13 included cohorts*

Characteristics	ACT	Brazilian BAS	CC75C	CFAS	Duke/UNC C-ADRC	HAAS	Mayo/MCSA	Nun study	ROS-MAP	The 90 + study	UKY-ADRC	Vantaa 85 +	VITA	Total or weighted average**
Country	USA	Brazil	UK	UK	USA	USA	USA	USA	USA	USA	USA	Finland	Austria	
Sample size	863	625	228	510	109	321	209	382	1620	402	318	302	307	6196
Average age at death, years	89.0	72.2	91.5	87.0	94.1	90.3	86.6	91.1	89.9	97.2	88.4	92.4	83.2	88.1**
% Female	59.4	48.7	70.6	60.4	67.0	0	40.0	100	69.2	69.7	63.2	83.1	58.6	62.3%**
# Known <i>APOE</i> status	831	306	216	289	109	303	209	369	1554	371	318	282	0	5157
% <i>APOE</i> ε4 allele	27.4	13.0	32.5	29.0	27.6	21.8	24.9	23.0	24.9	20.8	33.6	31.6	N/A	25.5%**
Final cognitive status														
# clinical normal	308	461	63	189	29	139	142	167	529	114	142	107***	37	2427
# clinical MCI/proxy	17	44	39	10	26	68	44	48	372	113	49	N/A	11	841
# clinical dementia	381	101	113	275	49	97	22	136	718	174	110	195	26	2397
Neuropathologic features														
% neocortical LBs	12.9	3.4	4.8	3.7	3.9	6.2	6.2	6.9	14.2	9.2	13.2	14.2	4.1	9.5%**
% Braak stages 0-II	25.6	67.6	17.6	36.7	12.8	49.8	41.1	44.5	15.7	7.2	43.4	17.6	59.5	31.5%**
% Braak stages III-IV	36.3	23.9	60.8	41.5	67.0	31.2	41.6	23.6	56.0	54.0	26.7	47.2	20.3	42.0%**
% Braak stages V-VI	38.1	8.5	21.6	21.9	20.2	19.0	17.2	31.9	28.3	38.8	29.9	35.2	20.3	26.5%**
% LATE-NC****	47.9	11.1	48.7	67.7	30.3	24.7	24.9	16.2	52.2	36.1	36.0	37.1	16.9	39.4%**

ACT adult changes in thought [58], BAS Brazilian biobank for aging studies of the University of Sao Paulo [106], CC75C Cambridge city over-75 s cohort [16], CFAS medical research council cognitive function and ageing study [115], Duke/UNC-ADRC Duke/University of North Carolina AD Research Center [36], HAAS Honolulu Asia-aging study [116], MCSA Mayo Clinic study of aging [91], Nun Study [112], ROS-MAP Rush University Religious Orders Study/Memory and Aging Project [10], The 90 + Study [50], UKY-ADRC University of Kentucky AD Research Center [98], Vantaa 85 + Study [52], and VITA Vienna Trans-Danube Aging study [55], APOE apolipoprotein E, MCI mild cognitive impairment, LBs lewy bodies, LATE-NC limbic-predominant age-related TDP-43 encephalopathy neuropathologic changes

* See text for more details on data missingness and cohort-specific operationalizations

** weighted average

*** number of subjects with no dementia according to the DSMIIIR criteria

**** these percentages are for cases with full CERAD neuritic plaque data

see Supplemental Table 3, online resource

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Table 2

Number of cases with definite frontotemporal dementia (FTD) in the nine cohorts where this diagnosis was evaluated (among $n = 3267$ participants)

Cohort	Sample size	Number of definite clinical FTD cases identified	Notes
ACT	863	0	
CC75C	228	0	Ascertained by post-mortem clinical consensus
CFAS	510	0	Ascertained by post-mortem clinical consensus; 2 with "lobar atrophy"
Duke ADRC	109	0	
HAAS	321	0	
Mayo/MCSA	209	0	
The 90 + Study (UC Irvine)	402	0	2 "possible" bvFTD, 1 turned out to have AD, the other vascular pathology
Uky ADRC	318	0	
VITA	307	0	
Total number with clinical workup	3267		
Total number of definite clinical FTD cases identified		0	

Table 3

Overall percentage of participants with LATE-NC, stratified by the neuropathologic workups and in the subset of cases with low/no ADNC

	Participants with Braak NFT staging*	Participants with CERAD neuritic amyloid plaque scores*	Participants with Braak NFT stages and all LATE-NC stages**
Number of cohorts providing relevant data	13	13	8
Total number of individual participants	5985	6125	3803
Overall LATE-NC% in this group	38.4%	39.4%	38.3%
Criteria for low/no ADNC	Braak NFT stages = 0-II	CERAD score = "none"	Thal A β phase = 0
Number of participants with low/no ADNC	1883	1935	787
LATE-NC% in low/no ADNC group	22.4%	27.0%	19.4%

* See Table 4

** See Table 5

Joint distribution of LATE-NC positivity with CERAD neuritic amyloid plaque ratings [66] and Braak NFT stages [4, 13]

Table 4

		Total without LATE-NC	Total with LATE-NC	LATE-NC, %
CERAD neuritic amyloid plaque density scores (<i>n</i> = 6125)	None	1413	522	27.0
	Sparse	702	376	34.9
	Moderate	976	759	43.7
	Frequent	621	756	54.9
Braak NFT stages (<i>n</i> = 5985)	0-II	1461	422	22.4
	III	812	381	31.9
	IV	717	604	45.7
	V	492	643	56.7
	VI	205	248	54.7

Table 5

Numbers of participants with complete data on LATE-NC stages, Braak NFT stages, and Thal A β phases, stratified according to all three pathologic readouts ($n = 3803$)*

LATE-NC stage 0	Braak NFT stages						Total		
	0	I	II	III	IV	V		VI	
Thal A β phases	0	110	136	176	128	80	4	0	634
	1	18	76	119	130	101	7	1	452
	2	16	23	72	54	37	5	2	209
	3	7	34	62	130	119	55	8	415
	4	2	10	15	58	106	115	17	323
	5	0	4	10	23	58	138	68	301
									2334

LATE-NC stage 1	Braak NFT stages						Total		
	0	I	II	III	IV	V		VI	
Thal A β phases	0	4	9	15	22	10	2	0	62
	1	1	8	23	34	31	2	0	99
	2	1	2	8	10	7	0	1	29
	3	2	7	8	28	40	28	3	116
	4	0	1	0	9	48	32	7	97
	5	0	0	1	6	19	69	21	116
									519

LATE-NC stage 2	Braak NFT stages						Total		
	0	I	II	III	IV	V		VI	
Thal A β phases	0	3	12	9	16	32	3	0	75
	1	1	6	21	22	45	7	1	103
	2	0	0	5	22	21	3	1	52
	3	0	2	11	20	54	40	7	134
	4	0	0	2	10	53	112	11	188
	5	0	0	1	6	31	131	61	230

782

LATE-NC stage 3	Braak NFT stages						Total		
	0	1	II	III	IV	V		VI	
Thal A β phases	0	2	2	3	4	5	0	0	16
	1	2	7	5	5	2	0	0	21
	2	0	1	1	2	1	0	0	5
	3	0	2	1	6	15	14	1	39
	4	0	2	1	2	11	14	3	33
	5	0	0	2	2	6	32	12	54
							168		

* For the numbers of cases contributory from each cohort, see Supplemental Table 3, online resource

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