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The CADA-MRIT: an MRI Inventory Tool for evaluating cerebral lesions in CADASIL across cohorts

Ruiting Zhang, MD, PhD ^{1,2} Chih-Hao Chen, MD, PhD ^{3,4} Sophie Tezenas Du Montcel, MD, PhD ⁵ Jessica Lebenberg, PhD ^{1,6} Yu-Wen Cheng, MD, PhD ^{3,7} Martin Dichgans, MD, PhD ^{8,9,10} Sung-Chun Tang, MD, PhD ³ Hugues Chabriat, MD, PhD ^{1,6}

1. Paris-Cité University, Inserm U1141 NeuroDiderot, Paris, France
2. Department of Radiology, the Second Affiliated Hospital of Zhejiang University, School of Medicine, Hangzhou, China
3. Department of Neurology, National Taiwan University Hospital, Taipei, Taiwan
4. Department of Clinical Neurosciences, University of Calgary, Calgary, Alberta, Canada
5. Sorbonne Université, Paris Brain Institute, INSERM, INRIA, CNRS, APHP, Paris, France.
6. Lariboisière University Hospital, APHP, Translational Neurovascular Centre and Department of Neurology, Reference Center for Rare Vascular Diseases of the Central Nervous System and the Retina (CERVCO), FHU NeuroVasc, Paris, France
7. Department of Neurology, National Taiwan University Hospital Hsinchu Branch, Hsinchu, Taiwan.
8. Institute for Stroke and Dementia Research, University Hospital, Ludwig Maximilian University, Munich, Germany.
9. German Center for Neurodegenerative Diseases (DZNE), Munich, Germany.
10. Munich Cluster for Systems Neurology (SyNergy), Munich, Germany.

Correspondence: Hugues Chabriat Centre NeuroVasculaire Translationnel, Hôpital Lariboisière, 2 rue Ambroise Paré, 75010 Paris, France Tel: +33-1-49952593 Fax: +33-1-49952596 E-mail: hugues.chabriat@aphp.fr

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Abstract

Background and objectives: Cerebral Autosomal Dominant Arteriopathy with Subcortical Infarcts and leukoencephalopathy (CADASIL) is the most frequent genetic cerebrovascular disease. The clinical aspects of the disease in relation to the various types of lesions on MRI vary widely within families but also between different cohorts reported worldwide. Many limitations prevent comparison of imaging data obtained with different scanners and sequences in different patients cohorts. We aimed to develop and validate a simple tool to inventory quickly the key MRI features in CADASIL to compare imaging data across different populations.

Methods: The Inventory Tool (CADA-MRIT) was designed by consensus after repeated expert meetings. It consisted of eleven imaging items to assess periventricular, deep, and superficial white matter hyperintensity (WMH), lacunes, cerebral microbleeds (CMB), centrum semiovale and basal ganglia dilated perivascular spaces (dPVS), superficial and deep atrophy, large infarcts, and macrobleeds. The reliability, clinical relevance, and time-effectiveness of CADA-MRIT were assessed using data from three independent patient cohorts.

Results: Imaging data from 671 CADASIL patients (440 from France, 119 from Germany and 112 from Taiwan) were analyzed. Their mean age was 53.4 ± 12.2 years, 54.5% were women, 56.2% had stroke, and 31.1% had migraine with aura. Any lacune was present in at least 70% of individuals, whereas CMB occurred in 83% of patients from the Asian cohort and in only 35% of European patients. CADA-MRIT scores obtained for WMH, CMB and dPVS were comparable regardless of the scanner or sequence used (weighted $\kappa > 0.60$). Intrarater and interrater agreements were good to very good (weighted $\kappa > 0.60$). Global WMH and atrophy scores correlated strongly with accurate volumetric quantification of WMH or brain parenchymal fraction (Pearson $r > 0.60$). Different imaging scores were significantly associated with the main clinical manifestations of the disease. The time for evaluating one patient was around 2-3 minutes.

Discussion: The CADA-MRIT is an easy-to-use tool for analyzing and comparing the most frequent MRI lesions of CADASIL across different populations. This instrument is reliable. It can be used with different imaging sequences or scanners. It also provides clinically relevant scores in a very short time for completion.

Introduction

Cerebral Autosomal Dominant Arteriopathy with Subcortical Infarcts and leukoencephalopathy (CADASIL) is the most common inherited cause of stroke and vascular dementia.¹ It is a severe condition that progresses with age,² and its' clinical worsening varies widely between affected individuals.^{3,4} The origin of this variability is partly related to the location of the mutation in the NOTCH3 gene^{5,6} or to the presence of cerebrovascular risk factors^{7,8} but remains largely undetermined. The extent of characteristic cerebral lesions of the disease and their distribution are undoubtedly central to this variability^{3,9}. Thus, understanding the variability of MRI signal changes and their determinants at a large scale within, but also between large CADASIL cohorts is crucial to improve prediction of clinical course and develop future clinical trials.

Regarding brain imaging to date, comparative studies between cohorts remain rare and are only based on historical data from the literature.^{10,11} This is mainly because quantitative imaging studies are extremely difficult to perform for rare diseases, particularly in a routine clinical setting and when MRI data have been collected over many years. Also, some pitfalls encountered in the automatic segmentation of MRI lesions in CADASIL have not yet been fully resolved, such as the difficulty of separating certain white matter lesions of variable signal in close contact with the cerebral cortex¹² or the isolation of lacunes from the frequent dilated perivascular spaces (dPVS). Moreover, while the segmentation of white matter hyperintensities (WMH) has benefited from extensive research efforts supported by medical image computing societies,¹³ the automatic segmentation of lacunes, microbleeds or PVS are still being studied and are far from being considered valid methods to evaluate heterogeneous and variable imaging findings as observed in CADASIL patients.¹⁴ Elsewhere, for analyzing imaging data from multiple cohorts, there are many other limitations to be overcome such as: 1) differences in imaging sequences or protocols across centers,¹⁵⁻¹⁷ 2) modifications related to advances in imaging techniques or acquisitions over a long period of recruitment (1.5-Tesla MRI replaced by 3-Tesla or T2* replaced by susceptibility weighted imaging (SWI)), 3) inconsistency between quantitative methods based on multiple algorithms or approaches,^{18,19} 4) difficulties related to the cost for sharing original imaging data between centers in a reliable, secure and anonymous way for a long time and within an ethical framework.

An alternative approach, especially for comparative cohort studies, could be to adopt already validated visual rating scales from the literature. Different tools were previously designed for rating MRI lesions related to sporadic cerebral small vessel disease (CSVD) in stroke patients or in the general population.²⁰⁻²³ However, most of these instruments have been validated to specifically assess one category of lesions such as the Fazekas scale for WMH, the Global Cortex Atrophy (GCA) scale for brain atrophy, or the Microbleed Anatomical Rating Scale for assessing the location and number of microbleeds.²⁴⁻²⁶ Some of these scales were validated based on specific imaging protocols.^{21,23} For other scales, such as the Fazekas scale whose maximal score is reached in more than half of CADASIL patients, a ceiling effect was observed due to the early emergence of extensive WMH.^{19,20,27} Finally, the assessment of specific CADASIL imaging features such as WMH in the anterior temporal lobes has never been considered.

In the present study, we aimed to develop and validate a practical tool that could quickly inventory the key MRI features observed in CADASIL and could be used easily by clinicians and researchers to compare their data across multiple patient populations even when studied with different MRI scanners and sequences.

METHODS

DATA COLLECTION METHODS

CADASIL patients from three independent cohorts were included in the present study. Details of these cohorts have been already reported.^{9,28,29} Briefly, consecutive CADASIL patients who were at least 18 years of age, were evaluated at Lariboisière hospital (Cohort 1), Ludwig-Maximilians-Universität hospital (Cohort 2), and National Taiwan University Hospital (Cohort 3) between October 2003 and February 2021 (Cohort 1), April 2003 and September 2010 (Cohort 2), January 2019 and August 2022 (Cohort 3). In all cases, the diagnosis was confirmed by genetic testing showing a typical NOTCH3 mutation altering the number of cysteine residues. Clinical and demographic data were collected by study investigators at the time of inclusion. All enrolled subjects underwent detailed baseline neurological examination, including a brief evaluation of cognitive deficits with the Mini-Mental State Examination (MMSE) and degree of disability based on the modified Rankin scale (mRS). As in the previous study, we defined disability as mRS ≥ 3 .⁹ MRI examination was done on the same day of baseline neurological examination in Cohort 1 and 2, and within 3 months after inclusion in Cohort 3.

The MRI acquisition parameters and the quantification methods of imaging data are detailed in the eMethods section.

STANDARD PROTOCOL APPROVALS, REGISTRATIONS, AND PATIENT CONSENTS

Informed consent was obtained from each subject or a close relative if the subject was too severely disabled to give written consent. This study was approved by independent ethics committees (Cohort 1: updated agreement CEEI-IRB-17/388; Cohort 2: the LMU medical faculty, No. 299/03; Cohort 3: NTUH: No. 201807044RIND) in all participating centers and conducted in accordance with the Declaration of Helsinki.

DESIGN OF THE INVENTORY TOOL

The different items composing the inventory tool were prepared after 4 working zoom meetings of 2-3 hours between the authors. The selection of the different items was established by consensus after repeated discussion and tests so that the final scale would be: 1) easy to use, 2) quick to assess, 3) usable on the MRI sequences most commonly used in clinical practice for CADASIL patients, 4) usable to summarize all lesions features currently observed in the disease (for example, WMH in the temporal lobes), 5) able to capture the differences between cohorts (frequent hemorrhagic lesions in Asian and ischemic lesions in European patients), 5) sensitive to the extension and number of lesions from the beginning to the late stage of the disease, 6) simple and as conservative as possible by adapting items already proposed in previous validated scales.

The final inventory included eleven items for evaluating the different lesions using the STRIVE criteria: 1) periventricular WMH (PVH), 2) deep WMH, 3) superficial WMH, 4) lacunes, 5) microbleeds (CMB), 6-7) dPVS in the Centrum Semi-Ovale (CSO) or in basal ganglia (BG), 8) superficial atrophy, 9) deep atrophy, 10) large infarcts and 11) macrobleeds (Figure 1). The levels of each item were chosen according to the literature, the distribution of data in different subsamples including extreme cases observed in the different cohorts, and the long clinical experience of some authors (HC, SCT). The final scoring adopted by consensus is illustrated in Figure 2. The method used for building the scoring system is detailed in eMethods.

ASSESSMENT AND VALIDATION OF THE INVENTORY TOOL

To validate the inventory tool, we look for 1) the agreement between the scoring obtained from different MR protocols, 2) the inter and intra-rater agreement of the composing scores, and 3) the external validity via the correlation between some items and quantitative lesions measures. The clinical correlates of items derived from the inventory tool and its time-effectiveness were also assessed. This validation procedure was based on different results obtained by 4 experienced raters [a neuroradiologist – RZ (rater 1) and 3 neurologists - CHC (rater 2), HC (rater 3); YWC (rater 4)] with 6 to 35 years of experience in clinical and imaging research on CSVD. Analysis was performed using fully anonymized MR images from the different cohorts and blinded to the clinical status of all individuals.

1) Agreement between scoring obtained from different MR acquisition protocols

The scores obtained from the inventory tool were compared when images were from a basic or an advanced MR imaging protocol. More specifically, the agreement between WMH scores obtained with 3D or 2D Fluid-attenuated inversion recovery (FLAIR) images, CMB scores obtained with T2* or SWI, or dPVS scores obtained with T1 or T2-weighted images, was analyzed. For WMH, 40 randomly selected 3D FLAIR images from Cohort 1 were re-sampled to 2D FLAIR images, and then the original 3D images and the 2D images re-sampled from the 3D images were evaluated by rater 1 during two separate sessions at 4 weeks interval. Similarly, for CMB, data from 83 patients from Cohort 1 who had both SWI and T2* images were also analyzed by rater 1 in two separate 4-week sessions. For dPVS, 40 patients having had both T1 and T2-weighted images from Cohort 3 were randomly selected and analyzed by rater 2 in two separate additional sessions. The agreement between the different imaging protocols was analyzed for each inventory item using the linear weighted κ measure of agreement. The weighted κ results were interpreted as poor (0–0.20), fair (0.21–0.40), moderate (0.41–0.60), good (0.61–0.80), or very good (0.81–1) agreement.³⁰

2) Intrarater and interrater agreements for the different inventory items

Forty patients with a basic MRI protocol (2D FLAIR, low-resolution 3D T1, T2*) and 40 others with an advanced MRI protocol (3D FLAIR, high-resolution 3D T1, SWI) were randomly selected from Cohort 1 among 440 patients by age of distribution in 10-year intervals. Images were then displayed using the ITK-SNAP 3.0 software (Cognitica, Philadelphia, PA, USA) and assessed both by rater 1 and rater 3. Rater 1 also analyzed the same data a second time after a 4-week interval. Each rater was blinded to the other rater's ratings. Before these independent rating sessions,

the two raters had together training sessions on 10 randomly selected patients, and cases of disagreement were solved by discussion. Appropriate guidance for using the inventory tool was prepared based on this experience.

For additional validation, 40 patients were also randomly selected from Cohort 3 and assessed by rater 2 and rater 4 separately. Similarly, 40 patients were randomly selected from Cohort 2 and assessed twice by rater 1 for intra-rater agreement. We also built a mixed database with 14 patients from Cohort 1, 13 patients from Cohort 2, and 13 patients from Cohort 3, who were randomly selected from the three cohorts. The mixed imaging database was then assessed by rater 1 and rater 2 for interrater agreement.

Intra-rater and interrater agreements were calculated using the linear weighted κ measure of agreement.

3) Correlations between the items from the inventory tool and quantification of lesions

We studied the correlations between the different WMH items. We also looked at the relationship between the total WMH score obtained as the sum of the 3 different WMH items and the quantitative measure of WMH calculated in the whole brain. Similarly, we studied the correlation between the different atrophy items. The sum of the two atrophy items was also compared to the brain parenchymal fraction. For this purpose, first we transformed the data to follow a normal distribution using cube (normalized WMH fraction) or quantile transformation (BPF). Then, the Pearson correlation between the inventory tool items and the respective quantification lesion was computed. We also used scatterplots to show the correlation between these scores and their respective quantitative measurements.

4) Association between the inventory tool items and the main clinical features of the disease

To address the clinical value of the different items derived from the inventory tool, we combined the three cohorts ($n = 671$ individuals) and assessed their association with the past occurrence of stroke (ischemic and hemorrhagic), attacks of migraine with aura, degree of disability ($mRS < \text{or} \geq 3$) and MMSE score. Multivariable logistic (stroke, migraine with aura, disability) or linear model (MMSE), adjusted for age and sex, were used. Additional adjustment for the level of education was considered for the association with MMSE. To test the association between clinical variables and each of the 11 inventory tool items, statistical significance was set at a probability value of < 0.0045 ($0.05/11$, Bonferroni adjustment). To search for an independent association between the inventory tool scores and the different clinical features, multivariate regression analysis was performed with statistical significance set at a probability value of < 0.05 (Model 2: all imaging items included simultaneously in the model without selection). Finally, the models obtained after selecting the significant items from the CADA-MRIT for predicting each clinical outcome were compared to the predictive models obtained using a global score as the total CSVD score.³¹ All analyses were done by R version 3.6.2 or SPSS 22.0.

5) Time effectiveness

The time requested for evaluating MR images using the inventory tool was recorded using a stopwatch in 80 patients from Cohort 1 and 40 patients from Cohort 3 (the same patients selected for intra- and interrater analysis). Rater 1 and 2 recorded the time spent to fulfill each item of the inventory, while rater 3 recorded the time spent to complete the entire inventory.

DATA AVAILABILITY

The data that support the findings of this study are available from the corresponding author and co-authors (HC, MD, SCT) upon reasonable request.

Results

1) Main characteristics of patients from the 3 cohorts

The main demographic and clinical characteristics of the 671 patients from the three CADASIL cohorts are presented (n = 440 from Cohort 1, n = 119 from Cohort 2, and n = 112 from Cohort 3) in Table 1. The mean age of the whole population was 53.4 ± 12.2 years, 366 (54.5%) individuals were female, 377 (56.2%) had stroke, and 209 (31.1%) patients had a history of migraine with aura. Patients of Cohort 3 (from Taiwan) included the oldest patients. In contrast with the other cohorts, migraine with aura was observed only in one individual in this cohort. The patients' imaging characteristics are detailed in Table 2. Any lacune was present in at least 71% (Cohort 1) to 81% (Cohort 3) of patients. CMB were present in 83% of patients in Cohort 3 but in only 33-36% in cohorts 1 and 2.

2) Agreements between scoring obtained from different MR acquisition protocols

Agreements between the different WMH scores obtained using the original 3D images and the 2D images re-sampled from the 3D images were very good [PVH: weighted $\kappa = 0.939$ (0.859-1.000); deep WMH: weighted $\kappa = 0.828$ (0.650-0.961); superficial WMH: weighted $\kappa = 0.839$ (0.727-0.926)]. Agreement between CMB scores obtained with SWI and T2* images was very good [weighted $\kappa = 0.848$ (0.751-0.932)]. Agreement between dPVS scores obtained using T1 and T2 were also good to very good [CSO dPVS: weighted $\kappa = 0.811$ (0.652-0.941); BG dPVS: weighted $\kappa = 0.631$ (0.399, 0.824)].

3) Intrater and interrater agreements for the different inventory items

As shown in eTable 1 and 2, 80 out of 88 intrater and interrater agreements for the different inventory tool items were good or very good (weighted κ between 0.647 and 1.000). The eight fair to moderate agreements included the interrater agreements for deep WMH from cohort 1 with advanced MRI protocol (weighted $\kappa = 0.516$), superficial atrophy from cohort 3 (weighted $\kappa = 0.556$), and the dPVS items especially when they were obtained using the basic MRI protocol based on low-resolution images (Interrater: weighted κ between 0.211 and 0.256; intrater: 0.388 to 0.589).

4) Correlations between the items from the inventory tool and quantifications of lesions

PVH, deep WMH and superficial WMH items were correlated with each other (PVH and deep WMH: Pearson $r = 0.645$, $p < 0.001$; PVH and superficial WMH: Pearson $r = 0.399$, $p < 0.001$; deep WMH and superficial WMH: Pearson $r = 0.425$, $p < 0.001$). The correlation between the WMH total score, calculated by summing the previous scores, and the cube root of WMH fraction was high in all three cohorts (Cohort 1: Pearson $r = 0.792$, $p < 0.001$; Cohort 2: Pearson $r = 0.897$, $p < 0.001$; Cohort 3: Pearson $r = 0.877$, $p < 0.001$; All patients: Pearson $r = 0.833$, $p < 0.001$) (Figure 3A).

Superficial atrophy and deep atrophy items were correlated with each other (Pearson $r = 0.712$, $p < 0.001$). The atrophy total score was correlated with the normal quantile transformed BPF in all three cohorts (Cohort 1: Pearson $r =$

-0.753, $p < 0.001$; Cohort 2: Pearson $r = -0.617$, $p < 0.001$; Cohort 3: Pearson $r = -0.572$, $p < 0.001$; All patients: Pearson $r = -0.707$, $p < 0.001$) (Figure 3B).

5) Association between the inventory tool scores and the main clinical features of the disease

The results of the association analysis between the different inventory tool items and the clinical variables are presented in Table 3. In the first model, stroke was positively associated with the WMH scores, lacunes, and CMB scores, but negatively associated with the CSO dPVS score. Migraine with aura was positively associated with the superficial WMH score but negatively associated with the CMB and superficial atrophy scores. Disability was positively associated with the PVH, deep WMH, lacune, CMB, and atrophy scores, but negatively associated with the CSO dPVS score. Lower MMSE was positively associated with the PVH, lacune, CMB, and atrophy scores, but negatively associated with the CSO dPVS score.

When all items were considered together (Model 2) for analysis, the results showed that scores of lacunes and macrobleeds were positively associated with stroke. CMB and superficial atrophy scores were negatively associated with migraine with aura, while deep atrophy score was positively associated with migraine with aura. PVH, lacune, superficial atrophy scores, large infarct, and macrobleeds were positively associated with disability. Higher PVH, deep WMH, atrophy and macrobleeds scores were associated with lower MMSE.

Finally, the models based on a combination of items derived from the inventory tool (Table 3) largely outperform those obtained using a global score as the CSVD total score for predicting the different clinical outcomes. Not only the R^2 was found always higher using items from CADA-MRIT but also two major clinical outcomes, migraine with aura and the MMSE score, were associated with scores derived from our tool but not with the total CSVD score (eTable 3).

6) Time effectiveness

As shown in eTable 4, the mean time for using the inventory tool for assessing MRI data from a CADASIL patient was around 2-3 minutes. The mean time needed for evaluating images obtained using an advanced MRI protocol including 3D MRI data was 50 seconds longer than with images obtained using the basic MRI protocol. The most time-consuming items were the lacunes and the dPVS.

Discussion

In this study, we have developed and evaluated an easy-to-use imaging inventory tool, for analyzing and comparing the most frequent MRI lesions observed in CADASIL across different cohorts or populations. This tool was developed based on data from three independent cohorts originating from Europe and Asia. The results showed that this instrument was not only reliable, but could also be used with different imaging sequences, and provide scores that were clinically relevant, all this in a very short time for completion.

This tool was designed to obtain a multifaceted assessment of cerebral MRI lesions in CADASIL, not to quantify a single imaging marker as proposed through different visual rating scales available in the literature. The inventory tool allows for evaluating the extent of the most characteristic imaging features observed in CADASIL patients, i.e. WMH, lacunes, microbleeds but also cerebral atrophy and the presence of large infarcts or large hemorrhages. All these markers have been repeatedly shown to vary considerably according to the progression of the disease and to be differently related to the clinical severity.^{4, 18} The results showed that its use allowed to detect easily the high frequency of hemorrhagic lesions and rarity of temporal damage in the Taiwanese cohort (Cohort 3) in contrast to the two other cohorts.³² (Figure 4) This is exactly one of the possible uses of such a tool which might be also employed for evaluating in future the influence of vascular risk factors, mutation locations of the NOTCH3 gene, or even additional genetic variants on the multiple imaging manifestations of the disease.

The results we obtained from different angles showed that this inventory tool was highly reliable. First, the items derived from the inventory were strongly correlated when images were acquired using different MR imaging sequences. These results are important since they suggest that the levels of items taken from this inventory in different centers and cohorts could be actually compared. Second, we observed that these imaging items were strongly related to the quantitative measures of lesions obtained using various algorithms and multiple data preparation steps. This demonstration was obtained after summing the different items of hyperintensities in the periventricular, deep, and superficial white matter that we compared to the global volumetric measures of lesions. Identical results were obtained after summing the superficial and deep atrophy items that we compared to the BPF. These results also suggest that such global scores of WMH or atrophy derived from the inventory tool could also be considered for estimating the total amount of white matter lesions or degree of cerebral atrophy across CADASIL patient groups. Elsewhere, because the rating of white matter lesions was developed from items originating from the long-validated Fazekas scale, the two subscores of the latter could be easily retrieved by only changing the PVH level 4 in level 3 in the inventory tool. Third, the different items derived from the inventory were also highly reproducible. For most of them, a high inter and intra-rater agreement was reached which supports that the inventory tool could be used by multiple observers and repeatedly when needed. These results were however obtained with MRI read by experienced neurologists or neuroradiologists and would need further evaluation for other categories of readers. Our experience during the study suggested however that meeting prior to the first evaluation for discussing how to use the method in detail, evaluating about ten cases in common, and using a clear illustration of rating as shown in figure 2, were presumably crucial to reach highly reproducible items. As expected with experiencing the method during the study, only the rating of dPVS did not reach a high level of inter-rater agreement, particularly when using images from basic MRI protocols without T2 images or low-resolution data from 1.5-T MRI. This was not surprising insofar as dPVS were the finest signal abnormalities to be detected on MRI³³. Thus, the quality of detection depended not only on the possibility to follow the corresponding vascular trajectories in 3D but also on the level of image resolution as already illustrated using extreme resolution with 7.0-Tesla MRI.^{34, 35}

The analysis of the association between the different imaging items and the main clinical manifestations of the disease showed results in perfect agreement with different links already established between quantitative MRI markers and the clinical severity in CADASIL. The significant association between stroke and the score of lacunes or that of macrobleeds which was not found with WMH items has been previously reported by measuring the load of lacunes and total volume of WMH.^{9,32} Also, the independent association between disability or MMSE score with the lacune and atrophy items are in line with previous reports showing that the accumulation of lacunes was not only a key driver of motor and cognitive decline during the progression of CADASIL but also promoted the development of cerebral and cortical atrophy.³⁶⁻³⁸ Finally, the association observed with the PVH item but not with the superficial WMH item is in agreement with the different types of white matter lesions of distinct nature and clinical impact already delineated in CADASIL.^{12,39} Interestingly, we observed that migraine with aura was negatively correlated with superficial atrophy and possibly related to WMH in the temporal lobes. Further analyses are needed to better understand the potential clinical links between lesions in the temporal white-matter relatively specific to the disease.^{40,41} Interestingly, although our tool was not primarily developed to reflect the clinical severity of the disease in terms of brain damage, scores that could be derived from our tool might also provide crucial information about the brain lesions variably associated with different clinical manifestations of the disease. Our results clearly showed that the CADA-MRIT items allowed, in any case, to predict the different clinical manifestations of CADASIL better than the use of a simple and global score as the total CSVD score developed in another context and for different purposes.³¹

Obviously, this study based on the development and validation of a specific tool to inventory multiple categories of MRI lesions in CADASIL presents many strengths. This tool was proposed based on a real need to compare easily and quickly patients' imaging data across countries or cohorts. It was developed both by CADASIL and MRI experts of CSVD from data collected from over 600 individuals and 3 distinct cohorts with a large spectrum of clinical and cerebral manifestations. Its reliability was extensively assessed from multiple angles and the derived imaging items were found clinically relevant. The tool was particularly easy to use and fast and could provide directly different imaging items at the patient bed and for clinical research. Based on this inventory, a rapid and more massive combination of data could be envisaged in future research. Some refinements could certainly improve it further for investigating other rare CSVD or answering very specific questions. For example, lacunes in the pons could be added as an additional item when evaluating pontine autosomal dominant microangiopathy and leukoencephalopathy (PADMAL).⁴² The location of lacunes or microbleeds which was not assessed in the inventory would also need additional items. We are also aware of some limitations of such an approach. Although we compiled different imaging protocols for validating the reproducibility of the inventory tool, we could not ascertain that other parameters from other MRI sequences not used in the present study could not influence the different imaging items. Also, the method can in no way be used to assess the progression of the disease over a limited time frame. In addition, it does not provide any continuous measure which limits the statistical power for detecting small differences. The evaluation of lesions remains also dependent on the quality of data and resolution of images, particularly for PVS. Moreover, as the different items of the inventory tool were variously associated with the clinical manifestations of the disease, we could not provide immediately a "total score" of lesions related to the clinical progression of the disease from this inventory. The weights of its multiple items might be probably adjusted in future to obtain a global score of severity according to the aim and design of future studies. In the meantime, the results from our inventory can be easily transformed to the CSVD total score³¹ or to the Fazekas score²⁰ when seeking for a simple comparison with the burden of lesions assessed in other databases or SVDs.

In conclusion, we believe that the inventory tool developed and validated in the present study is a new instrument for CADASIL research. It should first greatly facilitate the common assessment of large amounts of brain imaging data

and comparison between cohorts and multiple samples of patients. Further studies will be needed to determine how this tool could be also used to facilitate the instant evaluation of patients at bedside, prognostication, and triage for future therapeutic trials in CADASIL.

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Disclosure

The authors report no disclosures relevant to the manuscript.

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Table 1 The demographic and clinical characteristics of the cohorts

	Cohort 1 (n = 440)	Cohort 2 (n = 119)	Cohort 3 (n = 112)
Age, years, mean±SD	52.4 ± 12.0	48.3 ± 9.9	62.5 ± 10.6
Sex, female, n (%)	244 (55.5%)	69 (58.0%)	53 (47.3%)
Education level†, median (IQR)	5 (4-7)	5 (4-5)	6 (5-7)
Stroke, n (%)	230 (52.3%)	82 (68.9%)	65 (58.0%)
Migraine with aura, n (%)	163 (37.0%)	45 (37.8%)	1 (0.9%)
Disability, n (%)	66 (15.1%)	12 (10.1%)	24 (21.4%)
MMSE, median (IQR)	28 (26-30)	29 (27-30)	28 (24-29)
Normalized WMH fraction, median (IQR)	3.3% (1.8%-5.9%)	5.7% (3.7%-10.4%)	2.4% (1.2%-3.7%)
Lacune number, median (IQR)	5 (1-13)	3 (0-7)	3 (1-9)
CMB number, median (IQR)	0 (0-2)	0 (0-1)	9 (2-30)
BPF, median (IQR)	82.6% (78.6%-85.0%)	88.1% (84.3%-90.9%)	72.2% (68.9%-75.5%)

†Education level: 7 levels according to the number of education years. Level 1 = 0; Level 2 = 1-5; Level 3 = 6-8; Level 4 = 9; Level 5 = 10-11; Level 6 = 12; Level 7 = over 12 years.

MMSE = Mini-Mental State Examination; WMH = White matter hyperintensities; CMB = Cerebral microbleeds; BPF = Brain parenchymal fraction.

Table 2 Distribution of the different scores of the inventory in the 3 cohorts

		Cohort 1 (n = 440)	Cohort 2 (n = 119)	Cohort 3 (n = 112)
PVH	0	4 (0.9%)	0 (0%)	1 (0.9%)
	1	26 (5.9%)	4 (3.4%)	12 (10.7%)
	2	140 (31.8%)	20 (16.8%)	24 (21.4%)
	3	187 (42.5%)	69 (58.0%)	57 (50.9%)
	4	83 (18.9%)	26 (21.8%)	18 (16.1%)
Deep WMH	0	3 (0.7%)	2 (1.7%)	6 (5.4%)
	1	48 (10.9%)	11 (9.2%)	11 (9.8%)
	2	74 (16.8%)	19 (16.0%)	33 (29.5%)
	3	315 (71.6%)	87 (73.1%)	62 (55.4%)
Superficial WMH	0	30 (6.8%)	10 (8.4%)	85 (75.9%)
	1	177 (40.2%)	21 (17.6%)	17 (15.2%)
	2	87 (19.8%)	30 (25.2%)	4 (3.6%)
	3	95 (21.6%)	37 (31.1%)	3 (2.7%)
	4	51 (11.6%)	21 (17.6%)	3 (2.7%)
Lacune	0	126 (28.6%)	26 (21.8%)	21 (18.8%)
	1	123 (28.0%)	33 (27.7%)	51 (45.5%)
	2	79 (18.0%)	19 (16.0%)	24 (21.4%)
	3	112 (25.5%)	41 (34.5%)	16 (14.3%)
CMB	0	281 (63.9%)	80 (67.2%)	19 (17.0%)
	1	95 (21.6%)	21 (17.6%)	37 (33.0%)
	2	28 (6.4%)	10 (8.4%)	14 (12.5%)
	3	36 (8.2%)	8 (6.7%)	42 (37.5%)
BG dPVS‡	0	142 (32.3%)	50 (42.0%)	15 (13.4%)
	1	247 (56.1%)	65 (54.6%)	35 (31.3%)
	2	47 (10.7%)	3 (2.5%)	62 (55.4%)
CSO dPVS‡	0	235 (53.4%)	78 (65.5%)	60 (53.6%)
	1	150 (34.1%)	31 (26.1%)	32 (28.6%)
	2	51 (11.6%)	9 (7.6%)	20 (17.9%)
Superficial atrophy	0	290 (65.9%)	98 (82.4%)	20 (17.9%)
	1	89 (20.2%)	18 (15.1%)	45 (40.2%)
	2	57 (13%)	2 (1.7%)	32 (28.6%)
	3	4 (0.9%)	1 (0.8%)	15 (13.4%)
Deep atrophy	0	279 (63.4%)	77 (64.7%)	29 (25.9%)
	1	101 (23%)	29 (24.4%)	42 (37.5%)
	2	48 (10.9%)	9 (7.6%)	30 (26.8%)
	3	12 (2.7%)	4 (3.4%)	11 (9.8%)
Large infarct	0	429 (97.5%)	116 (97.5%)	109 (97.3%)
	1	10 (2.3%)	3 (2.5%)	3 (2.7%)
	2	1 (0.2%)	0 (0%)	0 (0%)
Macrobleeds	0	427 (97.0%)	119 (100%)	84 (75.0%)
	1	9 (2.0%)	0 (0%)	21 (18.8%)
	2	4 (0.9%)	0 (0%)	7 (6.3%)

‡ dPVS were evaluated in 436 patients in Cohort 1, 118 patients in Cohort 2 and 112 patients in Cohort 3. Four patients in Cohort 1 and one patient in Cohort 2 were excluded because of severe artifact on T1-weighted images.

PVH = Periventricular WMH; WMH = White matter hyperintensities; CMB = Cerebral microbleeds; CSO = Centrum semiovale; BG = Basal ganglia; dPVS = dilated perivascular spaces.

Table 3 Association between the different inventory tool items and different clinical outcomes

	Stroke, OR (95% CI)		Migraine with aura, OR (95% CI)		Disability, OR (95% CI)		MMSE, β (95% CI)	
	Model 1	Model 2	Model 1	Model 2	Model 1	Model 2	Model 1	Model 2
PVH	2.167 (1.708~2.749)*	1.141 (0.801~1.625)	1.117 (0.881~1.417)	0.961 (0.661~1.396)	4.305 (2.889~6.415)*	2.208 (1.252~3.896)**	<u>-1.418 (-1.849~-0.986)*</u>	<u>-1.211 (-1.794~-0.629)***</u>
Deep WMH	1.952 (1.523~2.502)*	1.173 (0.849~1.622)	1.432 (1.103~1.860)	1.303 (0.935~1.815)	2.316 (1.321~4.060)*	0.766 (0.379~1.546)	0.006 (-0.459~0.470)	0.928 (0.397~1.460)**
Superficial WMH	1.301 (1.145~1.480)*	1.078 (0.904~1.285)	1.355 (1.174~1.565)*	1.162 (0.964~1.402)	1.127 (0.959~1.324)	0.959 (0.753~1.221)	-0.029 (-0.276~0.217)	-0.121 (-0.405~0.163)
Total WMH	1.271 (1.177~1.374)*	/	1.160 (1.068~1.260)*	/	1.267 (1.131~1.419)*	/	-0.160 (-0.305~-0.016)	/
Lacune	2.454 (2.042~2.950)*	2.317 (1.854~2.895)***	0.996 (0.841~1.179)	1.036 (0.833~1.288)	2.759 (2.105~3.616)*	2.271 (1.591~3.241)***	<u>-0.835 (-1.144~-0.526)*</u>	-0.250 (-0.597~0.097)
CMB	1.329 (1.112~1.588)*	0.958 (0.770~1.193)	<u>0.659 (0.526~0.826)*</u>	<u>0.698 (0.536~0.909)**</u>	1.514 (1.232~1.861)*	1.175 (0.914~1.512)	<u>-0.674 (-1.013~-0.335)*</u>	-0.079 (-0.436~0.278)
CSO dPVS	<u>0.712 (0.567~0.894)*</u>	0.881 (0.677~1.146)	1.104 (0.863~1.412)	1.144 (0.871~1.501)	<u>0.562 (0.386~0.817)*</u>	0.885 (0.567~1.381)	0.829 (0.388~1.270)*	0.337 (-0.092~0.765)
BG dPVS	1.068 (0.808~1.410)	1.063 (0.766~1.473)	0.756 (0.556~1.030)	0.904 (0.642~1.274)	1.045 (0.704~1.550)	0.860 (0.521~1.420)	-0.042 (-0.596~0.513)	0.216 (-0.318~0.750)
Total dPVS	0.859 (0.730~1.010)	/	0.959 (0.802~1.147)	/	0.776 (0.609~0.990)	/	0.420 (0.100~0.741)	/
Superficial atrophy	1.090 (0.851~1.396)	0.908 (0.643~1.282)	<u>0.363 (0.25~0.526)*</u>	<u>0.344 (0.218~0.545)***</u>	2.570 (1.860~3.551)*	1.581 (1.046~2.392)#	<u>-1.886 (-2.346~-1.426)*</u>	<u>-1.145 (-1.697~-0.592)***</u>
Deep atrophy	1.290 (1.022~1.627)	0.861 (0.619~1.199)	0.787 (0.599~1.034)	1.487 (1.024~2.159)#	2.926 (2.178~3.932)*	1.352 (0.910~2.009)	<u>-1.885 (-2.311~-1.459)*</u>	<u>-0.719 (-1.254~-0.185)**</u>
Total atrophy	1.120 (0.978~1.283)	/	<u>0.698 (0.583~0.834)*</u>	/	1.981 (1.646~2.384)*	/	<u>-1.215 (-1.462~-0.968)*</u>	/
Large infarct	NA	/	1.387 (0.520~3.697)	1.338 (0.474~3.776)	3.822 (1.468~9.951)	3.158 (1.093~9.122)#	-1.141 (-2.999~0.717)	-0.261 (-1.972~1.450)
Macrobleeds	2.136 (1.147~3.980)	3.116(1.534~6.332)**	0.196 (0.051~0.747)	0.372 (0.102~1.360)	1.822 (1.072~3.097)	2.122 (1.062~4.239)#	<u>-1.652 (-2.637~-0.667)*</u>	<u>-1.248 (-2.218~-0.277)#</u>

Association between each inventory tool item and the different clinical outcomes, the analysis was performed separately for each item (Model 1, significance at *p < 0.0045).

Association between inventory tool item and the different clinical outcomes independently from the other items (Model 2, multivariate analysis, significance at ***p < 0.001; **p < 0.01, #p < 0.05). The total scores are not included in Model 2 to avoid collinearity with the raw items.

NA: All patients with large infarct had stroke.

PVH = Periventricular WMH; WMH = White matter hyperintensities; CMB = Cerebral microbleeds; CSO = Centrum semiovale; BG = Basal ganglia; dPVS = dilated perivascular spaces; MMSE = Mini-Mental State Examination.

Values in bold represented significant positive association between inventory tool scores and clinical outcomes, while values in italic and with underline represented significant negative association.

Figure legends

Figure 1: The CADASIL imaging inventory

Legend Figure 1: The items and scores composing the imaging inventory tool are presented on a single page sheet.

Figure 2 White matter hyperintensities (WMH), lacune, cerebral microbleeds (CMB), dilated perivascular spaces (dPVs) and atrophy scores from the inventory tool

Legend Figure 2: The different scores obtained from the imaging inventory tool are presented with the corresponding images to facilitate their interpretation.

Figure 3: Relationships between the total white matter hyperintensities (WMH) score (sum of periventricular, deep and superficial WMH scores) and the volume of WMH and between the total atrophy score (sum of deep and superficial atrophy scores) and the brain parenchymal fraction.

Legend Figure 3: Scatter plots showing (A) the significant association between the total WMH score and the cube root of normalized WMH fraction; (B) the significant association between the total atrophy score and the normal quantile transformed BPF.

Figure 4: Illustrative cases showing different scores derived from the Inventory tool obtained from various images

Legend Figure 4: Patient A: A 64-year-old woman, with predominantly superficial WMH [PVH, deep WMH, superficial WMH scores = 3, 3, 4, respectively] but no lacune, CMB or atrophy. She had no stroke history, mRS = 0 and MMSE = 29. Patient B: A 59-year-old man, with predominantly periventricular WMH [PVH, deep WMH, superficial WMH scores = 3, 3, 0, respectively], more than 10 lacunes (lacune score = 3) and significant brain atrophy (superficial and deep atrophy scores = 2, 1, respectively). He had stroke history, with mRS = 3 and MMSE = 26. Patient C: A 62-year-old man, from cohort 3, with moderate PVH and deep WMH and no superficial WMH [PVH, deep WMH, superficial WMH score = 3, 2, 0, respectively], more than 30 CMBs (CMB score = 3), one macrobleed and significant brain atrophy (superficial and deep atrophy score = 2, 1, respectively). He had stroke history, with mRS = 3 and MMSE = 30. Pv: Periventricular; Superf: Superficial. [§] 3D-T1 was used for assessment but not shown.

Appendix 1

Name	Location	Contribution
Ruiting Zhang, MD, PhD	Paris-Cité University, Inserm U1141 NeuroDiderot, Paris, France. Department of Radiology, the Second Affiliated Hospital of Zhejiang University, School of Medicine, Hangzhou, China	Drafting/revision of the manuscript for content, including medical writing for content; study concept or design; and analysis or interpretation of data
Chih-Hao Chen, MD, PhD	Department of Neurology, National Taiwan University Hospital, Taipei, Taiwan. Department of Clinical Neurosciences, University of Calgary, Calgary, Alberta, Canada	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data
Sophie Tezenas Du Montcel, MD, PhD	Sorbonne Université, INSERM, Unité Mixte de Recherche 1136, Institut Pierre Louis d'Épidémiologie et de Santé Publique, Paris, France. Sorbonne Université, INSERM, Institut Pierre Louis d'Épidémiologie et de Santé Publique, AP-HP, Hôpitaux Universitaires Pitié Salpêtrière - Charles Foix, F75013 Paris, France.	Drafting/revision of the manuscript for content, including medical writing for content; and analysis or interpretation of data
Jessica Leberberg	Paris-Cité University, Inserm U1141 NeuroDiderot, Paris, France. Lariboisière University Hospital, APHP, Translational Neurovascular Centre and Department of Neurology, Reference Center for Rare Vascular Diseases of the Central Nervous System and the Retina (CERVCO), FHU NeuroVasc, Paris, France	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data
Yu-Wen Cheng, MD, PhD	Department of Neurology, National Taiwan University Hospital, Taipei, Taiwan. Department of Neurology, National Taiwan University Hospital Hsinchu Branch, Hsinchu, Taiwan.	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data
Martin Dichgans, MD, PhD	Institute for Stroke and Dementia Research, University Hospital, Ludwig Maximilian University, Munich, Germany. German Center for Neurodegenerative Diseases (DZNE), Munich, Germany. Munich Cluster for Systems Neurology (SyNergy), Munich, Germany.	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data

Sung-Chun Tang, MD, PhD	Department of Neurology, National Taiwan University Hospital, Taipei, Taiwan	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data
Hugues Chabriat, MD, PhD	Paris-Cité University, Inserm U1141 NeuroDiderot, Paris, France. Lariboisière University Hospital, APHP, Translational Neurovascular Centre and Department of Neurology, Reference Center for Rare Vascular Diseases of the Central Nervous System and the Retina (CERVCO), FHU NeuroVasc, Paris, France	Drafting/revision of the manuscript for content, including medical writing for content; major role in the acquisition of data; study concept or design; and analysis or interpretation of data

The CADASIL MRI Inventory Tool (CADA-MRIT)

White matter hyperintensities (WMH)

Periventricular WMH (PVH) [In contact with the ventricles]

- 0 : Absence
- 1 : “Caps” or pencil-thin lining
- 2 : Smooth “halo”
- 3 : PVH extending into the deep white matter partly merging with deep WMH
- 4 : PVH extending into the deep white matter completely merging with deep WMH (PVH cannot be separated from deep WMH anywhere)

Deep WMH [Between PVH and superficial WMH]

- 0 : Absence
- 1 : Punctate foci
- 2 : Beginning confluence of foci
- 3 : Large confluent areas

Superficial WMH [In contact with U fibers; Evaluate frontal and temporal pole white matter specifically, if different in the 2 locations, use the most severe one]

- 0 : Absence
- 1 : Punctate foci
- 2 : Beginning confluence of foci
- 3 : Confluent WMH partly merging with deep WMH
- 4 : Confluent WMH completely merging with deep WMH

Lacunae

- | | | | |
|--------------------------------|----------------------------------|-----------------------------------|----------------------------------|
| <input type="checkbox"/> 0 : 0 | <input type="checkbox"/> 1 : 1-5 | <input type="checkbox"/> 2 : 6-10 | <input type="checkbox"/> 3 : >10 |
|--------------------------------|----------------------------------|-----------------------------------|----------------------------------|

Cerebral microbleeds (CMB)

- | | | | |
|--------------------------------|---|---|--|
| <input type="checkbox"/> 0 : 0 | <input type="checkbox"/> 1 : 1-5(T2*)/1-10(SWI) | <input type="checkbox"/> 2 : 6-10(T2*)/11-20(SWI) | <input type="checkbox"/> 3 : >10(T2*)/>20(SWI) |
|--------------------------------|---|---|--|

Dilated perivascular spaces (dPVS)

[Count in the slice with the highest number. T1 – sum of both sides, and T2 – the higher score on one side]

<i>Centrum semiovale (CSO)</i>	<i>Basal ganglia</i>
--------------------------------	----------------------

- | | |
|----------------------------------|----------------------------------|
| <input type="checkbox"/> 0 : <10 | <input type="checkbox"/> 0 : <10 |
|----------------------------------|----------------------------------|

- | | |
|------------------------------------|------------------------------------|
| <input type="checkbox"/> 1 : 10-20 | <input type="checkbox"/> 1 : 10-20 |
|------------------------------------|------------------------------------|

- | | |
|----------------------------------|---|
| <input type="checkbox"/> 2 : >20 | <input type="checkbox"/> 2 : >20, innumerable or status cribrosum |
|----------------------------------|---|

Atrophy

<i>Superficial [Dilation of the sulcus; Evaluate central sulcus and lateral sulcus specifically, if different in the 2 locations, use the most severe one]</i>	<i>Deep [Dilation of the lateral ventricles]</i>
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- | | |
|---------------------------------------|---------------------------------------|
| <input type="checkbox"/> 0 : Absence | <input type="checkbox"/> 0 : Absence |
| <input type="checkbox"/> 1 : Mild | <input type="checkbox"/> 1 : Mild |
| <input type="checkbox"/> 2 : Moderate | <input type="checkbox"/> 2 : Moderate |
| <input type="checkbox"/> 3 : Severe | <input type="checkbox"/> 3 : Severe |

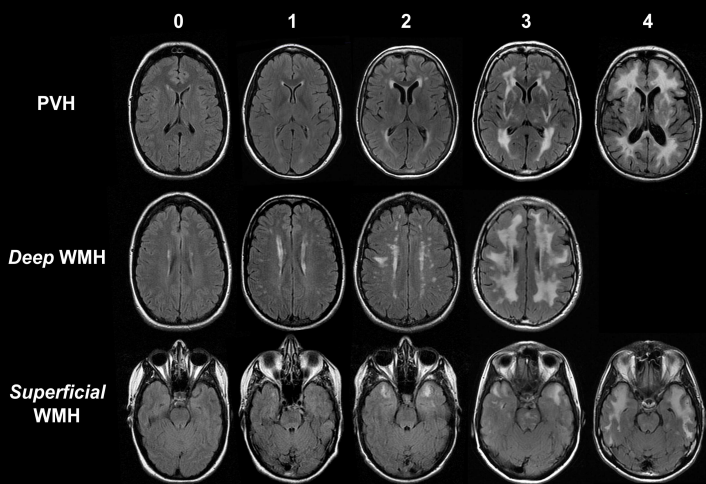
Large infarct (>20mm)

- | | | |
|--------------------------------|--------------------------------|---------------------------------|
| <input type="checkbox"/> 0 : 0 | <input type="checkbox"/> 1 : 1 | <input type="checkbox"/> 2 : >1 |
|--------------------------------|--------------------------------|---------------------------------|

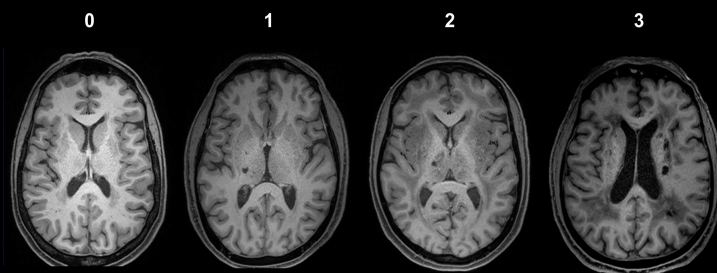
Macrobleeds (>15mm)

- | | | |
|--------------------------------|--------------------------------|---------------------------------|
| <input type="checkbox"/> 0 : 0 | <input type="checkbox"/> 1 : 1 | <input type="checkbox"/> 2 : >1 |
|--------------------------------|--------------------------------|---------------------------------|

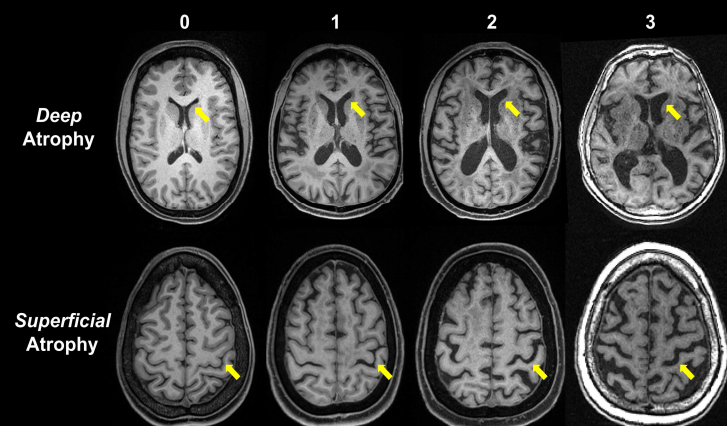
WMH scores



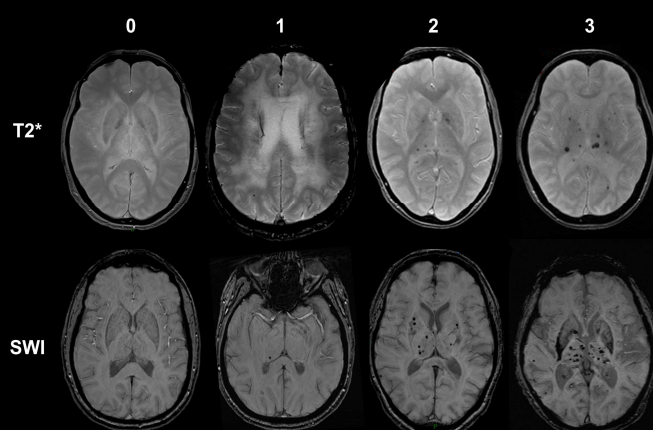
Lacune scores



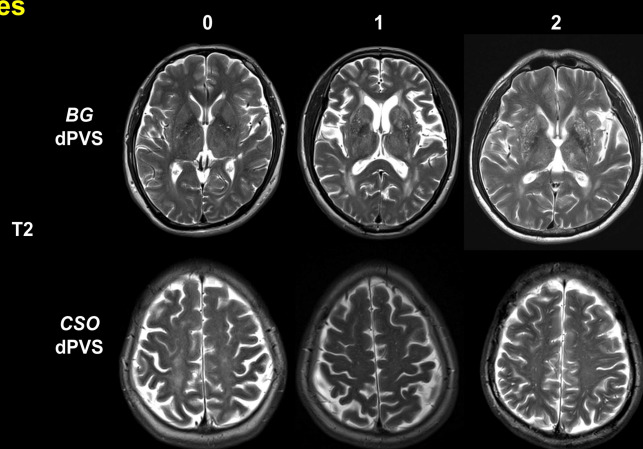
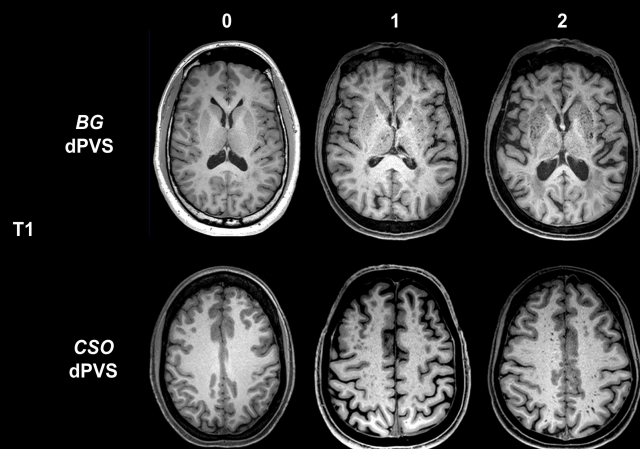
Atrophy scores

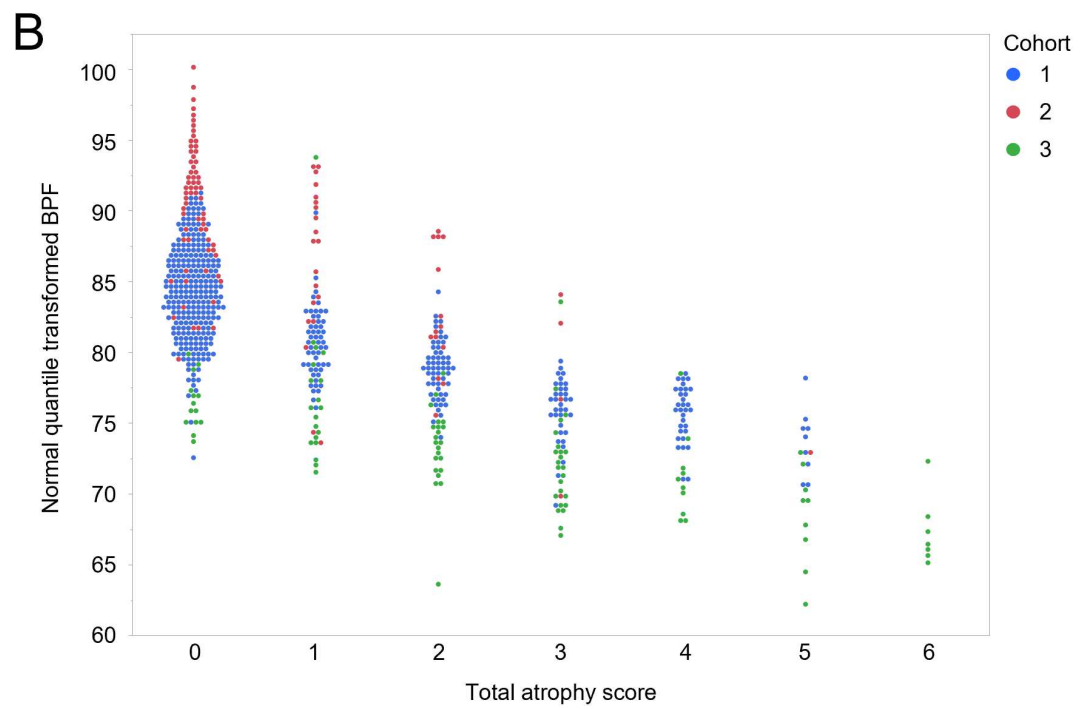
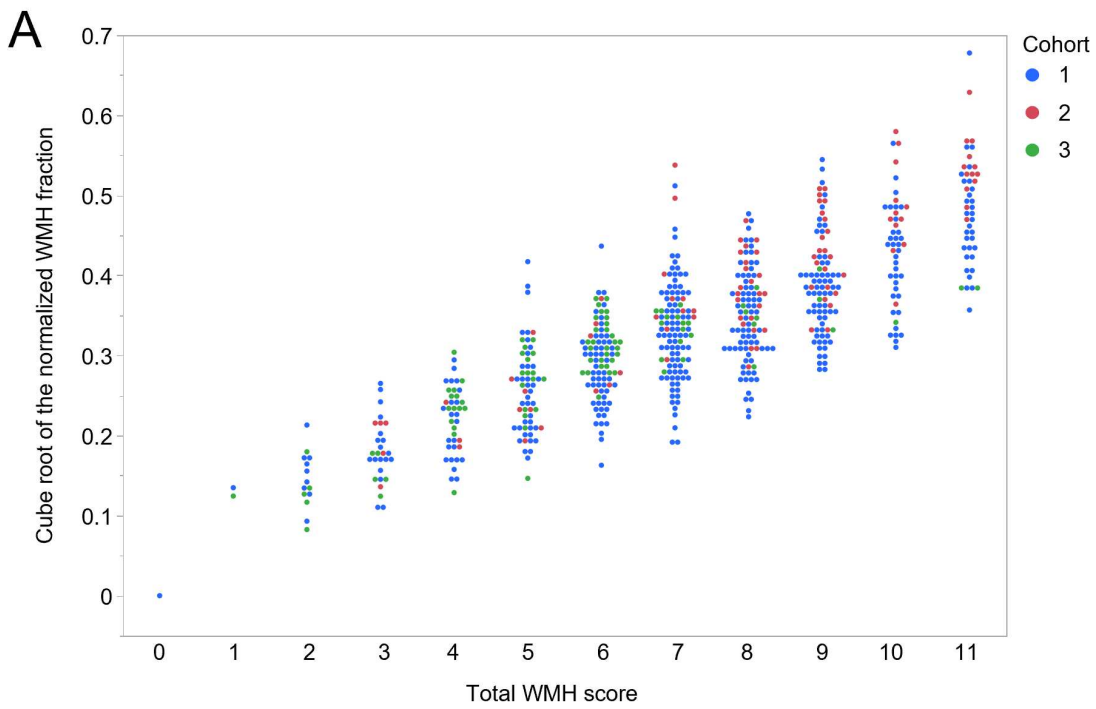


CMB scores



dPVS scores





Patient A

Female, 64y, mRS = 0, MMSE = 29

Inventory scores:

WMH: Pv-3, Deep-3, Superf-4

Lacune: 0; CMB: 0

§ Atrophy: Superf-0, Deep-0

Patient B

Male, 59y, mRS = 3, MMSE = 26

Inventory scores:

WMH: Pv-3, Deep-3, Superf-0

Lacune: 3; CMB: 1

§ Atrophy: Superf-2, Deep-1

Patient C

Male, 62y, mRS = 3, MMSE = 30

Inventory scores:

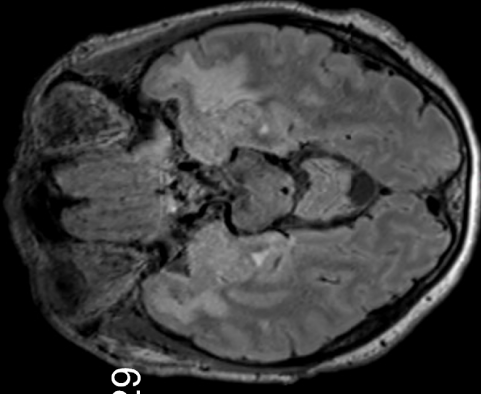
WMH: Pv-3, Deep-2, Superf-0

Lacune: 2; CMB: 3

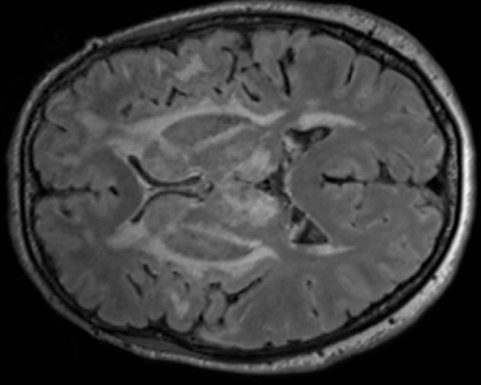
§ Atrophy: Superf-2, Deep-1

Macrobleeds: 1

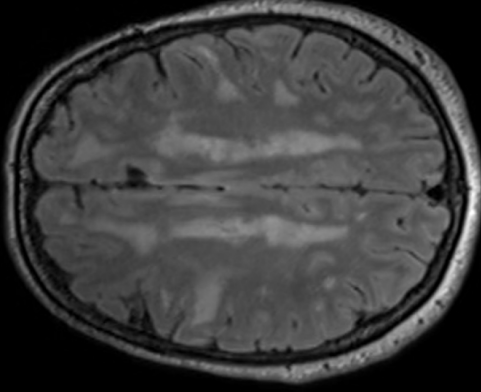
T2FLAIR



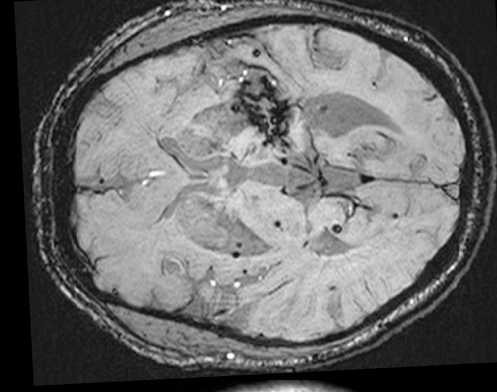
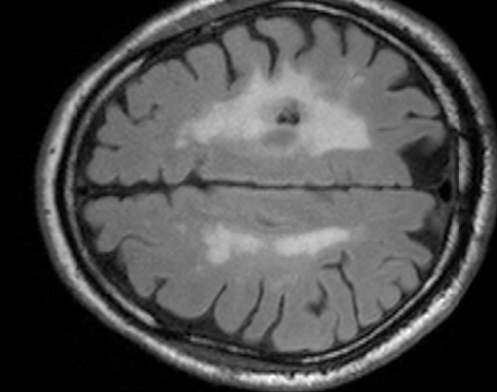
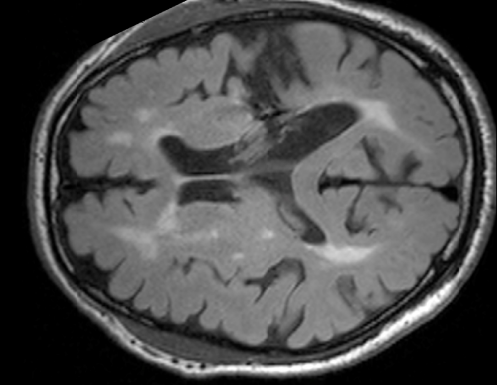
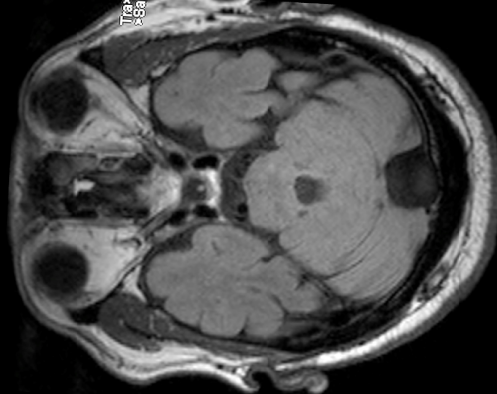
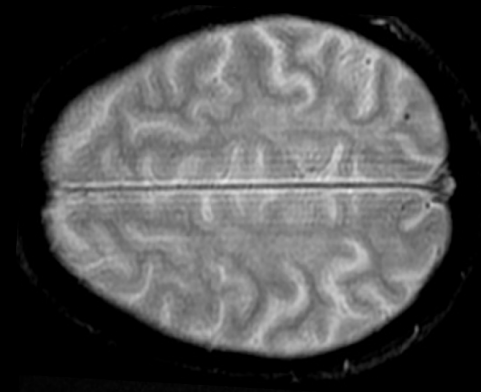
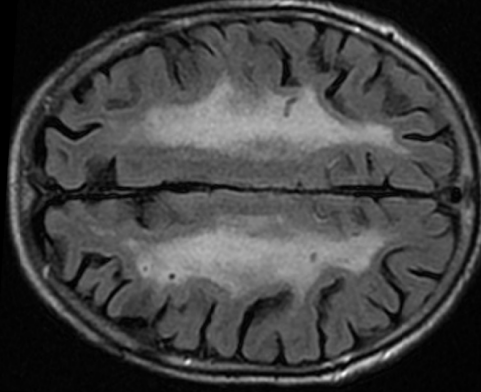
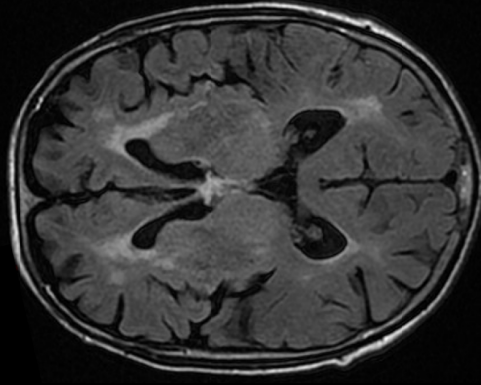
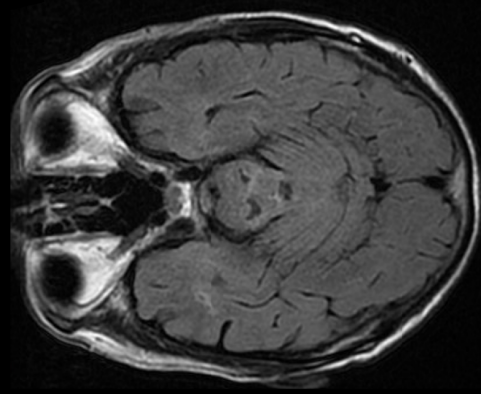
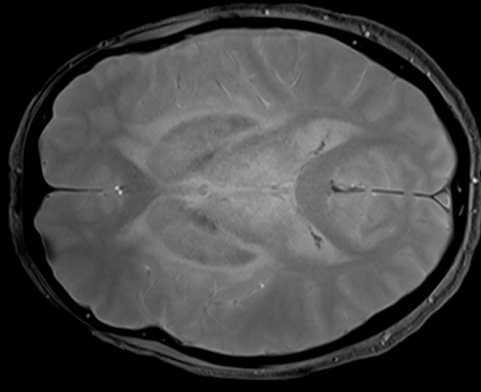
T2FLAIR



T2FLAIR



T2*/SWI



eMethods

MRI acquisition parameters

As the MRI scans were performed in a clinical cohort setting over a period of more than 15 years, modifications in MRI parameters and sequences occurred from time to time. They are detailed below.

COHORT 1: All MRI sequences were obtained with no interslice gap. From September 2003 to April 2014, MRI acquisition was performed on a 1.5-T MRI machine (Signa General Electric Medical Systems). The sequences and corresponding parameters were as follows: 1) three-dimensional T1-weighted (3D-T1): TR/TE 9/2 ms, slice thickness 1.60 mm; 2) Fluid-attenuated inversion recovery (FLAIR): TR/TE/TI 8402/161/2100 ms, slice thickness 5.5 mm; 3) T2*-weighted imaging (T2*): TR/TE 500/15 ms, slice thickness 5.5 mm. From May 2014 to November 2021, MRI scans were obtained by the use of a 3.0-T system (Skyra SIEMENS) as follows: 1) 3D-T1: TR/TE 2200/2.43 ms, slice thickness 1 mm from May 2014 to July 2015; TR/TE 1800/2.35 ms, slice thickness 0.85 mm from July 2015 to September 2017; TR/TE 2000/3.15 ms, slice thickness 0.84 mm from September 2017 to November 2021; 2) FLAIR: TR/TE/TI 6000/387/2200 ms, slice thickness 0.90 mm from May 2014 to July 2018; TR/TE/TI 5000/393/1600 ms, slice thickness 1 mm from July 2018 to February 2020; TR/TE/TI 5000/378/1650 ms, slice thickness 1.13 mm from February 2020 to November 2021, 3) SWI: TR/TE 27/20 ms, slice thickness 1.5 mm from May 2014 to September 2017; TR/TE 22/15 ms, slice thickness 2 mm from September 2017 to February 2021, 4) T2*: TR/TE 600-720/10-18 ms, slice thickness 4 mm from May 2014 to May 2018.

COHORT 2: MRI scans were obtained by the use of a 1.5-T system (Signa General Electric Medical Systems), always without interslice gap between images. The parameters were as follows. 1) 3D-T1: TR/TE 11.4/4.4 ms, slice thickness 1.19 mm; or TR/TE 22/6 ms, slice thickness 1.20 mm; 2) FLAIR: TR/TE/TI 4284/110/1428 ms, slice thickness 5 mm; or TR/TE/TI 8402/150.6/2100 ms, slice thickness 5 mm; 3) T2*: TR/TE 1056/22 ms, slice thickness 5 mm.

COHORT 3: MRI scans were acquired on three 3.0-T scanners (Aera, Verio or TIM Trio, Siemens) with the same imaging settings. The imaging protocol included 1) 3D-T1: TR 1850-2530 ms, TE 2.27-4.6 ms, TI 720-1100 ms, slice thickness 1mm; 2) FLAIR: TR 10000 ms, TE 90-100 ms, TI 2500-2600 ms; 3) T2-weighted imaging: TR 3500-5500 ms, TE 88-105 ms, slice thickness 5mm; 4) SWI: TR 28-48 ms, TE 20-40 ms, flip angle 15-20°, slice thickness 1.6-2.5 mm.

Construction of the different inventory items

Scoring of WMH was built based on the Fazekas scale^{e1} to take into account the large areas of white matter lesions, particularly when periventricular and deep WMH were totally merged as often seen in CADASIL. For PVH, we added a level 4 as “PVH extending into the deep white matter completely merging with deep WMH (PVH cannot be separated from deep WMH anywhere)” and modified the score 3 as “PVH extending into the deep white matter partly merging with deep WMH”. The level of the deep WMH item did not differ from that of the Fazekas scale. In the inventory tool, we also added an item for the superficial WMH to take into account lesions particularly detected in CADASIL in close contact with the cortex detected in the temporal lobes or in the anterior part of the frontal lobes.^{e2} These superficial WMH

were scored as 0; absence; 1: punctate WMH; 2: foci of WMH becoming confluent; 3: confluent WMH partly merging with deep WMH; 4: confluent WMH completely merging with deep WMH.

Scoring of lacunes was obtained as follows: 0: no lacune; 1: 1 to 5 lacunes; 2: 6 to 10 lacunes; 3: >10 lacunes. We chose 1, 5, and 10 as thresholds based on the quartile distribution of the number of lacune number in all patients [1, 4, 11], along with the consideration of using numbers easy to remember, such as multiples of five.

Scoring of CMB was obtained as follows: 0: no CMB; score 1: 1 to 5 CMB for T2*, 1-10 CMB for SWI; score 2: 6 to 10 CMB for T2*, 11-20 CMB for SWI; score 3: >10 CMB for T2*, >20 CMB for SWI. The cut-offs 0 and 1 were chosen to be able to easily evaluate the presence or absence of CMB. The cut-offs of 5 and 10 for T2* and of 10 for SWI based on the quartile distribution of CMB numbers in the patients with CMB [T2*: 1, 3, 12, n=151; SWI: 3, 10, 36, n = 157], and also on a consensus to facilitate the evaluation using numbers easy to remember and similar to those used for lacunes. The highest level for SWI was initially > 30 according to the 75 percentiles but modified to > 20 in order to shorten the time spent on counting CMB after the discussion in the panel of authors.

Scoring of atrophy was derived from the GCA scale.^{e3} We kept the four levels (from 0 to 3) of this scale as absent, mild, moderate, and severe atrophy but reduced the thirteen brain regions initially proposed to only two areas in the present inventory tool. An item was chosen for superficial cortical atrophy based on the aspect of the central and lateral sulcus at the cortex level. Another item was proposed for deep cerebral atrophy based on the dilation of the cerebral lateral ventricles. For patients with large infarcts or macrobleeds, atrophy was evaluated on the hemisphere contralateral to the largest lesion.

The scoring of dPVS was based on two previous scales already widely used for dPVS observed on T1 and T2-weighted images, either on both sides or on one side of the brain using a single slice.^{e4,5} We combined the first two levels of the previous scales as the first level of the previous scales was hardly seen in our patients, and we modified the last level of the T1-weighted image-based scale from innumerable to >20 for BG region to avoid inconsistency between observers. As shown in Figure 1, CSO and BG dPVS items each had three levels (from 0 to 2), and both were evaluated in the slice containing the greatest number of dPVS. The numbers referred to the sum number of dPVS counted on both sides of the slice for T1-weighted images, and on one side for T2-weighted images (the side with the highest number).

Finally, the occurrence of large infarcts was defined as an area of significant signal changes presumably related to an ischemic lesion of diameter > 20mm. Three levels (from 0 to 2) were given, 0 in the absence of a large infarct, one for a single, and 2 for multiple large infarcts. Macrobleeds were defined as hemorrhagic lesions of diameter >15mm. Three levels (from 0 to 2) were also given in the absence, or presence of one or multiple macrobleeds.

Quantitative imaging data

COHORT 1: The segmentation of the whole brain tissue and the intracranial cavity (ICC) were

obtained in processing T1w-MRI with ANTs (<http://www.picsl.upenn.edu/ANTs>).^{e6} The volume of WMH was calculated in each subject using the BIANCA (Brain Intensity AbNormality Classification Algorithm, <https://fsl.fmrib.ox.ac.uk/fsl/fslwiki/BIANCA>) segmentation tool (using T1w and FLAIR as inputs).^{e7} For 78 patients with insufficient quality of results from BIANCA, the FLAIR images of each subject were further segmented with the Lesion Segmentation Toolbox in SPM12 by applying the best threshold to the probabilistic lesion map to derive the binarized WMH image (<https://www.applied-statistics.de/lst.html>). The segmented lesions of these patients were then visually checked and manually corrected by an experienced neuroradiologist (RZ).

COHORT 2: Determination of the global brain volume was performed as previously detailed.^{e8} A mask of WMH lesions was generated from FLAIR images after the application of the ICC mask by applying a threshold on signal intensity derived from the signal intensities histogram. Two trained neurologists using drawing tools then validated each discrete lesion and corrected its delimitation if necessary.^{e9}

COHORT 3: The segmentation of the whole brain tissue and the intracranial cavity was done by Freesurfer. FLAIR images of each subject were segmented with the Lesion Segmentation Toolbox in SPM12 by applying the best threshold to the probabilistic lesion map to derive the binarized WMH mask. The segmented lesions were visually checked and manually corrected by an experienced neurologist (CHC).

In all cohorts, the brain parenchymal fraction (BPF) was calculated for each individual as the ratio of the brain tissue volume to the ICC volume. The total volume of WMH was normalized to the ICC in each patient [normalized WMH fraction = (WMH volume / ICC volume) * 100%].

eTable 1 Intrarater agreements for each inventory item in different cohorts

Intrarater agreement, weighted κ (95% CI)	Cohort 1 – “basic” protocol	Cohort 1 – “advanced” protocol	Cohort 2	Cohort 3
PVH	0.944 (0.857-1.000)	0.821 (0.684-0.929)	0.968 (0.885-1.000)	0.886 (0.740-0.975)
Deep WMH	0.873 (0.711-1.000)	0.869 (0.699-1.000)	0.856 (0.639-0.969)	0.820 (0.614-0.932)
Superficial WMH	0.877 (0.786-0.961)	0.857 (0.746-0.932)	0.877 (0.763-0.949)	1.000 (1.000-1.000)
Lacune	0.868 (0.734-0.944)	0.871 (0.770-0.958)	0.783 (0.650-0.975)	0.801 (0.654-0.915)
CMB	0.854 (0.675-0.952)	0.820 (0.666-0.930)	0.802 (0.653-0.927)	0.958 (0.882-1.000)
CSO dPVS	0.388 (0.135-0.677)	0.847 (0.726-0.966)	0.677 (0.439-0.856)	0.774 (0.612-0.893)
BG dPVS	0.589 (0.310-0.769)	0.757 (0.554-0.898)	0.371 (0.126-0.604)	0.913 (0.737-1.000)
Superficial atrophy	0.754 (0.622-0.862)	0.809 (0.541-0.941)	0.647 (0.374-0.863)	0.715 (0.528-0.852)
Deep atrophy	0.791 (0.641-0.903)	0.910 (0.814-1.000)	0.856 (0.660-1.000)	0.768 (0.587-0.898)
Large infarct	1.000 (1.000-1.000)	1.000 (1.000-1.000)	1.000 (1.000-1.000)	1.000 (1.000-1.000)
Macrobleeds	1.000 (1.000-1.000)	1.000 (1.000-1.000)	NA	1.000 (1.000-1.000)

NA: None of the patients had macrobleeds in the selected sample of cohort 2.

PVH = Periventricular WMH; WMH = White matter hyperintensities; CMB = Cerebral microbleeds; CSO = Centrum semiovale; BG = Basal ganglia; dPVS = dilated perivascular spaces.

eTable 2 Interrater agreements for each inventory item in different cohorts

Interrater agreement weighted κ (95% CI)	Cohort 1 – “basic” protocol	Cohort 1 – “advanced” protocol	Cohort 3	Mixed
PVH	0.706 (0.545-0.874)	0.657 (0.493-0.813)	0.807 (0.625-0.913)	0.673 (0.506-0.804)
Deep WMH	0.609 (0.387-0.771)	0.516 (0.331-0.688)	0.840 (0.652-0.960)	0.765 (0.568-0.909)
Superficial WMH	0.699 (0.517-0.819)	0.759 (0.662-0.863)	0.837 (0.634-0.958)	0.807 (0.706-0.907)
Lacune	0.786 (0.674-0.880)	0.726 (0.513-0.856)	0.677 (0.481-0.810)	0.761 (0.608-0.880)
CMB	0.863 (0.742-0.975)	0.812 (0.653-0.949)	0.713 (0.556-0.808)	0.885 (0.762-0.961)
CSO dPVS	0.256 (0.010-0.438)	0.733 (0.511-0.867)	0.667 (0.472-0.820)	0.488 (0.217-0.730)††
BG dPVS	0.211 (0.044-0.379)	0.653 (0.394-0.802)	0.657 (0.479-0.832)	0.690 (0.469-0.892)††
Superficial atrophy	0.695 (0.524-0.833)	0.616 (0.380-0.781)	0.556 (0.401-0.721)	0.652 (0.424-0.838)
Deep atrophy	0.651 (0.503-0.768)	0.828 (0.626-0.953)	0.631 (0.476-0.778)	0.771 (0.579-0.892)
Large infarct	1.000 (1.000-1.000)	1.000 (1.000-1.000)	1.000 (1.000-1.000)	NA
Macrobleeds	1.000 (1.000-1.000)	1.000 (1.000-1.000)	0.958 (0.869-1.000)	1.000 (1.000-1.000)

Interrater agreement was not studied in cohort 2.

††This was calculated in patients with advanced imaging data (n = 21). NA: None of the patients had large infarct in the mixed sample.

PVH = Periventricular WMH; WMH = White matter hyperintensities; CMB = Cerebral microbleeds; CSO = Centrum semiovale; BG = Basal ganglia; dPVS = dilated perivascular spaces.

eTable 3 Association between the CSVD total score, inventory items and the different clinical outcomes

Dependent variable	Model	Independent Variables	OR (95% CI)	p	R ²
Stroke	Model I CSVD total score	<i>Age</i>	0.992 (0.975~1.009)	0.352	0.181
		<i>Sex</i>	0.654 (0.468~0.914)	0.013	
		CSVD total score	2.101 (1.714~2.574)	< 0.001	
	Model II Inventory†	<i>Age</i>	1.005 (0.984~1.027)	0.652	0.297
		<i>Sex</i>	0.728 (0.513~1.090)	0.131	
		Lacune	2.317 (1.854~2.895)	< 0.001	
		Macrobleeds	3.116(1.534~6.332)	0.002	
Disability	Model I CSVD total score	<i>Age</i>	1.053 (1.027~1.079)	< 0.001	0.182
		<i>Sex</i>	0.709 (0.449~1.121)	0.141	
		CSVD total score	1.577 (1.188~2.095)	0.002	
	Model II Inventory	<i>Age</i>	1.020 (0.985~1.056)	0.258	0.429
		<i>Sex</i>	1.280 (0.735~2.228)	0.384	
		PVH	2.208 (1.252~3.896)	0.006	
		Lacune	2.271 (1.591~3.241)	< 0.001	
		Superficial atrophy	1.581 (1.046~2.392)	0.030	
		Large infarct	3.158 (1.093~9.122)	0.034	
		Macrobleeds	2.122 (1.062~4.239)	0.033	
Migraine with aura	Model I CSVD total score	<i>Age</i>	0.950 (0.932~0.968)	< 0.001	0.148
		<i>Sex</i>	1.684 (1.176~2.412)	0.004	
		CSVD total score	0.896 (0.734~1.094)	0.282	
	Model II Inventory	<i>Age</i>	0.966 (0.944~0.987)	0.002	0.263
		<i>Sex</i>	1.570 (1.055~2.335)	0.026	
		CMB	0.698 (0.536~0.909)	0.008	
		Superficial atrophy	0.344 (0.218~0.545)	< 0.001	
		Deep atrophy	1.487 (1.024~2.159)	0.037	
MMSE	Model I CSVD total score	<i>Age</i>	-0.096 (-0.129~-0.063)	< 0.001	0.222
		<i>Sex</i>	0.301 (-0.340~0.941)	0.356	
		<i>Education level</i>	0.902 (0.697~1.107)	< 0.001	
		CSVD total score	-0.244 (-0.615~0.126)	0.196	
	Model II Inventory	<i>Age</i>	-0.013 (-0.048~0.022)	0.467	0.368
		<i>Sex</i>	-0.623 (-1.243~-0.002)	0.049	
		<i>Education level</i>	0.799 (0.607~0.991)	< 0.001	
		PVH	-1.211 (-1.794~-0.629)	< 0.001	
		Deep WMH	0.928 (0.397~1.460)	0.001	
		Superficial atrophy	-1.145 (-1.697~-0.592)	< 0.001	
		Deep atrophy	-0.719 (-1.254~-0.185)	0.004	
		Macrobleeds	-1.248 (-2.218~-0.277)	0.012	

†Large infarct was not included in the model, since all patients with large infarct had stroke.

Bold: Only significant items from CADA-MRIT and results obtained with CSVD total score were presented in the Table.

Italics: Age, sex and education (for MMSE only) were forced into the models.

CSVD = Cerebral small vessel disease; CMB = Cerebral microbleeds PVH = Periventricular WMH; WMH = White matter hyperintensities; MMSE = Mini-Mental State Examination.

eTable 4 Mean time for evaluating the inventory items

Inventory items	Cohort 1 – “basic” protocol, time (s)	Cohort 1 – “advanced” protocol, time (s)	Cohort 3, time (s)
WMH	35.0 ± 14.8	31.6 ± 13.7	21.2 ± 8.0
Lacune	66.2 ± 37.0	57.2 ± 40.3	21.9 ± 10.3
CMB	16.9 ± 11.5	27.1 ± 11.5	20.3 ± 9.3
dPVS	26.6 ± 7.1	43.0 ± 23.7	39.2 ± 13.2
Atrophy	14.5 ± 6.6	13.5 ± 9.7	20 ± 8.2
Large infarct	1.0 ± 0.0	1.1 ± 0.6	2.9 ± 3.6
Macrobleeds	1.7 ± 3.4	2.5 ± 7.4	2.6 ± 4.2
Total time – rater 1	161.8 ± 52.9	175.8 ± 71.6	/
Total time – rater 3	153.8 ± 52.4	201.1 ± 77.9	/
Total time – rater 2	/	/	128.2 ± 28.2

WMH = White matter hyperintensities; CMB = Cerebral microbleeds; dPVS = dilated perivascular spaces.

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