Proximity to dementia onset and multi-modal neuroimaging changes: The prevent-dementia study

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\section*{A R T I C L E   I N F O}

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\section*{A B S T R A C T}

\textbf{Background:} First-degree relatives of people with dementia (FH+) are at increased risk of developing Alzheimer’s disease (AD). Here, we investigate "estimated years to onset of dementia" (EYO) as a surrogate marker of pre-clinical disease progression and assess its associations with multi-modal neuroimaging biomarkers.

\textbf{Methods:} 89 FH+ participants in the PREVENT-Dementia study underwent longitudinal MR imaging over 2 years. EYO was calculated as the difference between the parental age of dementia diagnosis and the current age of the participant (mean EYO = 23.9 years). MPRAE, ASL and DWI data were processed using Freesurfer, FSL-BASIL and DTI-TK. White matter lesion maps were segmented from FLAIR scans. The SPM Sandwich Estimator Toolbox was used to test for the main effects of EYO and interactions between EYO, Time, and APOE-ε4+. Threshold free cluster enhancement and family wise error rate correction (TFCE\textsubscript{FWE}) was performed on voxelwise statistical maps.

\textbf{Results:} There were no significant effects of EYO on regional grey matter atrophy or white matter hyperintensities. However, a shorter EYO was associated with lower white matter Fractional Anisotropy and elevated Mean/Radial Diffusivity, particularly in the corpus callosum (TFCE\textsubscript{FWE} p < 0.05). The influence of EYO on white matter deficits were significantly stronger compared to that of normal ageing. APOE-ε4 carriers exhibited hyperperfusion with nearer proximity to estimated onset in temporo-parietal regions. There were no interactions between EYO and time, suggesting that EYO was not associated with accelerated imaging changes in this sample.

\textbf{Conclusions:} Amongst cognitively normal midlife adults with a family history of dementia, a shorter hypothetical proximity to dementia onset may be associated with incipient brain abnormalities, characterised by white matter disruptions and perfusion abnormalities, particularly among APOE-ε4 carriers. Our findings also confer biological validity to the construct of EYO as a potential stage marker of preclinical progression in the context of sporadic dementia. Further clinical follow-up of our longitudinal sample would provide critical validation of these findings.

1. Introduction

It is now established from clinicopathological and in vivo imaging studies that the neuropathological process of Alzheimer’s disease (AD) and dementia begins decades before the earliest manifestation of cognitive or behavioural symptoms (Ritchie et al., 2016). A greater understanding of brain alterations occurring in the preclinical phase might identify novel biomarkers, aid earlier detection and assist in establishing treatment targets for AD. One approach to delineate early alterations is to evaluate longitudinal in vivo neuroimaging markers in asymptomatic individuals who are at increased risk for developing dementia.

Parental family history of dementia (FH+) is a recognised risk factor linked with increased cumulative lifetime risk for dementia, earlier age of onset as well as cognitive impairment (Scarabino et al., 2016). Previous neuroimaging studies have hinted at the presence of brain abnor-

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malities in FH+ individuals, such as reduced functional connectivity, hyperperfusion and grey matter atrophy (Adluru et al., 2014; Honea et al., 2010). However, our systematic reviews of structural and functional imaging studies in preclinical dementia have highlighted numerous discordant findings in the literature (Habib et al., 2017; Mak et al., 2017). Methodological differences, study designs and different age ranges are all plausible explanations for the discrepancy of findings in the FH literature. Yet an alternative possibility is that the presence of brain abnormalities may only be reliably detected amongst individuals who are on the precipice of symptom onset. Indeed, studies in cohorts of persons with familial AD mutations have proposed the estimated years to onset (EYO) as a stage marker of preclinical severity (Bateman et al., 2012). By exploiting the near certainty of dementia onset in carriers of familial AD mutations, EYO has been derived based on the difference between an individual’s age and the parental age of symptom onset. Its association with deficits in functional connectivity (Ringman et al., 2011), alterations in white matter tracts (Ringman et al., 2007), white matter lesions (Vilqar et al., 2015) and grey matter atrophy (Cash et al., 2013) has been investigated. A brief summary of key imaging findings on familial AD mutation carriers in relation to EYO is shown in Supplementary Table 1.

Data from the Swedish Twin Registry has demonstrated a strong heritable component of AD for both men and women (Gatz et al., 2006; Pedersen et al., 2004). A previous study also showed that heritable factors were associated with intergenerational age of symptom onset (Day et al., 2019). Collectively, these lines of evidence imply that EYO may be generalizable to some extent in sporadic AD and thus represents a surrogate index of preclinical disease progression. Our group first extrapolated the construct of EYO to a longitudinal cohort of cognitively normal midlife adults and found that FH+ individuals with a shorter EYO had worse visual working memory scores (Ritchie et al., 2017). Recent reports from an ‘at risk’ cohort have revealed that EYO was associated with temporo-parietal brain atrophy (Vogel et al., 2018) and lower levels of cerebrospinal fluid (CSF) Aβ1-42, particularly among APOE ε4 carriers (Villeneuve et al., 2018). Nevertheless, the mean age of their cohort was 63 years, and consequently the mean EYO was shorter (11 years to onset) relative to our sample in the PREVENT-Dementia study (Ritchie et al., 2013, 2017).

To extend previous cross-sectional studies (Caballero et al., 2018; Villeneuve et al., 2018; Vogel et al., 2018) and address the current gaps in our understanding of brain changes occurring during the preclinical stages, we mapped out an in-depth characterisation of EYO-related changes in (1) grey matter, (2) white matter microstructure, (3) white matter lesions, and (4) cerebral perfusion in a longitudinal cohort of FH+ individuals (mean follow-up of 2 years). We tested the hypotheses that individuals approaching their parental age of dementia diagnosis would show more severe deficits on multi-modal neuroimaging measures, expressed by a lower FA, higher MD and RD, more severe burden of white matter lesions, cortical and hippocampal atrophy, as well as reduced cerebral perfusion.

2. Materials and methods

2.1. Participants and clinical data

The protocol for the PREVENT study has been described in detail previously (Ritchie and Ritchie, 2012). Participants aged 40 to 59 years old with or without a parent with dementia were recruited through multiple sources, such as the register database held at West London Mental Health National Health Service Trust, part of the UK National Health Service. Other participants were recruited via the Join Dementia Research website through information about the study on the Internet and public presentations. Consented participants were seen at the West London Cognitive Disorders Treatment and Research Unit, West London Mental Health NHS Trust, where they were given a standardised neuropsychiatric interview and life-style questionnaires. Cognitive assessments were performed with the Addenbrooke’s Cognitive Examination (ACE-III) during the follow-up visit. Blood was taken for APOE ε4 genotyping with all members of the research and clinical teams remaining blind to the results.

2.2. Ethics statement

Approval for the study has been given by the NHS Research Ethics Committee London Camberwell St-Giles. EYO was estimated as the difference between the age of the participant and the age of dementia diagnosis of the earliest affected parent (if more than one parent had dementia) (Ritchie et al., 2017). Global cognition and domain-specific cognition was also evaluated using the Addenbrooke’s Cognitive Examination (ACE-III) at Visit 2.

2.3. Multi-modal imaging acquisition

All subjects were scanned using a 3T Siemens Verio scanner. The following MR scans were acquired: A 3D T1-weighted magnetization prepared rapid gradient-echo (MPRAGE) image (TR = 2.3 s, TE = 2.98 ms, 160 slices, flip angle = 9°, voxel size = 1 mm3 isotropic); a diffusion weighted image (DWI) with 64 directions (b = 1000 s/mm2) in addition to a single non-diffusion weighted volume (TR = 11.7 s, TE = 90 ms, 63 slices, voxel size = 2 × 2 × 2 mm, flip angle = 90°); a Fluid-attenuated inversion recovery (FLAIR) scan (TR = 9 s; TE = 94 ms, 27 slices, voxel size = 0.4 × 0.4 × 4 mm, TI = 2500 ms; flip angle = 150°) and an arterial spin labelling (ASL) dataset (PICORE Q2TIPS, 50 pairs of control/tag images, one calibration image, TR = 2.5 s / TE = 11 ms, inversion time = 1.8 s, bolus duration = 700 ms, voxel size 3.0 × 3.0 × 6.0 mm, 14 slices, flip angle = 90°). Details about the number of utilised scans per modality can be found in Supplementary Fig. 1.

2.4. Image preprocessing

2.4.1. Cortical thickness

The T1-MPRAGE scans were analysed using the automated longitudinal stream of Freesurfer version 6.0 (http://surfer.nmr.mgh.harvard.edu/) pipeline. The full details have been previously described (Reuter et al., 2012, 2010). Briefly, the pipeline applies motion correction, skull stripping, correction for field inhomogeneities, registration to and from the Talairach coordinate space, cortical parcellation and subcortical structure segmentation (Fischl, 2012). Three subjects were excluded due to failure of preprocessing steps of the Freesurfer recon-all pipeline (i.e. excessive misclassification of pial and white matter surfaces or severe skull stripping errors).

2.4.2. MNI normalisation

The individual T1-MPRAGE images were segmented with SPM12 to generate grey matter, white matter and CSF tissue probability maps. The generated grey matter images were inspected and when necessary manually corrected when parts of dura or residual skull were erroneously included in the grey matter cluster. Subsequently, the DARTEL pipeline was used to create a study specific template based on the grey and white matter masks from both study time-points. DARTEL makes use of non-linear diffeomorphic registration to register individual grey matter masks to the study-specific template and subsequently an affine registration to register the study specific template to the MNI space (Ashburner, 2007). The derived flow fields were used in the subsequent analyses to spatially normalise the T1 MRI to the MNI space. The grey matter mask for subsequent analyses was derived from the average grey matter maps and thresholded at 25% probability.

2.4.3. Perfusion measures

ASL post-processing was performed using FSL’s Bayesian Inference for ASL MRI (BASIL) toolbox (Chappell et al., 2009). The acquired ASL
scans were motion corrected using FSL’s MCFLIRT and calibrated based on a proton density acquisition. Spatial regularization was applied prior to cerebral blood flow (CBF) calculation. CBF was quantified using the Buxton ASL kinetic model (Buxton et al., 1998) according to the model suggested in the ASL white paper (Alsp et al., 2015). The generated CBF images were corrected for the presence of partial volume effects using FSL’s NILM’s adaptive spatial prior approach (Chappell et al., 2011). To proceed to partial volume correction (PVC), the MPRAGE images were registered to the ASL calibration scan using FSL’s FLIRT. The same transformation was applied to coregistere the high-resolution partial volume maps to the ASL resolution. PVC grey matter perfusion maps and grey matter CBF mean values were recorded for voxels with grey matter > 10%. Subsequently, the grey matter CBF maps were registered to the T1 images using the inverse transform generated in the previous step and normalized to the MNI space using SPM’s DARTEL. To proceed to voxel-wise comparisons a grey matter CBF mask was created for each subject including voxels with at least 5 ml/100 g/min. All individual masks were summed up and a threshold of 95% was used for voxel retention. Finally, this mask was multiplied by the grey matter mask generated in the previous step. Consistent with previous methodology (Dolui et al., 2017), the temporal SNR (tSNR) of the ASL acquisition was determined as follows: the time-series were motion corrected using FSL MCFLIRT. Subsequently, individual perfusion weighted images (control-labelled) were generated for all 50 repetitions. A mean value for each of the 50 perfusion weighted images within areas of grey matter >10% was calculated, and tSNR was measured as the mean over the standard deviation of these values (2.11 ± 0.9).

2.4.4. White matter microstructure

The diffusion datasets were visually inspected for optimal coverage and to ensure minimal eddy-current distortions by 2 authors (EM and EMK). 9 subjects were excluded due to excessive EPI distortions, while 1 subject was excluded due to poor field-of-view coverage. DWI data were stripped of nonbrain tissue using the Brain Extraction Tool (BET). Eddy currents and head movements were corrected with “eddy” in FSL (Version 6.0.1). Quantitative identification of slices with signal loss was performed in “eddy” and these volumes were replaced by non-parametric predictions using the Gaussian process (Andersson et al., 2016). In addition, we also corrected for intra-volume movement (Andersson et al., 2017). The b-matrix was subsequently reoriented by applying the rotational part of the affine transformation used during eddy correction (Leemans and Jones, 2009). Diffusion tensors were generated from the eddy-corrected DWI datasets using DTIFIT in FSL 6.0.1. We pursued an unbiased longitudinal approach for capturing normalisation of diffusion datasets using the Diffusion Tensor Imaging Toolkit (DTI-TK). Firstly, tensor images at baseline and follow-up were created by converting the diffusion eigenvectors and eigenvalues using the function fsl_to_dttk in DTI-TK. Secondly, for each participant, an unbiased intra-subject template was generated using the tensor datasets from his or her baseline and follow-up after an iterative process of rigid, affine and non-linear diffeomorphic registrations. Next, the within-subject templates were used to create a study-specific population template using the same iterative process. Each participant’s DWI data was normalised to the group-specific template via a single interpolation step. Fractional anisotropy (FA), mean diffusivity (MD) and radial diffusivity (RD) maps were generated using the normalised tensor images. Following the tensor-based registration, a mean FA map was derived from the group template and skeletonised using tbsk skeleton in FSL to create the mean FA skeleton (thresholded at FA > 0.2). FA, MD and RD images from each participant were projected onto the mean FA skeleton by searching for maximum FA values perpendicular to the skeleton. To complement our voxel-wise analyses, mean global FA, RD and MD values were extracted at baseline and follow-up from each participant’s white matter skeleton using the fslnetx command.

2.4.5. Quantification of WMH

WMH lesion maps were obtained using an automated script on the Statistical Parametric Mapping 8 (SPM8) suite (http://www.fil.ion.ucl.ac.uk/spm/) on FLAIR MRI; details on the procedures involved have been described previously (Firbank et al., 2003; Smart et al., 2011). All lesion maps from both baseline and follow-up visits were reviewed by a single experienced rater (AL) while blinded to all clinical information. Lesion maps obtained from the segmentation procedure were used as starting points for manual WMH delineation. Baseline and follow-up FLAIR images were evaluated side by side during delineation to ensure consistency. WMH volumes were normalised by total intracranial volume to account for individual differences in head size. To generate a descriptive voxelwise map representing the spatial topography of WMH occurrence in our sample, we normalised the binarized lesion maps to the MNI152 template using the forward transformations from T1-MPRAGE to MNI152 space. Next, the normalised lesion maps were concatenated using false. A mean image was created to represent the lesion probability map, in which the intensity of each voxel represents the likelihood of WMH occurrence across our sample.

2.5. Statistical analyses

A summary of the key imaging outcomes is illustrated in Fig. 1. To characterise brain imaging changes in relation to EYO, we used all available longitudinal datasets for all participants. Demographic variables were compared between the APOE-ε4 groups using t-tests, Wilcoxon Rank sum tests, chi-square tests where appropriate. To increase robustness of our correlations, EYO was transformed with the inverse normal transformation due to its skewed distribution, increasing the robustness of correlations (Miller et al., 2016). Voxelwise associations of EYO with imaging changes were investigated using the longitudinal Sandwich Estimator (SwE) toolbox (http://www.nisox.org/Software/SwE) (Guillaume et al., 2013). This marginal model describes expected variability as a function of predictors in a design matrix, while additionally accounting for correlations due to repeated measurements and unexplained variations across individuals. The main models featured the main effects of EYO as well as its interactions with time (i.e. centred within-subject age of participant). The latter metric allowed us to assess whether a hypothetical proximity to dementia may have an accelerating effect of imaging changes over time, while the main effect of EYO indicates whether a hypothetical proximity to dementia is related to imaging differences across the sample, independent of other covariates. The covariates included are (i) the demeaned mean age of a subject over both time points, (ii) the demeaned quadratic term of age to account for potential non-linear age associations (Kodiweera et al., 2016; Salami et al., 2012), (iii) sex, (iv) demeaned years of formal education and time. The demeaned average WMH volume and its interaction with time were included as additional covariates in all DTI analyses to ensure that white matter changes are not confounded by small vessel disease. In addition, we examined both potential moderating and interactive effects of APOE-ε4 by modelling the main effects of APOE-ε4 and its interactions with EYO and time into the SwE models. Non-parametric statistical inference was based on the Wild Bootstrap procedure (Guillaume et al., 2015), and corrections for multiple comparisons were performed with Threshold-Free Cluster Enhancement (TFCE) and Family-Wise Error Rate (TFCE P_FWER < 0.05) (Smith and Nichols, 2009). To report the coordinates of significant clusters in standard MNI152 space, the mean FA of the group-specific template was affine registered to the MNI152 atlas using FLIRT and visually inspected. The affine transformation was then applied onto the statistical maps for anatomical localisations in MNI152 space. To eliminate noise due to averaging and normalization, significant clusters with < 50 voxels were excluded from the table of results. Non-voxelwise analyses of global FA, MD, RD, mean grey matter perfusion, and mean cortical thickness were performed using Linear Mixed Effect models (LME), implemented with the lmer function of the lme4.
package in R (Bates et al., 2015). Consistent with our SwE models, the LME models included a random intercept for each subject, and fixed effects of demeaned averaged age, demeaned quadratic age, sex, demeaned years of education and time (i.e. the demeaned within-subject age). WMH volumes were also included as additional covariates in our analyses on FA, MD and RD.

As the EYO was derived based on the participant’s age, both measures were expected to be correlated ($r = -0.3$, $p < 0.01$; Supplementary Fig. 2). We also took extra caution to verify the absence of multicollinearity in our models by calculating variance inflation factors (VIF) using the CAR package (Fox and Monette, 1992; Fox and Weisberg, 2019). Finally, two sets of sensitivity analyses were conducted to examine the robustness of the significant EYO associations against chronological age. First, we repeated all significant LME and SwE models while substituting the EYO variable with age. Secondly, to compare the relative contributions of age, EYO and APOE-ε4 status on imaging changes, we conducted group comparisons of imaging data between median-split age groups, median-split EYO groups, and APOE-ε4 status using ANCOVA.

2.6. Data availability

Data will be made available upon reasonable requests and for non-commercial reasons. The statistical maps, group templates and other relevant files are available to view on Neurovault (https://identifiers.org/neurovault.collection:9099).

3. Results

3.1. Sample characteristics

Demographic details about the sample with diffusion imaging datasets are summarised in Table 1. The EYO ranged from 38 years before the parental onset to $-3$ years after onset. There were no significant differences with regards to age, gender, education years or ACER between the APOE-ε4 carriers and non-carriers.

| Table 1 | Participant characteristics of subjects with diffusion weighted imaging datasets. |
|---------|---------------------------------|
| Age (years) | APOE-ε4 – (N=42) | APOE-ε4 + (N=37) | P-value |
| Mean (SD) | 53.4 (4.34) | 51.8 (4.83) | 0.086 |
| Median [Min, Max] | 54.0 [41.0, 60.0] | 52.0 [40.0, 60.0] |  |
| Gender | | | |
| Female | 31 (73.8%) | 27 (73.0%) | 0.99 |
| Male | 11 (26.2%) | 10 (27.0%) |  |
| Education (years) | | | |
| Mean (SD) | 15.7 (2.92) | 15.7 (3.12) | 0.561 |
| Median [Min, Max] | 16.0 [11.0, 22.0] | 17.0 [11.0, 21.0] |  |
| ACE-III | | | |
| Mean (SD) | 94.3 (4.73) | 95.1 (3.93) | 0.644 |
| Median [Min, Max] | 95.0 [80.0, 100] | 97.0 [82.0, 100] |  |
| Missing | 2 (4.8%) | 0 (0%) |  |
| ACE-III Memory | | | |
| Mean (SD) | 24.6 (2.07) | 24.8 (1.91) | 0.974 |
| Median [Min, Max] | 26.0 [18.0, 26.0] | 25.0 [18.0, 26.0] |  |
| Missing | 2 (4.8%) | 0 (0%) |  |
| Time interval (years) | | | |
| Mean (SD) | 2.00 (0.122) | 2.01 (0.131) | 0.957 |
| Median [Min, Max] | 2.00 [1.72, 2.32] | 1.98 [1.86, 2.46] |  |
| EYO (years) | | | |
| Mean (SD) | 24.6 (6.73) | 23.5 (7.86) | 0.345 |
| Median [Min, Max] | 26.0 [4.00, 33.0] | 25.0 [-3.00, 38.0] |  |

Abbreviations: ACE-III = Addenbrooke’s Cognitive Examination, EYO = Estimated years to onset.

3.2. Reliability of imaging data across time

The sjstats package in R was used to calculate the intra class coefficients. ICC was calculated as the proportion of between-subject variance to total variance. The dependent variable for these models was the global imaging measurement for each subject on two separate visits, fixed effects included the average age and time. Subject-specific intercept was included as a random effect. Across all imaging measurements,
ICC was moderate to excellent (Mean cortical thickness: 0.87, FA: 0.73, MD: 0.74; RD: 0.74; ASL: 0.55).

3.3. Grey matter atrophy

There were no significant main effects of EYO on regional cortical thickness and hippocampal volumes that survived false discovery rate correction for multiple comparisons. No regions showed a significant interaction between EYO × APOE4 status, or three-way interactions between EYO × Time × APOE4 status after false discovery rate correction for multiple comparisons.

3.4. White matter microstructural changes

The association between white matter changes and EYO are illustrated in Fig. 2. There were no significant effects of APOE-ε4 status on white matter changes in terms of FA (T = -1.13, p = 0.26), MD (T = 1.11, p = 0.27) or RD (T = 1.16, p = 0.25). LME models showed significant main effects of EYO on FA (T = 2.58, p = 0.01), MD (T = -2.55, p = 0.01) and RD (T = -2.66, p = 0.01). There were no significant interactions between time and EYO (T = -1.12-1.43, p = 0.16-0.43). Voxelwise SwE revealed widespread and significant main effects of EYO on FA, MD, and RD maps, after adjusting for averaged age, age², gender, education years, averaged white matter hyperintensity volume, APOE-ε4 status, and within-subject age (TFCE FWER p < 0.05). The EYO-associated white matter changes were predominantly found in the forceps minor of the corpus callosum, inferior longitudinal fasciculi, superior longitudinal fasciculi, cingulum bundles, anterior thalamic radiation, inferior fronto-occipital fasciculus, uncinate fasciculus (see Supplementary Table 3 for MNI152 coordinates of peak clusters).

3.5. White matter hyperintensities

We illustrated the baseline WMH incidence, voxel-wise, across our sample and revealed the expected spatial distribution of WMHs in normal ageing (Fig. 4). The highest probabilities of WMH were concentrated around the periventricular areas and in deep white matter regions. There were no significant main effects of EYO (T = -0.74, p = 0.46) or EYO – Time interactions (T = -1.06, p = 0.29) on WMH volumes.

3.6. Cerebral perfusion

LME and SwE voxel-wise analyses revealed a significant interaction of EYO with APOE-ε4 on grey matter perfusion (LME: T = -2.87; p = 0.01; SwE: TFCE FWER p < 0.05) such that APOE-ε4 carriers showed increased perfusion with nearer proximity to onset while APOE-ε4 non-carriers did not (Fig. 5). The difference in effects between APOE-ε4 carriers and non-carriers was qualified by a significant interaction term (LME: T = -2.87; p = 0.01). Simple slope analyses and slope difference tests further confirmed a significant difference in the relationship between EYO and ASL for the APOE-ε4 carriers and non-carriers, such that a significant association was found only amongst the APOE-ε4 carriers (Slope analysis: T = -3.27; p < 0.01) and not for the non-carriers (Slope analysis: T = 0.74; p = 0.462). Similarly, SwE voxel-wise analyses revealed that the EYO × APOE-ε4 interaction was localised within tempo-parietal regions, such as the hippocampus, precuneus/posterior cingulate cortices (TFCE FWER p < 0.05; see Supplementary Table 4 for MNI152 coordinates of peak clusters). There were no significant three-way interactions between EYO, APOE-ε4 and time.

3.7. Cross-modal relationships between ASL and DTI data

Scatter plots of the relationships between grey matter perfusion and DWI metrics are shown in Fig. 6. Post-hoc robust linear regressions indicated a significant negative correlation of FA (T = -2.95, p < 0.01) and positive correlations of MD (T = 2.29, p = 0.03), and RD (T = 2.66, p = 0.01) with grey matter perfusion after adjusting for age, age², gender, education years, APOE-ε4 status, and WMH volumes.

3.8. Sensitivity analyses

3.8.1. Substituting EYO with age

Given that age is correlated with EYO, we performed two sets of sensitivity analyses to evaluate the age-independence of the EYO results. First, the EYO term was substituted by the chronological age of the participant. These analyses are informative in terms of showing whether the spatial extent of proximity-related white matter changes could be recapitulated from the participant’s age itself. First, age was not significantly correlated with global mean FA (T = -1.88, p = 0.06), MD (T = 1.31, p = 0.19) or RD (T = 1.64, p = 0.11). Akaike Information Criteria analysis also indicated that including EYO resulted in a better model fit compared to a reduced model with age alone (p = 0.007). Second, in our voxelwise SwE models without the EYO term, older age was only associated with lower FA but not MD or RD (TFCE FWER p < 0.05). The spatial extent of the age-FA associations was markedly smaller to that of the EYO (Dice volume overlap = 17.4%). The voxels that are unique to either EYO, age, or overlapping effects are illustrated in Fig. 7, showing that EYO is sensitive to microstructural changes over and beyond that of normal ageing. Similarly, in contrast to EYO, there was no
significant interaction between age and APOE-ε4 status on mean grey matter perfusion ($T = 0.46, p = 0.65$).

### 3.8.2. Group comparisons of white matter across binarized subgroups

We also attempted to disentangle the contributions of ageing, APOE-ε4 and EYO through a series of group comparisons (Figure 8). The dependent variable was the imaging metric at baseline (i.e. FA). The categorical covariates are age groups (Age $\geq 53$ and Age $< 53$), APOE4 (Carriers and Non-Carriers), and EYO proximity groups (EYO $\geq 25$ and EYO $< 25$). WMH volumes were also included to account for white matter alterations due to small vessel diseases. This parsimonious stratification of the sample enabled a comparison of the relative impact of each risk factor on imaging changes. ANOVA indicated a significant main effect that was exclusive to EYO ($F = 6.52, p = 0.01$). Furthermore, the effect size was the largest for EYO (Cohen $f = 0.3$) relative to age (Cohen $f = 0.06$) and APOE-ε4 (Cohen $f = 0.09$). The predominant effect of EYO was also evident in MD and RD analyses.

### 4. Discussion

Our study revealed an assortment of brain changes as cognitively normal FH+ individuals approach their expected onset of dementia in midlife. A shorter hypothetical proximity to dementia was associated with widespread white matter disruption (↑ FA ↑ MD ↑ RD), in the absence of grey matter atrophy. In addition, Moreover, our data showed that hyperperfusion with shorter EYO was strongly predicated on the presence of APOE-ε4 status.

We did not find a robust effect of EYO on regional grey matter atrophy. This finding was not entirely unexpected, as cortical atrophy detected on MRI is widely believed to represent the terminal end-product of neuronal loss, cell shrinkage, reductions in dendritic extent and synaptic loss (McEwen, 1997). In contrast, extensive white matter deficits were revealed as FH+ individuals approached their estimated age of onset, involving major white matter bundles that are characteristically compromised in mild cognitive impairment and AD (Acosta-Cabronero et al., 2010). Cross-sectional studies in late-onset AD and familial AD have demonstrated preferential vulnerability of the corpus callosum, the superior and inferior longitudinal fasciculi, and cingulum bundles of the limbic projections, and our findings extend those reports by documenting a similar profile of white matter degeneration in a cognitively normal cohort of midlife adults. To the extent that EYO is a valid stage marker of preclinical disease progression, it is conceivable that the pattern of white matter deficits in our data may reflect incipient neuropathologies (i.e. amyloid or early build-up of neurofibrillary tau tangles). While the EYO correlations preclude causal interpretations, our hypothesis is aligned with a recent study that included both ante-mortem DTI and histological analyses in cognitively normal subjects (Kantarci et al., 2017). In that study, Kantarci et al., demonstrated associations of temporo-parietal white matter tracts with amyloid burden and tau accumulation (Kantarci et al., 2017). Despite the lack of a
prior finding in the FH+ literature, our findings are corroborated by recent data from the Dominantly Inherited Alzheimer’s Network (DIAN) project, which found elevated MD with shorter EYO (Caballero et al., 2018). Interestingly, there were pronounced EYO-related changes in the genu of the corpus callosum, coinciding with the striking convergence of proximity-related FA, MD, and RD changes in our sample (Fig. 3).

In contrast to the anticipated trajectory of ↓ FA ↑ MD ↑ RD on white matter tracts, a shorter EYO was accompanied by increased perfusion particularly amongst APOE-ε4 carriers. Hyperperfused regions encompassed the temporo-parietal regions (i.e. precuneus, posterior cingulate, medial temporal regions and hippocampus). At first glance, these results may appear to contradict the wealth of evidence showing ~40% global deficit of CBF in people with AD in the same aforementioned regions (Alsop et al., 2010; Wierenga et al., 2014). However, our data are also aligned with other reports of hyperperfusion amongst APOE-ε4 carriers (Bangen et al., 2012; Wierenga et al., 2013): increased glucose uptake from FDG-PET studies have been found in FAD mutation carriers as early as 25 years before expected onset (Benzinger et al., 2013).
while Alsop et al. have reported hippocampal hyper-perfusion in cases with AD (Alsop et al., 2008). Taken together, our current perfusion data are consistent with a model of disease progression in the asymptomatic period of dementia, in which APOE-ε4 carriers display elevated cerebral perfusion very early during the presymptomatic phase as a possible attempt to maintain an adequate blood supply at rest. Hyperperfusion could also be a response to neuroinflammation (Alsop et al., 2008), alterations in the capillary transit time (Østergaard et al., 2013) or associated with amyloid accumulation (Fazlollahi et al., 2020). Another possible explanation is that the higher ASL signal is driven by delayed intravascular signal. However, our subjects were all healthy participants hence we do not expect large differences in the arterial transit time. The inversion time used in our protocol is also aligned with previous recommendations (Alsop et al., 2015). In a study by Wierenga et al. (2013) the interaction of age and APOE-ε4 was investigated in relation to CBF and it was found that young APOE-ε4 (mean age: 24) demonstrated hyper-perfusion, whereas older APOE-ε4 carriers (mean age: 75) demonstrated hypo-perfusion compared to non-carriers. Some plausible explanations for the observed associations include the (a) uncoupling of the flow from metabolic demands in young adults, in turn leading to cerebrovas-
cular dysregulation, (b) a differential effect of APOE-ε4 on neurovascular function in aging, (c) and hypo-perfusion as a result of amyloid accumulation or even underlying structural alterations in the vasculature in older carriers. These hypotheses are especially pertinent to this study, considering the intermediate mean age of our cohort relative to the aforementioned age groups (Wierenga et al., 2013). Future studies with additional measurements of cerebral hemodynamics using arterial transit time sensitive ASL acquisitions or diffusion weighted ASL may be well-suited to reveal further insights into the hemodynamic changes and identify the inevitable inflection point associated with the emergence of the AD-signature pattern of hypoperfusion.

A key strength of the present study is the integration of multi-modal MR imaging, enabling several inferences regarding the evolution of distinct pathological mechanisms along the preclinical spectrum. First, the coupling of EYO with DTI instead of cortical thickness provides further support to the argument that microstructural changes – typically not visible on T1-MPRAGE – predate macroscale neurodegeneration/brain atrophy, and thus could predict subsequent cognitive decline with greater sensitivity (Kantarci et al., 2005; Müller et al., 2005; Ringman et al., 2007; Weston et al., 2015). Secondly, against the backdrop of widespread white matter degeneration, it is tempting to speculate that elevated perfusion may reflect the harnessing of greater neural resources to help circumvent the deteriorating efficiency of cognitive networks. There is some tangential support for this hypothesis from a recent study showing a paradoxical relationship between elevated cerebral perfusion and increased amyloid burden in a cognitively normal elderly cohort (Fazlollahi et al., 2020). Likewise, a compensatory hypothesis would have the corollary that elevated cerebral perfusion is accompanied by microstructural impairments (↓ FA ↑ MD ↑ RD), as was confirmed in our post-hoc cross-modal associations between DTI and ASL. Interestingly, a recent study found the inverse relationship in patients with chronic traumatic brain injury (TBI), but not in those with less remote TBI (Clark et al., 2017). Taken together, these findings provide preliminary evidence for a dynamic association between CBF and white matter integrity that may exert deleterious effects on cognitive outcomes.

Fig. 8. Proximity to dementia is the predominant contributor to white matter changes compared to age and APOE-ε4 status. FH+ individuals who are less than 25 years to estimated onset showed significantly reduced white matter FA compared to FH+ individuals who are ≥ 25 years away from estimated onset. EYO group had the largest effect size relative to age group and APOE-ε4 status. Abbreviation: FA = Fractional Anisotropy, EYO = Estimated years to onset.

E. Mak, M.-E. Dounavi, A. Low et al. NeuroImage 229 (2021) 117749
Further validation of the EYO in larger samples of cognitively normal APOE-ε4 carriers/FH+ is warranted and will be performed as more subjects in PREVENT-Dementia complete their follow-up assessments.

Credit author statement

EM and MD performed the imaging analysis, conducted the statistical analysis, and drafted the manuscript. GTM provided statistical advice and critical feedback on the manuscript. AL, SFC, EMK, GBW, PSJ, JC, KR, CR, LS and JOB reviewed and provided critical feedback on the manuscript. CR, JOB, LS also helped obtain funding, supervised this study, was involved with its design.

Data availability

Data will be made available upon reasonable requests and for non-commercial reasons. The statistical maps, group templates and other relevant files are available to view on Neurovault (https://identifiers.org/neurovault.collection:9099).

Financial disclosures

John T. O’Brien has no conflicts related to this study. Unrelated to this work he has received honoraria for work as DSMB chair or member for TauRx, Axon, Eisai, has acted as a consultant for Roche, has received research support from Alliance Medical and Merck. Elijah Mak, Maria-Eleni Dounavi, Stephen Carter, Audrey Low, Li Su, Elizabeth McKiernan, Karen Ritchie, Graciela Muniz, Isabelle Carriere, Guy Williams, Simon Jones and Craig Ritchie have no conflicts of interest to declare.

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Supplementary materials

Supplementary material associated with this article can be found, in the online version, at doi:10.1016/j.neuroimage.2021.117749.

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