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Age-related changes of Peak width Skeletonized Mean Diffusivity (PSMD) across the adult life span: a multi-cohort study

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36

37 Abstract

Parameters of water diffusion in white matter derived from diffusion-weighted imaging (DWI), 38 such as fractional anisotropy (FA), mean, axial, and radial diffusivity (MD, AD and RD), and 39 40 more recently, peak width of skeletonized mean diffusivity (PSMD), have been proposed as 41 potential markers of normal and pathological brain ageing. However, their relative evolution 42 over the entire adult lifespan in healthy individuals remains partly unknown during early and late adulthood, and particularly for the PSMD index. Here, we gathered and meta-analyzed 43 44 cross-sectional diffusion tensor imaging (DTI) data from 10 population-based cohort studies in order to establish the time course of white matter water diffusion phenotypes from post-45 46 adolescence to late adulthood. DTI data were obtained from a total of 20,005 individuals aged 47 18.1 to 92.6 years and analyzed with the same pipeline for computing DTI metrics. For each individual MD, AD, RD, and FA mean values were computed over their FA volume skeleton, 48 PSMD being calculated as the 90% peak width of the MD values distribution across the FA 49

50 skeleton. Mean values of each DTI metric were found to strongly vary across cohorts, most 51 likely due to major differences in DWI acquisition protocols as well as pre-processing and DTI 52 model fitting. However, age effects on each DTI metric were found to be highly consistent across cohorts. RD, MD and AD variations with age exhibited the same U-shape pattern, first 53 54 slowly decreasing during post-adolescence until the age of 30, 40 and 50, respectively, then 55 progressively increasing until late life. FA showed a reverse profile, initially increasing then continuously decreasing, slowly until the 70's, then sharply declining thereafter. By contrast, 56 57 PSMD constantly increased, first slowly until the 60's, then more sharply. These results 58 demonstrate that, in the general population, age affects PSMD in a manner different from that of other DTI metrics. The constant increase in PSMD throughout the entire adult life, including 59 60 during post-adolescence, indicates that PSMD could be an early marker of the ageing 61 process.

62 1 Introduction

63 Parameters of water diffusion in white matter derived from diffusion-weighted imaging (DWI), 64 such as fractional anisotropy (FA), mean, axial, and radial diffusivity (MD, AD and RD) are 65 well-established markers of normal brain maturation(1)(2)(3)(4)(5)and ageing 66 (6)(7)(8)(9)(10)(11)(12) and have been proposed as potential tools for the investigation of various brain disorders (13)(14)(15)(16)(17)(18)(19). 67

68 More recently, peak width of skeletonized mean diffusivity (PSMD, (20)), a new phenotype of 69 white matter microstructure that can be derived from DWI, has been proposed as an imaging 70 biomarker of small vessel disease (SVD, (20)(21)) and a correlate of cognitive impairment, 71 particularly processing speed (20)(22)(21). So far, our knowledge of the PSMD distribution in 72 healthy individuals has been limited to these three previously mentioned studies that all 73 included people aged over 50 years. In addition, none of these studies addressed the issue of changes in PSMD across lifespan, which is critical for establishing whether PSMD could 74 75 be used as an imaging marker of brain aging as well as an early predictor of age-related disorders or to serve as a tool to monitor outcomes in clinical trials. Here, we gathered and 76 77 meta-analyzed cross-sectional diffusion tensor imaging (DTI) data from 10 population-based 78 cohort studies in order to establish the time course, from post-adolescence to late adulthood, 79 of the PSMD distribution and compare it with that of more commonly used white matter water 80 diffusion phenotypes in white matter.

81 2 Materials and Methods

82 2.1 Participants

83 Ten independent datasets coming from cross-sectional cohort studies were gathered in the present study, namely MRi-Share, BIL&GIN, SYS, LIFE-Adult, 1000 BRAINS, UKBiobank, 84 85 ASPSF, OATS, LBC1936, MAS; (see acronym definition in Table 1 caption). All but three 86 (LIFE-Adult, 1000BRAINS, and UKBiobank) were part of the BRIDGET Consortium (BRain 87 Imaging, cognition, Dementia and next generation GEnomics: a Transdisciplinary approach to search for risk and protective factors of neurodegenerative disease), supported by EU-88 89 JPND (European Union Joint Programme – Neurodegenerative Disease Research). The 10 datasets included a total of 20,005 individuals (age range: 18.1 to 92.6 years, 10,807 women 90 91 and 9,198 men). Tables 1 and 2 detail sample size and age distribution for the 10 cohorts that 92 were all of cross.

93 2.2 Diffusion-weighted image acquisition and preprocessing

Tables 3 and 4 summarize the main characteristics of the DWI acquisition and preprocessing 94 95 for the 10 cohorts. Overall, there was considerable variability between studies regarding almost all acquisition parameters, including scanner manufacturer, field strength, gradient 96 97 strength, diffusion pulse sequence, resolution and number of directions. For the present work, it was not possible to access raw DWI data at different sites in order to harmonize processing 98 from the initial DICOM data. For this reason, DWI datasets were pre-processed with 99 100 procedures specific to each site, including exclusion of data upon visual detection of major 101 artifacts due to eddy current distortions or head motion. AD, RD, MD and FA maps were 102 computed by fitting the DTI model parameters in each voxel from these preprocessed DWI 103 volumes. Additional details on DWI preprocessing and DTI parameter map computation for 104 each dataset are provided in the Supplementary Material section.

105 2.3 Derivation of DTI metrics

106 Various metrics were derived from the DTI data using a script developed by Baykara et al.
107 (<u>http://www.psmd-marker.com, (20)</u>). This original script was designed to extract PSMD, an
108 index of the dispersion of MD values across the white matter skeleton. Briefly, the
109 computation included two steps: 1-WM skeletonizing using the FA map, and 2- analyzing the

110 voxel value distribution histogram in the MD volume masked by the WM-FA skeleton. The FA volume of each individual was skeletonized using the FSL-TBSS software, part of the FMRIB 111 112 Software Library (FSL) (23)(24), using the FMRIB 1mm FA template and applying a 0.2 threshold on FA maps. Then the MD volume of the same individual was masked, keeping 113 114 only voxels within the FA skeleton. Furthermore, in order to reduce contamination of the 115 skeleton by CSF voxels, the FA-masked MD volumes were further masked by both a standard FA skeleton with a threshold of 0.3 on FA values and a custom mask (provided with the PSMD 116 software tool) designed so as to exclude regions adjacent to the ventricles, such as the fornix. 117 Finally, PSMD was computed as the difference between the 95th and 5th percentiles of the so-118 masked MD volume voxel value distribution. Here, we extended this script in order to obtain, 119 120 in addition to PSMD values, estimates of the mean values of axial, radial, and mean diffusivity 121 (AD, RD, MD, respectively) as well as of fractional anisotropy (FA) over the same customized 122 skeleton. All 10 cohorts were processed separately with this customized script and the results sent to the Bordeaux site where they were combined for further statistical analysis. 123

124 2.4 Statistical analyses

125 2.4.1 Age category definition

Due to previously reported non-linear effects of age on DTI metrics (1)(3)(8), we divided each cohort sample into subsamples of 10-years age range starting at 18 years of age, the last subsample (i.e. [78 to 98]) including all subjects aged over 78 years, as there was only a small number of individuals aged over 88 years. Table 1 and Figure 1 detail the contribution of each cohort to each age category. Because we planned running analyses at the age category by cohort by sex level, we discarded subsamples of small sizes, namely containing less than 30 individuals.

133 2.4.2 Assessing age-related changes of PSMD and other DTI metrics

For each of the five DTI metrics and each age category, we performed an analysis of variance including "Age" as the main effect, and "Sex", "TIV" (total intracranial volume), and "Cohort" as confounding factors. The Cohort effect was included in order to account for apparent large differences in DTI metric average values across cohorts contributing to the same age category dataset (see Figure 2). In order to document differences of age effects between cohorts contributing to the same age category, we also performed an analysis of variance for

140 each age category and each cohort, including "Age", "Sex" and "TIV" as effects. Moreover,

141 we analyzed the effects of age on DTI metrics in men and women separately.

142 2.4.3 Assessing the effects of Sex and TIV on PSMD and other DTI metrics

143 For each of the five DTI metrics and each age category, we also performed an analysis of144 variance including "Sex" and "TIV" as main factors and "Cohort" as confounding factors.

145 All statistical analyses were performed using the JMP Pro Software (version 14.3.0, SAS146 Institute Inc.).

147 3 Results

148 3.1 Descriptive statistics

Tables 3a to 3e provide basic statistics across age categories and cohorts for PSMD and the four other DTI metrics, while Figure 2 illustrates their respective profiles across the adult lifespan. From both Table 3 and Figure 2 it is apparent that there is considerable variability in all five DTI parameter values, and that within a given age category the variability across cohorts is larger than the variability between individuals of the same cohort (see the extreme case of AD and FA values for the LBC1936 study performed at 1.5T, for example).

Figure 3 compares the inter-individual variability of PSMD (in terms of its coefficient of variation, CV in %) within each cohort to those of the other DTI metrics, again for each age category and each cohort, revealing that PSMD CV's is in the order of 10 to 15% (with values as high as 20% for later ages) while those for AD, RD, MD and FA are in the order of 2 to 5%. Note also that the CV's of all DTI metrics increase with age, more for PSMD than for the other parameters.

161 3.2 Effects of age on PSMD and other DTI metrics

The evolution of PSMD across the adult life is different from that of the other metrics (Figure 2). Indeed, PSMD seems to increase monotonically with age whereas AD, RD, and MD exhibit a similar J-shape profile, initially slightly decreasing during post-adolescence before later increasing during adulthood. As for FA, it shows a reverse profile to that of AD, RD and MD, with an initial small increase followed by a later decrease.

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167 This apparent specific lifespan profile of PSMD was confirmed by the quantitative estimates of the effects of age on PSMD and other DTI metrics provided by the between-cohort ANOVA 168 169 (see Table 4) and their profiles of evolution across age categories as shown in Figure 4. 170 Estimates of the age effect on PSMD were indeed positive for all age categories and 171 significant for all but the [28 to 38] age period. This increase in PSMD accelerated during late 172 life periods, its value being multiplied by a factor of 3 between the [58 to 68] and the [78 to 173 98] periods. By contrast, AD, RD, and MD age variation profiles were characterized by an 174 initial small but significant decrease (negative age effect), followed by a stable period (non-175 significant age effect) and a final significant increase (positive age effect). What distinguished 176 AD, RD and MD profiles was their respective timings, the initial decrease in RD and MD being 177 significant only for the [18 to 28] subsample, while it extended over the [38 to 48] period for 178 AD. Note that since the stable period covered two periods of 10 years for all three metrics, 179 the increase in AD was delayed by 10 years compared with RD and MD. At the same time, 180 FA exhibited the reverse contrast consisting of an initial increase during the [18 to 28] period 181 followed by a stable period ([28 to 38]) before an accelerated decrease during the rest of the 182 adult life course. The age effects on DTI metrics were not significantly different between men 183 and women as revealed by the separate sex-specific analyses of age effects.

184 3.3 Effects of Sex and TIV on DTI

Amplitude of sex effects on DTI metric average values were found to be quite variable across the various cohorts for the different age categories (see Figure 5). When pooling all datasets, we found that sex had significant effects on all DTI parameters except on FA. Women had higher mean AD and MD values than men, who conversely had higher PSMD values than women (see Table 5). As for TIV, we found that it had positive effects on all DTI parameters (see Table 5), including PSMD, these effects being very significant (p < 10^{-4}) in all cases but RD (p=0.53).

192 4 Discussion

We will first discuss methodological issues and potential limitations in interpreting results of our study. Second, we will discuss what the present study adds to the already existing literature on age effects on classical DTI parameters. In the third part, we will focus on the original findings regarding PSMD distribution and evolution over the adult life.

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197 **4.1** Methodological issues and potential limitations of the present study

198 In order to study the effects of age over the entire adult lifespan, we gathered DTI data from 20,005 individuals scanned at 10 different sites with the major objective of maximizing 199 200 statistical power. By so doing, we were aware that the variability of DTI parameters derived 201 from the entire dataset would be much larger than for a single site study because of between-202 site differences in data acquisition protocols such as scanner manufacturer, field strength, 203 number and strength of diffusion-encoding gradients, image voxel size, and post-processing 204 of raw DWI data, and so on (23)(24)(25)(26)(27)(28). Accordingly, a key issue was to reduce 205 as much as possible the variability due to sources that were controllable so that we could 206 maximize benefits of the very large sample size.

207 Concerns with DTI data meta-analysis have been emphasized by several authors (29)(30) 208 and strategies have been proposed for harmonizing DWI data before proceeding to statistical 209 analysis of DTI parameter maps. However, it has also been reported that DTI parameters 210 estimated over the whole white matter or large regions of interest actually exhibit high intra-211 and inter-scanner reproducibility, making them suitable for multisite studies without extensive 212 harmonization (17)(31)(32). This is the approach that we implemented here as it was not 213 possible to access raw DWI data at different sites in order to harmonize processing from the 214 very earliest stages (see Supplementary Materials). In our study, harmonization was limited 215 to the post-processing of DTI maps pre-computed by each site using the same script for 216 generating individual skeletons and computing individual DTI parameter values across these 217 skeletons. However, our findings indicate that it is possible to draw meaningful conclusions 218 from such minimal harmonization by focusing on the effects of interest (age in our case) rather 219 than the absolute values of the measurements, an approach previously used by others (33).

Multi-site studies do have some advantages as compared with single-site investigations, namely statistical power due to very large sample size, and the ability to recruit a sufficient number of individuals over the entire life or adult lifespan. As emphasized by others, these are important features in the context of clinical studies that are often multisite in nature (17)(31). In fact, there have been several previous reports of multisite studies on DTI parameters in adults but they dealt with issues other than age effects such as methodology (26)(17)(29) and genetic effects (34)(35), for example.

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227 In this study, we focused on the mean DTI metrics over a white matter skeleton, rather than over the entire white-matter compartment, since our primary metric of interest was PSMD, 228 229 which is defined over the FSL-TBSS white matter skeleton derived from FA data. This 230 approach should in theory minimize any partial volume effects by limiting the measurement 231 over the core of white matter tracts. Indeed, when we compared the mean DTI metrics over 232 the white matter skeleton with those measured over the whole white-matter compartment in two cohorts of the study (MRi-Share and BIL&GIN), mean FA values were higher and 233 234 diffusivity values lower when averaged over the white matter skeleton than when averaged 235 over the entire white-matter compartment. However, the cross-subject variability measured 236 as the CV of the mean DTI metrics decreased only marginally (a fraction of a percent for CVs 237 ranging from 2% to 4%) when mean values were computed over the white matter skeleton 238 rather than the whole white matter. In contrast, the CV of PSMD decreased markedly (on 239 average from 24% to 8%) when its computation was performed on the white matter skeleton 240 rather than on than the global white-matter mask. This demonstrates the importance of 241 choosing a measure of MD dispersion values over a white matter skeleton for controlling 242 between subject variability.

243 In the present work, we restricted the analysis to classical DTI metrics as only two of the 244 contributing cohorts had high angular resolution and/or multi-shell acquisition schemes that 245 could be used for estimating advanced white-matter microstructural parameters with more 246 sophisticated models (25). Here, DWI data processing was solely based on the classical DTI 247 model. The DTI model limitations are well-known (24) and it has been shown, for example, 248 that correction for free water has a major impact on classical DTI parameter values (36)(37). 249 However, although investigating advanced white-matter microstructural parameters is highly 250 desirable, it was beyond the scope of our study: it would require additional datasets with multi-251 shell acquisition, especially for individuals aged 30 to 50 years or over 70 years, in order to 252 supplement existing data (8) on the adult lifespan trajectory of these microstructural 253 parameters.

We included "Sex" and "TIV" as covariates in our statistical analyses. Mixed results have been reported regarding the impact of sex on DTI measures ((38)(39)(40)(12)(41), see review in (3)). Here we also observed mixed results across the different cohorts, although very significant sex effects on all DTI parameters, except FA, were uncovered when combining the 258 entire dataset. Note, however, these sex effects were of very small size (for PSMD, for example, $\text{omega}^2 = 6.7 \times 10^{-3}$ for the sex effect to be compared with 7.6×10^{-2} for the age effect). 259 which could explain the mixed findings in the literature, and suggests further investigations 260 261 are required in order to understand their biological origins. TIV effects on DTI parameters are not well established in the literature. In our study, we found that TIV was positively correlated 262 with all DTI parameters except for RD. Similar to sex, TIV effects when significant were very 263 small (for PSMD again, omega²= 1.5×10^{-3} for the TIV effect). Here again additional 264 investigations are needed to understand the origins of these effects. 265

266 Finally, and importantly, it should be stressed that interpretation of the results of the present 267 study should be taken with caution because of the cross-sectional nature of the data that we 268 analyzed. Numerous reports have indeed pointed out the caveats of cross-sectional design 269 for assessing effects of age and demonstrated how such design may lead to spurious findings 270 when compared to those obtained with longitudinal data ((12)(42)(43)). In the present work, 271 there was no attempt to use a single model to describe the variation of DTI parameters with 272 age over the entire adulthood period. Rather, we selected a piecewise linear model to 273 examine/compare age-related changes in 10-year duration consecutive time bins, thereby 274 minimizing the generation bias between cohorts of nearby categories. Understandably, such 275 an approach does not eradicate the intrinsic limits of our cross-sectional study. But it should 276 be reminded that a fully longitudinal design is guasi impossible to implement in the context of 277 lifespan research, since, in practice, measures in an individual can be repeated only a few 278 times and at short duration intervals. As a consequence, such longitudinal studies suffer from 279 some of the limitations of cross-sectional ones. This may explain why the results of the 280 present study are compatible with those a previously published longitudinal study (12) in 281 which individuals aged between 20 and 84 years were observed twice 3 years apart.

4.2 Adult life span profiles of variation of classical DTI parameters AD, RD, MD and FA

Effects of age on white-matter microstructure assessed with DTI have been intensively investigated over the past decade from a developmental perspective (4) as well as in a lifespan/ageing framework (1)(11)(40)(3)(12)(11)(5)(8)(36). Briefly, and considering only DTI metrics estimated at the global level, AD, RD and thus MD were reported to follow similar Uor better J-shape age variation patterns, initially decreasing during childhood and

adolescence (see (4) for review) then exhibiting an accelerated increase during the adult life (12)(8), while FA followed a reverse profile. Our own findings agree with this body of results during the adult life course. Raw data plots show J-shape profiles for AD, RD and MD, and the reverse profile for FA, as well as acceleration of these changes during late life.

293 Maximum global FA values and minimum global MD, RD, and AD values have been reported 294 to occur before the age of 40 (1)(40)(11)(5)(12), although large variations were found when 295 considering individual tracts (44)(39)(45)(5). Here, we found extreme values for RD, MD and 296 FA occurring between 28 and 38 years, well in line with these previous findings. In addition, 297 we found that the decrease of AD in the post-adolescence period extended into adulthood by 298 about 10 years more than for RD and MD, thereby uncovering heterochrony of AD and RD 299 variations during adulthood. Such a heterochrony during adulthood was not detected in a 300 previous longitudinal study (12) possibly due to an insufficient sample size and large DTI 301 metric variability between individuals (as can be seen in Fig. 7 of the mentioned report). Note 302 that two recent studies (46)(47) have reported opposite age effects for AD (decrease) and 303 RD (increase) with stable MD during the 18 to 55 year age period; however, as both studies 304 used simple linear modeling due to small sample sizes, no age at extreme value could be 305 observed. Rather, our findings are compatible with the AD-RD variation heterochrony that has 306 been noticed earlier during childhood and adolescence at the individual tract level with 307 stronger decrease for RD than for AD (44)(30). According to these and our findings, the AD 308 decrease / RD increase profile (6) would occur only during mid-adulthood.

309 4.3 PSMD is a diffusion imaging phenotype with a profile of variation across the adult 310 life span that differs from that of other DTI parameters

311 The distribution of PSMD values observed for the different cohorts and age categories of the 312 present study are consistent with the few comparable data reported in the literature for older participants (no data are available in young adults). For example, Baykara et al. reported in 313 their pioneering article a PSMD median value around 3.0 (in $mm^2s^{-1} \times 10^{-4}$, range [2.5, 4.9]) 314 in a sample of healthy individuals aged 60 to 80 years drawn from the ASPF cohort (see Table 315 316 2 of Baykara et al (20)), values that are comparable to those reported in Table 3d of our study in subsamples of other cohorts of similar age category. Similarly, Wei et al (21) recently 317 reported a PSMD average value of 2.4 x 10⁻⁴ mm²s⁻¹ in a sample of healthy controls aged 318 around 60 years. In both studies, PSMD CVs were close to 10%, a value again similar to 319

those observed in our own study. That the CV of PSMD is 2 to 3 times larger than the CVs of other DTI metrics could be expected since PSMD is a dispersion rather than a central tendency statistic. Moreover, the larger increase in PSMD CV as age advances (as compared to the other DTI metrics) indicates that this phenotype should be used with caution especially during the late life period. However, it is important to note that the CV of PSMD was found to be quite stable across cohorts with similar age ranges.

326 The main goal of the present study was to document the profile of PSMD evolution across 327 age bands during adulthood. In this respect, and the proviso that the data we gathered were 328 not longitudinal, our results show that PSMD increases continuously from post-adolescence 329 to late adult life, that this increase is accelerating at later ages, and that this acceleration is 330 larger than for the other DTI metrics. As there are no available data of PSMD in childhood 331 and adolescence, it is not possible to decide whether the lifetime PSMD evolution profile is 332 similar to those of AD, RD, MD, i.e. with a decrease during childhood that reaches the 333 minimum value before adulthood, or if it shows continuous increase throughout the lifespan. 334 Nevertheless, it remains the case that the continuous and accelerating increase of PSMD 335 during adulthood is an indication that it is an adequate and potentially valuable marker of white matter ageing. In particular, it is notable that PSMD increases during early adulthood 336 337 when the other DTI metrics variations appear to be still undergoing late maturational 338 processes.

339 The biological mechanisms of the origin of PSMD evolution with age are at present unknown, 340 but one can think of several reasons why PSMD may be more prone to increase with age as 341 compared with the other metrics. First, it is important to remember that PSMD is a measure 342 of MD values dispersion across a skeleton of white matter. As such, it will be directly affected 343 by differences across MD values of the individual tracts. Consequently, regional 344 heterogeneity as well as heterochrony in MD values of the fiber tracts will result in higher 345 PSMD values more than in average MD values. Second, MD itself is a weighted average of 346 AD (1/3) and RD (2/3) values, and thus MD value dispersion will also be affected by 347 heterochrony in AD and RD variations with age. Overall, what possibly makes PSMD an early 348 and sensitive imaging marker of ageing is that it captures multiple sources of heterogeneity 349 in white matter water diffusion parameters. With this regard, it would be interesting to 350 investigate variations in the pattern of MD dispersion at the regional level using tract-based

351 DTI metrics since it is well established that heterochrony is a major feature of the development 352 and aging of the different fiber tracts (see for example (44)(39)(8)(5)). Accordingly, variations 353 of PSMD value provide only a gross and possibly biased estimate of the white matter-354 microstructure dynamics. We did not implement regional analysis as our study focused on 355 PSMD that is by definition a dispersion statistic over the entire white matter skeleton. 356 Nevertheless, a regional approach would certainly be interesting and feasible since peak 357 width of MD values could be measured on a white-matter skeleton at the tract level in the 358 same manner as it has been done for other DTI metrics (see (30)(46) for example).

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378 6 Author Contributions

379 Study conception (BM, SD), data collection (GB, CT, SC, ZP, TP, RS, PS, HB, NK, JT, ID, 380 VW, AV, BM), data analysis (GB, LP, SC, JS, YP, LP, PS, WW, NA, MB, SMM, VW, MD,

BM), drafting (GB, AT, BM), revising the manuscript (LP, CT, SC, JS, ZP, YP, TP, R, LP, PS,
WW, NA, ID, MB, JW, SMM, VW, AV, MD, SD, BM).

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544 **8 Tables**

545 **Table 1.** Basic statistics for the 10 contributing datasets. MRi-Share: Magnetic Resonance imaging subcohort of Internet-based Students HeAlth Research Enterprise;

546 BIL&GIN: Brain Imaging of Lateralization study at Groupe d'Imagerie Neurofonctionnelle; SYS cohort: Saguenay Youth Study; ASPSF: Austrian Stroke Prevention

547 Study Family; OATS: Older Australian Twin Study; LBC 1936: Lothian Birth Cohort; MAS: Memory and Ageing Study. TIV: total intracranial volume estimated with

548 Freesurfer 5.3 version except "*" with FreeSurfer 6.0 version.[†] data subsample size < 30 (not included in the age category statistical analysis)

| Dataset | | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|------------------------|------------------|--------------|----------------|----------------|--------------|--------------|-------------------|------------------|-------------------|--------------|--------------|
| Sample size | 20,005 | 1,824 | 410 | 512 | 1,906 | 1,209 | 12,397 | 277 | 386 | 672 | 412 |
| | (% females) | 72% | 51% | 54% | 51% | 44% | 53% | 60% | 64% | 47% | 53% |
| Age (years) | mean (SD) | 22.1 (2.3) | 26.9 (8.0) | 49.5 (5.0) | 58.1 (15.1) | 60.8 (13.4) | 62.6 (7.4) | 65.0 (11.1) | 71.1 (5.1) | 72.7 (0.7) | 80.3 (4.6) |
| | range | [18.1, 34.9] | [18.1, 57.2] | [36.4, 65.4] | [19.0, 82.0] | [18.5, 85.4] | [45.1, 80.3] | [38.5, 85.6] | [65.4, 89.5] | [71.0, 74.2] | [72.7, 92.6] |
| TIV (cm ³) | females | 1515 (104)* | 1323 (93) | 1348 (164) | 1637 (115) | 1401 (104) | 1466 (203)* | 1396 (110) | 1369 (133) | 1353 (100) | 1267 (141) |
| mean (SD | males | 1703 (118)* | 1503 (127) | 1622 (131) | 1831 (131) | 1570 (118) | 1576 (147)* | 1572 (118) | 1574 (138) | 1537 (112) | 1460 (163) |
| N per age | 18 to 28 (2,200) | 1,778 | 285 | 0 | 101 | 36 | 0 | 0 | 0 | 0 | 0 |
| category | 28 to 38 (340) | 46 | 84 | 3^{\dagger} | 142 | 65 | 0 | 0 | 0 | 0 | 0 |
| | 38 to 48 (707) | 0 | 24^{\dagger} | 176 | 250 | 93 | 126 ^{††} | 38 | 0 | 0 | 0 |
| | 48 to 58 (4,287) | 0 | 17^{\dagger} | 311 | 218 | 176 | 3,525 | 40 | 0 | 0 | 0 |
| | 58 to 68 (6,535) | 0 | 3† | 22^{\dagger} | 511 | 431 | 5,375 | 52 ^{††} | 144 ^{††} | 0 | 0 |
| | 68 to 78 (5,515) | 0 | 0 | 0 | 656 | 358 | 3,351 | 129 | 195 | 672 | 154 |
| | 78 to 98 (421) | 0 | 0 | 0 | 28† | 50 | 20^{\dagger} | 18 [†] | 47 | 0 | 258 |

549

Table 2. Diffusion-weighted imaging acquisition the 10 contributing datasets. (see Table1 legend for the meaning of dataset abbreviated names and supplementary

552 material section for a more detailed description of these datasets). *: multi-shell acquisition.

| Dataset | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|---------------------------------|------------------------|--------------------|-----------|------------------|---------------------|------------------------|---------------------|-----------------------------|--------------------------|--------------------------------|
| MRI scanner | SIEMENS PRISMA | PHILIPS ACHIEVA | SIEMENS | SIEMENS VERIO | SIEMENS TIM TRIO | SIEMENS SKYRA | SIEMENS TIM TRIO | PHILIPS (x2) SIEMENS(x2) | GE Signa Horizon HDxt | PHILIPS ACHIEVA Quasar Dual |
| B0 strength | 3Т | 3T | 1.5T | 3T | 3T | 3Т | 3T | 1.5T & 3T | 1.5T | 3T |
| b-gradient (s/mm ²) | 1000* | 1000 | 1000 | 1000 | 1000 | 1000* | 1000 | 1000 | 1000 | 700 or 1000 |
| Nr of directions | 32 | 2 x 21 | 64 | 60 | 30 | 50 | 6 or 12 | 32 | 64 | 6 or 32 |
| Voxel size (mm ³) | 1.7x1.7x1.8 | 2x2x2 | 2.3x2.3x3 | 1.7x1.7x1.7 | 2x2x2 | 2x2x2 | 1.8x1.8x2.5 | 2.5x2.5x2.5 | 2x2x2 | 2.5x2.5x2.5 |
| TR (ms) | 3540 (multiband x3) | 8500 | 8000 | 13,800 | 7800 | 3600 (multiband x3) | 4900 or 6700 | 7800 or 8600 | 16500 | 7800 |
| TE (ms) | 75 | 81 | 102 | 100 | 83 | 92 | 81 or 95 | 68 or 96 | 98 | 68 |
| Acquisition duration | 9'45 | 7'45 | 10' | 16'08 | 4'30 | 9'45 | 4'51 or 6'10 | 4'40 or 4'56 | 19'30 | 4'40 |

- 555 **Table 3a.** Basic statistics of axial diffusivity (AD, in mm²sec⁻¹ x 10⁻⁴) average across a white matter skeleton for each age category and each contributing dataset.
- 556 Values are mean (S.D. and range [min, max] across the age category sample). See Table1 legend for the meaning of dataset abbreviated names

| Age range | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|-----------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|----------------|---------------|----------------|
| 18 to 28 | 11.80 (0.19) | 12.70 (0.18) | | 12.39 (0.23) | 11.60 (0.20) | | | | | |
| | [10.87, 12.45] | [12.26, 13.18] | | [11.70, 13.10] | [11.09, 11.96] | | | | | |
| 28 to 38 | 11.68 (0.21) | 12.58 (0.17) | | 12.33 (0.20) | 11.41 (0.27) | | | | | |
| | [11.10, 12.12] | [12.14, 13.04] | | [11.80, 12.90] | [10.71, 11.95] | | | | | |
| 38 to 48 | | 12.45 (0.21) | 11.58 (0.22) | 12.25 (0.21) | 11.33 (0.25) | 12.21 (0.27) | 11.88 (0.24) | | | |
| | | [12.07, 12.88] | [11.03, 12.29] | [11.60, 12.80] | [10.52, 11.89] | [10.76, 12.85] | [11.50, 12.46] | | | |
| 48 to 58 | | | 11.58 (0.23) | 12.25 (0.24) | 11.31 (0.25) | 12.21 (0.25) | 11.89 (0.27) | | | |
| | | | [11.01, 12.38] | [11.40, 13.10] | [10.59, 12.02] | [10.18, 13.03] | [11.42, 12.68] | | | |
| 58 to 68 | | | 11.67 (0.24) | 12.34 (0.26) | 11.37 (0.30) | 12.25 (0.26) | 12.03 (0.30) | 11.74 (0.49) | | |
| | | | [11.24, 12.21] | [11.60, 13.20] | [10.30, 12.32] | [10.52, 13.24] | [11.41, 12.76] | [10.61, 12.80] | | |
| 68 to 78 | | | | 12.48 (0.28) | 11.50 (0.32) | 12.34 (0.29) | 12.06 (0.37) | 11.83 (0.58) | 9.79 (0.43) | 11.83 (0.31) |
| | | | | [11.40, 13.40] | [10.36, 12.58] | [10.70, 13.53] | [9.69, 12.88] | [10.69, 13.18] | [8.41, 10.94] | [11.06, 12.56] |
| 78 to 98 | | | | 12.49 (0.30) | 11.64 (0.33) | 12.44 (0.27) | | 12.12 (0.62) | | 11.96 (0.32) |
| | | | | [11.90, 13.20] | [11.07, 12.47] | [11.66, 12.83] | | [10.80, 13.49] | | [10.99, 13.05] |

- **Table 3b.** Basic statistics of radial diffusivity (RD, in mm²sec⁻¹ x 10⁻⁴) average across a white matter skeleton for each age category and each contributing dataset.
- 560 Values are mean (S.D. and range [min, max] across the age category sample. (see Table1 legend for the meaning of dataset abbreviated names)

| Age range | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|-----------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|
| 18 to 28 | 4.74 (0.18) | 4.81 (0.19) | | 4.83 (0.19) | 4.66 (0.18) | | | | | |
| | [4.15, 5.53] | [4.26, 5.44] | | [4.33, 5.29] | [4.26, 5.19] | | | | | |
| 28 to 38 | 4.72 (0.21) | 4.79 (0.20) | | 4.83 (0.20) | 4.60 (0.18) | | | | | |
| | [4.35, 5.10] | [4.45, 5.56] | | [4.42, 5.36] | [4.20, 5.17] | | | | | |
| 38 to 48 | | 4.83 (0.22) | 5.45 (0.24) | 4.87 (0.22) | 4.64 (0.20) | 4.95 (0.22) | 5.11 (0.29) | | | |
| | | [4.49, 5.27] | [4.88, 6.36] | [4.35, 5.59] | [4.28, 5.27] | [4.19, 5.64] | [4.54, 5.79] | | | |
| 48 to 58 | | | 5.51 (0.25) | 4.99 (0.24) | 4.72 (0.22) | 4.97 (0.23) | 5.22 (0.30) | | | |
| | | | [4.93, 6.54] | [4.47, 6.01] | [4.21, 5.38] | [4.17, 6.22] | [4.57, 6.05] | | | |
| 58 to 68 | | | 5.58 (0.31) | 5.13 (0.29) | 4.84 (0.25) | 5.08 (0.26) | 5.47 (0.37) | 5.27 (0.38) | | |
| | | | [5.02, 6.27] | [4.29, 6.36] | [4.21, 5.57] | [4.34, 6.40] | [4.73, 6.59] | [4.36, 6.05] | | |
| 68 to 78 | | | | 5.31 (0.33) | 4.99 (0.31) | 5.21 (0.29) | 5.49 (0.35) | 5.40 (0.46) | 5.21 (0.38) | 5.66 (0.34) |
| | | | | [4.48, 6.73] | [4.16, 6.29] | [4.29, 7.10] | [4.85, 6.69] | [4.44, 6.77] | [4.28, 8.87] | [5.05, 6.78] |
| 78 to 98 | | | | 5.37 (0.33) | 5.19 (0.37) | 5.36 (0.20) | | 5.75 (0.48) | | 5.81 (0.40) |
| | | | | [4.87, 6.31] | [4.69, 5.99] | [5.03, 5.69] | | [4.80, 7.08] | | [4.88, 7.17] |

- **Table 3c.** Basic statistics of mean diffusivity (MD, in $mm^2 \sec^{-1} x \, 10^{-4}$) average across a white matter skeleton for each age category and each contributing dataset.
- 564 Values are mean (S.D. and range [min, max] across the age category sample. (see Table1 legend for the meaning of dataset abbreviated names)

| Age range | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|-----------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|
| 18 to 28 | 7.09 (0.16) | 7.44 (0.17) | | 7.35 (0.18) | 6.97 (0.17) | | | | | |
| | [6.54, 7.76] | [7.02, 7.98] | | [6.89, 7.75] | [6.54, 7.34] | | | | | |
| 28 to 38 | 7.04 (0.18) | 7.39 (0.17) | | 7.33 (0.18) | 6.87 (0.19) | | | | | |
| | [6.64, 7.38] | [7.08, 7.97] | | [6.92, 7.78] | [6.43, 7.39] | | | | | |
| 38 to 48 | | 7.37 (0.20) | 7.50 (0.21) | 7.33 (0.20) | 6.87 (0.19) | 7.37 (0.22) | 7.37 (0.25) | | | |
| | | [7.05, 7.80] | [6.98, 8.17] | [6.81, 7.84] | [6.42, 7.39] | [6.38, 8.05] | [6.97, 8.01] | | | |
| 48 to 58 | | | 7.53 (0.23) | 7.41 (0.22) | 6.92 (0.21) | 7.39 (0.22) | 7.45 (0.27) | | | |
| | | | [6.97, 8.45] | [6.89, 8.31] | [6.35, 7.58] | [6.17, 8.42] | [6.91, 8.26] | | | |
| 58 to 68 | | | 7.61 (0.27) | 7.53 (0.26) | 7.01 (0.25) | 7.47 (0.24) | 7.66 (0.34) | 7.43 (0.41) | | |
| | | | [7.09, 8.20] | [6.82, 8.64] | [6.35, 7.78] | [6.41, 8.67] | [6.98, 8.58] | [6.55, 8.18] | | |
| 68 to 78 | | | | 7.70 (0.30) | 7.16 (0.29) | 7.59 (0.27) | 7.68 (0.33) | 7.54 (0.48) | 6.74 (0.35) | 7.72 (0.31) |
| | | | | [6.93, 8.92] | [6.30, 8.39] | [6.43, 9.23] | [6.51, 8.74] | [6.57, 8.91] | [5.79, 7.86] | [7.14, 8.71] |
| 78 to 98 | | | | 7.74 (0.31) | 7.34 (0.34) | 7.72 (0.20) | | 7.87 (0.51) | | 7.86 (0.36) |
| | | | | [7.23, 8.61] | [6.84, 8.15] | [7.34, 8.01] | | [6.92, 9.21] | | [6.96, 9.06] |

- **Table 3d.** Basic statistics of peak width skeletonized mean diffusivity (PSMD, in mm²sec⁻¹ x 10⁻⁴) for each age category and each contributing dataset. Values are
- 568 mean (S.D. and range [min, max] across the age category sample. (see Table1 legend for the meaning of dataset abbreviated names.

| Age range | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|-----------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|
| 18 to 28 | 1.54 (0.14) | 2.13 (0.17) | | 2.02 (0.19) | 2.26 (0.28) | | | | | |
| | [1.20, 2.13] | [1.75, 2.78] | | [1.68, 2.53] | [1.86, 3.47] | | | | | |
| 28 to 38 | 1.58 (0.17) | 2.18 (0.22) | | 2.05 (0.19) | 2.32 (0.24) | | | | | |
| | [1.33, 2.01] | [1.75, 2.85] | | [1.65, 2.63] | [1.75, 3.00] | | | | | |
| 38 to 48 | | 2.24 (0.18) | 2.68 (0.30) | 2.17 (0.22) | 2.49 (0.29) | 2.10 (0.22) | 2.63 (0.39) | | | |
| | | [1.89, 2.58] | [2.07, 4.44] | [1.64, 3.00] | [1.94, 3.46] | [1.60, 3.11] | [1.95, 3.53] | | | |
| 48 to 58 | | | 2.82 (0.34) | 2.31 (0.28) | 2.67 (0.32) | 2.12 (0.25) | 2.69 (0.36) | | | |
| | | | [1.99, 5.12] | [1.70, 3.63] | [2.03, 4.07] | [1.49, 4.58] | [2.18, 3.70] | | | |
| 58 to 68 | | | 2.92 (0.30) | 2.53 (0.38) | 2.94 (0.38) | 2.27 (0.31) | 2.82 (0.49) | 2.96 (0.54) | | |
| | | | [2.51, 3.57] | [1.83, 4.85] | [2.14, 5.03] | [1.56, 5.71] | [2.15, 4.54] | [2.11, 4.90] | | |
| 68 to 78 | | | | 2.78 (0.47) | 3.26 (0.47) | 2.48 (0.39) | 3.17 (0.49) | 3.10 (0.64) | 3.18 (0.53) | 3.84 (0.61) |
| | | | | [1.88, 5.26] | [2.23, 5.53] | [1.61, 6.19] | [2.15, 4.60] | [2.17, 6.66] | [2.09, 5.72] | [2.56, 6.02] |
| 78 to 98 | | | | 2.96 (0.45) | 3.68 (0.58) | 2.74 (0.33) | | 3.77 (0.90) | | 4.29 (0.77) |
| | | | | [2.15, 3.94] | [2.83, 5.32] | [2.10, 3.45] | | [2.47, 6.51] | | [2.95, 7.96] |

- **Table 3e.** Basic statistics of fractional anisotropy (FA, unitless) average across individual FA skeleton for each age range category and each contributing dataset.
- 572 Values are mean (S.D. and range [min, max] across the age category sample. (see Table1 legend for the meaning of dataset abbreviated names)

| Age range | MRi-Share | BIL&GIN | SYS | LIFE | 1000 BRAINS | UKBiobank | ASPSF | OATS | LBC 1936 | MAS |
|-----------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|--------------|
| 18 to 28 | 0.53 (0.01) | 0.56 (0.01) | | 0.54 (0.02) | 0.54 (0.01) | | | | | |
| | [0.47, 0.58] | [0.51, 0.60] | | [0.49, 0.58] | [0.49, 0.56] | | | | | |
| 28 to 38 | 0.53 (0.02) | 0.55 (0.02) | | 0.54 (0.02) | 0.54 (0.01) | | | | | |
| | [0.49, 0.56] | [0.50, 0.58] | | [0.50, 0.58] | [0.49, 0.57] | | | | | |
| 38 to 48 | | 0.55 (0.02) | 0.46 (0.02) | 0.54 (0.02) | 0.53 (0.02) | 0.53 (0.01) | 0.51 (0.02) | | | |
| | | [0.52, 0.57] | [0.39, 0.50] | [0.48, 0.58] | [0.48, 0.56] | [0.49, 0.56] | [0.46, 0.55] | | | |
| 48 to 58 | | | 0.45 (0.02) | 0.53 (0.02) | 0.52 (0.02) | 0.52 (0.02) | 0.49 (0.02) | | | |
| | | | [0.40, 0.50] | [0.46, 0.57] | [0.47, 0.58] | [0.44, 0.57] | [0.42, 0.54] | | | |
| 58 to 68 | | | 0.45 (0.02) | 0.52 (0.02) | 0.51 (0.02) | 0.52 (0.02) | 0.48 (0.02) | 0.48 (0.02) | | |
| | | | [0.41, 0.49] | [0.44, 0.58] | [0.46, 0.57] | [0.44, 0.57] | [0.41, 0.53] | [0.43, 0.55] | | |
| 68 to 78 | | | | 0.51 (0.02) | 0.51 (0.02) | 0.51 (0.02) | 0.48 (0.02) | 0.48 (0.03) | 0.41 (0.03) | 0.45 (0.02) |
| | | | | [0.40, 0.56] | [0.44, 0.56] | [0.37, 0.57] | [0.41, 0.53] | [0.38, 0.54] | [0.31, 0.47] | [0.36, 0.49] |
| 78 to 98 | | | | 0.50 (0.02) | 0.50 (0.02) | 0.50 (0.01) | | 0.46 (0.02) | | 0.45 (0.03) |
| | | | | [0.45, 0.54] | [0.44, 0.53] | [0.48, 0.52] | | [0.40, 0.51] | | [0.34, 0.51] |

Table 4. ANOVA effect of age (estimate (standard error) and significance p-value) on five DTI metrics (AD, RD, MD, PSMD, and FA evaluated across individual FA

577 skeletons). Values are in $mm^2s^{-1}year^{-1}x10^{-6}$ for AD, RD and MD, in $mm^2s^{-1}year^{-1}x10^{-7}$ for PSMD, and in $year^{-1}x10^{-3}$ for FA.

| | | AD | RD | | MD | | PSMD | | FA | |
|-------------------|--------------|----------|--------------|----------|--------------|----------|-------------|----------|--------------|----------|
| Age range (years) | estimate | р | estimate | р | estimate | р | estimate | р | estimate | р |
| 18 to 28 | -2.01 (0.20) | < 0.0001 | -1.06 (0.18) | < 0.0001 | -1.38 (0.17) | < 0.0001 | 0.41 (0.15) | 0.0058 | 0.30 (0.15) | 0.041 |
| 28 to 38 | -1.32 (0.43) | 0.0023 | -0.22 (0.41) | 0.59 | -0.58 (0.37) | 0.12 | 0.60 (0.42) | 0.15 | -0.27 (0.32) | 0.40 |
| 38 to 48 | -0.68 (0.37) | 0.065 | 0.40 (0.37) | 0.27 | 0.047 (0.33) | 0.89 | 1.84 (0.41) | < 0.0001 | -0.54 (0.27) | 0.046 |
| 48 to 58 | -0.11 (0.13) | 0.41 | 0.61 (0.12) | < 0.0001 | 0.37 (0.12) | 0.0019 | 1.11 (0.14) | < 0.0001 | -0.56 (0.09) | < 0.0001 |
| 58 to 68 | 0.95 (0.12) | < 0.0001 | 1.46 (0.11) | < 0.0001 | 1.29 (0.11) | < 0.0001 | 2.18 (0.14) | < 0.0001 | -0.91 (0.08) | < 0.0001 |
| 68 to 78 | 1.75 (0.19) | < 0.0001 | 1.94 (0.18) | < 0.0001 | 1.87 (0.17) | < 0.0001 | 3.67 (0.24) | < 0.0001 | -0.97 (0.12) | < 0.0001 |
| 78 to 98 | 2.30 (0.60) | 0.0002 | 4.01 (0.64) | < 0.0001 | 3.44 (0.59) | < 0.0001 | 6.99 (0.12) | < 0.0001 | -2.36 (0.39) | < 0.0001 |

- 580 **Table 5.** ANOVA effect of sex and TIV (estimate (standard error) and significance p-value) on five DTI metrics (AD, RD, MD, PSMD, and FA evaluated across
- 581 individual FA skeletons). TIV: total intracranial volume. Values for the effects of sex are in $mm^2s^{-1}x10^{-6}$ for AD, RD, MD, and PSMD and in $x10^{-3}$ for FA, and for the
- 582 effects of TIV are in $mm^2s^{-1}cm^{-3} \times 10^{-6}$ for AD, RD MD, and PSMD and in $cm^{-3} \times 10^{-5}$ for FA.

| | A |) | RD | | MD | | PSI | ٨D | FA | |
|-----------|-------------|----------|--------------------------|-------|-------------|----------|--------------|----------|-------------|----------|
| Effect | estimate | р | estimate | р | estimate | р | estimate | р | estimate | р |
| Sex (F-M) | 1.13 (0.21) | < 0.0001 | 0.45 (0.21) | 0.032 | 0.68 (0.19) | 0.0005 | -5.34 (0.26) | < 0.0001 | 0.04 (0.14) | 0.74 |
| TIV | 1.41 (0.12) | < 0.0001 | 0.07 (0.12) ⁻ | 0.53 | 0.52 (0.01) | < 0.0001 | 1.43 (0.15) | < 0.0001 | 0.54 (0.08) | < 0.0001 |

585 9 Figure legends

586 **Figure 1.** Box-plots and histogram distribution of the age of the participants of the 10 587 contributing datasets.

Figure 2. Mean and standard deviation (bars) of the five DTI metrics for each age subcategory and each dataset. The dotted lines connect values for the different age categories of the same dataset.

Figure 3. Coefficients of variation of the five DTI metrics for each age subcategory and eachdataset.

593 **Figure 4.** ANOVA estimates and 95% confidence intervals of the effects of age on the five

594 DT metrics for each age subcategory. Units are mm²/s/year x10⁻³ for AD, RD and MD,

595 mm²/s/year x10⁻⁴ for PSMD, /year x10⁻³ for FA.

Figure 5. ANOVA estimates and 95% confidence intervals of the effects of age on the fiveDT metrics for each age subcategory and each dataset.

599 **10 Conflict of Interest**

- 600 The authors declare that the research was conducted in the absence of any commercial or
- financial relationships that could be construed as a potential conflict of interest.

602 11 Data Availability Statement

603 Requests to access these datasets should be directed to:

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