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What is normal in normal aging? Effects of aging, amyloid and Alzheimer's disease on the cerebral cortex and the hippocampus



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ABSTRACT

What can be expected in normal aging, and where does normal aging stop and pathological neurodegeneration begin? With the slow progression of age-related dementias such as Alzheimer's disease (AD), it is difficult to distinguish age-related changes from effects of undetected disease. We review recent research on changes of the cerebral cortex and the hippocampus in aging and the borders between normal aging and AD. We argue that prominent cortical reductions are evident in fronto-temporal regions in elderly even with low probability of AD, including regions overlapping the default mode network. Importantly, these regions show high levels of amyloid deposition in AD, and are both structurally and functionally vulnerable early in the disease. This *normalcy-pathology homology* is critical to understand, since aging itself is the major risk factor for sporadic AD. Thus, rather than necessarily reflecting early signs of disease, these changes may be part of normal aging, and may inform on why the aging brain is so much more susceptible to AD than is the younger brain. We suggest that regions characterized by a high degree of life-long plasticity are vulnerable to detrimental effects of normal aging, and that this age-vulnerability renders them more susceptible to additional, pathological AD-related changes. We conclude that it will be difficult to understand AD without understanding why it preferably affects older brains, and that we need a model that accounts for age-related changes in AD-vulnerable regions independently of AD-pathology.

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1. Introduction

The major risk factor for Alzheimer's disease (AD) is age, with a sharp increase in incidence after 60 years (Kawas et al., 2000). This has inspired many researchers to propose that to understand AD, we must understand its inherent relationship to aging (Herrup, 2010). Why is the aging brain so susceptible to AD, compared to the middle-aged or young brain? What features distinguish normal brain changes from those seen in early AD? How should we understand the fact that three of the major symptoms of AD observed in vivo – disruption of episodic memory function (Koivisto et al., 1995; Nyberg et al., 2012), brain atrophy (Raz et al., 2005; Driscoll et al., 2009; Fjell et al., 2009a) and accumulation of amyloid protein (Morris et al., 2010) – are also found in many presumably healthy elderly? Given these commonalities, it can be argued that AD cannot be understood separately from its major risk factor – age. However, we suggest that this statement can also be reversed: if we understand why the older brain is susceptible to AD, we may have a better chance of understanding brain aging itself. With the aging of the population, a comprehensive understanding of normal, non-demented changes in brain and cognition is arguably as important as understanding AD.

How the link between aging and AD should be understood is thus a major question in contemporary neuroscience. However, it is not obvious that studying the relationship between the two is the best starting point for understanding either phenomenon. Some argue that AD should be viewed as a disease with distinct etiology and neuropathology, separate from normal aging, and that it is less fruitful to view AD in light of normal age changes (Nelson et al., 2011). AD may be driven by factors less related to aging per se, for instance differences in amyloid precursor protein expression (APP) (Nelson et al., 2011), from which the presumably most toxic form of amyloid (A β 42) originates. However, we have still not understood the role of amyloid in brain atrophy and cognitive decline. Current models of the role of amyloid in AD, as for instance reflected in the proposed diagnostic guidelines from the National Institute of Aging – Alzheimer's Association (NIA-AA) (Jack et al., 2011; Sperling et al., 2011a) and the popular 'dynamic biomarker model' (Jack et al., 2010a, 2013), suggest that the influence of amyloid is greatest in very early phases – at a stage where cognitive and clinical symptoms are not yet detected. When accelerated brain atrophy and cognitive decline become evident, the therapeutic window for anti-amyloid drugs may very well be closed. Thus, it is absolutely necessary to study the relationship between amyloid, brain integrity and memory in healthy elderly if the role of amyloid in neurodegeneration and cognitive decline is to be understood. Animal models of AD are not characterized by the massive brain atrophy that correlates with memory problems in AD patients, and therefore can provide only limited insight into relationships between amyloid, brain integrity and episodic memory decline in non-demented older adults.

In the present paper, we review recent research on cortical and hippocampal changes in normal aging, the relationship between changes in normal aging vs. early AD, and the role played by amyloid. First, we will discuss the characteristics of presumably normal brain aging. What kind of macroscopic brain changes can be expected in older adults without dementia, and what consequences do these brain changes have for cognitive function?

We try to identify and evaluate some of the proposed major organizing principles for brain aging, such as the theory of retrogenesis or the principle of "last in, first out". In the cognitive domain, we focus especially on episodic memory, which is of interest because it is affected both in normal aging and very early in AD. Second, we investigate similarities and differences in the pattern of brain atrophy between normal aging and AD, with a special focus on comparisons between AD patients and elderly individuals with low AD-risk. Finally, we discuss the role of amyloid in brain atrophy and cognition, and evaluate current available knowledge related to the question of why brain aging is associated with the dramatic increase in AD-risk.

2. What is normal in normal aging?

2.1. Magnitude, pattern and timing of change

Reductions in specific cognitive abilities like mental speed (Salthouse, 1996), executive function (Connelly et al., 1991; Schretlen et al., 2000; Rabbitt et al., 2001) and episodic memory (Salthouse, 2003; Buckner, 2004; Nyberg et al., 2012) are commonly experienced in aging, while verbal abilities and world knowledge are typically maintained (Park and Reuter-Lorenz, 2009). However, there is disagreement about whether longitudinal changes in older adults reflect continuous ongoing processes starting in young adulthood or whether changes begin in middle age or beyond. Closely related to this is an unsettled discussion about whether changes observed in cross-sectional studies reflect real ongoing change within individuals (Salthouse, 2009) or arise from methodological artifacts (Schaie, 2009; Nyberg et al., 2012). Cross-sectional studies may suffer from cohort effects, and, potentially more seriously, different recruitment bias across age-groups. For instance, most studies are based on convenience samples, and there may be systematic differences in individuals who are recruited for, and agree to participate in, research at younger ages and older ages. For example, young participants are often college students, middle-aged participants may be more likely to be unemployed or under-employed, and older participants may be recruited from senior centers. The older age group may be biased by participants motivated to volunteer due to concerns about their cognitive abilities, or, alternatively, only the superior functioning older adults volunteer or are accepted into the study due to strict exclusion criteria. Thus these three age groups may differ in critical characteristics, which may affect the estimated age-trajectories. This is referred to as covariance between age and sampling bias. Longitudinal studies are, in principle, not affected by this bias and can measure change over time within individuals. However, longitudinal studies too have limitations that can influence the results – such as selective attrition and test-retest effects that may be larger than the change across time points (Salthouse, 2012). Adding to this, few longitudinal studies sample the entire adult age-span, precluding estimations of change rates as a function of age.

Despite the limitations of cross-sectional and longitudinal study designs, there is a consensus that episodic memory, which is of special focus in the present review, declines from about the age of 50–60 years on a population basis (Nyberg et al., 2012), although earlier decrements cannot be ruled out. In a very interesting study, longitudinal trajectories in episodic memory were mapped over

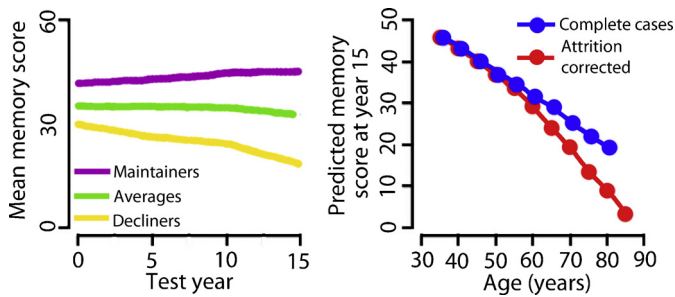


Fig. 1. Longitudinal episodic memory decline in aging. Left panel: By using a random-effects pattern-mixture model, [Josefsson and colleagues \(2012\)](#), found individual differences both in initial memory scores (offset at time 0) and change over time in their large sample ($n > 1500$) of initially healthy participants, with some maintaining their functional level (18%), some showing age-average decline (68%), while some declined (13%). The researchers were able to identify environmental and genetic factors predicting group membership. Right panel: Correcting for selective attrition reveals a much steeper decline in episodic memory with increasing age than analyzing the full sample of available data. Figures modified from [Josefsson et al. \(2012\)](#).

15 years in more than 1500 participants from the Betula study ([Josefsson et al., 2012](#)). Using a random-effects pattern-mixture model, the authors were able to divide the participants into three groups in terms of memory change – maintainers (18%), those who showed age-typical decline (68%) and decliners (13%) (see [Fig. 1](#)). Physical activity, being female and living with someone increased the likelihood of being a maintainer, while lower education, not being employed and being male increased likelihood of being a decliner. Further, the authors corrected the age-memory reduction function for the effects of selective attrition. Correcting for selective attrition revealed a much steeper decline in episodic memory with increasing age than merely analyzing the available raw data. This indicates that studies may actually under-estimate the true age-decline due to selective attrition in longitudinal designs, or, likely, covariance between selection biases and age in cross-sectional. Carefully conducted longitudinal studies yield opportunities for correcting for these biases, revealing truer age-functions.

A popular view is that at least some portion of the age-changes in cognitive function is caused by structural brain changes. Cross-sectional correlations between brain volume and cognitive function cannot alone be used to justify such claims, and other types of cross-sectional evidence, e.g. cross-sectional mediation analysis, also have caveats ([Rabbitt, 2011](#); [Raz and Lindenberger, 2011](#); [Salthouse, 2011](#)). Detailed mapping of regional longitudinal brain changes, which then can be correlated with longitudinal cognitive changes, is a more promising avenue for understanding the neurobiological foundation for age-related decrements in cognitive function.

Compared to cognitive testing, an advantage with MRI is that re-test effects are likely to be minute, and follow-up examinations at short intervals are highly feasible. With high-quality MR-scans and state-of-the-art analysis tools ([Reuter et al., 2010](#); [Holland and Dale, 2011](#); [Holland et al., 2012b](#); [Reuter et al., 2012](#)), brain changes can be detected reliably and monitored over intervals as short as one year or less even in healthy elderly ([Fjell et al., 2009a](#); [Murphy et al., 2010](#)). Although longitudinal studies cannot fully replace cross-sectional studies in brain aging research for a number of practical reasons, such as the relatively short life-expectancy of MR-scanners and continual advances in technology, repeated MRIs yield a unique tool to monitor brain changes naturally occurring among older adults.

Reductions in gross brain volume over time are now well-documented, with annual decreases on the order of 0.2–0.5% ([Scahill et al., 2003](#); [Ezekiel et al., 2004](#); [Enzinger et al., 2005](#);

[Fotenos et al., 2005](#); [Hedman et al., 2012](#)). Significant longitudinal change has been observed across almost the entire cerebral cortex ([Driscoll et al., 2009](#); [Fjell et al., 2009a](#)), but with substantial heterogeneity across regions. Traditionally, the frontal lobes have been regarded as especially vulnerable to normal age-changes, inspiring frontally based theories of cognitive aging ([West, 1996](#); [Robbins et al., 1998](#); [Rabbitt, 2005](#)). Evidence is accumulating, however, that in healthy older adults, temporal areas undergo reductions over time of comparable magnitude to the frontal changes ([Driscoll et al., 2009](#); [Fjell et al., 2009a](#); [Raz et al., 2010](#); [Pfefferbaum et al., 2013](#)). Annual atrophy rates of 0.79–2.0% for the hippocampus ([Jack et al., 1998](#); [Scahill et al., 2003](#); [Ezekiel et al., 2004](#); [Raz et al., 2005](#); [Du et al., 2006](#); [Fjell et al., 2009a](#)) and of 0.3–2.4% for the entorhinal cortex ([Du et al., 2003, 2006](#); [Ezekiel et al., 2004](#); [Raz et al., 2005](#); [Fjell et al., 2009a](#)) have been reported. Thus, age-related atrophy is not restricted to the frontal lobes.

Age-related changes in cortical volume are illustrated in [Fig. 2](#), which shows one-year percentage change in healthy elderly participants from the Alzheimer's Disease Neuroimaging Initiative (ADNI) (see www.adni-info.org). These participants remained free of a mild cognitive impairment (MCI) or AD diagnosis for three years following the baseline scan. As can be seen, cortical volume reductions are found across most of the brain surface, with annual change rates of around 0.5% in most regions. [Fig. 2](#) also shows this annual change expressed in standard deviations, highlighting the relative degree of change across different regions (Z-transformed maps). The frontal and the temporal lobes stand out with the highest degree of relative change, with substantial change observed also in the medial parietal area (precuneus and the adjacent retrosplenial and posterior cingulate cortices). This predominant fronto-temporal pattern of atrophy is in accordance with the results of several independent longitudinal studies using different methods for quantification of brain change. As we discuss below, this has implications for how we should understand the macro-structural correlates of decline in specific cognitive abilities

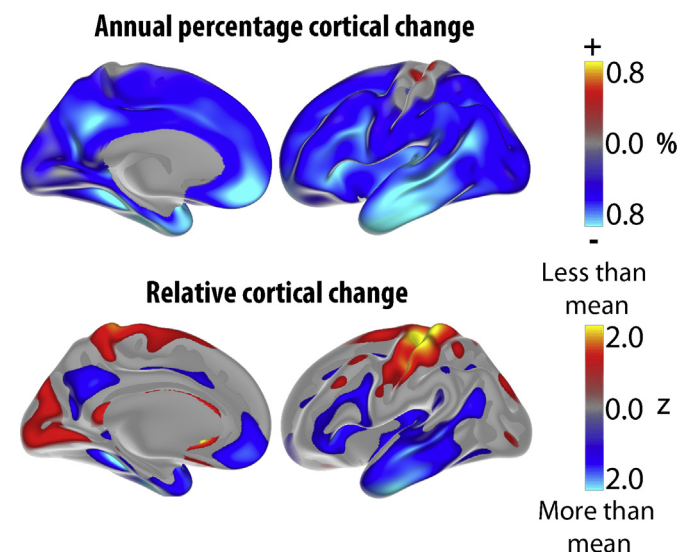


Fig. 2. Longitudinal cortical volumetric reductions in aging. Upper panel shows annual percent volume reduction of the cerebral cortex in a longitudinal sample of 132 healthy elderly (55–91 years at baseline) from ADNI (Alzheimer's Disease Neuroimaging Initiative). Lower panel shows the same data, but the values have been re-scaled to yield a mean of zero and a standard deviation of one, to allow better visualization of regional distribution of change. Blue–cyan colors represent areas that are reduced more in one year than the rest of the cortex, while red–yellow colors represent areas of less than average reduction. The often-observed fronto-temporal pattern of relative increased atrophy is evident, with medial parietal cortex/posterior cingulate as additional regions with high rates of atrophy in aging. Data from [Fjell et al. \(2013a\)](#).

in normal aging, and how we should conceptualize the differences between normal aging and age-related neurodegenerative disease like AD.

Longitudinal studies can track within-person changes in brain structures over time. However, no longitudinal MR-studies can follow the same participants over extended time-periods. Thus, to investigate the effect of age across the life-span on brain changes, combinations of longitudinal and cross-sectional designs seem to be the optimal solution. Among such studies, several have found modest acceleration of regional decline with increasing age (Raz et al., 2004a,b, 2005, 2010; Du et al., 2006; Driscoll et al., 2009; Fjell et al., 2009a; Schuff et al., 2010; Thambisetty et al., 2010; Holland et al., 2012a; Taki et al., 2013; Pfefferbaum et al., 2013). Thus, there is a tendency that the rate of atrophy in specific brain regions is higher in older adults than in middle-aged adults. However, very few longitudinal studies have sampled the entire adult life-span, and this hinders direct comparisons between young and older adults, although important exceptions exist (e.g. Raz et al., 2005; Taki et al., 2013; Pfefferbaum et al., 2013). This is the major strength of the cross-sectional design. Dozens of cross-sectional studies have reported that different macro-structural measures of the cerebral cortex are negatively correlated with age across the adult life-span. Negative correlations between cortical thickness and age have been reported in all regions (Fjell et al., *in press-b*), exceeding $-.70$ in the most vulnerable areas. As long as the MR scans are of good quality, the method used to quantify cortical reductions is identical and the sample size is reasonably large, high consistency can be observed across studies (Fjell et al., 2009b). In Fig. 3, Pearson correlations between age and cortical thickness in a large sample of healthy adults are shown. Strong correlations are seen in the frontal cortex, both laterally and medially, but also in other regions, such as the lateral temporal cortex.

Cross-sectional studies of cortical thickness and volume usually find mainly linear age-relationships, although tendencies toward accelerating or reduced estimated decline with increasing age are sometimes reported for specific regions, such as accelerated decline of the entorhinal and the lingual cortex with age, and decreasing decline in anterior parts of the cingulate (Fjell et al., *in press-b*)

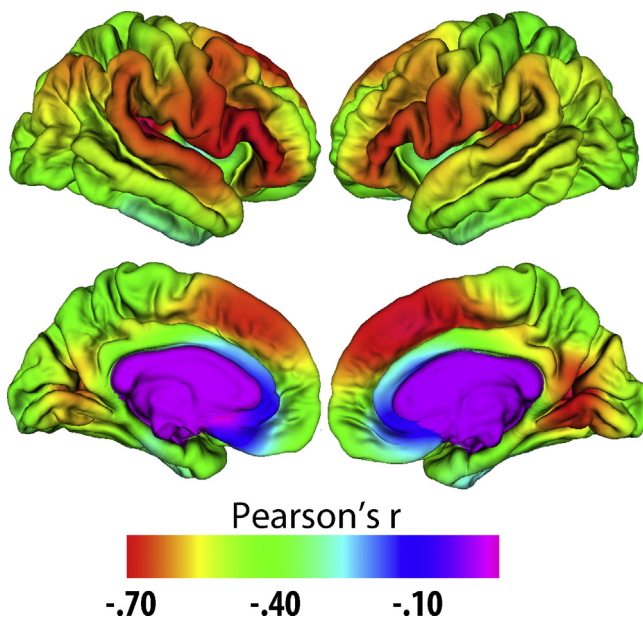


Fig. 3. Age-thickness correlations. Correlations between age and cortical thickness in a healthy multi-site adult life-span sample ($n = 1100$, age 18–94). Correlations exceed $-.40$ in large regions, approaching $-.70$ in the prefrontal and lateral temporal cortex.

Data from Fjell et al., *in press-b*.

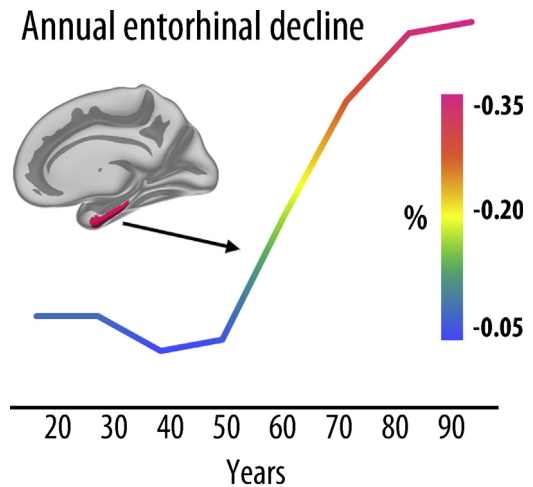


Fig. 4. Accelerated entorhinal decline in aging. Cross-sectional estimates of annual rate of cortical thinning in the entorhinal cortex across the adult life indicate a marked increase in atrophy from 50 years. Age-related acceleration of decline in elderly has been confirmed with longitudinal data.

Data from Fjell et al., *in press-b*.

(see Fig. 4). This is in contrast to the trajectory for hippocampal volume, which almost invariably shows accelerated decline with higher age across studies. In Fig. 5, typical age-trajectories for the volume of the hippocampus and for the rest of the cerebral cortex observed in cross-sectional studies are illustrated. While total cortical volume declines almost linearly from 20 years, hippocampus is relatively stable until about the age of 60 years, after which the trajectory indicates prominent loss.

More longitudinal studies including participants from the entire adult lifespan are expected, and will likely advance our knowledge about this critical question over the next few years. A challenge, however, is that the potential cross-sectional fallacy of covariance between sampling bias and age, i.e. that the oldest participants are not representative of the younger population, will carry over to the combined cross-sectional and longitudinal design.

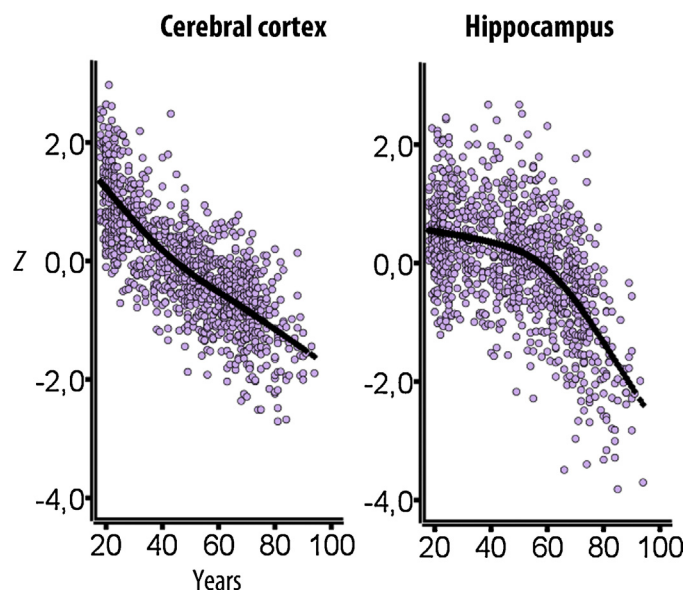


Fig. 5. Life-span trajectories of volumetric reductions. Cross-sectional estimates of adult life-span trajectories of total cerebral cortex volume and total hippocampal volume. Volume is expressed in units of standard deviations.

Data from Fjell et al. (2013).

However, a way to account for this effect by use of multiple samples has been suggested (see e.g. [Schaie and Hofer, 2001](#)).

Two recent studies have used longitudinal data to demonstrate flaws in apparent cross-sectional trends. Nyberg et al. followed a sample of elderly over six years, measuring brain activation during a semantic categorization task ([Nyberg et al., 2010b](#)). The cross-sectional analyses indicated frontal over-recruitment in aging. In contrast, the longitudinal analyses revealed frontal reductions over time, and showed that the cross-sectional over-recruitment could be accounted for by a select sample of high-performing elderly. We recently made a similar observation in a combined longitudinal and cross-sectional study of cortical thickness change in aging ([Fjell et al., in press-b](#)). In a sample of healthy elderly adults from ADNI, we observed a positive correlation between thickness and age in the cingulate cortex, especially the anterior parts. This has been found previously in several independent cross-sectional samples ([Salat et al., 2002](#); [Fjell et al., 2009b](#)). However, when we examined the follow-up data for the same participants, consistent thinning was found throughout the cerebral cortex, including the entire anterior cingulate. The results are illustrated in [Fig. 6](#). In addition, there was a general down-scaling of estimated change in the cross-sectional compared to the longitudinal results. This indicates that cross-sectional studies may under-estimate rather than exaggerate real change, an observation also made previously ([Raz et al., 2005](#)). Similar observations have been made for cognitive functions like episodic memory, where cross-sectional studies have found almost linear decline from 20 to 80 years of age but practice-adjusted longitudinal data have shown preservation of function until 60 years of age ([Nyberg et al., 2012](#)). Actually, if the true age-trajectory is non-linear, cross-sectional results may over-estimate the age-effect in the first part of adult life and under-estimate it in the second part.

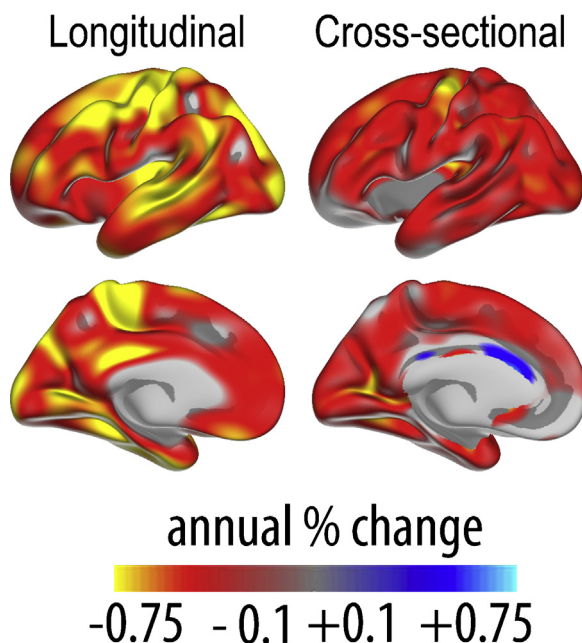


Fig. 6. Cross-sectional vs. longitudinal results. Left panel shows percentage annual change in cortical thickness measured longitudinally in a sample of healthy elderly ($n = 207$, 60–93 years). The right panel shows percentage annual change in cortical thickness estimated cross-sectionally in the same sample of participants. As can be seen, the apparent thickening of the anterior cingulate cortex in the cross-sectional analyses is not confirmed by the longitudinal results, indicating that this likely arises from issues with selective sampling. Also, estimated change is smaller in the cross-sectional compared to the longitudinal results. In other respects, the results are more similar.

Data from [Fjell et al., in press-b](#).

With such discrepancies between cross-sectional and longitudinal results in mind, it is important to be cautious about inferences from one to the other. Still, the differences are not always as prominent as in these examples, and cross-sectional and longitudinal results often converge ([Fjell et al., in press-b](#)). Thus, confirming results by use of both cross-sectional and longitudinal analyses will strengthen conclusions.

2.2. How best to quantify brain change?

Surface-based segmentation techniques allow structural cortical changes to be measured along several different dimensions, such as thickness, area, volume, gyrification and signal intensity or gray matter–white matter contrast. In recent years, more researchers have been conscious of the differences between these measures. Volume is the more complicated measure to interpret, since it is the product of two genetically independent metrics (thickness and area) ([Panizzon et al., 2009](#)), and is affected by fundamentally different neurobiological properties during development and evolution ([Rakic, 2009](#); [Rakic et al., 2009](#); [White et al., 2010](#)). Area has expanded tremendously during evolution, without comparable increase in thickness: a twofold difference in cortical thickness between rodents and primates does not measure up to the 1000-fold difference in total cortical surface area between mice and humans ([Rakic, 2009](#)). In human adults, weak ([Winkler et al., 2010](#)) or even negative ([Hogstrom et al., 2012](#)) correlations between thickness and area have been observed. It has been suggested that myelin growth, both intracortically and in subcortical regions, may stretch the cortex tangentially to the surface ([Seldon, 2005](#)). This is hypothesized to contribute to disentangling of neighboring neuronal columns and enabling the relevant parts of the cortex to better differentiate afferent signal patterns increasing functional specialization ([Seldon, 2007](#)). This model provides one possible mechanistic and functional account for a negative relationship between cortical thickness and surface area. Few studies have systematically compared differences in age-vulnerability between cortical thickness, area and volume, but there are indications that thickness is more affected than area ([Hogstrom et al., 2012](#)) (see [Fig. 7](#)). In sum, although volume may be a sensitive measure of atrophy and age-related differences, the underlying neurobiological causes leading to volume reduction are difficult to identify due to the independence of the two constituting measures of cortical thickness and area.

There is discussion about whether changes in glucose metabolism in aging as evidenced by [18 F]fluorodeoxyglucose positron emission tomography (FDG-PET) occurs prior to, at the same time as, or is a consequence of atrophy. While this discussion is outside the scope of this paper, it can be noted that glucose metabolism is reduced in normal aging, e.g. in anterior regions such as the anterior cingulate ([Yoshizawa et al., 2014](#)), and it is suggested that hypermetabolism in specific regions, e.g. anterior cingulate, as well as functional connectivity, are positively related to cognitive reserve ([Arenaza-Urquijo et al., 2013a](#)). However, the relationship between degeneration, cognitive function and metabolism is not simple, as other studies have found hypermetabolism in amyloid positive non-demented elderly, e.g. in the bilateral superior temporal gyrus ([Johnson et al., 2014](#)). In a comprehensive study of age-changes in atrophy and metabolism based on the ADNI database, age- and symptom-severity dependent glucose metabolism both in temporal, parietal and precuneus regions was found in a sample of AD patients and healthy controls ([Dukart et al., 2013](#)). The authors further suggested a dissociation with larger effects on metabolism compared to atrophy in early phases of AD and the opposite pattern at more advanced stages. Targeted investigations into the relationship between changes in metabolism and atrophy have

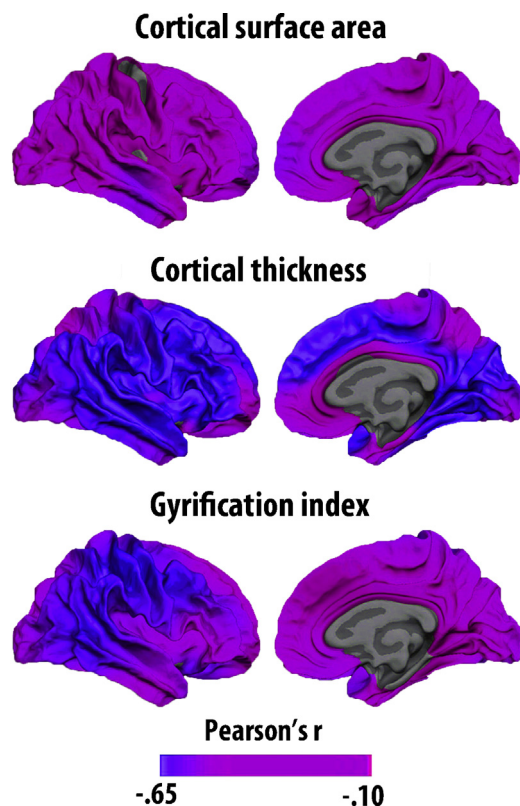


Fig. 7. Different effects of age on cortical surface area, thickness and gyrification. Different structural features of the cerebral cortex have different genetic architecture and are reflecting different neurobiological events. Cortical surface area, thickness and gyrification are all negatively related to age, but to a different degree and with somewhat different regional distribution of effects. The results are displayed on the white matter surface of the brain. Gyrification index is defined as the ratio between the buried cortex and the visible, outer surface cortex. Data from [Hogstrom et al. \(2012\)](#).

the potential to yield important insights into the cerebral changes in normal aging and their cognitive correlates.

2.3. Principles of regional cortical age-changes: development and evolution

As reviewed above, cortical reductions are relatively global in normal aging, not being confined to specific regions. Still, substantial heterogeneity is seen, with several studies identifying fronto-temporal areas as especially prone to the effects of aging. Attempts have been made to explain the regional differences in age-vulnerability based on different overarching principles. While some of these proposed principles capture critical aspects of brain aging, all also have substantial weaknesses. One popular view is that late-maturing regions are most vulnerable to age-changes, often referred to as the “last in, first out” or “retrogenesis” hypothesis. These areas tend to have a more complex cortical architecture than areas that develop early. The finding that late-maturing medial frontal areas, such as the medial orbitofrontal cortex ([Gogtay et al., 2004](#); [Tamnes et al., 2012](#)), are highly vulnerable to aging has led to the hypothesis of an anterior-to-posterior gradient of age vulnerability, which has been invoked to explain the decline in executive function often observed in healthy elderly samples ([Gunning-Dixon and Raz, 2003](#); [Head et al., 2009](#)). For instance, while degeneration of the medial temporo-parietal memory network is characteristic for AD, change in a fronto-striatal network supporting executive functions has been proposed as a hallmark of healthy aging ([Buckner, 2004](#); [Head et al., 2005](#)) (see [Fig. 8](#)). Decline of executive functioning can impact memory

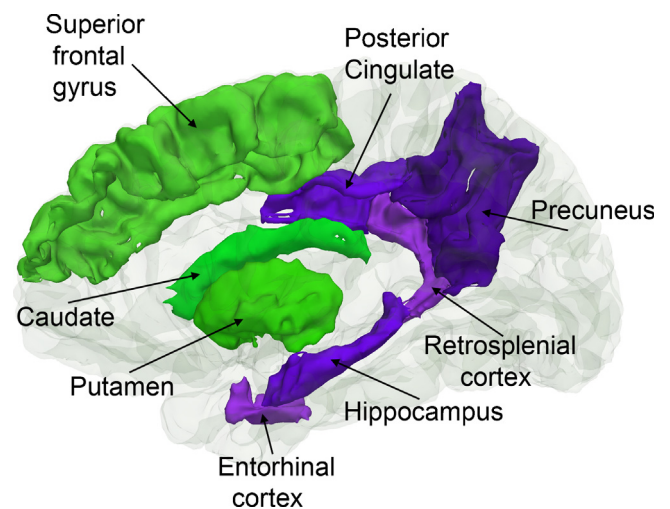


Fig. 8. Fronto-striatal vs. temporo-parietal networks. While normal aging affects a fronto-striatal network important for cognitive control and executive function (green structures), AD has additional effects on a temporo-parietal network important for episodic memory function (purple structures).

function, and contribute to memory problems in non-demented elderly. This can occur even in MCI patients. It has been shown that MCI patients with higher executive function perform better on episodic memory tests, with higher memory performance related to thicker cortices in regions outside the medial temporal lobe, including prefrontal regions ([Chang et al., 2010](#)). From this view, even though reduction in episodic memory function is seen in both AD and healthy aging, this reduction could have partially different brain structural causes. Thus, the “last in, first out” model has many similarities to the traditional “frontal theory of aging” ([West, 1996](#); [Robbins et al., 1998](#); [Rabbitt, 2005](#)).

Although the “last in, first out” model is consistent with some of the data on brain aging, there are important discrepancies between the available data on cortical maturation and aging. This becomes especially evident when maturation and aging of the medial temporal lobes are compared. Medial temporal lobe structures are typically found to undergo relatively modest maturation after school age is reached ([Shaw et al., 2008](#); [Ostby et al., 2009](#); [Tamnes et al., 2012](#); [van Soelen et al., 2012](#)), yet are strongly impacted by age, especially after 60 years (see [Figs. 4 and 5](#)). Thus, in this respect, “the last in, first out” theory of brain maturation does not account for one of the most important features of cortical brain aging – the vulnerability of medial temporal regions.

The “last in, first out” hypothesis is based on a view that more complex brain areas, supporting higher-order cognitive functions such as executive functions, take longer to develop, follow more complex developmental trajectories ([Shaw et al., 2008](#)) and are more vulnerable to the negative effects of normal aging. A version of this hypothesis takes human evolution as a starting point instead of ontogenetic development, applying the same principles of cortical complexity and age-vulnerability. Human cortical surface area has expanded tremendously during evolution ([Kaas, 2008](#)), more than what can be explained by body size alone – a phenomenon referred to as encephalization ([Fonseca-Azevedo and Herculano-Houzel, 2012](#)). The adaptive benefits of improved general intellectual function likely underlie this expansion ([Gibson, 2002](#); [Sherwood et al., 2008](#)). Cortical expansion during evolution is spatially heterogeneous, with large scaling effects in some regions and minute effects in others ([Van Essen and Dierker, 2007](#)). Interestingly, Hill et al. recently demonstrated similarities between cortical expansion in evolution and development ([Hill et al., 2010](#)) – evolutionary high expanding cortical areas tended to show high developmental expansion in childhood, suggesting that

evolutionary factors have shaped ontogenetic cortical development. To a certain extent, cortical regions supporting mental capacities in which humans excel compared to other primates have expanded the most (Haug, 1987; Sherwood et al., 2008; Fjell et al., in press-a), and it has been suggested that human-specific cognitive adaptations are correlated with enlargement of the neocortex (Sherwood et al., 2008). Thus, it is possible that high-expanding cortical regions during evolution will have a protracted ontogenetic developmental course, tend to support higher-order late-maturing cognitive functions, and be more vulnerable to decline in normal aging.

In Fig. 9, cortical area expansion maps from macaque to humans, previously published in (Bardet et al., 2007; Hill et al., 2010), and longitudinal atrophy maps from healthy ADNI participants, are projected onto the same cortical surface and displayed on the same scale (Z scores). As can be seen, there is overlap between high-expanding cortical areas during evolution and areas that undergo rapid decline during normal aging. Contrasting annual percentage decline in regions of high, medium or low evolutionary expansion revealed clear differences in a dose-dependent manner. Annual percentage change in cortical volume increase from 0.23 to 0.41 and 0.53 from low to medium to high expansion regions. Overlap between evolutionary expansion and cortical age-vulnerability was most prominent on the lateral surface, with high expansion and high rate of atrophy in the lateral temporal cortex, inferior regions of the parietal cortex and most of the lateral prefrontal cortex.

Medially, however, there are important differences. High expansion and high rate of atrophy are seen in parts of the superior frontal cortex and the anterior cingulum. Interestingly, the medial temporal lobes, including entorhinal and parahippocampal cortices, as well as posterior cingulate/retrosplenial cortex and the precuneus, show a high degree of decline in normal aging, but relatively low expansion during evolution. Also, the most anterior part of the cingulate cortex, the subgenual cortex (Brodmann area 25) shows substantial age-decline without high evolutionary expansion. Several of these areas belong to the allocortex or archicortex, which is phylogenetically older and has a cellular organization that deviates from the typical six-layered structure of most of the rest of the neocortex. For instance, parts of the cingulate cortex have a less distinct laminar differentiation (Vogt et al., 1995), lacking the internal granular layer (IV). These evolutionary low-expanding, age-vulnerable cortical regions overlap well with the Papez circuit, important in normal memory function, and also with areas previously found to show simpler developmental trajectories (Shaw et al., 2008).

A note of caution must also be included in a discussion of phylogenetic brain expansion, because it is challenging to draw inferences about brain evolution from inter-species comparisons. For instance, it has been argued that comparison of proportional size of different brain areas across species conflate selective enlargement with allometric scaling, and consequently that it is invalid to infer that humans are specialized for frontal lobe functions in particular compared to other species (Barton and Venditti, 2013). Due to allometric constraints on connectivity caused by the fast increase in white matter during evolution, prefrontal volume is expected to take up a higher proportion of the brain in humans than in non-human primates with smaller brains. In a recent paper, Barton and Venditti argued that the widely held assumption that human brain evolution involved relative expansion of the frontal lobes may lack support, and that "... unless one is willing to take seriously the hypothesis that lemurs have more of the qualities bestowed by the frontal cortices than do humans, or that llamas possess more than monkeys, it must be concluded that testing the hypothesis that any species is specialized for frontal functions (...) requires scaling to be taken into account." (Barton and Venditti, 2013, pp. 9001–9002). Further, it should also be noted that all living species have their own evolutionary history, and thus cross-species comparisons cannot be equated with evolutionary adaptations.

2.4. Principles of regional cortical age-changes: default mode network

Interestingly, the more recently described default mode network (DMN) is almost completely encapsulated in the high-declining regions in Fig. 2. DMN (Raichle et al., 2001) has received much attention recently (Snyder and Raichle, 2012), and its robust appearance across studies indicates that it plays a major role in human brain function (Buckner, 2012). For a delineation of some important DMN regions, see Fig. 10. The DMN consists of a specific set of brain areas that decrease activity during performance of a wide range of tasks, and that are typically also active also during periods of rest or introspection (Snyder and Raichle, 2012). The different DMN structures are densely interconnected to each other and limbic structures by polysynaptic connections, with few connections to sensory and motor areas (Binder et al., 2009; Buckner, 2012). Structural and functional aspects of DMN are affected both in normal aging (Lustig et al., 2003; Andrews-Hanna et al., 2007; Addis et al., 2011) and AD (Greicius et al., 2004; Walhovd et al., 2010; Jones et al., 2011). DMN maps onto a network of core brain areas involved in episodic memory and imagination (Buckner and Carroll, 2007; Schacter et al., 2007), and is therefore

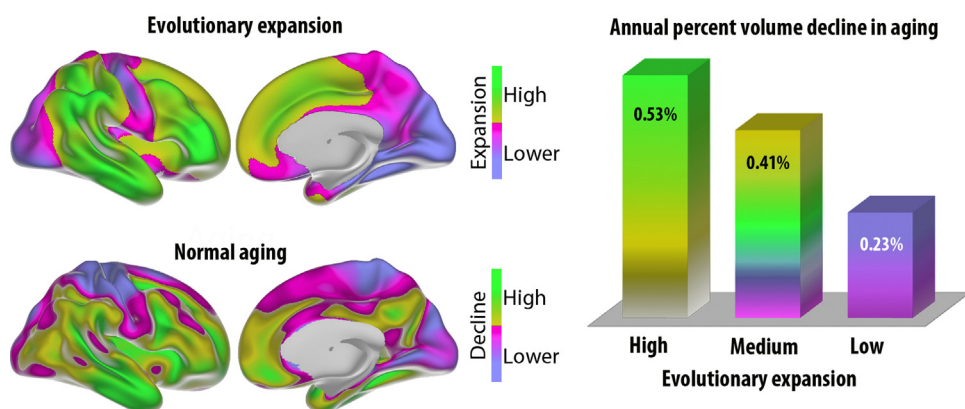


Fig. 9. Cortical expansion across primates and volume decline in aging. Upper panel shows regions of high vs. lower cortical expansion from macaque monkeys to humans (maps re-computed from Hill et al., 2010). Lower panel shows regions of high vs. lower volumetric reduction in aging (based on the data from Fig. 2). There is overlap between evolutionary high-expanding regions in temporal and frontal cortex and regions with high decline in aging. The right panel shows annual volumetric decline in regions of high ($z > 0.5$ SD), medium ($-0.5 < z < 0.5$) and low ($z < -0.5$) evolutionary expansion. As can be seen, annual decline in aging is related to degree of expansion during evolution.

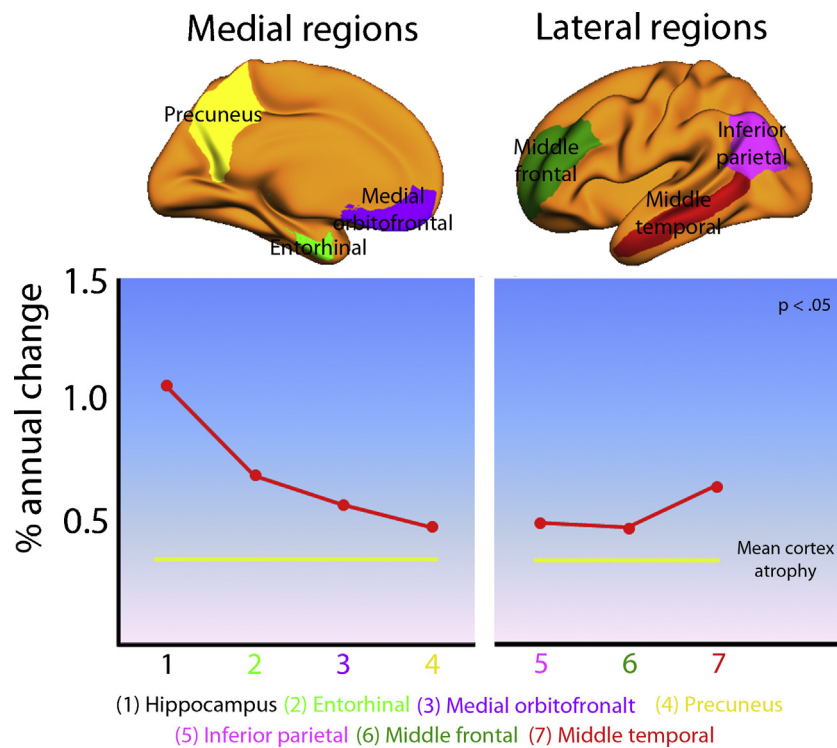


Fig. 10. Aging of the default mode network. Annual percentage change in brain volume for selected areas of the default mode network (DMN) in 132 healthy elderly (based on data from Fig. 2). Both the medial and the lateral DMN changed significantly more than the whole brain rate of 0.44% (mean atrophy in the medial DMN: 0.70%, $t[131] = 4.75$, $p < 10^{-5}$; mean atrophy in the lateral DMN: 0.53%, $t[131] = 2.10$, $p < .05$). Compared to the average cortical change (not including hippocampus) of 0.39%, the age-vulnerability of DMN is larger.

highly relevant for the understanding of episodic memory problems in aging. Recent normative studies (Laird et al., 2009; Andrews-Hanna et al., 2010) have pointed to at least seven important regions within the default mode and episodic memory/simulation network: rostral middle frontal, inferior parietal cortex and middle temporal cortex on the lateral side, and medial orbitofrontal cortex, precuneus, entorhinal cortex and the hippocampus on the medial side (although the roles played by hippocampus and entorhinal cortex are debated (see Nyberg et al., 2010a; Squire et al., 2010; Cooper et al., 2011; Yeo et al., 2011)). Early on it was suggested (Andreasen et al., 1995) that the network primarily consists of association cortices that have undergone especially large changes from nonhuman primates to humans, have the most complex columnar organization and are slow to reach mature levels of myelin (Catani and Ffytche, 2005). However, while this is true for some of the structures involved, especially on the lateral surface as shown above, a characterization of the DNM structures overall as unique to higher-order evolution would not be correct. Rather, some of these structures, such as the hippocampus, the entorhinal cortex and the retrosplenial cortex, share considerable similarities, likely both structurally and functionally, across several species. These structures belong to the medial temporal lobe DMN subsystem (Andrews-Hanna et al., 2010), critical for the ability to navigate in time and space in many species. At the same time, this is a capability that is affected relatively early in AD. Also the medial parietal cortex, including the precuneus, belonging to the midline core of the DMN in the same classification, has expanded relatively little during evolution (see Fig. 9). Thus, there are considerable areas anatomically not overlapping across the DMN and evolutionary cortical expansion maps. Although DMN likely serves important functions for human cognition and mental life, resting-state networks overlapping the midline core of DMN have also been identified in monkeys. In a recent study, functional networks were directly compared

between humans and monkeys, and both inter-species corresponding and human-specific networks were identified (Mantini et al., 2013). In sum, the theory of DMN as localized in evolutionary recent cortical areas does not seem justified.

Naturally, DMN is a prime target for investigation into the structural correlates of reduced cognitive function in both normal aging and AD. As can be seen from Fig. 10, rates of yearly cortical volume reduction in these areas are higher than in most other cortical regions, indicating that the DMN is vulnerable to normal aging. This is also true for the posterior cingulate/retrosplenial/precuneus and MTL, which have been relatively low-expanding in evolution. A speculative suggestion may be that these areas support fundamental aspects of episodic memory that would have large adaptive benefits during early evolution, such that they are relatively well developed even in non-human primates. Recent studies have compared resting state (Hutchison and Everling, 2012) and task-related functional networks between humans and non-human primates (Mantini et al., 2012; Miyamoto et al., 2013), showing functionally correspondent networks across species. Corballis recently argued for the existence of evolutionary continuity in the capacity for mental time travel (Corballis, 2013a,b), which is closely connected to hippocampal and medial parietal cortical function. Although the discussion of whether this capacity exists in non-human primates is far from settled (Suddendorf, 2013), a natural conclusion from available evidence is that DMN is vulnerable to age-changes for reasons not grounded in evolutionary expansion.

Several studies have shown that longitudinal brain changes especially in MTL are correlated with changes in cognitive capacities, episodic memory function in particular (Rodrigue and Raz, 2004; Murphy et al., 2010; Persson et al., 2012; Fjell et al., in press-b; Nyberg et al., 2012). Thus, the DMN seems to provide a better account of the distribution of structural cortical changes in normal aging than the frontal or the “last in, first out” theories. Of

special interest is the vulnerability of DMN to AD (Lustig et al., 2003), and the high overlap between DMN and areas of A β deposition and PET-FDG hypo metabolism (Buckner et al., 2005). We discuss this in more detail in Section 4.

3. Normal aging vs. Alzheimer's disease: how to tell the difference?

Age-related dementia typically has a slow and gradual onset, with atrophy manifest years in advance of clinical symptoms (Davatzikos et al., 2009; Jack et al., 2010a, 2013). Some have even suggested that cortical decline in aging is caused by undetected disease and is not a feature of normal aging (Burgmans et al., 2009). An important implication of the apparent temporal gap between detectable cerebral and cognitive expression of dementia is that cases with undetected disease in presumably normal samples could bias inferences about normal brain aging (Sliwinski and Buschke, 1999). This is a particular concern since the proportion of elderly with undetected neurodegenerative disease is expected to increase with the age of the population, potentially leading to invalid conclusions of accelerated age-related decline in cortical areas vulnerable to disease pathology, especially the entorhinal cortex and hippocampus (Braak and Braak, 1985; Jack et al., 1997; McDonald et al., 2009). However, it is also possible that stringent screening criteria could result in an over-representation of very high-functioning elderly in convenience samples (Nyberg et al., 2010b), thereby reducing the amount of incipient dementia in aging-studies of the oldest old. The contribution of latent pathology to age-related decline in apparently normal aging thus remains an open question. However, differences in AD biomarkers, such as patterns of atrophy and amyloid deposition may shed light on this issue.

The pattern of atrophy in AD is not random, but usually evolves slowly, following a specific pathway that first involves the entorhinal cortex and the hippocampus, then spreads out to association areas in medial parietal, lateral temporal and frontal regions, eventually affecting all regions of cortex. In Fig. 11, rate of atrophy in healthy controls vs. patients with MCI or AD from ADNI is shown. Even at the stage of MCI, annual atrophy is several times higher than in normal aging, with further increases in atrophy rate with progression to a full AD diagnosis.

The organized pattern of brain changes in early AD can be used to differentiate between non-demented elderly with lower vs. higher risk of progression to MCI or AD (Driscoll et al., 2009; McEvoy et al., 2009, 2011). For instance, Driscoll and colleagues identified a pattern of brain changes typical for AD patients, including temporal and orbitofrontal cortex, and argued that the differential patterns of decline in participants who subsequently developed MCI differed from that of participants who remained cognitively stable. The finding that both those who progress to AD and those who remained cognitively stable showed accelerated tissue loss indicates that atrophy does not necessarily signify impending dementia, and thus may not necessarily reflect latent AD. Likewise, McEvoy and colleagues used linear discriminant analysis to identify a pattern of regional atrophy by comparing healthy controls with AD patients, and found accelerated atrophy in the temporal cortex, isthmus of the cingulate and orbitofrontal cortex. These results could later be used to predict conversion to AD in ADNI's MCI participants.

These two examples show that magnitude of atrophy in collections of brain areas can be used to distinguish normal aging from AD. Still, when looking at atrophy maps in normal aging, e.g. Fig. 2, it is evident that temporal regions, both medial and lateral, are among those that decline the most. Even though hippocampus and entorhinal cortex are the areas that best distinguish AD from controls in terms of atrophy rates (Fjell et al., 2010b), and these

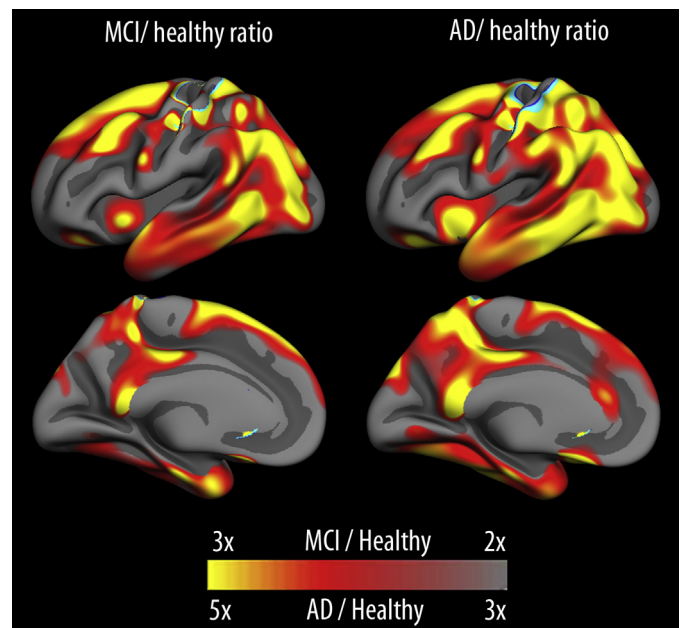


Fig. 11. Differences in cortical atrophy rates between healthy elderly and Mild Cognitive Impairment/Alzheimer's Disease. In mild cognitive impairment (MCI), rates of cortical volumetric reductions are more than double that seen in healthy aging across large areas of the cerebral cortex, and rates more than three times larger are seen in AD. Please note that the scale is different between MCI/healthy and AD/healthy to allow appreciation of the regional patterns of effects across groups.

Data from the Alzheimer's Disease Neuroimaging Initiative.

temporal areas figured prominently in the AD risk patterns in the studies described above, low-risk elderly also show accelerated change in these areas. Similar results have been reported for hippocampus in several independent studies (Driscoll et al., 2009; Fjell et al., 2009a; Raz et al., 2010; Pfefferbaum et al., 2013). Thus, MTL atrophy is common in normal aging and AD. This is illustrated in Fig. 12 (Fjell et al., 2013a). When the large differences in magnitude of atrophy are removed by standardizing the atrophy maps within each diagnostic group, two interesting features become apparent. First, both the low-risk healthy elderly and MCI/AD patients are characterized by elevated atrophy in the temporal lobes, medially and laterally. Atrophy in the temporal lobe exceeded two standard deviations above mean cortical atrophy in all groups. A second interesting feature is the relatively high level of atrophy in the orbitofrontal cortex in the normal controls. In the patients, these areas do not stand out as more atrophic than the rest of the cortex. Thus, the fronto-temporal pattern of atrophy in the low-risk elderly is replaced by an even more distinct temporal pattern of atrophy in the patients. Relatively high levels of atrophy in the medial parietal cortex, i.e. precuneus and posterior parts of the cingulate, were also seen across groups. Of interest, the accelerated prefrontal atrophy in the healthy group overlaps with one of the cortical regions that has expanded the most during evolution, while not being among the regions most vulnerable to early AD. Thus, it seems that evolutionary expansion, even with the discrepancies found in the medial temporal cortex, characterizes the brain changes in healthy aging as good as or better than the atrophy seen in early AD.

The vulnerability of the temporal lobes in aging and AD naturally prompts the question of whether brain-changes observed in non-demented older adults are in part caused by undetected neurodegenerative processes of an Alzheimer type. At age 79, the probability of significant AD neuropathological changes in the brain is 30–40%, but the likelihood of having been diagnosed

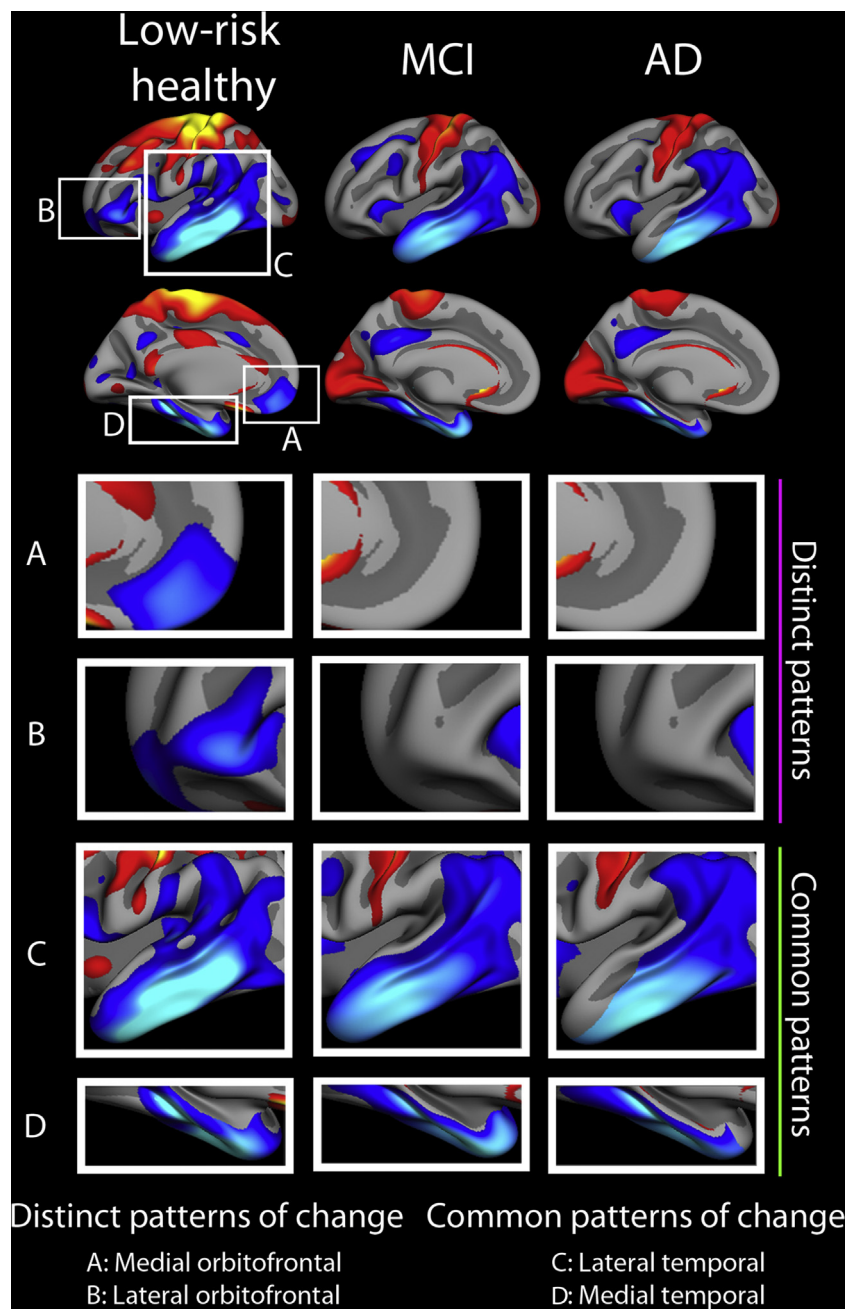


Fig. 12. Comparison of aging and Mild Cognitive Impairment/Alzheimer's Disease. Comparison of the standardized pattern of atrophy in a group of APOE $\epsilon 4$ negative elderly with normal levels of CSF A β , mild cognitive impairment (MCI) and AD. Atrophy maps are standardized within each group by Z-transformation, yielding maps showing areas of more (blue–cyan) vs. less (red–yellow) atrophy for each group in terms of standard deviations. Thus, atrophy is scaled within group, and changes are relative to group means. Across groups, we see common patterns of standardized change in the lateral and medial temporal lobe (including the hippocampus, not shown), and a distinct pattern characterizing only healthy elderly in the prefrontal cortex, especially the orbitofrontal part. Figure from Fjell et al. (2013a). Atrophy across the cortex in the healthy elderly and the AD patients correlated .81, showing substantial overlap in regional vulnerability.

with dementia due to AD is only 15% (Nelson et al., 2012). Subtle cognitive symptoms have been identified more than 10 years before dementia diagnosis (Elias et al., 2000), and the currently popular hypothetical biomarker model suggests that brain changes are detectable before the first cognitive symptoms (Jack et al., 2010a, 2013). Thus, the question of whether brain atrophy in typical AD areas, i.e. the temporal lobes, can be attributed to preclinical manifestations of the disease is not a trivial one. However, there are several pieces of evidence indicating that these brain changes indeed constitute a part of normal, non-disease specific aging.

First, Resnick and colleagues followed 92 healthy elderly from the Baltimore Longitudinal Study of Aging for four years, and found

significant reductions in brain volume in all cortical lobes (Resnick et al., 2003). Of most interest, these changes were also seen in a subgroup of 24 very healthy participants who experienced no medical condition or cognitive impairment four years after the initial examination. The authors argued that the uniformity of tissue loss across individual participants indicated that these were not pathological changes associated with preclinical dementia, unless all participants had been in such a preclinical stage, which is highly unlikely. Thus, these results constitute evidence that tissue loss in older adults does not need to be caused by undetected, preclinical dementia. A similar conclusion was drawn in the later study by the same group where participants were followed for

periods up to ten years after baseline, furthering ruling out the possibility that all participants were in a preclinical stage of AD (Driscoll et al., 2009).

Second, in our own work, we have focused specifically on the fate of AD-prone areas in healthy controls, and attempted to disentangle normal age-changes from changes related to preclinical AD. We quantified one-year cortical thinning in 138 normal elderly from ADNI, and found the expected reductions in the temporal lobe, including the medial temporal lobes and the entorhinal cortex (Fjell et al., *in press-b*). From the full sample of non-demented participants, subsamples with very low probability of AD were identified. A group of cognitively “super-stable” participants ($n = 18$) were selected based on a lack of decline over two years on scores of two clinical measures and seven neuropsychological tests. Another group was selected based on their negative CSF A β test ($n = 28$). A final group consisted of those that were both amyloid negative and negative for APOE $\epsilon 4$ ($n = 22$). Interestingly, rate of cortical thinning in the entorhinal cortex did not differ between the full sample of non-demented older adults and the different low-risk groups. This further supports the view that brain changes in older adults are not necessarily related to early AD-processes, but can be part of the normal aging process, even in areas prone to early AD. Interestingly, in the 94 participants with no change over two years in the two clinical measures, the entorhinal changes correlated with changes in memory performance over time, suggesting that non-AD mechanisms in AD-prone areas may still be causative for cognitive decline within the normal range. With longer follow-up data from ADNI more recently available, investigation of the same participants up to four years after the initial exam has shown that significant entorhinal and hippocampal atrophy can be detected in groups of elderly at very low risk of AD defined by clinical neuropsychological, genetic and biomarker criteria (Fjell et al., 2013a). An independent study also identified small hippocampal volumes in a group of amyloid negative elderly (Knopman et al., *in press*). Further, Oh et al. found correlations between age and cortical volume and thickness, including in the medial and lateral temporal lobe, in healthy elderly even when amyloid status as indexed by PIB PET was used as covariate (Oh et al., *in press*). Such results can be used to argue that brain atrophy in AD-prone regions can be a part of the normal aging-process. Of course, there is no guarantee that such changes are due to aging or non-AD pathology, and it cannot be ruled out

that a component of the net loss may be due to very early AD in individuals stable over several years. Also, a danger associated with low-risk group analyses is that the power to detect differences in decline diminishes with the reduction in sample size. Still, demonstration of cortical atrophy in low-risk older adults makes a strong case that preclinical AD is not the only factor driving brain change in aging.

4. Neuroplasticity in the aging brain – a critical vulnerability factor?

If atrophy in normal aging does not reflect latent AD pathology, a critical question that must be addressed is *why* it is that AD-prone areas are also highly vulnerable to normal age-changes? These areas include not only the temporal regions, but regions around the medial parietal cortex (the precuneus/posterior cingulate/retrosplenial cortex) – an area that is an important component of the DMN. One speculation is that areas characterized by high degree of life-long plasticity are those most vulnerable to detrimental effects of normal and pathological aging (Neill, 1995; Mesulam, 1999; Buflin and Carbonell, 2004; Rapoport and Nelson, 2011; Neill, 2012; Buflin et al., 2013). The pattern of effects may be explained by the special role of the medial temporal lobes and other parts of DMN in learning and memory, with high demands for life-long plasticity required for these cognitive functions (Aimone et al., 2010; Deng et al., 2010). For instance, neurogenesis in the hippocampus may be important for structural plasticity and network maintenance, and it has been suggested that altered neurogenesis in the hippocampus is an early critical event in AD (Mu and Gage, 2011). Down-regulation of neurogenesis in the aged mouse brain has also been found (Lazarov et al., 2010). As neurogenesis in the adult brain is likely limited to the dentate gyrus of the hippocampus as well as the olfactory bulb, it is unlikely to be a main factor in the changes in brain and cognition associated with aging. However other mechanisms related to neuroplasticity could be at play in other age-vulnerable regions. Dendritic spines may represent a primary site of structural plasticity in the adult brain, and spine plasticity seems to play an important role in long-term memory in rodents (Sanders et al., 2012). Dendritic spine plasticity in the prefrontal cortex is reduced in aged rodents (Bloss et al., 2011), and spine density is likely reduced in aging (Jacobs et al., 1997; Esiri, 2007; Freeman et al., 2008; Benavides-Piccione et al., 2013) (see Fig. 13).

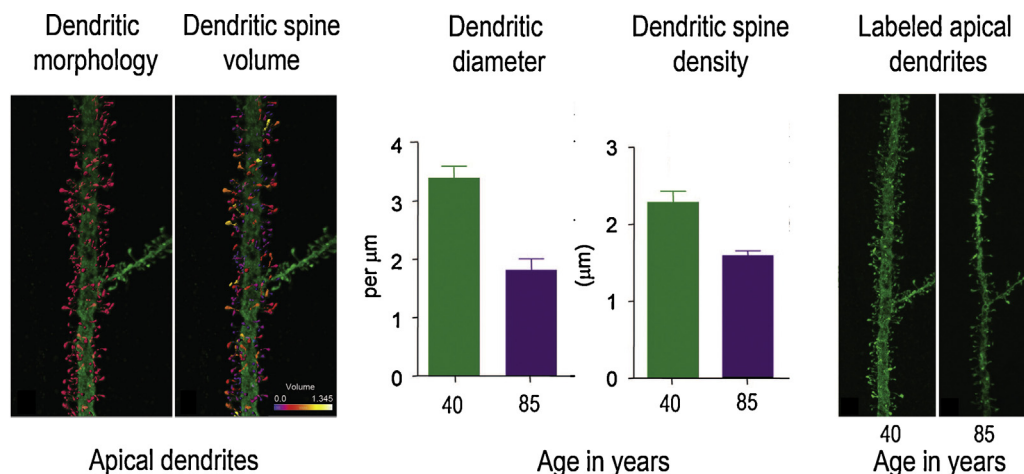


Fig. 13. Dendritic spine morphology in aging. Benavides-Piccione et al. (2013) 3D reconstructed 8900 individual dendritic spines from layer III pyramidal neurons in the cingulate cortex from two male humans of age 40 and 85 years, using intracellular injections of Lucifer Yellow in fixed tissue. The left two panels show the 3D reconstruction of the complete morphology of each spine of an apical dendritic segment at 100 μm distance from the soma, and the estimation of the spine volumes shown in color codes (0–1.345 μm^3). The middle two panels show mean dendritic diameter and dendritic spine density in apical dendrites in the middle-aged and the older participant. The right two panels show examples of apical dendritic segments from the middle-aged and the older participants. The differences in spine morphology are easily seen. Figure modified with permission from Benavides-Piccione et al. (2013).

Another interesting feature is that the medial temporal lobe structure dentate gyrus is the only site of adult human neurogenesis besides the olfactory bulb (Eriksson et al., 1998; Lotsch et al., in press). This is different from other primate species, where it is possible that new neurons can be added throughout life in select regions of the neocortex (Gould et al., 2001; Bernier et al., 2002; Vessel and Darian-Smith, 2010). Thus, the medial temporal lobe is an evolutionary old and at the same time a neuroplastic area within an otherwise much more developed and expanded neocortex. The downside of being lone areas involved in or surrounding this type of plasticity in the human brain may be increased vulnerability and accumulation of negative impacts through life. Lesions in the highly neuroplastic limbic system could then have consequences for association cortices with which they are reciprocally interconnected (Mesulam, 1999).

Thus, although still highly speculative, it may be that certain brain regions are characterized by increased demands for life-long plasticity mechanisms, and that this makes them especially vulnerable to subtle lesions and pathology accumulating through life. The cost of maintained plasticity may be increased vulnerability to factors which can trigger cognitive decline (Buftill et al., 2013). Perturbation of neuroplasticity has previously been proposed as a fundamental principle of a unitary theory that could potentially account for all clinical and neuropathological features of AD, on the assumption that amyloid depositions and neurofibrillary tangles are manifestations of the same underlying phenomenon (Mesulam, 1999). The potential for neuroplasticity is higher in the limbic system than in other parts of the cerebral cortex, which increases its vulnerability to neurofibrillary degeneration. A consequence of this view may be that late-onset AD is not a disease at all, but “*the inevitable manifestation of a failure to keep up with the increasingly more burdensome work of plasticity*” (Mesulam, 1999, p. 526). This is in line with a systems vulnerability view on aging and AD, where critical networks are subject to various negative impacts, aging in particular, rather than being selectively targeted in AD (Jagust, 2013).

Somewhat related to the concept of plasticity is an idea based on the observed overlap between regional distribution of deposition of amyloid and the DMN that the level of brain activity itself may be causally related to amyloid deposition (Jagust and Mormino, 2011; Buckner, 2012). Amyloid depositions are part of the histopathological definition of AD, but the role of amyloid in normal aging and in the transition from normal aging to neurodegeneration is still poorly understood, and the target of several large research initiatives.

5. Amyloid and brain changes in non-demented older adults: preclinical dementia or a role in normal aging?

Amyloid depositions are part of the histopathological definition of AD, and thus much weight is put on in vivo biomarkers of amyloid in contemporary AD research, as reflected in the NIA-AA proposed diagnostic guidelines for Alzheimer's disease and its preclinical stages (Jack et al., 2011; Sperling et al., 2011a). The role of amyloid in neurodegeneration is envisioned to be very early in the cascade of detrimental events that culminates with dementia and a clinical diagnosis of AD (Jack et al., 2010a, 2013). Current models posit that brain atrophy in AD progresses years before clinical symptoms (Jack et al., 2011; Sperling et al., 2011a), but that nerve cell degeneration is downstream from even earlier causative events, especially abnormal processing of amyloid- β peptides (Goedert and Spillantini, 2006; Jack et al., 2010a). It has been suggested that, once initiated, the neurodegeneration in AD progresses independently of its amyloid-trigger, leading to the commonly expressed concern that the therapeutic window for anti-amyloid drugs may close when neurodegeneration has started

to accelerate (Sperling et al., 2011b). Thus, it is absolutely necessary to better understand the relationship between amyloid, brain integrity and memory in healthy elderly. It has been suggested that older adults with normal cognitive function despite AD-related pathology should be the focus of intense investigation, because they may be in the earliest phases of AD (Jagust, 2013). The difficult question is then: What amyloid-related changes in brain and cognition represent AD-related pathology, and what, if any, such changes can be expected as part of the “normal” aging-process?

A β -atrophy correlations have been demonstrated in MCI/AD patients (de Leon et al., 2006; Fjell et al., 2010b) and normal controls (Storandt et al., 2009; Fjell et al., 2010a; Becker et al., 2011; Fortea et al., 2011; Arenaza-Urquijo et al., 2013b), eliciting debate about whether brain atrophy related to amyloid in presumably healthy elderly signifies early AD-related changes (Becker et al., 2011; Fortea et al., 2011; Shim and Morris, 2011; Sperling et al., 2011a) or whether they may be part of the normal aging process (Fjell and Walhovd, 2012). This pertains directly to the question of whether the factors that drive atrophy in AD also play a role in normal aging. In ADNI, relationships between CSF levels of A β_{1-42} and cortical atrophy have been found in MCI patients (Schuff et al., 2009; Fjell et al., 2010b; Jack et al., 2010b; Tosun et al., 2010) and controls (Fjell et al., 2010a; Schott et al., 2010). In the controls, however, CSF A β_{1-42} predicts atrophy only in the subgroup of participants with very low A β_{1-42} levels (an indication of a high level of A β deposition in the brain). In these participants, strong A β_{1-42} – atrophy correlations can be observed in large areas of the cortex, but not MTL. This is shown in Fig. 14. In MCI, the correlations are weaker, and restricted to the temporal lobes, medial parietal cortex and posterior cingulate. Thus, the pattern of correlation is almost non-overlapping between the healthy older adults and the MCI patients. In contrast, no relationship between atrophy and CSF p-tau levels can be seen in the healthy older adults, while the expected correlation between p-tau and temporal atrophy can be seen in MCI patients. Interestingly, recent studies from ADNI have shown interactions between p-tau levels and A β_{1-42} in clinical decline (Desikan et al., 2012) and entorhinal atrophy (Desikan et al., 2011) in non-demented elderly. In general, the anatomical distribution of tau pathology, initially restricted to MTL, fits better with the spreading of atrophy in AD, and also with the earliest cognitive symptoms of memory and navigation problems. Histopathological data indicate that accumulation of tau-related pathology is an earlier event than A β deposition (Braak and Braak, 1997a,b). An interesting feature is that A β is only modestly related to age, while at least levels of total tau increases substantially. As age is the number one risk factor for AD, understanding brain events that also increase in intensity with age would set us in a position to better understand why aging plays such a prominent role in AD development. A recent study found that MTL tangle formation in non-demented elderly was independent of neuritic plaques, and proposed an age-related, non-amyloid process contributing to this (Mungas et al., 2014). In addition, it was suggested that a primary amyloid-based AD process would make the largest contribution to MTL tangles. Other studies have also demonstrated MTL tangles without neuritic amyloid plaques (Nelson et al., 2009). Thus, there may be both direct amyloid-independent and indirect amyloid-dependent processes causing tangle formation, leading to cerebral and cognitive decrements in aging. Age-related changes in tangle formation in MTL could contribute to explain the high levels of MTL atrophy observed even in healthy elderly at low AD-risk and the consequent decline in episodic memory.

The relationship between A β and tangle formation is also closely related to the discussion of which brain changes in aging result from early AD-related processes and which are AD-independent. A β -atrophy correlations in cognitively stable older

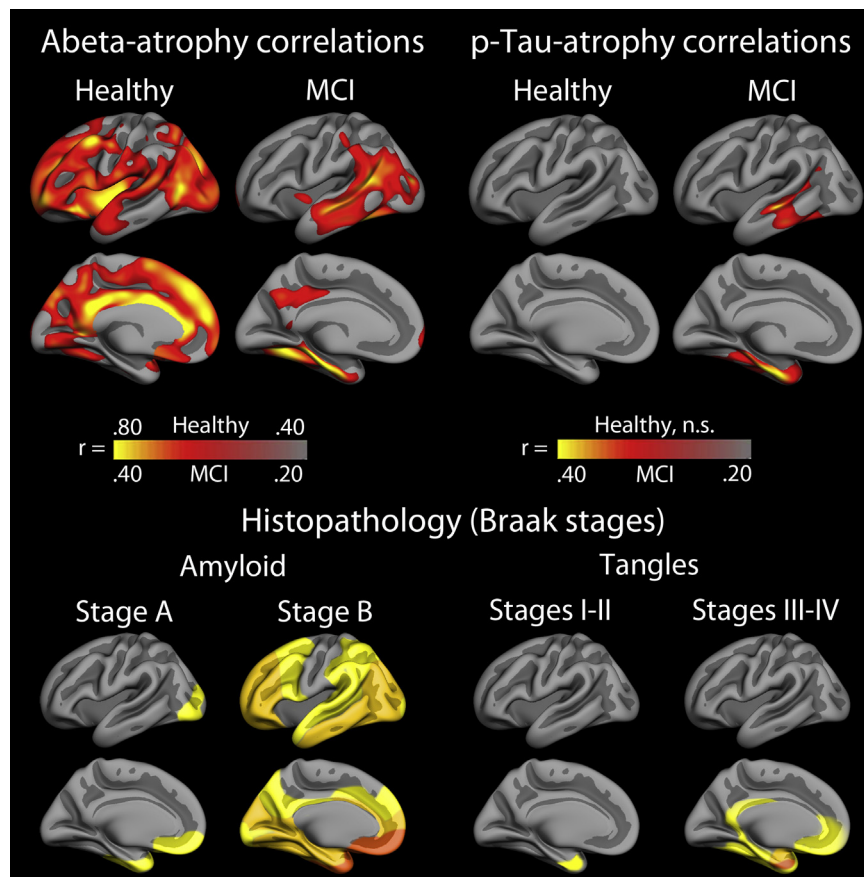


Fig. 14. CSF biomarkers and regional atrophy in aging and MCI. Upper panel: CSF biomarkers of A β and p-Tau, indexing brain levels of amyloid and tangle load, respectively, correlate with temporal atrophy in stable MCI patients from ADNI ($n = 213$). In contrast, the correlations between atrophy and A β in amyloid positive healthy elderly (A $\beta_{1-42} < 175$ pg/ml; Fjell et al., 2010a) are seen in other cortical areas almost exclusively. This may indicate that the relationship between atrophy and amyloid is different in healthy aging and MCI, and that the selective lack of relationship between amyloid beta and atrophy in the medial temporal lobe in controls may represent a key to understand why these participants have not progressed to AD. Lower panel: Histopathological maps from Braak and Braak, projected onto the same template brain. The topographical pattern of accumulated amyloid in AD brains resembles the A β -atrophy correlations seen in MCI to a much larger extent than the A β -atrophy correlations seen in non-demented elderly. Note, both the medial wall and the insula are masked out from the histology maps.

adults have been interpreted as reflecting early AD (Tosun et al., 2010; Becker et al., 2011; Fortea et al., 2011; Shim and Morris, 2011; Sperling et al., 2011a). However, the healthy older adults shown in Fig. 14 are not on the verge of converting to MCI or AD, as evidenced by their status as normal controls two years later, and their normal levels of atrophy and normal memory function (Fjell

et al., 2010a). The few other studies testing A β -atrophy correlations in healthy elderly in vivo generally show the same lack of medial temporal lobe-relationships in the cerebral cortex, especially in entorhinal cortex (Storandt et al., 2009; Fjell et al., 2010a; Tosun et al., 2010; Becker et al., 2011; Oh et al., 2011; Arenaza-Urquijo et al., 2013b) (see Fig. 15 for examples), even

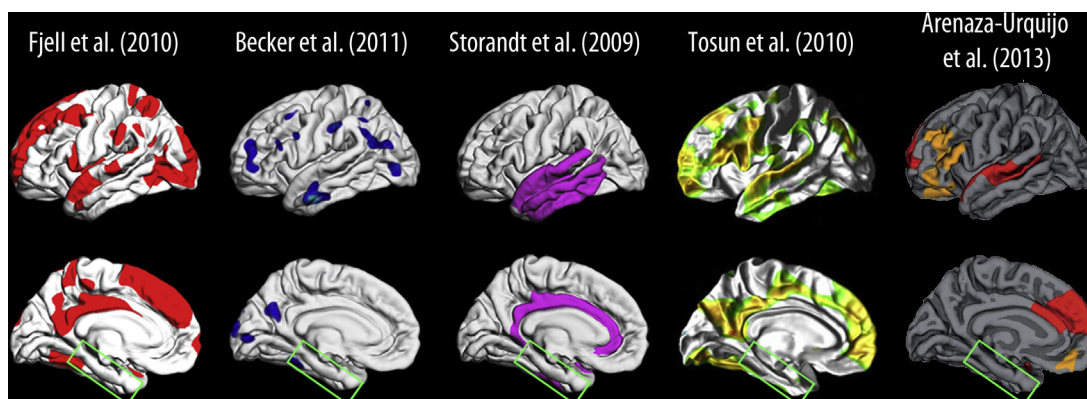


Fig. 15. Selected studies on amyloid and cortical structure in non-demented older adults. Representative studies of the relationship between amyloid levels (PiB PET or CSF A β) and cortical thickness (baseline) or cortical atrophy (longitudinal) in healthy elderly (Storandt et al., 2009; Fjell et al., 2010a; Tosun et al., 2010; Becker et al., 2011; Arenaza-Urquijo et al., 2013b). Common for these studies was the use of anatomically unbiased surface-based cortical analyses using FreeSurfer (surfer.nmr.mgh.harvard.edu). We extracted the effect sites from the published figures, and projected them onto the same standard brain to allow visual comparison of the results. For some, the colors of the effects were changed to aid discriminability between the different studies. The main conclusions are that A β levels are not related to cortical thickness/atrophy in cognitively healthy elderly in typical AD areas in the medial temporal lobe (green box).

though the results sometimes are complex (Bourgeat et al., 2010; Chételat et al., 2010a,b). In contrast, A β -atrophy correlations in MCI are found mainly in the temporal lobe, laterally and medially. Correlations between A β and hippocampal volume (Mormino et al., 2009; Arenaza-Urquijo et al., 2013b) or atrophy (Fjell et al., 2010a) have been observed, but when compared to correlations with other subcortical structures, A β -atrophy relationships often do not stand out as especially strong (Fjell et al., 2010a). This was also found in a longitudinal study from the Australian Imaging, Biomarker and Lifestyle Study of Aging (AIBL). Healthy controls with higher levels of amyloid deposition showed more hippocampal atrophy, but also more total gray matter loss (Villemagne et al., 2013). As recently reviewed, significant hippocampal atrophy in amyloid-positive cases has been inconsistently observed (Chételat et al., 2013).

One way to reconcile these findings could be to hypothesize that A β reflects processes related to brain changes in elderly without AD-symptoms, but that it is only when these processes impact MTL that cognitive decline of the sort observed in early AD/MCI becomes evident. As described above, MTL plays a special role in learning and memory, with high demands for life-long plasticity and neurogenesis required for these cognitive functions. Rodent research has shown that A β , although usually in extremely high doses, can affect synaptic function by modulation of maturation and plasticity of newborn neurons in hippocampus (Bischoff et al., 2009; Lilja et al., 2013). Thus, the impact of A β on the temporal lobes may cause malfunction that cannot be compensated for by mechanisms of neural plasticity elsewhere in the brain. Although the atrophy-driving mechanisms may be related to A β , the causal role of amyloid in neurodegeneration in healthy older adults remains an open question. Such an account may call for a revision of the prevailing view of amyloid biomarkers in normal aging and AD, in that healthy aging and AD may have partly common neural mechanisms for brain atrophy (see also Glass and Arnold, 2012). A critical issue may thus be the regional distribution of A β , not only the amount (see Fig. 16).

Several authors have pointed out the overlap between the DMN and the distribution of amyloid (for a review, see Buckner, 2012),

and it has been suggested that lifelong patterns of neural activity lead to increased A β production and deposition (Jagust and Mormino, 2011). It has been demonstrated that synaptic activity increases amyloid-beta levels in mice (Cirrito et al., 2005), and that neural activity regulates the regional concentration of A β , which again drives local A β aggregation (Bero et al., 2011). Thus, the cognitive importance of DMN has been envisioned to cause its susceptibility to A β (Buckner, 2012, p. 11): “AD may be so devastating to higher cognitive function specifically because it targets brain systems expanded in the human brain and important to the default network, robbing the ability to remember and imagine. The clinical insight from this connection is that downstream consequences of normal basal activity and metabolism may facilitate AD pathology.” Further support for the neural activity-amyloid mechanism has come from recent research showing a relationship between brain aerobic glycolysis and A β (Vaishnavi et al., 2010; Vlassenko et al., 2010). Education is an often-used proxy for cognitive reserve, which is defined as the ability to maintain cognitive function despite high levels of AD pathology (Stern et al., 1999; Stern, 2006, 2012). Higher education is typically regarded as facilitating the development or use of alternate strategies to cope with pathology. However, according to the activity-dependent amyloid accumulation view, the situation may be the opposite. Higher education itself could lead to the build-up of amyloid pathology through its effect of promoting life-long intellectual activity.

This theory is intriguing, but prompts difficult questions: First, both targeted, experimental cognitive interventions (Engvig et al., 2010; Zatorre et al., 2012) and lifespan mental activity (Valenzuela et al., 2008) are shown to have positive effects on brain atrophy and cognitive function in elderly. One study even found a relationship between self-reported lifespan cognitive activity and lower amyloid deposition as measured by Pittsburgh Compound-B (PiB) PET (Landau et al., 2012), and the authors suggested that “(...) lifestyle factors found in individuals with high cognitive engagement may prevent or slow deposition of β -amyloid, perhaps influencing the onset and progression of AD.” A recent study by Selkoe and colleagues found that enriched environments increased synaptic plasticity in the hippocampus and yielded better

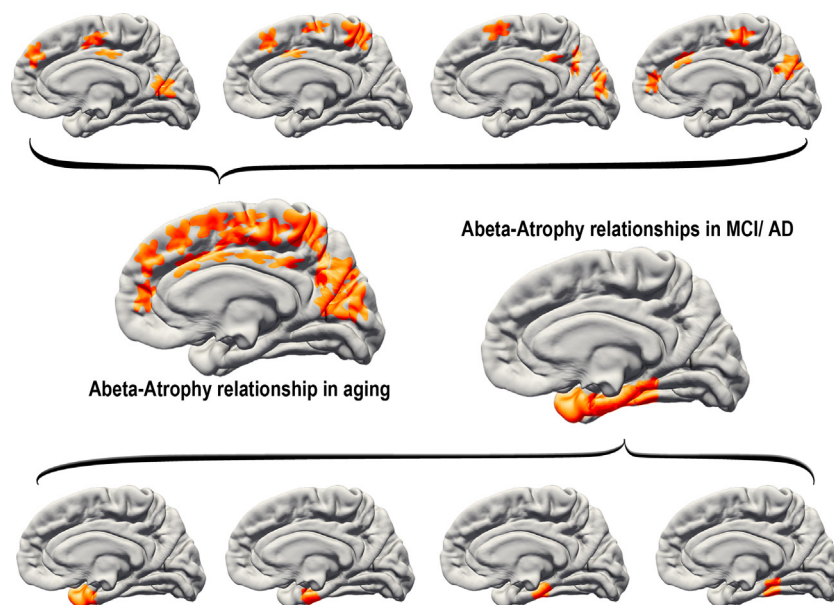


Fig. 16. A speculative model of A β -atrophy relationships in normal aging and MCI. One speculative explanation for the discrepant A β -atrophy relationships in healthy aging vs. MCI (see Fig. 14) would be to assume that lesions related to amyloid levels can occur in several places in the cortex in aging. Lesions targeting the temporal lobe will tend to yield memory problems that are difficult to compensate for, increasing the likelihood of an MCI/AD diagnosis. In contrast, lesions in other parts of the cortex are less directly related to memory problems and may also be easier to compensate for, thus reducing the likelihood of an MCI/AD diagnosis. This would cause the observed discrepancy in regional distribution of amyloid-atrophy relationships between clinically normal older adults and MCI patients.

resistance to toxic effects of A β oligomers in wild-type mice (Li et al., 2013). The researchers then went one step further, and showed that blocking the β 2-adrenergic pathway hindered the positive effects of environmental exploration while stimulating β 2-adrenergic signaling yielded the same benefits as environmental enrichment, thus providing a clue to one possible mechanism behind the often-reported positive association between intellectual activity and cognitive function. The idea that neural activity itself may initiate a cascade of neurodegenerative events needs to be reconciled with such findings.

Second, even if the overlap between DMN and early A β deposition, as evidenced by amyloid imaging, is causally related to the high activity of certain parts of the DMN (Jagust and Mormino, 2011; Buckner, 2012), it yet not clear whether the life-long pattern of brain activity is higher in the areas with higher A β depositions, e.g. the medial posterior parietal cortex, than in the areas with less, such as the medial temporal lobes. The hippocampus and surrounding cortices are involved in encoding of all new information, consolidation of memories, episodic memory retrieval and spatial navigation (Buzsaki and Moser, 2013), and even show a high level of activity during sleep (Grosmark et al., 2012). As can be seen in Fig. 14, A β deposition is detected in histopathological studies in the medial temporal lobes before advanced AD (Braak stage A), but is usually not found to be among the first regions to show increased amyloid deposition by amyloid PET imaging in non-demented elderly (Sojkova et al., 2011; Mormino et al., 2012). If cognitive activity is directly related to amyloid production and later deposition, we need to understand why amyloid deposition is not more evident in these areas in early phases. One explanation could be that cognitive activity increases brain efficiency – people who perform tasks as experts or who are better trained actually may use fewer neural resources. Thus, people who remain cognitively active throughout life develop more efficient ways of performing mental tasks, and therefore produce less A β (Jagust, personal communication). This could possibly unite the seemingly contradictory findings of increased A β production with synaptic activity combined with possibly lower levels of A β deposition with sustained intellectual activity. Another possibility is that amyloid accumulation in non-demented elderly actually is higher in the medial temporal lobes than in most other parts of the brain. This was suggested by a recent study of network properties of amyloid accumulation in healthy elderly with low amyloid levels, where it was argued that accumulation begins in the amygdala–orbitofrontal–hippocampus formation, before extending to entorhinal cortex and the posterior medial and lateral temporal pole, the medial parietal cortex and the medial and lateral prefrontal cortex (Sepulcre et al., 2013). Recent results from AIBL also indicated substantial longitudinal change in amyloid deposition in the temporal lobe in healthy elderly, even in those with initially little or no detectable amyloid burden (Villain et al., 2012). Still, amyloid deposition as detected by PIB PET generally seems to be most intense in posterior medial parietal cortex and medial frontal cortex.

Finally, although intellectual activity has been shown to have positive effects on brain biomarkers in some studies, there are discrepant findings, and a firm relationship between intellectual activity and amyloid deposition has not yet been established. Results from the large EClipSE (Epidemiological Clinicopathological Studies in Europe) initiative did not provide evidence in favor of any relationship between education and amyloid deposition. A relationship was found between education, reduced dementia risk and larger brain weight at death, but with no correlation between education and neurodegenerative or vascular pathologies (Members et al., 2010). Interestingly, higher education levels reduced dementia risk largely independent of severity of pathology, by mitigating the impact of pathology on the clinical expression of

dementia before death, consistent with the cognitive reserve hypothesis. A recent publication from the Mayo Clinic Study of Aging did not find any relationship between amyloid deposition, glucose metabolism or hippocampus volume and lifetime or current intellectual activities (Vemuri et al., 2012). The relationships may be complex and affected by several variables. One study found that the relationship between cognitive function and AD histopathological markers was modified by the participants feeling of purpose in life (Boyle et al., 2012).

Another issue is that there is growing agreement that A β accumulation in sporadic AD is more related to decreased clearance than increased production (DeMattos et al., 2004; Mawuenyega et al., 2010; Castellano et al., 2011), in contrast to increased A β production seen in familial AD (Scheuner et al., 1996). Thus, it may not be increased A β production that is the most relevant factor for amyloid deposition, but rather the ability of the brain to keep the A β peptides soluble (Robakis, 2010). In sum, we still lack convincing evidence for a relationship between amyloid deposition and intellectual activity in humans.

To summarize, regions with relatively high mental activity are parts of the heteromodal cortex, supporting complex cognition, with a complex architecture and high functional connectivity, with some being late maturing in development and high-expanding in evolution. These areas also show higher levels of atrophy in aging (see Fig. 9). Interestingly, prefrontal cortex is high-expanding in evolution and shows high levels of atrophy in normal aging, but is not among the most vulnerable regions in AD. Degeneration in these different age-vulnerable regions likely has a multitude of causes that may affect deposition of amyloid, through increased production or through processes causing reduced ability to prevent oligomerization. The chain of causation is still unknown, however. For instance, with cerebral amyloid angiopathy, amyloid deposition is found in the inner walls of blood vessels in the brain. Cerebrovascular diseases, e.g. small vessel disease with accompanying cerebral microbleeds, thus increase the levels of deposited amyloid in the brain detected by PiB PET (Park et al., 2013). However, the cause of atrophy and degeneration could be directly related to the vascular problems, not amyloid deposition per se. The causal direction between vascular factors and amyloid accumulation is not yet established, and both are related to APOE ϵ 4 (Song et al., 2004). Vascular disease risk factors, such as hypertension are also risk factors for AD and have been shown to affect brain structure (Raz et al., 2005; Salat et al., 2012; Jacobs et al., 2013) and amyloid accumulation (Rodrigue et al., 2013) in non-demented elderly and AD patients (Grimmer et al., 2012). Atrophy in these areas in normal aging may be caused by amyloid-independent events, while a gradual increase in deposition of amyloid may still be seen over time in the same areas. This may or may not be the same mechanisms observed in AD. It has been argued that the important task is not to decide whether the age-changes observed in these areas are disease-related, but to attempt to understand the fundamental mechanisms by which they occur (Jagust and Mormino, 2011). This will have implications for a number of diseases and conditions associated with aging.

Since increased amyloid deposition is observed in AD, related to brain atrophy in both patients and healthy elderly, and has a preference for deposition in regions supporting complex cognitive functions like episodic memory (Buckner, 2012), one could expect amyloid levels to be related to cognitive function in older adults. Testing the relationship between amyloid levels and cognitive function has been a very active area of research, but the results so far have been conflicting, with some studies finding modest relationships while others do not. In a recent meta-analysis, Hedden et al. analyzed amyloid-cognition relationship across 34 independent studies of almost 3500 subjects (Hedden et al., 2013).

Episodic memory function was related to amyloid burden, with a correlation of .12, while executive function and global cognition correlated 0.08 and 0.09. The authors concluded that amyloid levels have small but nontrivial associations with cognitive function in older adults.

An important consideration in the interpretation of the in vivo biomarkers of amyloid is that they may not be sensitive to the most relevant form of amyloid. Both CSF levels of $A\beta_{1-42}$ and amyloid imaging with PIB PET reflect fibrillar $A\beta$ levels and amyloid plaque load in the brain (Blennow et al., 2010). Soluble oligomeric forms of $A\beta$ may be more toxic than fibrillar $A\beta$ (McLean et al., 1999), but are very difficult to measure reliably in CSF (Rosen et al., 2013). Consequently, to be amyloid negative based on in vivo PET imaging or CSF does not guarantee that there are not malign amyloid-processing ongoing in the brain. It has been suggested that there actually could be several interacting pools of amyloid, with a large insoluble pool that is derived from a constantly turning over smaller soluble pools (McLean et al., 1999).

6. Integration, summary and conclusion

As argued above, we believe that a key element in understanding the borders between normal aging and the earliest stages of AD may be related to neuroplasticity; cognitive decline results when the brain is not able to compensate for accumulation of insults. The major genetic risk factor for AD, apolipoprotein $\epsilon 4$ (APOE), is related to neural plasticity. Higher levels of cognitive reserve, as indicated by education level and socio-economic status, are protective of AD diagnosis (Stern, 2012), as individuals with greater reserve can maintain cognitive function in the face of higher levels of brain amyloid (Members et al., 2010). The weak correlations between amyloid level and cognitive function suggest that other mechanisms, such as functional compensation, impact cognitive ability. The view that cognitive function can be maintained in aging by compensatory cognitive processes, and that decline is seen when a person is no longer able to compensate for reduced workings of primary brain structures and circuits, has many supporters in contemporary research (Cabeza et al., 2002; Park and Reuter-Lorenz, 2009; Grady, 2012). According to such a view, the magnitude of brain insults that can be accommodated without cognitive decline and progression to AD varies significantly between individuals, and a key to understanding why may be individual differences in neural and cognitive plasticity. Thus, it is important to understand the mechanisms that cause some older adults to develop AD and some to maintain cognitive function in spite of accumulated brain amyloid. To reach this aim, we need knowledge about individual differences in the relationship between the major neurocognitive events in AD. For instance, it is likely that the relationship between brain atrophy and amyloid load will vary substantially between different groups of participants (Fjell et al., 2010a), and this variation is likely not random. A key to improved diagnosis and prognosis, as well as to a better understanding of the disease mechanisms themselves, is to systematically map individual differences in the relationships between the major events in early AD. For instance, one approach to close in on the borders between healthy aging and AD is to study the characteristics of participants who show

There is a dearth of studies focusing directly on the relationship between brain plasticity and aging- and AD-related biomarkers, such as intervention studies with systematic mapping of structural and functional brain characteristics in combination with amyloid imaging. Such studies would have the potential to elucidate the relationship between cognitive function, brain integrity and mechanisms of cognitive and neural plasticity on the borders between normal aging and the earliest phases of AD.

Several overarching principles of age-vulnerability have been proposed, including the “last in, first out” hypothesis. These principles only partially fit existing evidence, as for instance the medial temporal lobes are vulnerable to age-changes but relatively low-expanding in evolution. So far, the best overlap seems to be between cortical age-vulnerability maps and the DMN. As shown in the present review, cortical age-related reductions are often found in AD-prone areas, even in those individuals at low risk of AD. We argue that these structural changes are an inevitable part of the normal aging process, and not necessarily related to neurodegenerative disease. It is likely that the vulnerability of these areas to normal age-changes contributes to their vulnerability to pathological processes involved in AD. The background for this idea is that age affects the brain independently of AD, and there is a growing number of proponents of the view that it is important to understand age-related changes to better understand AD: Herrup has developed a patho-physiological model for AD starting with age as the prime risk factor (Herrup, 2010), Jagust suggested that brain systems may be vulnerable to aging in “in the absence of even subtle disease”, and that research should thus focus on fundamental causes of neural system vulnerability rather than solely on AD-specific degeneration (Jagust, 2013), and Khachaturian proposed that we should focus less on single etiological factors and rather study the full spectrum of pathogenesis in favor of a system-level approach (Khachaturian, 2011). One idea may be that regions characterized by a high degree of plasticity are vulnerable to normal aging, which renders them even more susceptible to AD-pathology. With the *normalcy-pathology homology* as starting point, studying the brain regions that are vulnerable to both aging and AD may allow us to disentangle disease-specific mechanisms from normal age-related events.

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- (a) Normal cognition and structural brain integrity as measured by MRI despite elevated levels of AD-biomarkers such as brain amyloid
- (b) Reduced cognition and brain integrity in AD-vulnerable regions with normal amyloid biomarkers
- (c) High levels of brain maintenance as measured by structural and functional neuroimaging (Nyberg et al., 2012)

support ADNI clinical sites in Canada. Private sector contributions are facilitated by the Foundation for the National Institutes of Health (www.fnih.org). The grantee organization is the Northern California Institute for Research and Education, and the study is coordinated by the Alzheimer's Disease Cooperative Study at the University of California, San Diego. ADNI data are disseminated by the Laboratory for Neuro Imaging at the University of Southern California.

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