

Review

Does the Interplay Between Aging and Neuroinflammation Modulate Alzheimer's Disease Clinical Phenotypes? A Clinico-Pathological Perspective

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Abstract.

Alzheimer's disease (AD) is a chronic neurodegenerative disorder and is the most common cause of dementia worldwide. Cumulative data suggests that neuroinflammation plays a prominent and early role in AD, and there is compelling data from different research groups of age-associated dysregulation of the neuroimmune system. From the clinical point of view, despite clinical resemblance and neuropathological findings, there are important differences between the group of patients with sporadic early-onset (<65 years old) and late-onset AD (>65 years old). Thus, it seems important to understand the age-dependent relationship between neuroinflammation and the underlying biology of AD in order to identify potential explanations for clinical heterogeneity, interpret biomarkers, and promote the best treatment to different clinical AD phenotypes. The study of the delicate balance between pro-inflammatory or anti-inflammatory sides of immune players in the different ages of onset of AD would be important to understand treatment efficacy in clinical trials and eventually, not only direct treatment to early disease stages, but also the possibility of establishing different treatment approaches depending on the age of the patient. In this review, we would like to summarize what is currently known about the interplay between "normal" age associated inflammatory changes and AD pathological mechanisms, and also the potential differences between early-onset and late-onset AD taking into account the age-related neuroimmune background at disease onset.

Keywords: Aging, Alzheimer's disease, inflammation, microglia, phenotype

INTRODUCTION

Alzheimer's disease (AD) is a chronic neurodegenerative disorder and is the most common

cause of dementia worldwide. The two major neuropathological hallmarks of the disease are senile plaques, which are mainly composed of extracellular deposits of amyloid- β ($A\beta$) and neurofibrillary tangles, which consist of intracellular aggregates of aberrantly phosphorylated tau protein. This is accompanied by neuronal and synaptic loss, dendritic and axonal changes, and inflammatory reaction

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37 lesions [1, 2]. Cumulative data suggests that neuroinflammation plays a prominent and early role in AD [3–8]. Microglia cells are the predominant resident immune cells in the central nervous system (CNS) [9]. Recently, some studies highlighted the biological process of age-related changes associated with microglial cells [10–12] and suggest that microglial senescence can be directly associated to neurofibrillary degeneration [13]. From the clinical point of view, despite clinical resemblance and neuropathological findings, there are important differences between the group of patients with sporadic early-onset (<65 years old, EOAD) and late-onset AD (>65 years old, LOAD). Thus, it seems important to understand the age-dependent relationship between neuroinflammation and the underlying biology of AD in order to identify potential explanations for clinical heterogeneity, interpret biomarkers, and promote the best treatment to different clinical AD phenotypes.

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56 In this article, we will discuss the current knowledge regarding the interplay between “normal” age associated inflammatory changes and AD pathological mechanisms. In addition, we will discuss the potential differences between EOAD and LOAD taking into account the age-related neuroimmune background at disease onset. We will give particular emphasis to microglia due to their predominant role in the immunological process within the CNS.

66 BRAIN IMMUNE SYSTEM

67 Microglia are the resident immune cells of the CNS and considered the tissue-resident macrophages. These cells were first described by Nissl in 1899, who distinguished microglia from other neural cells based on the shape and their nuclei [14]. Microglia cells arise from myeloid precursors and constitute an autonomous population distinct from the peripheral circulating mononuclear phagocytes [15]. These cells account for up to 16% of total cell CNS population and this is dependent on the brain region [9]. There is limited replication and turnover of microglia, suggesting that microglia are a very long-lived and stable cell population [9, 12]. Microglia can provide several macrophage-related activities that provide an innate immune response as the first and main form of active immune defense in the brain [9]. The term microglial activation encloses the process where microglia change shape, molecular signature, and cellular physiology in order to

86 respond to injury or disease [16]. Resting microglia are characterized by a small cell body, highly ramified processes with weak expression of associated cell surface marker antigens [17]. In contrast, activated microglia display shortened and extensively branched processes and hypertrophy of cell body [18]. The definition of resting microglia does not mean a passive spectator in the healthy adult CNS. *In vivo* two-photon microscopy imaging studies showed that microglia survey the brain parenchyma by constantly extending and retracting their processes, and react rapidly to brain injury or insult, and are more properly termed “surveillant” [19–21]. The functions of microglia in the normal healthy brain beyond immune surveillance are unclear, but recently more sophisticated functions were described such as participating actively in the maintenance and plasticity of neuronal circuits and contributing to the protection and remodeling of synapses [22, 23].

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105 Microglial activation states have been classically described as activated (M1) or alternatively activated (M2) [24]. The M1 phenotype is characterized by production of proinflammatory cytokines, such as IL-1 β , tumor necrosis factor alpha (TNF α), and IFN- γ , whereas in the M2 phenotype microglia secrete anti-inflammatory cytokines, such as IL-4, IL-10, and transforming growth factor- β , which downregulate inflammation and promote tissue remodeling/repair and angiogenesis [25]. However, this categorizing system relays on peripheral macrophages studies, which do not recapitulate all microglial functions and is likely an oversimplification [21].

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118 The second type of neuroimmune cells is the perivascular macrophages [26]. They seem to be derived from circulating macrophages, and are able to perform all the known functions of peripheral macrophages; they undergo complete turnover approximately every 3 months [27, 28]. Finally, the circulating blood monocyte can enter the CNS, but it is not clear how often it happens under non-inflammatory conditions. In conditions of disrupted blood-brain barrier, and when properly stimulated, they can differentiate into microglia-like cells or perivascular macrophages morphologically and phenotypically [26].

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137 Astrocytes are the most abundant glial cells in the CNS and their function is critical for the support of neuronal homeostasis. The term astrogliosis describes a wide range of both molecular and functional changes in astrocytes aimed to neuroprotection and repair of injured neural tissue [29, 30]. Recently it has been shown that that reactive astrogliosis and glial

scar formation play essential roles in regulating CNS inflammation [29]. Reactive astrocytes in response to different kinds of insult can produce molecules with either pro- or anti-inflammatory potential. Additionally, reactive astrocytes can exert both pro- and anti-inflammatory effects on microglia [31, 32].

NEUROINFLAMMATION IN BRAIN AGING

There is clinical and experimental evidence that neuroinflammation in the aged brain is characterized by a shift toward a pro-inflammatory state [9, 33]. *In vivo* imaging studies using ^{11}C -R-*PK1195* PET ligand, which is upregulated in activated microglia cells, showed an increase in the specific binding with age in several cortical and subcortical structures, indicating that activated microglia gradually appear in the aging human brain [34]. In parallel, age senescent alterations can contribute to a dysfunctional microglia [12, 35, 36]. In the next paragraphs, we will address these apparent competitive perspectives of age-related neuroinflammation.

Inflammation in the brain is defined by upregulated astrocyte and microglial cell reactivity in association with increased levels of circulating cytokines such as $\text{TNF}\alpha$, $\text{IL-1}\beta$, and $\text{IFN-}\gamma$ [37–39]. With aging, microglia phenotype shifts progressively toward the activated form, together with enhanced sensitivity to inflammatory stimuli (priming phenomena) [9, 40]. In normal human brain aging, microglia is characterized by upregulation of glial activation markers such as $\text{IL-1}\alpha$ [41] and major histocompatibility complex II (MHC II) [42]. MHC II is important because it is conserved across species and is interpreted to indicate microglial priming [9]. There is compelling evidence from different research groups and aging models, that following different types of challenge (bacteria, virus, stress, surgical intervention), aged animals exhibited a clear and exaggerated neuroinflammatory response, when compared to young adult animals [33, 43–46]. These studies provided evidence that during lifespan, episodes of systemic inflammation and cytokine stimulation can “instruct” microglia and increase their reactivity [23, 33]. Interestingly, some of these sensitized neuroinflammatory responses are specific to the hippocampal formation, which is important for memory function [33]. Microglia from the aged CNS could be described as hyper-vigilant to disturbances in central homeostasis with less capability of shifting among functional states.

Proteins expressed in CNS microenvironment, which are known to inhibit microglia activation or pro-inflammatory immune responses, were implicated in the mechanism how microglia becomes chronically sensitized during normal aging [47]. In fact, some lines of research describe various proteins that activate anti-inflammatory signals following ligand and receptor interactions [48], particularly CD200 [49–51] and fractalkine (CX3CL1) [51–53]; interestingly, both are preferentially expressed in neurons. These proteins inhibit microglia through their cognate receptor, which is expressed predominantly in myelomonocytic cell types. During aging, the expression of levels of these ligands decreases concurrently with increases in microglial activation status. More recently, another line of research suggests that significant and prolonged elevation in hippocampal corticosterone (the endogenous glucocorticoid in rodents) leads to microglial priming [51]. However, the simplistic view that aging CNS shifts microglial polarization from alternative M2 state to the classical, proinflammatory state, should be interpreted cautiously because many studies found that both M1 markers and M2 markers are increased in aged mice [12]. For example, active microglia from aged mice actually had higher levels of IL-10 production (an anti-inflammatory cytokine) than those of adult mice and lower expression of $\text{TGF}\beta$ (an inflammatory cytokine) [54]. In this case, the maintenance of inflammatory response could be attributed to an impaired response to IL-10 in the aged brain [9]. Furthermore, primed microglia phenomena have been described mainly in mouse models [9, 55], and less in human brain research [56]. More recently, research studies showed that the cerebrospinal fluid (CSF) levels of YKL-40 (a microglial marker) increase in normal aging [57–59].

Together with this perspective that microglia becomes primed and more reactive with age, others showed that microglia becomes senescent and less reactive with age [10, 11, 13]. In the healthy young CNS, microglia have a typical ramified morphology and are distributed throughout the neural parenchyma in a “space-filling” manner [60]. Due to the prolonged lifespan of CNS microglia, they are more susceptible to accumulate aging-related changes [61], such as in the distribution, morphology, and behavior [12, 60] (Table 1). Many microglial cells in the aged brain show dystrophic features indicative of age-related alterations. This dystrophic microglia have de-ramification or decrease arborization of their processes, loss of finely branched cytoplasmic process,

Table 1

Summary of principal changes associated to microglial aging (adapted from Wong [60] and Wyss-Coray [6])

Changes in aged microglia**Changes in microglial distribution**

Replicative senescence (reduced mitotic activity in response to CNS injuries)

Decreases in regularity in distribution

Changes in morphology

Decrease in individual microglial ramification (dendritic arbor area, branching, and total process length)

Appearance of morphological changes suggestive of increase activation state (shortened and extensively branched processes and hypertrophy of cell body)

Appearance of dystrophic microglia (deramified, fragmented, or tortuous processes, cytoplasmic beading/spheroid formation)

Changes in microglial dynamic behavior and function

Decrease in the motility and migration process

Changes in intercellular signaling and marker expression (MHC II, CD11b)

Impaired phagocytosis

Impaired proteostasis

cytoplasmic beading/spheroid formation, and shortened and twisted cytoplasmic processes, and in some instances there is partial or complete cytoplasmic fragmentation [38]. The meaning of these morphological changes or why they happen is still to be understood.

Age-related changes were also described in astrocytes, particularly emphasizing that aged astrocytes show characteristics of the senescence-associated secretory phenotype, which involves increased secretion of inflammatory components [62].

In summary, aged microglia are primed with exaggerated and prolonged responses to inflammatory stimuli and also display dysfunctional dystrophic age associated features. Yet, it is still to be determined if microglia activation is the cause of neurodegeneration or a secondary reactive (beneficial) process; or if the neurodegeneration is actually secondary to microglia senescence and associated loss of microglial protection.

NEUROINFLAMMATION IN ALZHEIMER'S DISEASE

After two decades of the amyloid cascade hypotheses proposed by Hardy and Higgins [63], multiple lines of research still support the A β aggregation as the critical step that initiates AD pathology. However, despite required, it seems that A β aggregation is not sufficient for the development of the neuropathological and clinical syndrome of AD [64]. Several research studies report links between AD and genes regulating immunity as well as the expression of immune factors in blood, CSF, and brain tissue [8, 65–68]. There is compelling data that neuroinflammation in AD is not a passive mechanism

activated by senile plaques and neurofibrillary tangles, but instead contributes, as much or even more, to pathogenesis as do plaques and tangles [65, 68, 69]. Epidemiological studies indicate that systemic markers of the innate immunity are risk factors of LOAD [70–73] and more recently, inflammation in AD gained strong support from genome-wide association studies that identified genes involved in inflammation that are associated with increased risk of developing AD [74], including TREM2 [75, 76] and CD33 [77, 78]. Prospective cohorts' studies suggested that elevations in inflammatory mediators may be present years before clinical disease onset [70, 79, 80]. However, other longitudinal studies did not report associations between inflammation and AD risk [81, 82]. Furthermore, non-steroidal anti-inflammatory drug (NSAID) epidemiology and clinical trials showed mostly negative results, playing against the importance of inflammation in AD pathogenesis. However, these disappointing results are no surprising taking into account that normal physiological cytokine regulation of glia activation and microglial phenotypes are highly dependent of the context and the disease stage [65]. More recently, studies have consistently found an increase in CSF YKL-40 levels in AD. They also found a correlation between CSF YKL-40 levels with markers of neurodegeneration, such as tau, and with at-risk ϵ 4 carriers during mid middle age [57–59].

Neuropathological studies have shown the presence of a broad variety of inflammation-related proteins (complement factors, acute-phase proteins, proinflammatory cytokines) and clusters of activated microglia around amyloid plaques (Fig. 1) in AD subjects and also AD mice models [8], and these findings have been implicated in the neurodegeneration process [4, 83]. Neuropathological studies also showed

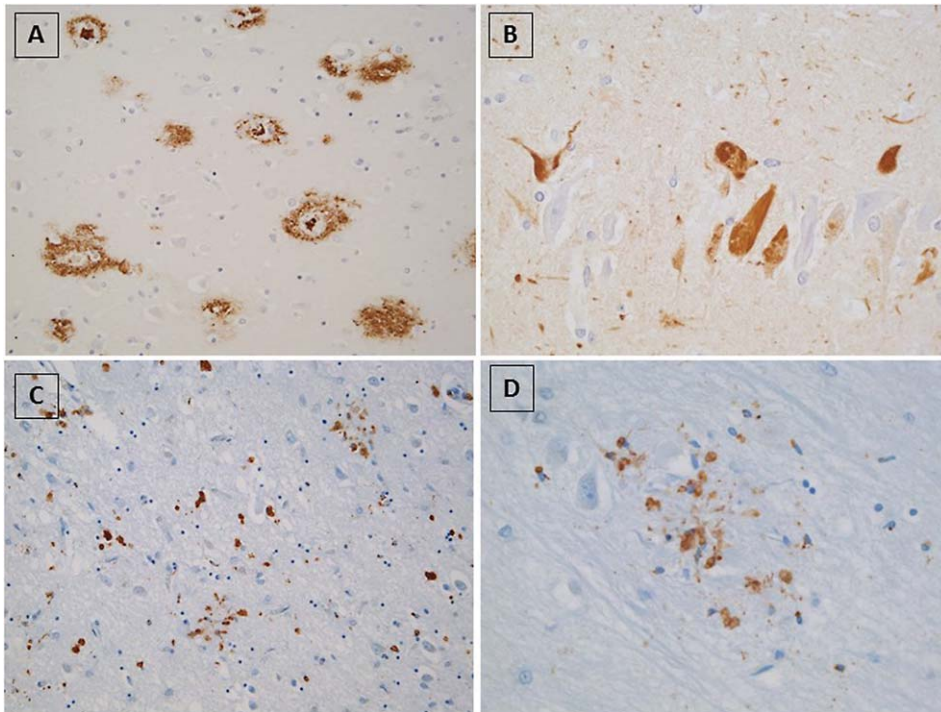


Fig. 1. Alzheimer's disease neuropathology. A) Senile plaques and globose diffuse deposits demonstrated with anti-A β antibody (M 0804, Dako). B) Neurofibrillary tangles demonstrated by phosphorylated tau protein immunohistochemistry (PHF-Tau; AT8, Thermo Scientific). C) Diffuse distribution of activated microglia in the cortex with clustering within and around amyloid plaques. D) Higher magnification of amyloid plaque with activated microglial in the CA4 region of the hippocampus (C and D: CD68 immunohistochemistry; PGM1 clone, Dako).

310 that the neuroinflammatory response in the neocortex is present in the early stages of AD pathology and precedes the late stage, tau-related pathology [84].
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 312 Furthermore, microglial activation has been shown
 313 to progress with the clinical stage of dementia, with
 314 neuropathological stage of disease severity, and with
 315 stage of progression of A β plaques [67, 85, 86].
 316 *In vivo* imaging studies, using ¹¹C-R-PK1195 PET
 317 ligand, showed that activated microglia accumulate
 318 near the amyloid plaque pathology, and that activated
 319 microglia burden correlates with cognitive decline
 320 [87].
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322 The pathological accumulation of A β is considered
 323 the key factor that drives neuroinflammation
 324 responses in AD [65]. The chronic deposition of
 325 A β stimulates the persistent activation of microglial
 326 cells in AD [88]. Microglia undergoes a progres-
 327 sive switch from a neuroprotective M2 status to a
 328 classically activated phenotype M1, characterized
 329 by production of proinflammatory cytokines [89].
 330 The persistent microglia activation and consequently
 331 microglia-derived cytokine overexpression, caused
 332 by continuous formation of A β and positive feedback
 333 loops between inflammation and amyloid- β protein

334 precursor processing, can increase A β production
 335 and decrease A β clearance, ultimately causing neu-
 336 ronal damage [65, 86, 89]. In addition, ongoing
 337 exposure to A β , chemokines, cytokines, and other
 338 inflammatory mediators can be responsible for the
 339 functional impairment of microglial cells seen at
 340 plaque sites [11, 90] and thus impede the protec-
 341 tive role of microglia in A β clearance [91]. Recently,
 342 Kim et al. [92] showed that soluble A β oligomers
 343 impair synaptic plasticity and cause synaptic loss in
 344 mouse AD models and brains of AD patients bind-
 345 ing to the murine PirB (paired immunoglobulin-like
 346 receptor B) and its human ortholog LILRB2 (leu-
 347 cyte immunoglobulin-like receptor B2) receptors,
 348 respectively. The PirB receptor was first described
 349 exclusively in the immune system but is now know
 350 to be expressed by neurons.
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352 Microglia can have different roles and effects
 353 depending on the particular disease stage and which
 354 brain region is affected in each model [65]. AD
 355 mouse models studies showed that in younger ages,
 356 together with the appearance of the first A β plaques,
 357 the microglia is activated toward the alternative
 358 state and at older ages, together with the increased
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358 accumulation of extracellular oligomeric A β , there
359 is a widespread microglial activation toward the clas-
360 sic phenotype [93]. Recently, Sudduth et al. described
361 that in the early-stage AD brains there is an apparent
362 polarization toward either M1 or M2 brain inflam-
363 matory states [94]. The M2 polarized group had
364 great number of neuritic plaques, eventually reflect-
365 ing disease progression. The heterogeneity found in
366 the early stage AD can influence the response to
367 therapeutic agents that act on immune system and
368 inflammation [94]. The neuropathological study of
369 AD patients that had undergone active A β vaccina-
370 tion as part of the AN1792 trial showed significantly
371 reduced levels of A β and reduction of aggregated
372 tau in neural processes (not in neurofibrillary tan-
373 gles), and, although there was no difference on total
374 microglial load, there were reduced levels of a range
375 of activated microglial species when compared to
376 patients who died from AD without treatment [95,
377 96]. These findings suggest that downregulation of
378 microglial activation through A β immunotherapy
379 possibly reduces the inflammatory component of the
380 neurodegeneration of AD [95]. However, a different
381 line of research supports that aging-related microglial
382 degeneration and loss of microglial neuroprotection
383 rather than microglial activation contributes to the
384 onset of sporadic AD [11]. A role for peripheral-
385 derived macrophage cells in AD pathophysiology
386 have recently come under attention [97]. There is
387 extensive evidence that blood-derived monocytes can
388 phagocytose A β [98] and that these cells can be
389 recruited to the AD brain, albeit in low numbers [99].

390 Reactive astrocytes tend to accumulate around fib-
391 rillar amyloid plaques [100]. Similar to microglia,
392 astrocytes release cytokines and other potentially
393 cytotoxic molecules after exposure to A β thus aggra-
394 vating the neuroinflammatory response [65]. Glial
395 cell activation can be an early event in AD process,
396 even preceding A β deposition. Recently, Rodriguez-
397 Vieitez and colleagues [101], using a PET tracer
398 for astrocytes (^{11}C -deuterium-L-deprenyl), showed
399 prominent initially high and then declining astrocy-
400 tosis in autosomal dominant AD carriers, contrasting
401 with the increasing A β plaque load during disease
402 progression. This study provided *in vivo* evidence that
403 astrocyte activation is a very early feature of, at least
404 familial, AD pathology [101]. Other lines of research
405 have linked senescent astrocytes to the increase risk
406 of sporadic AD [102].

407 In summary, the role of microglia remains con-
408 troversial in AD pathogenesis and the question
409 of whether activated microglia aids in promoting

410 clearance of toxic A β species or if their proinflamma-
411 tory profile exacerbates pathology is currently a topic
412 of debate [103]. Although there is broad evidence of
413 a large immune response component in AD, the issue
414 of which activation phenotype affects the onset or
415 progression of the disease and, consequently, which
416 should be the therapeutic target remains to be deter-
417 mined [104]. Furthermore, the questions regarding
418 the role of excessive astrogliosis or astrocyte senes-
419 cent loss of function in AD pathogenesis remains to
420 be solved [100].

421 EARLY AND LATE-ONSET 422 ALZHEIMER'S DISEASE

423 Regardless of the clinical resemblance and
424 neuropathological findings, important differences
425 between EOAD and LOAD patients have been
426 reported. The separation of EOAD from LOAD at 65
427 years old is a conventional cutoff point indicative of
428 a sociological partition in terms of employment and
429 retirement, but there is no specific biological signifi-
430 cance to use this specific age, and there is a range of
431 disease features that do not respect this arbitrary divi-
432 sion [105, 106]. However, this arbitrary cutoff point
433 has been used widely by different research groups
434 and allowed the uniform study of AD patients with
435 different ages of onset.

436 *Clinical presentation*

437 Whether age of onset defines the clinical presen-
438 tation of AD has been a matter of debate for decades
439 and reports on this issue are often contradictory.
440 Nonetheless, some differences have been consis-
441 tently recognized. Earlier onset is associated with a
442 worse prognosis and a faster progression. Younger-
443 onset patients have comparatively worst outcomes
444 in the Mini-Mental State Examination at baseline,
445 show a steeper cognitive and functional decline, and
446 seem to have higher mortality risks when compared
447 to older-onset patients [107–109]. In addition, dif-
448 ferent patterns of cognitive deficits are apparent;
449 non-amnesic presentations are more often found in
450 early-onset disease, described in 33–64% of EOAD
451 compared to 6–12.5% of LOAD patients [105, 110].

452 Earlier neuropsychological studies have shown
453 that younger patients have more language disability
454 when compared to older-onset patients [111–113].
455 The risk of having language difficulties detected by
456 caregivers has also been shown to nearly duplicate
457 for each 10-year decrease in AD patients' age [114].

458 Other groups have recognized a greater impairment in
459 measures of attention, praxis, and visuo-contruction
460 tasks in EOAD [115–117]. On the other hand, LOAD
461 patients seem to consistently have preferential mem-
462 ory involvement [118–120]. To explore the relation
463 between this clinical duality and pathologic features,
464 Murray et al. [121] divided a cohort of AD patients
465 into “hippocampal sparing”, “limbic predominant”,
466 and “typical AD” according to neurofibrillary pathol-
467 ogy distribution. They have shown that a younger age
468 of onset (mean 63 years) was associated with greater
469 neurofibrillary tangle burden in cortical association
470 areas and that older age (mean 76 years) was more
471 often associated with limbic predominant pathology.
472 The hippocampal sparing group had greater preva-
473 lence of atypical presentations and a faster cognitive
474 decline, similar to what has been described in EOAD.

475 Seizures and extrapyramidal features seem to be
476 more frequent in EOAD [111, 122]. There are con-
477 tradictory reports in other symptoms in both groups.
478 For example, there are reports of higher anxiety lev-
479 els in EOAD [123], while others have shown greater
480 neuropsychiatric and behavioral symptoms in LOAD,
481 including anxiety, depression, agitation, hallucina-
482 tions, and delusions [124, 125].

483 Limited research has been reported into sex differ-
484 ences in brain aging, particularly neuroinflammation
485 process. However, gender effect is an interesting
486 issue due to the differences of the neuroendocrine
487 milieu and its possible relation to inflammation cas-
488 cades (particularly steroid-related pathways). The
489 dynamic change in hormonal status in women during
490 the menopause transition may promote a dysregula-
491 tion of cellular processes involved in hypothalamic-
492 pituitary-adrenal axis and thus have potential implica-
493 tions on stress-mediated neurotoxicity [126]. It is also
494 important to recognize the importance of immunolog-
495 ical differences in males and females within the CNS
496 at different development time points and their possi-
497 ble relevance for the susceptibility in the development
498 of neurological conditions later in life [127]. A recent
499 work in mice by Mangold and colleagues showed a
500 greater induction of MHC class I components and
501 receptors with aging with this finding being greater
502 in females than in males [128]. However, despite the
503 prevalence of AD being greater in women, the pre-
504 vailing view has been that this difference is due to
505 the fact that women live longer than men on average,
506 and older age is the greatest risk factor for AD. Many
507 studies of incidence of AD have found no significant
508 difference between men and women in the proportion
who develop AD at any given age [129].

Biomarkers

509
510 Magnetic resonance imaging (MRI) studies show
511 that younger-onset patients have greater general-
512 ized neocortical atrophy than LOAD subjects when
513 compared to healthy controls [118, 130]. This is
514 in accordance with glucose metabolism studies,
515 which demonstrate a premature decline in glucose
516 metabolism and a more severe and widespread
517 hypometabolism in EOAD [131]. Regarding regional
518 differences, older patients tend to have a more circum-
519 scribed involvement, with preferential reduction in
520 the hippocampus and related structures, including the
521 amygdala [132] and retrosplenial and temporopari-
522 etal junction volumes [130], while younger patients
523 tend to have a greater temporoparietal and parietooc-
524 cipital grey matter atrophy [115, 120]. White matter
525 atrophy mimics this pattern [133]. Moreover, both per-
526 fusion and glucose metabolism studies have shown
527 a predilection for temporo-parietal-occipital associ-
528 ation areas in EOAD versus medial temporal cortex
529 susceptibility in LOAD [119, 134]. Interestingly,
530 another study has shown no significant difference in
531 total or regional amyloid burden, measured by Pitts-
532 burg compound-B PET, despite showing decreased
533 glucose metabolism in bilateral temporoparietal and
534 occipital cortex in EOAD. This finding suggests that
535 both early A β and increased susceptibility to pathol-
536 ogy in younger onset patients might be responsible
537 for cortical dysfunction in EOAD [135]. The greater
538 involvement of hippocampal-related structures in
539 LOAD is also apparent in functional connectivity
540 studies that have shown that older patients have
541 decreased activation of the anteromedial temporal net-
542 work, correlating with poorer performance in memory
543 tasks; EOAD was associated with less activation of the
544 dorsolateral prefrontal network, manifested by worse
545 performance on executive function tasks [118].

546 CSF pathophysiological markers for AD include
547 decrease levels of A β_{1-42} and increase levels of
548 total tau and hyperphosphorylated tau. The use of
549 these biomarkers combined is associated with sig-
550 nificant sensitivity and specificity in the diagnosis
551 of AD [136]. There is some evidence that EOAD
552 patients have a greater reduction of A β_{1-42} (and
553 corresponding greater elevation of tau) than LOAD
554 patients when compared to young and old controls,
555 respectively, although no differences emerge in the
556 direct comparison between EOAD and LOAD [137].
557 Others have reported lower levels of A β_{1-42} in EOAD
558 [138] or no differences [120, 139]. A study compar-
559 ing CSF biomarkers along different EOAD subtypes,

560 including amnesic, logopenic progressive aphasia
561 and posterior cortical atrophy found no differences in
562 the A β levels, but showed that posterior cortical atro-
563 phy had lower levels of total tau and phosphorylated
564 tau [140].

565 *Genetics*

566 Amyloid precursor protein, presenilin 1, and pre-
567 senilin 2 mutations can cause autosomal dominant
568 AD, and although they may be present in up 71%
569 of familial cases, they account for only about 1–5%
570 of all AD patients. These patients typically have an
571 early or very early-onset disease (<45 years) [136,
572 141, 142]. A well-recognized genetic risk factor for
573 AD is the APOE ϵ 4 allele. It is usually associated
574 with greater hippocampal atrophy and a poorer per-
575 formance in memory based tasks [121, 142] and it
576 decreases the age of onset by up to 2.45 years for
577 each copy of the allele [142, 143]. Conversely, non-
578 APOE ϵ 4 patients tend to have greater structural and
579 clinical involvement of non-hippocampal, neocortical
580 areas [121]. ApoE ϵ 4 allele carriers among AD
581 patients are most frequently found in the 60–69-year-
582 old range [144], therefore including both older EOAD
583 patients and younger LOAD patients. The ApoE ϵ 2
584 allele is less frequently found in AD patients than in
585 normal controls and there seems to be no difference
586 in its prevalence between EOAD and LOAD [144].
587 Genome wide association studies have identified sev-
588 eral other risk genes for LOAD. The association
589 between nine of them (PICALM, CLU, CR1, BIN1,
590 CD2AP, EPHA1, MS4A4A, CD33, and ABCA7) has
591 been shown to account for 1.1% of age of onset vari-
592 ation, versus 3.9% of variation provided by ApoE.
593 The most significant association was found for the
594 CR1, BIN1, and PICALM genes [143]. Another can-
595 didate gene that may have an impact on age of onset
596 is DCHS2, a gene expressed in the cerebral cortex
597 [145]. Yet, and surprisingly, these genetic variants do
598 not seem to bring significant value for the distinction
599 between EOAD and LOAD, as they simply seem to
600 anticipate pathology.

601 **INTERPLAY BETWEEN BRAIN AGING, 602 NEUROINFLAMMATION, AND AD 603 PHENOTYPES**

604 AD prevalence is strongly associated with increas-
605 ing age and aging changes in microglia have been
606 hypothesized to play a prominent role in disease
607 pathogenesis [60]. Recently, the consistent pattern

608 of increases in YKL-40 level with aging supports
609 the concept that neuroinflammation is a process that
610 occurs normally with aging [57–59]. The additional
611 finding of a stronger association with at-risk ϵ 4
612 carriers during mid middle age suggests that this age-
613 related process may be further exacerbated in the
614 presence of insults including amyloid deposition and
615 neuronal injury [59]. There are important clinical dif-
616 ferences between sporadic EOAD and LOAD. Taking
617 into account the data regarding the importance of
618 neuroinflammation in the pathogenesis of AD, par-
619 ticularly the role of microglia, and the differences
620 of the neuroimmunological milieu of the aged brain,
621 it is conceivable that the neuroinflammation associ-
622 ated to the AD can, at least in the beginning, differ
623 between these two groups and contribute for the clinical
624 differences. Not many studies have addressed this
625 issue.

626 Hoozemans et al. [146] compared the presence
627 of microglia and astrocytes, in clinically and patho-
628 logically confirmed AD and non-demented control
629 cases in relation to age. In their study they suggested
630 that the association between neuroinflammation and
631 AD is much stronger in relatively young patients as
632 compared to the older patients (age at death <80 ver-
633 sus >80 years old). Microglial activation increases
634 with the neuropathological stage and disease severity
635 [67, 85]. A key issue would be to know if inflamma-
636 tion differs between these two groups (EOAD versus
637 LOAD) at different pathological and clinically AD
638 stages.

639 Another remarkable finding is that, in contrast
640 to AD, activated microglia is not found in the
641 similar-appearing A β diffuse deposits of the brains of
642 neurologically normal elderly individuals [147]. One
643 of the possibilities is that for those unusual elderly
644 individuals with only diffuse A β deposits there is an
645 inherent difference in the responsiveness of microglia
646 [86]. Interestingly, plaque-associated microglia were
647 not seen in diffuse plaque-only young Down's syn-
648 drome brain [148]. This subgroup of cases was from
649 very young patients (between 12 and 29 years old),
650 possible supporting the notion that A β inflammatory
651 response can also differ in the very young. More
652 recently, a study showed that in Down's syndrome
653 patients with AD pathology (>40 years old), there is
654 a distinct neuroinflammatory phenotype compared to
655 sporadic AD due to microglia bias toward an M2b
656 phenotype [149]. Clinicopathological studies from
657 brain donation programs showed that the presence of
658 moderate and severe AD type pathology changes is
659 more associated to dementia in younger old persons
660

660 than in older old persons [150]. These findings sug-
 661 gest that additional factors are involved in the clinical
 662 expression of dementia in the oldest old, such as vari-
 663 able tolerance to neuropathological lesions [150]. We
 664 speculate that different neuroinflammation apparatus
 665 in this age can partial explain this discrepancy.

666 The study of inflammatory cytokines in CSF as
 667 biomarkers of AD has shown very different and con-
 668 tradictory results between different research groups
 669 [89]. The analysis of different neuroinflammation-
 670 related proteins in the blood, including several
 671 interleukines (IL-1 α , IL-1 β , IL-6, IL-10), α 2-
 672 macroglobulin, brain-derived neurotrophic factor
 673 (BDNF), complement factor H, and heat shock pro-
 674 tein 90 (Hsp90) has not shown significant differences
 675 between EOAD and LOAD, but studies are scarce and
 676 with small samples [151, 152]. TNF α levels have
 677 been shown to be both higher and lower in EOAD
 678 [152, 153].

679 Some of the risk loci in modifying age of disease
 680 onset identified in genome wide association stud-
 681 ies have recognized roles in the immune system,
 682 including phagocytosis and immune cell traffick-
 683 ing [154]. Both CLU and CR1 encode for proteins
 684 that regulate complement activation; EPHA1, mostly
 685 expressed in leukocytes, is involved in T cell regula-
 686 tion; ABCA7 is highly expressed in the hippocampal
 687 neurons and in microglia and is involved in A β pro-
 688 cessing; and CD33, overexpressed in AD patient's
 689 microglia, encodes for an endocytic receptor that
 690 takes part in cell-cell interactions and in immune cell
 691 regulation [154, 155]. TREM2, another loci associ-
 692 ated to increase risk for AD identified, is involved
 693 in immune response [75]. There are studies that
 694 found a significantly earlier symptom of onset in
 695 patients with TREM2 variants [156], but others found
 696 only an association to shortened disease duration
 697 and not to age of onset [76]. A β cerebral amyloid
 698 angiopathy (CAA) and particularly A β related angio-
 699 itis (ABRA), is other AD related clinical feature that
 700 bridges AD, inflammation and age. CAA describes a
 701 group of biochemically and genetically diverse dis-
 702 orders, which have in common the deposition of
 703 amyloid in media and adventitia of cortical and lep-
 704 tomeningeal vessels [157]. Sporadic CAA and AD
 705 have overlapping biology with shared risk factors
 706 [158]. A β vascular deposition affects about 30% of
 707 the otherwise normal elderly and over 90% of those
 708 with AD, in whom CAA tends also to be more severe
 709 [157, 159]. ABRA is characterized by a vasculitic
 710 transmural, often granulomatous, inflammatory infil-
 711 trates affecting leptomeningeal and cortical vessels

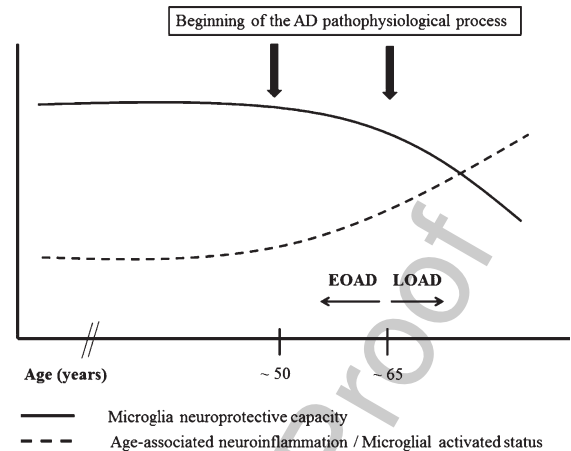


Fig. 2. Diagram illustrating age associated microglia dynamics and temporal Alzheimer's disease onset. Arrows exemplify two time points for the beginning of AD biomarkers [A β accumulation (CSF/PET), sequentially followed by tau-mediated neuronal injury (CSF)] at the preclinical stage.

712 that have abundant A β deposition within the vessel
 713 walls [159, 160]. The recent finding of autoantibod-
 714 ies against A β ₁₋₄₀ and A β ₁₋₄₂ forms of amyloid in
 715 the CSF of two patients with ABRA and inflamma-
 716 tion associated to CAA [161, 162], together with the
 717 description of meningoencephalitis caused by active
 718 or passive immunotherapeutic approaches to reduce
 719 A β burden in AD [163], suggests that an immune
 720 response directed against A β may represent a com-
 721 mon disease mechanism shared by ABRA and in
 722 complications of therapy for AD [160]. The mean
 723 age of presentation of ABRA is lower than that
 724 of sporadic non-inflammatory A β -related CAA (66
 725 versus 76 years, respectively) [159, 160]. These find-
 726 ings support a role for the interactions between age,
 727 and inflammation in AD related pathophysiology and
 728 clinical expression.

729 In summary, the pathophysiological mechanisms
 730 underlying the clinical differences between EOAD
 731 and LOAD are still not well known, but the dif-
 732 ferences of neuroinflammation characteristics with
 733 aging can help to partially explain it (Fig. 2).

734 CONCLUSION

735 Understanding both sides of microglial and astro-
 736 cytosis inflammation process at functional and
 737 molecular level will be necessary for the development
 738 of treatment strategies for AD and aging [12].

739 Additionally, the study of this delicate balance in
 740 the different ages of onset of AD would be important

to understand treatment efficacy in clinical trials and eventually, not only direct treatment to early disease stages, but also the possibility of establishing different treatment approaches in light of the age of the patient. The boost on AD diagnostic biomarkers will increase diagnostic certainty in life for the diagnosis of dementia with AD pathology. This refinement will allow the increased recognition of the more often atypical clinical presentations in EOAD and thus increase the knowledge (epidemiology, clinical progression, biomarkers studies, neuroinflammation associated process, etc.) for a possible better understanding of this complex disorder.

DISCLOSURE STATEMENT

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