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Mild Cognitive Impairment: Pathology and mechanisms

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Abstract

Mild cognitive impairment (MCI) is rapidly becoming one of the most common clinical manifestations affecting the elderly. The pathologic and molecular substrate of people diagnosed with MCI is not well established. Since MCI is a human specific disorder and neither the clinical nor the neuropathological course appears to follow a direct linear path, it is imperative to characterize neuropathology changes in the brains of people who came to autopsy with a well-characterized clinical diagnosis of MCI. Herein, we discuss findings derived from clinical pathologic studies of autopsy cases with various subtypes of MCI antemortem. The heterogeneity of clinical MCI imparts significant challenges to any review of this subject. The pathologic substrate of MCI is equally complex and must take into account not only conventional plaque and tangle pathology but also a wide range of cellular biochemical and molecular deficits many of which relate to cognitive decline as well as compensatory responses to the progressive disease process. The multifaceted nature of the neuronal disconnection syndrome associated with MCI, suggests that there is no single event, which precipitates this prodromal stage of AD. In fact, it can be argued that neuronal degeneration initiated at different levels of the central nervous system drive cognitive decline as a final common pathway at this stage of the dementing disease process.

Keywords

Alzheimer's disease; amyloid; cholinergic; dementia; MCI; neurofibrillary tangles; neuropathology; molecular; neurotrophins; synapses

Mild Cognitive Impairment

The development of the concept of mild cognitive impairment (MCI) was derived from memory clinics, which attracted milder cases of dementia than Alzheimer's disease (AD) and epidemiological studies of elderly populations followed over time, in which subjects were evaluated annually for cognitive status. Memory clinic researchers noticed that many of the earlier, milder cases of cognitive loss in the elderly did not meet the "two cognitive domains impaired" criteria required for an NINDS/ADRDA diagnosis of AD established by McKhann and coworkers [98]. In the 1990's such cases, were found most frequently but not always to be characterized by an amnesic disorder and the term "Mild Cognitive Impairment" or MCI, was popularized by Petersen [133] becoming the category in which these 'single domain impaired' individuals (who did NOT meet criteria for dementia/AD) were placed. The term MCI, however, was devoid of any pathological implications and was more descriptive than specific. Thus, while it emerged from the study of AD and the implication was that "all MCI was a precursor to AD", increasing studies led to the identification of multiple types of MCI-some static (e.g., because of an injury, most commonly vascular), and some progressive, most commonly due to AD but possibly to vascular or other neurodegenerative disease processes, such as FrontoTemporal Dementia (FTD) [129].

While amnesic MCI was the most common form of MCI, especially in Memory Disorders Clinics, it was clear that MCI with other affected single cognitive domains were a small, but significant, component of this clinical presentation. While some presented with rare neurodegenerative disorders (e.g., dysexecutive or aphasic dysfunction as the sole cognitive problem in the MCI phase of FTD), some were vascular and many progress to AD. Moreover, the nature of MCI might differ by the nature of the research entity. Thus. Memory Disorders clinics report amnesic MCI (aMCI) as the most common form of MCI. In epidemiological populations, in which subjects have vascular disease, chronic medical diseases (e.g., congestive heart failure) and require medications for chronic illness, which may affect cognition, do not have aMCI as the predominant condition [90, 91]. However, people with a clinical diagnosis of MCI comprise a heterogeneous cohort of which, those who present solely with memory deficit are classified as single domain amnesic MCI (aMCI), while those who have a deficit in memory as well as other cognitive domains are categorized as multi-domain aMCI [76, 130]. Those with aMCI are at a higher risk of developing AD [76, 130]. Those with impairment in cognitive domains other than memory are designated as either single or multi-domain non-amnesic MCI (naMCI) [131, 188].

If the pathobiology of MCI is to be determined, autopsies of subjects with well-characterized MCI of different types are a necessity. Our goal, in this review, is to extend beyond the many excellent neuropathologic reviews of MCI [64, 137, 148]. Here, in addition to discussing the gross brain features of MCI, we will concentrate on pathomechanistic signature(s) of this clinical construct including biochemical and neuroanatomical alterations, synaptodegeneration, cell loss, neurotrophic failure, cellular genetics, neuronal selective vulnerability and other factors that occur in the MCI brain. Since, to date, there are no true animal models of MCI, we emphasize relevant research derived from clinical pathologic investigations of the human condition, the gold standard for the field. Understanding the complete neuropathological status of individuals with MCI will be essential for appropriate therapeutic interventions and realistic expectations for slowing or stopping the clinical decline.

Gross Morphologic features of MCI

Although there have been many advances in defining the clinical definition of MCI [5, 43, 168], distinguishing brains at autopsy from people who died with a clinical diagnosis of NCI and MCI is challenging. For example, the cortical gyral and sulcal patterns appear similar between NCI (Fig. 1a) and nonamnestic MCI (naMCI) (Fig. 1b) brains. However, we have observed a widening of sulci, such as the ventral ramus of the lateral fissure as well as a blunting of the anterior tip of the temporal pole in aMCI (Fig. 1c) and mild AD (Fig. 1d) compared to the NCI or naMCI brain. These gross morphological changes are magnified and extended to other cortical regions in late stage AD (Fig. 1e). Brains harvested from those who came to autopsy with a clinical diagnosis of MCI are providing the material needed to unravel the structural and cellular pathobiologic substrate of this prodromal stage of the dementia.

Amyloid Pathology in MCI

Brain amyloid beta ($A\beta$) plaques, are a hallmark lesion of people with a clinical diagnosis of MCI. The distribution of $A\beta$ deposits changes with time and reflects the spread of $A\beta$ deposition in the diseased brain [181]. Senile plaques first appear as diffuse and “fleecey” [180] plaques throughout the neocortex and extend hierarchically into other brain regions. In a second stage, $A\beta$ plaques occur in allocortical areas (e.g. entorhinal cortex and subiculum/CA1 region). In the third phase, plaques appear in the basal ganglia, thalamus, and hypothalamus. In the fourth phase, amyloid reaches the midbrain and medulla oblongata. Finally, in a fifth stage, senile plaques appear in the pons and cerebellum. Later stages feature neuritic (Fig. 2a) and amyloid cored (Fig. 2b) plaques. Whether cored plaques evolve from dying neurons and their role in the onset of dementia remain unclear. Furthermore, whether amyloid plaque pathology defines the substrate underlying MCI is still unanswered. The distribution of amyloid deposits in MCI appears to be intermediate between the changes seen in the NCI and AD brain [93]. Many aged control and MCI brains exhibit a similar degree of $A\beta$ deposition as shown in postmortem brain tissue [137] as well as in clinical imaging studies [4], limiting the use of these lesions as a true pathologic marker for the distinction between normal aging and MCI. Early clinical pathological studies revealed that people with clinical dementia rating (CDR) scores of 0.5 (questionable dementia/MCI) displayed significant increases in the density of diffuse plaques (DPs) in the temporal cortex [105]. The density of plaques increased with increasing dementia and the proportion of plaques shifted from diffuse to neuritic. By contrast, a recent study using samples from the University of Kentucky Alzheimer’s Disease Research Center (UKADRC) revealed that the brains of people without cognitive impairment and with a Braak score of II showed no differences in the number of DPs in the neocortex or limbic medial temporal lobe (MTL) structures compared to MCI cases [94] as defined by the CDR criteria [105, 131]. Neuritic plaque (NP) counts were significantly greater in the medial frontal cortex and amygdala in MCI compared to normal controls, whereas no differences were found in the number of NPs within the hippocampal CA1 region, subiculum or entorhinal cortex (ERC). Numbers of DPs were unchanged in most cortical regions between controls and MCI, whereas NP number was significantly increased in the above limbic cortical regions when MCI and early AD were compared suggesting that NPs mark this transition. A clinical pathologic study using a mixed cohort of MCI (aMCI and mdMCI) cases from the Rush Religious Orders Study (RROS), a clinical pathologic longitudinal study of aging and dementia [15, 109] found that ERC $A\beta$ -amyloid peptide load in MCI cases appeared intermediate between controls and early AD, but the difference was not statistically significant [109] lending support to the suggestion that β -amyloid deposition is not the major pathological factor defining the transition to MCI [93]. Moreover, age-adjusted group comparisons of cortical insoluble $A\beta$ 40 and $A\beta$ 42 levels correlated with neuropathologic AD status (CERAD and

Braak staging) but were not predictive of a clinical diagnosis of MCI [46]. In the precuneus region of the medial parietal cortex [Brodmann area 7, (BA7)], soluble A β 42 concentration and [3H] PiB binding to insoluble A β aggregates were elevated in MCI cases; both measures correlated with lower Mini-Mental State Exam (MMSE) scores [71]. Although MCI and early AD are distinguishable biochemically from aged controls in terms of the A β 42/A β 40 ratio (both soluble and insoluble) [183], removal of amyloid plaques following A β vaccination therapy failed to prevent further cognitive decline in people with a clinical diagnosis of early AD [67]. Interestingly, decreased levels of A β ₁₋₄₂ and increased phospho-tau levels detected in cerebral spinal fluid may provide a biomarker for those destined to develop dementia [107, 173], but this remains to be more fully validated. Therefore, solely targeting this protein for the treatment of dementia may not be entirely feasible, and anti-amyloid therapies may need to be initiated prior to dementia onset. In light of the recent finding that A β normally functions as an antimicrobial peptide in the innate immune system in response to clinically relevant pathogenic micro-organisms, its removal may result in increased vulnerability to infection [167] and continued brain dysfunction. Perhaps the field requires a major paradigm shift to go beyond the problems of A β protein cleavage, processing and aggregation [78, 142] in MCI.

Recent studies suggest that A β monomers, dimers and higher order oligomeric forms may be an early and toxic form of amyloid [83]. In this regard, A β oligomers were found to accumulate in the frontal cortex of people with a clinical diagnosis of MCI (CDR= 0.5) and mild to moderate AD (CDR=1 or 2) compared to age matched controls (CDR=0) from the Alzheimer's Disease Research Center at the University of California, San Diego [134]. Increased A β oligomer levels correlated with the severity of cognitive impairment (MMSE and Blessed Information Memory Concentration score), AD neurodegeneration (Braak staging), and lower levels of the presynaptic vesicle protein synaptobrevin and the postsynaptic density protein PSD-95 [134]. Perhaps A β monomers and oligomers underlie amyloid toxicity, causing a disruption of synaptic function in MCI, which initiates cognitive decline.

During the past several years, genomic analysis has revealed genetic risk factors not only for the development of dementia but which affect amyloid deposition during the early phase of the disease. The seminal findings of Roses and colleagues revealed that apolipoprotein E (APOE) e4 carriers are at increased risk for AD [32]. Imaging studies of presymptomatic e4 carriers have shown reduced cerebral glucose metabolism [162], increased cerebral amyloid deposition [139] and enhanced cortical atrophy [31] as well as accelerated memory decline [26]. Although these changes are milder, they qualitatively resemble those seen in AD suggesting that they may represent a very early stage that precedes the symptomatic expression of MCI. In fact, the total amyloid load in AD cases is higher in APOE e4 carriers than non-carriers [42]. Two studies, one examined a large Finnish cohort [85] and another a population from the Sun Health Brain Donation program [27], found that APOE e4 carriers without premortem diagnosis of dementia, including those with an MCI diagnosis at death displayed higher amyloid (but not NFT) burdens than non-carriers. PiB imaging of amyloid deposition seems to be predictive of decline in MCI cases, but not that predictive in pre-MCI cases [45, 166].

PiB-PET imaging data derived from four cortical regions known to have high amyloid deposition were used as a quantitative phenotype to identify genes related to fibrillar amyloid burden in 103 Alzheimer's Disease Neuroimaging Initiative subjects [174]. Of the various genes examined only the intronic single nucleotide polymorphisms (SNP) of the DHCR24 gene was significantly associated with a lower average PiB uptake. The DHCR24 gene appears to play a neuroprotective role and has been shown to confer resistance to A β and oxidative stress induced apoptosis by inhibiting caspase-3 activation [55]. Another SNP

located in the amyloid-associated LR11 gene (*SORL1*) is also associated with an increased risk for developing AD [16, 86], although this association is controversial [102]. LR11 (also termed sorLA) is a multi-functional neuronal receptor that binds apolipoprotein E (ApoE), plays a crucial regulatory role in the processing of the amyloid precursor protein (APP) and may maintain low levels of the pathological form of the A β peptide [6, 7, 41, 123]. A quantitative immunohistochemical report found that neuronal LR11 levels in MCI were intermediate between NCI and AD in superior frontal cortex (SFC) tissue obtained from the RROS [147]. MCI subjects with low LR11 levels displayed significant cognitive impairment compared to high LR11 level cases, suggesting that the loss of LR11 is an early alteration in the cascade of events underlying the development of MCI by regulating APP processing.

Neurofibrillary Tangle Pathology in MCI

NFTs are composed of aggregates of hyperphosphorylated forms of the protein tau [17] (Fig. 2c), display a beta pleated sheet conformation (Fig. 2d) and are argyrophilic (Fig. 2a). Recent biochemical reports suggest that NFT development progresses according to a linear sequence of molecular and conformational changes in the tau molecule before and during NFT formation and maturation [18, 47]. The Braaks [21] described the pathologic topographic spread of NFTs from the MTL to the neocortex, which composed six stages according to the location of the tangle-bearing neurons and the severity of changes (transentorhinal stages I–II: clinically silent cases; limbic stages III–IV: incipient AD; neocortical stages V–VI: fully developed AD). Clinical subtypes of MCI as well as many NCI cases have variable Braak staging scores ranging from a complete absence of NFTs (Stage 0) to stages V/VI representing AD [109, 112, 114, 132, 137]. Although these observations indicate that there is no clear demarcation between some control, MCI and AD brains using the Braak staging criteria, a study suggests that Braak scores are better at identifying aMCI from naMCI [138].

A broader evaluation of the distribution of NFTs indicate that they are significantly increased in the amygdala, entorhinal cortex (ERC), subiculum and the inferior parietal cortex (IPC) in MCI compared to controls suggesting that NFTs are critical for the transition to MCI [21, 58, 94, 118]. Alterations in the phosphorylation or conformation state of tau are related to MCI [64]. Clinical pathologic investigations using tissue from the RROS have shown a significant increase in tau positive NFTs and neuropil treads in the ERC and perirhinal cortex between MCI and NCI cases [103] coincident with a significant reduction in the number and atrophy of ERC layer 2 stellate neurons [87], which display prominent NFTs [69]. An examination of brains from two large autopsy populations, the Nun Study [140, 164, 165] and the National Alzheimer's Coordinating Center (NACC) registry [13] failed to document an example of a truly end-stage NFT pathology coexisting with intact cognition [2]. These findings suggest a continuum of tau induced NFT pathology underlying the transition between normal aging, MCI and AD. A parsimonious corollary to the formation of NFTs within neurons, suggest that areas containing these lesions should exhibit significant neuronal loss in the MCI brain.

Neuronal loss in MCI

Numerous studies have examined changes in neuron numbers in MCI, focusing on structures in the MTL. Within MTL, the ERC, a major paralimbic cortical multimodal sensory relay region to the hippocampus undergoes cell loss in MCI [21, 40, 54, 69, 87]. Cases from the Washington University cohort showed no significant age related decrease in numbers of Nissl-stained ERC neurons or volume in healthy non-demented individuals [136]. Few or no differences were observed between healthy controls and what was termed “preclinical AD” or cases with normal cognition (CDR 0) despite plaques and NFTs at autopsy [136].

However, neuronal numbers were significantly decreased in the ERC and hippocampal CA1 subfield in very mild AD (CDR 0/0.5 or CDR 0.5) suggesting that neuronal atrophy and death has already occurred at a time when patients begin to manifest clinical symptoms [54]. Unbiased quantitative stereology revealed a significant loss of NeuN-immunoreactive ERC lamina II neurons and volume in MCI cases harvested from the RROS [87] in agreement with previous seminal findings [54, 136]. Furthermore, ERC atrophy correlated with impairment on MMSE and clinical tests of declarative memory [87, 106], indicating that ERC neurodegeneration impacts the progression of MCI. A likely consequence to neuronal loss is that it should lead to a decrease in the number of synapses.

Synaptodegeneration in the hippocampus in MCI

The hippocampus, an integral component of a trisynaptic memory circuit, displayed a significant decline in the plasticity related post-synaptic density protein 95 (PSD 95) in a small aMCI cohort compared to subjects without dementia [172]. Quantitative immunoblotting found that synaptophysin, a key player in membrane trafficking events which precede exocytosis, as well as synaptotagmin, a Ca²⁺ microsensor that modulates activity-dependent exocytosis [172], were relatively preserved in aMCI cases from the RROS [33]. On the other hand, drebrin, an f-actin postsynaptic binding protein [66] involved in synaptic plasticity [66], was decreased in the hippocampus of MCI cases and correlated with cognitive decline [65]. Taken together, these findings suggest that the hippocampus displays decreased synaptic plasticity in MCI. Whether synaptic protein changes in hippocampus are reflected in changes to synapse number has great relevance to the level of cognitive decline seen in MCI.

Studies aimed at defining synapse loss in the hippocampus have been performed using tissue harvested at the UKADRC [158] or RROS [15] and processed for unbiased stereological sampling coupled with transmission electron microscopy. These investigations revealed no significant differences in synapse number in the outer molecular layer of the dentate gyrus between cognitively intact subjects and aMCI, whereas a milder AD group showed a significant loss of synapses compared to both NCI and aMCI [154]. A subsequent investigation using the same cases showed synaptic changes in the stratum radiatum region of the hippocampal CA1 subfield [153]. Unlike the outer molecular layer of the dentate gyrus, estimation of total CA1 synaptic numbers was more pronounced in aMCI. Importantly, synaptic counts strongly correlated with antemortem cognitive scores. It would be anticipated that synaptic defects are reflected in a change in the gross structure of these limbic regions.

Structural changes in the MTL in MCI

Support for the involvement of the ERC and hippocampus early in the disease process also comes from structural MRI investigations, which demonstrate atrophy of these regions in aMCI compared to cognitively healthy elderly controls [81, 170, 176]. Recent, voxel based morphometry studies demonstrated a significant decrease in the volume of the parahippocampal white matter a region that includes the perforant pathway projections to the hippocampus in aMCI compared to healthy control subjects [171]. Furthermore, the rate of MTL atrophy seems to be related to the extent of declarative memory dysfunction [171]. These *in vivo* results are in line with postmortem tissue investigations suggesting that AD-related pathology affects the entorhinal region before the hippocampus [169, 176] and that white matter volume change reflects not only loss of afferent and efferent fibers in the region of the parahippocampal gyrus, but may also be due to partial demyelination in remaining fibers in MCI [77, 143].

Synaptodegeneration in the cortex in MCI

Although neocortical synapse loss is a prominent feature of AD [97, 151], very few studies have reported changes in the MCI neocortex. Evaluation of cortical tissue from individuals with CDR scores of 0.5–1.0 containing probable MCI and early AD cases compared to aged non-demented controls revealed a 25% loss in synaptophysin immunostaining in the frontal cortex but no change in synaptotagmin and GAP-43 [95]. Quantitative protein immunoblotting using tissue from clinically diagnosed MCI and either mild/moderate or severe AD from the RRCS revealed decreased levels of the presynaptic vesicle marker, synaptophysin selectively in the superior temporal cortex (STC) and inferior parietal cortex (IPC) in AD compared to NCI supporting findings in mild or severe AD [68, 96, 160, 179]. Synaptophysin protein levels in the STC and IPC correlated with MMSE scores, suggesting that the loss of this presynaptic protein within select neocortical areas mark a clinical progression of the disease. However, levels of the predominantly presynaptic protein alpha-synuclein (α -syn), which is involved in neuronal plasticity, was significantly reduced in the SFC only in AD compared to MCI and NCI [186]. In contrast, cortical presynaptic synaptotagmin levels remained stable from NCI to MCI to AD independent of region with no association with cognition. Taken together, these findings suggest that select differential alterations of presynaptic proteins contribute to early perturbations in synaptic homeostasis in MCI cortex.

With respect to postsynaptic cortical proteins, studies using tissue from the RRCS revealed that levels of drebrin were significantly reduced in the STC in MCI compared to controls. On the other hand, SFC drebrin levels were increased in MCI compared to NCI and AD. Drebrin is localized to postsynaptic dendritic spines at excitatory synapses [8, 66, 161], where it may regulate spine morphogenesis and postsynaptic densities [66, 175]. Since drebrin levels in the anterior cingulate, STC and visual cortex correlated with cognitive impairment, the progression of dementia may, in part, be related to a loss of neocortical excitatory postsynaptic contacts in the cortex similar to the hippocampus.

A recent ultrastructural study demonstrated a significant loss of synaptic contacts from NCI to aMCI in the inferior temporal cortex (ITC, BA 20) in samples from UKADRC and RRCS (Fig. 3) [155]. There was a strong association between the number of synapses in the ITC and the individual performance on both the MMSE and tests of verbal fluency [155]. Immediately adjacent tissue showed a significant loss of different pre and post-synaptic proteins including synaptophysin and PSD-95 in aMCI, which correlated with MMSE score, supporting a role for the ITC in the early stages of disease progression [155]. Despite MTL synaptodegeneration, many other cellular processes are likely to play important roles in MCI-associated neuronal dysfunction.

Cholinergic basal forebrain systems in MCI

The MTL receives input from various neurotransmitter systems including cholinergic innervation from the basal forebrain (Fig. 4a, b) [100]. Biochemical [37] and histologic [187] investigations revealed a reduction in cortical cholinergic activity and loss of cholinergic basal forebrain (CBF) neurons (Fig. 4c), which led to the cholinergic hypothesis of AD [11]. Sustaining the function of the impaired central cholinergic system is the primary mechanism of action for currently effective FDA-approved drugs for AD [110]. This emphasizes the importance of defining the status of the cholinergic system in subjects with MCI as well as cognitively normal elderly people who are at risk for developing AD. However, whether or not cholinergic system degeneration is an early or late feature of AD is still an open question. In MCI, CBF neurons also contain early and late tau conformational markers of NFTs [99, 185]. Furthermore, cell expression profiling studies indicated that the

ratios of 3-repeat tau (3Rtau)/4-repeat tau (4Rtau) are significantly decreased due to a reduction in expression of 3Rtau rather than an increase in 4Rtau tau in CBF neurons in both MCI and AD [52]. Such a shift in ratio may lead to an increased propensity of tau to form toxic aggregates; this is suggested by *in vitro* studies indicating that 4Rtau is more prone to aggregate than is 3Rtau [82]. Hence, the CBF degenerative process(es) appear to be initiated early in the disease. However, quantitative stereological studies revealed that the number of CBF nucleus basalis (NB) perikarya expressing either choline acetyltransferase (ChAT, the synthetic enzyme for acetylcholine) or the vesicular acetylcholine transporter was stable in MCI in cases from the RROS (Fig. 4d) suggesting that frank NB cell loss is a later stage event despite the early appearance of NFTs [50]. Furthermore, George and colleagues [48] found that the volume of the substantia innominata (SI), which contains the cholinergic neurons of the NB, was significantly reduced in participants from the Rush Alzheimer's Disease Center clinic and RROS diagnosed with AD, as reported previously [61–63, 150]; however, the NCI and aMCI groups did not differ from each other. The lack of change in SI volume in individuals with MCI is consistent with postmortem tissue investigations showing the preservation of SI cholinergic neuron number early in the disease [50]. However, others found significant MRI-derived atrophy of the SI in patients with MCI [116]. These apparent differences may be due to the region of the SI examined or methodologies used by different investigators. Whether deficits in forebrain cholinergic projection neurons are mimicked in their cortical and hippocampal projection sites is an active area of investigation.

Biochemical analyses of ChAT [39] and acetyltransferase (AChE) [38] activities in neocortical regions from cases from the Jewish Home and Hospital in New York or the RROS, found no changes in MCI while significant deficits were detected in severely demented AD cases. By contrast, ChAT activity is increased in the MCI superior frontal [39] but reduced in the primary visual cortex, an area relatively spared in late stage [49] and mild-moderate AD cases [72]. Interestingly, the visual cortex displays significant loss of cholinergic fibers even in the absence of frank synaptic pathology [12]. The precuneus is another cortical area found to display a significant cholinergic deficit in AD cases from the RROS [71]. Collectively these studies demonstrate that early stages of AD dementia are associated with cholinergic deficits, however, these changes are detectable only in selected brain regions.

Of great interest are findings showing elevated ChAT activity in the hippocampus (Fig. 6a) and superior frontal cortex (SFC) in MCI subjects from the RROS [39, 73]. It appears that select hippocampal and cortical cholinergic projection systems are capable of compensatory responses during prodromal dementia. Increased hippocampal and SFC ChAT activity in MCI may be important in promoting biochemical stability, or compensating for neurodegenerative defects, which may delay the transition of these subjects to AD. Interestingly, hippocampal ChAT activity was increased selectively in MCI cases scored as a Braak III/IV stage, suggesting that upregulation of ChAT occurs, at least in part, due to the disconnection of glutamatergic ERC input to the hippocampus early in the disease process (Fig. 6b). The source of these compensatory responses may be the cholinergic neurons, which preserve their cholinergic phenotype in MCI [50] (Fig. 4d) resulting in a biochemical upregulation of ChAT protein or enzyme activity, which accommodates for reduced regional cholinergic fibers and axon varicosities [71]. The biochemical cortical cholinergic upregulation in MCI [39] is not paralleled by a structural plasticity of cortical cholinergic fibers [70], suggesting increased SFC ChAT activity precedes the loss of cholinergic axonal input to this brain region during the clinical progression of AD. Further evidence of cortical plasticity is reflected by a paradoxical increase in the number of glutamatergic synapses in the MCI midfrontal gyrus versus NCI in cases from the RROS [14].

The relationship between changes in cholinergic activity and regions of the brain, which display high levels of amyloid burden defined by PiB-PET binding in MCI has not been examined extensively. Although both frontal and precuneus cortex develop amyloid pathology early in the disease process [84, 182], ChAT activity is only upregulated in the former [39] and stable in the latter cortical region in MCI [71]. This disparity may reflect the differential ability of these two regions to launch a compensatory response. While the progression of amyloid pathology does not appear to affect regional cholinergic enzyme activity in the frontal cortex [39], there is a strong association between reduced ChAT activity levels and increased A β load in the precuneus as well as an association with impaired cognitive performance on the MMSE [71]. An analysis of frontal and parietal cortex demonstrated a reduction in ChAT activity and the coupling of M1 muscarinic acetylcholine receptor (mAChR) to G-proteins in non-demented elderly subjects with plaques compared to those without plaques using tissue from the Banner Institute Brain Bank, Phoenix, AZ, suggesting that cortical cholinergic dysregulation is initiated at the preclinical stage in parallel with increased amyloid plaque deposition [135]. Since anticholinesterase drugs are widely prescribed for early AD, it is imperative to determine the extent to which acetylcholinesterase (AChE), the cholinergic degrading enzyme, is affected in MCI. PET imaging using [11C] MP4A or [11C]PMP, has been used to measure alterations in acetylcholinesterase (AChE) activity *in vivo*. These studies revealed a reduction in the hippocampus and cortex with the most pronounced reduction in temporal polar region in MCI (CDR 0.5) [59] and early AD [141] (Fig. 5). A parsimonious event that most likely occurs in the cholinergic system would be impairments to cholinergic receptors in MCI.

Acetylcholine receptors (nAChR) in MCI

Currently there is no clear consensus on whether presynaptic or post-synaptic (or both) receptor components of cholinergic neurotransmission are defective in MCI. The nicotinic acetylcholine receptor (nAChR) system is altered in AD [122], however, these receptors have not been studied extensively in MCI. Using an *in vitro* binding assay with [H-3] methyllycaconitine (MLA, a potent ligand for α -bungarotoxin sensitive nAChR), α 7 receptor binding was examined in the superior frontal cortex of RROS cases and found to be unchanged in MCI and mild/moderate AD compared to NCI subjects [74]. Interestingly, this study indicated an association between increased α 7 nAChR binding and cortical neuritic plaques. This finding is in agreement with a previous observation that α -bungarotoxin binding correlates positively with A β plaque density in AD brains [128], and may reflect up regulation of α 7 nAChR mRNA in cortical-projecting cholinergic neurons in early AD [34]. These results support a role for α 7 nAChR in the development of cortical A β pathology [117, 177] and potential functional interaction between α 7nAChR and A β [89]. The role of α 4 β 2 nAChRs in MCI and AD is somewhat more controversial. A postmortem study of nAChRs, using samples from the Sun Health Research Institute Brain Donation Program, found that [3H](+/-) epibatidine binding was stable in the midfrontal cortex suggesting the preservation of cortical α 4 β 2 nAChRs [146]. Likewise, a PET study using 2-[(18)F]fluoro-3-(2(S)-azetidylmethoxy) pyridine (2-[(18)F]FA-85380) on 15 mild AD (CDR<2) and healthy control subjects found no evidence of α 4 β 2 reduction in early AD [44]. In contrast, another (18)FA-85380 PET study reported that MCI (CDR=0.5) and mild-moderate AD patients had significant reductions in cortical and hippocampal α 4 β 2 nAChR binding; interestingly, only those MCI who converted later to AD had deficient binding [80]. In this regard, α 4 β 2 nAChRs were also examined using 123I-5-IA-85380 SPECT confirming reductions of these receptors in temporal lobe [178] or their preservation in the MCI brain [104]. Therefore, it appears that reductions in the predominant brain nAChR subtypes are altered early during disease progression and likely impact cholinergic function.

It is generally accepted that M1 mAChR densities are unchanged in AD; however, details about their role in disease progression are limited [184]. [(35)S]GTPgammaS binding/immunocapture assay revealed that M1 mAChR function in MCI frontal (BA 10) cortex samples from the RROS were unchanged, but increased significantly in AD relative to controls [124]. Another study measured postsynaptic M1 mAChR coupling, using displacement of ³H-pirenzepine binding by oxotremorine-M in the presence and absence of GppNHp, and ChAT activity in frontal and parietal cortices from cognitively normal (preclinical aged subjects with either, the presence, or absence of amyloid plaques at postmortem examination) and clinical AD cases [135]. Both cholinergic markers were reduced in preclinical cases with amyloid burden, and this reduction was accelerated further in clinical AD. Therefore during AD pathogenesis, cholinergic deficits may emerge early, together with the development of amyloid plaques and prior to the onset of clinical symptoms. The mechanisms driving degeneration of the cholinergic system in MCI have recently received a great deal of attention.

Neurotrophic abnormalities in MCI

A central concept underlying the survival of CBF neurons is their dependence upon the neurotrophic substance nerve growth factor (NGF), which is synthesized from its precursor (proNGF) molecule and interacts with the cognate NGF receptor (TrkA) principally for pro-survival and the pan-neurotrophin receptor (p75^{NTR}) principally for pro-cell death functions [110, 111]. Examination of cases from the RROS revealed that the number of NB perikarya expressing either the TrkA or the pan-neurotrophin receptor p75^{NTR} was reduced ~50% in MCI and mild AD compared to NCI (Fig. 4d) and this deficit correlated with impaired performance on the MMSE and tests of working memory and attention [113, 114]. Many cholinergic NB neurons appear to undergo a phenotypic silencing of NGF receptor expression (Fig. 4e–g) in the absence of frank neuronal loss [149] during the early stages of cognitive decline. Moreover, TrkA gene expression, but not p75^{NTR}, was reduced in NB neurons in MCI and AD (Fig. 7a,c) [53]. These alterations may signify an early deficit in neurotrophic support during the progression to MCI. In MCI, NB neurons also contain early and late tau markers of NFTs [99, 185].

Although mature NGF (mNGF) levels are preserved in neocortex CBF neuron projections sites in MCI [112], proNGF is elevated relative to non-demented controls in MCI [127] and TrkA levels increase in mild AD, whereas p75^{NTR} levels are stable in tissue harvested from the RROS [35]. Recent studies suggest that the putative pro-apoptotic effect(s) of p75^{NTR}-mediated proNGF signaling is dependent on interactions between p75^{NTR} and the neurotensin receptor sortilin, a Vps10p domain trafficking protein that acts as a cell surface co-receptor with p75^{NTR} to mediate proNGF-induced cell death [92, 144]. Evaluation of cortical sortilin protein levels revealed no changes between NCI and MCI cases from the RROS [115]. These alterations in cholinergic activity may favor a shift in the pro-apoptotic proNGF:p75^{NTR}:sortilin trimeric ratio during the earliest stages of AD (Fig. 8). Increased cortical proNGF levels in MCI also suggest potentially pathogenic alterations in metabolic pathways mediating the maturation and degradation of mNGF [23]. In MCI cases from the RROS, there was up-regulation of MMP-9 protein levels and activity, that correlated inversely with cognitive status [24] suggesting yet another defect in the NGF biosynthetic pathway in MCI (Fig. 8), which may contribute to the selective vulnerability of CBF neurons in MCI. However, neurotrophic dysfunction is not the only and not the earliest biochemical defect associated with CBF cellular dysfunction in MCI adding to the complexity of the disease process underlying neuronal selective vulnerability.

Endosomal and oxidative stress in MCI

In this regard, several studies indicate that endosomal and oxidative stress dysregulation are among the earliest pathological changes observed in cortical, hippocampal CA1 pyramidal and NB CBF neurons [28–30, 119–121, 163], precede clinical symptoms and appear prior to substantial deposition of cerebral amyloid, vascular amyloid and tau pathology in AD [30, 120]. With regard to endosomal pathways, laser capture microdissection (LCM) coupled with microarray analysis of CA1 pyramidal neurons and NB cholinergic neurons revealed an upregulation of markers of endosomal activation, including select rab GTPases in MCI and mild/moderate AD tissue from the RR0S (Fig. 7b,c) [34, 51]. In NB cholinergic neurons [51, 53], select rab GTPase expression levels were selectively increased as antemortem measures of cognition declined and occurred in parallel with a down regulation of TrkB and TrkC, the cognate receptors for brain-derived neurotrophic factor (BDNF) and NT3, respectively, both members of the NGF neurotrophin family [9]. However, unlike the NGF/proNGF complex, proBDNF and mature BDNF are both reduced in the MCI cortex [126]. Hence, increased endocytic pathway activity, driven by elevated rab GTPase expression, may result in long-term deficits in hippocampal and basal forebrain neurotrophic signaling and represent a key pathogenic mechanism underlying the onset of MCI.

With regards to oxidative stress, heme oxygenase (HO-1, an indirect marker of oxidative stress), together with biliverdin reductase-A (BVR-A, a pleiotropic enzyme that plays a pivotal role in antioxidant defense against free radicals) are elevated in MCI hippocampus and temporal neocortex samples obtained from the RR0S [156, 157] and UKADC [10]. Astroglial HO-1 immunoreactivity in the temporal neocortex, but not hippocampus, correlated with the burden of neurofibrillary pathology [157]. In addition, there was a significant reduction in protein carbonyl-derivatives of BVR-A and an up regulation of inducible nitric oxide synthase in the MCI and AD hippocampus [156, 157]. MCI individuals also show increased levels of lipid peroxidation and nucleic acid oxidation in hippocampus [79]. Redox proteomics revealed excessive oxidative modification of the oxidative proteins A-enolase, glutamine synthetase, pyruvate kinase M2 and peptidyl-prolyl cis/trans isomerase 1 in MCI hippocampus compared to control samples accrued from the UKADRC [25]. These findings indicate the functional interaction of these oxidative proteins in energy metabolism, synaptic plasticity and mitogenesis/proliferation pathways and strengthen the suggestion that oxidative stress is an early cellular event, perhaps inducing cells to compensate for the increase of intracellular oxidative events during the prodromal stages of dementia [19]. Interestingly, Smith and coworkers [88] proposed that tau phosphorylation represents a compensatory neuronal response against oxidative stress, which acts as a neuroprotective event during the early phases of cellular insult. This mechanism may explain reports indicating that tangle bearing neurons survive for at least two decades [108], suggesting that NFT are not initiators of cell death [88] but may play a role in altering nuclear transcription factors.

An intriguing report using western blot analysis of isolated cytosolic and nuclear fractions prepared from postmortem human hippocampi harvested from the UKADRC with a premortem clinical diagnosis of NCI and MCI [131], demonstrated a shift of nuclear factor of activated T-cells (NFAT) 1 and 2 to nuclear compartments at different stages of AD neuropathology and cognitive decline, whereas NFAT2 remained unchanged [1]. Changes in NFAT3 were directly correlated to soluble amyloid- β ($A\beta_{1-42}$), and oligomeric $A\beta$ levels in hippocampus. These findings add to the growing number of factors that are dysregulated in MCI and suggest that NFAT signaling may play an important role in driving $A\beta$ -mediated neurodegeneration early in the development of dementia.

Concomitant Pathogenic Factors in MCI

Several clinical pathological studies have reported that some patients with aMCI had other concomitant neuropathologic features, which may contribute to the clinical presentation of the subjects. For example, aMCI cases also display argyrophilic grain disease [20, 132], hippocampal sclerosis and vascular disease [132, 159]. Macroscopic cerebral infarcts without a pathologic diagnosis of AD, were found to be more common in naMCI (18.6%) compared to aMCI (13.3%) [159]. Certain vascular dementia subtypes, particularly those related to subcortical microvascular disease, may be preceded by MCI, which exhibit similar domains of cognitive impairment and an overlapping progressive course that may mimic AD [101]. A recent report demonstrated a small number of MCI brains that also displayed neocortical Lewy bodies; however, no person displayed a combination of AD pathology, macroscopic infarcts and neocortical Lewy bodies [101, 159]. Therefore, as with other human disorders, for example, cardiovascular disease, there may be many contributing pathologies that create a vulnerability state for MCI and neurodegeneration as well as in other neurologic disorders.

Neuropathology in Parkinson's disease with MCI

A clinical study revealed that patients with Parkinson's disease (PD) and MCI, defined by the Peterson criteria, had a higher risk of developing dementia than cognitively intact PD subjects, suggesting that MCI in PD is an early manifestation of dementia [75]. However, the pathology of MCI in PD is only now receiving attention. In a preliminary report, the majority of PD-MCI cases were found to be Braak AD stages III–IV (two amnesic MCI cases being stage IV) [3]. However, the major pathologic features were limbic and/or neocortical Lewy body and AD histopathology and possibly cerebrovascular pathology. This is an area that requires extensive clinical pathologic investigation using well-characterized patient populations.

Monoaminergic and Serotonergic Pathology in MCI

Although most reviews of MCI have concentrated on the pathology of cortical and basal limbic forebrain regions, there is mounting evidence suggesting that the brainstem harbors the earliest cellular degenerative events, even before those seen in neo and limbic cortex. Clinical pathological investigations of the norepinephrine (NE) containing locus coeruleus (LC) demonstrate NFTs occur during aging and in the earliest Braak pathological stages [22]. In MCI, LC neurons display sequential early and late tau conformational epitopes linked to NFT formation [22, 33, 57]; abnormal tau aggregates occur within proximal axons of LC projection neurons in the absence of either NFTs or neuropil threads in the transentorhinal cortex [22]. Since LC cytopathology correlates with overall cognition, noradrenergic dysfunction should be considered among the earliest cytopathologic lesions mediating the onset of cognitive decline in the aging-MCI continuum [57]. Most likely this occurs by disrupting ascending LC noradrenergic input to the thalamus, hippocampus and cortex. [22]. Similar to the LC, clinical pathologic investigations reveal the involvement of the brain stem serotonergic raphe cortical projection neurons in AD [36, 60, 125]. In the early Braak stages 0, II, and III, phospho-tau cytoskeletal changes occur in the supratrochlear subnucleus of the dorsal raphe nucleus (ST-DR) [56, 145], suggesting that the ST-DR plays a key role in the induction and spread of AD-related cytoskeletal pathology. Currently, information on the state of raphe neurons in MCI is lacking. Once these regions of the brain are fully investigated in MCI, it may be that the currently accepted stages of the cytopathology of MCI may require reclassification based on the emerging concept of a neuron-to-neuron transynaptic propagation, which maybe initiated in the brainstem and over time spreads to the telencephalon [22].

Summary and conclusions

The data presented in this review indicate that the neuropathologic substrate(s) of MCI is complex and must take into account not only senile plaque and NFT pathology but also cellular dysfunction and the initiation of neuroplastic responses. The wide range of cellular dysregulation that occurs prior to and during the prodromal MCI stage of the disease process suggests that there is no “silver bullet” at this time, which best fits the diverse pathologic, molecular and cellular constellation of events that occur in the MCI brain. Just as there are subtypes of clinical MCI, there may be multiple pathological entities that drive the onset of MCI. It will clearly require much more investigation to tease the pathology underlying these states apart. Emerging data suggest that MCI pathology is initiated via a trans-synaptic neuron-to-neuron disconnection syndrome affecting multiple levels within the central nervous system. In any case the pathologic mechanism underlying MCI most likely begins years before the onset of cognitive decline. The data presented herein suggest that it is too simplistic to attribute a single event to the precipitation of MCI. Instead, the clinical pathologic data suggest that the brain undergoes multisystem dysfunction and that in some instances the disease process triggers cellular and biochemical repair mechanisms in an attempt to slow the disease process. Understanding the molecular pathogenesis of these compensatory processes will provide novel clues about how the brain naturally responds to the signals that propagate the onset of MCI pathology, and give insight on how to potentially treat this complex, prodromal disease state with novel therapeutic approaches and cognitive remediation.

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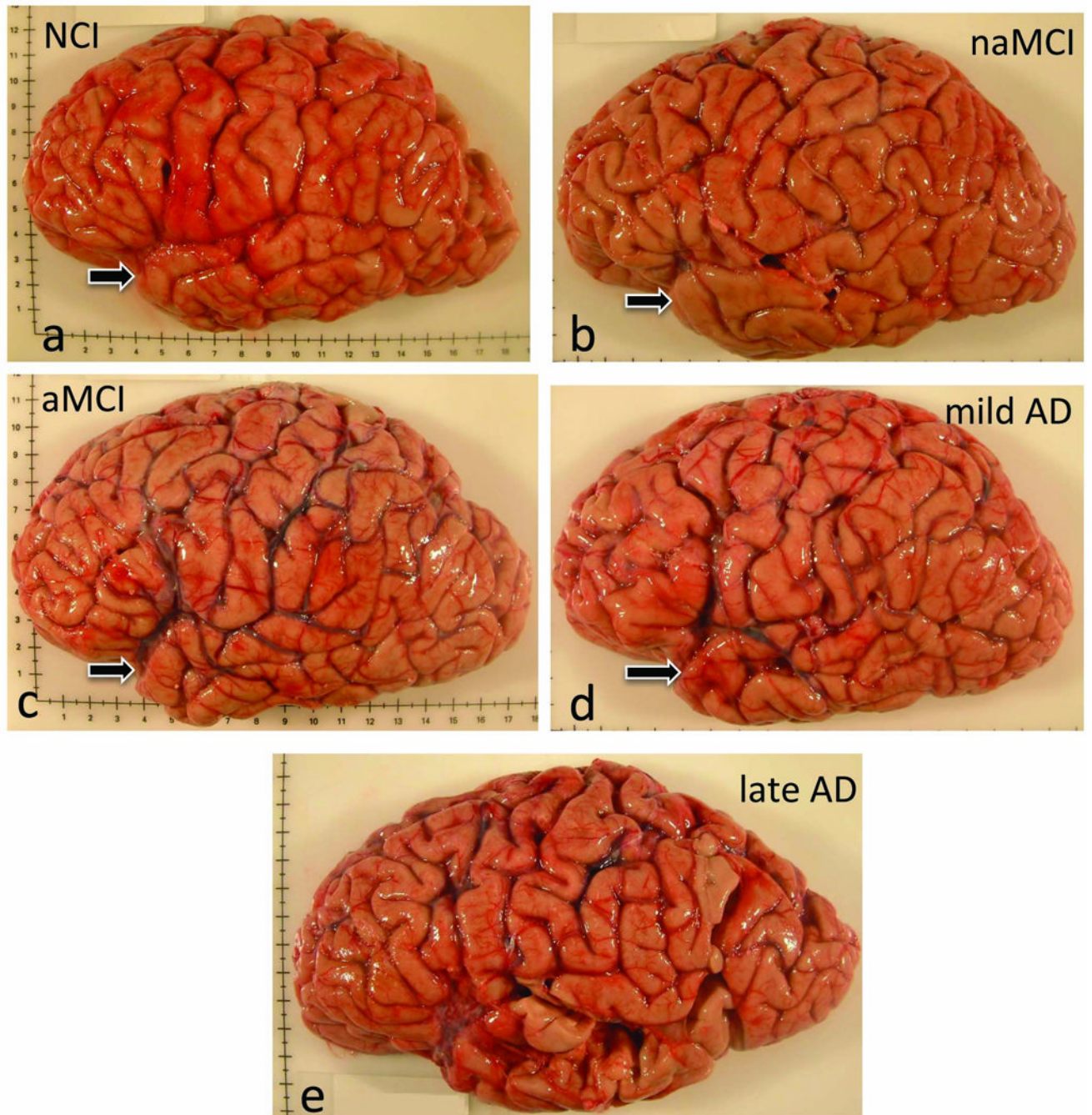


Fig. 1. Gross anatomy showing similarities and differences between the brain of a person who dies with a clinical diagnosis of (a) no cognitive impairment (NCI), (b) non amnesic mild cognitive impairment (naMCI), (c) amnesic MCI (aMCI), (d) mild and (e) severe Alzheimer's disease

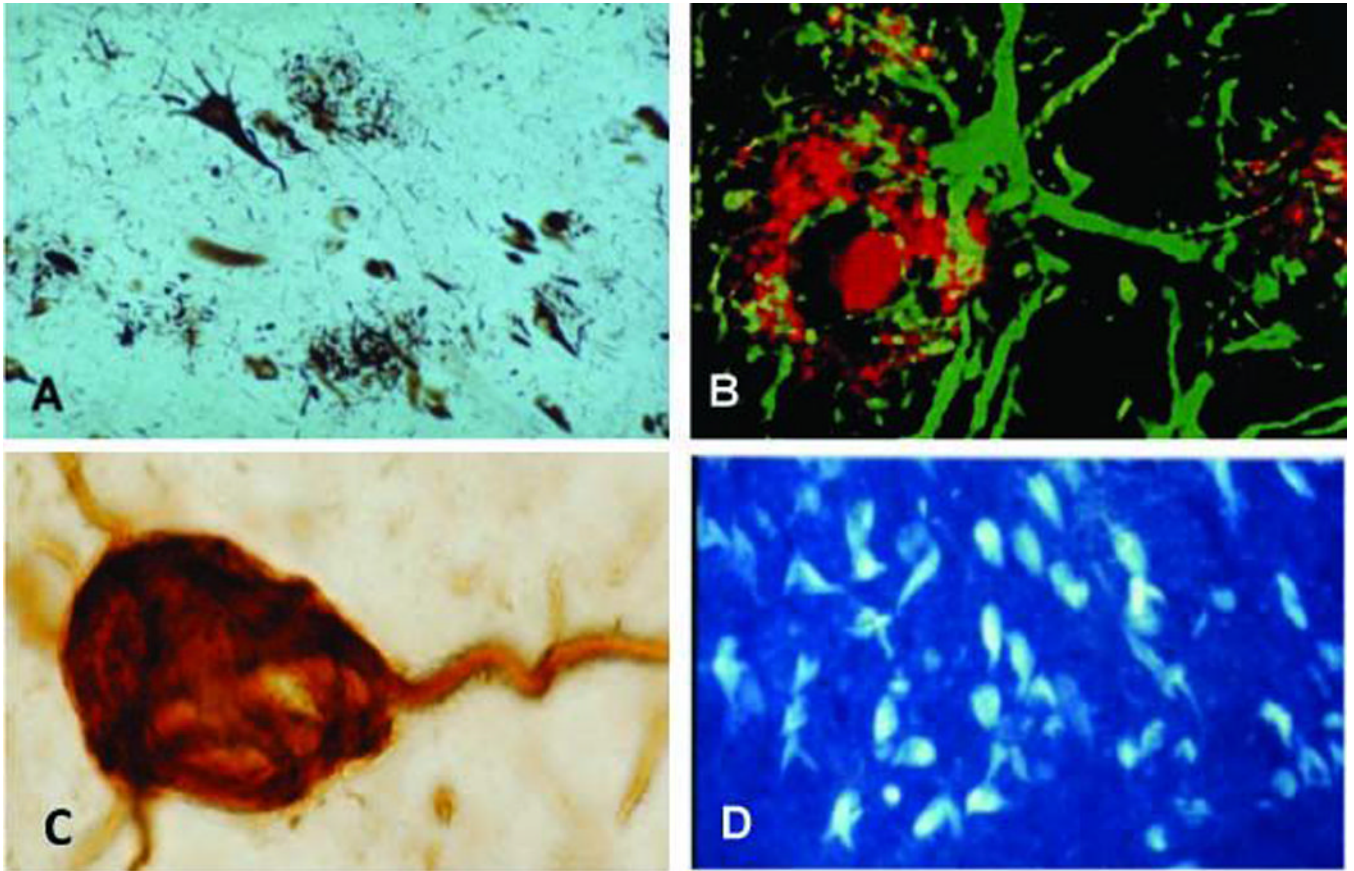


Fig. 2.
 (a) Neuritic plaques (arrows) and neurofibrillary tangle (star) visualized by the Bielschowsky silver method in AD. (b) Fluorescent image of a cored amyloid beta immunostained plaque (red) surrounded by a GFAP positive glia cell (green) in the AD frontal cortex. (c) NFT containing the tau epitope Alz-50 in the frontal cortical in the AD frontal cortex. (d) Thioflavin-S positive staining of NFTS revealing the beta fibrillar configuration within entorhinal cortex layer 2 stellate neurons in AD

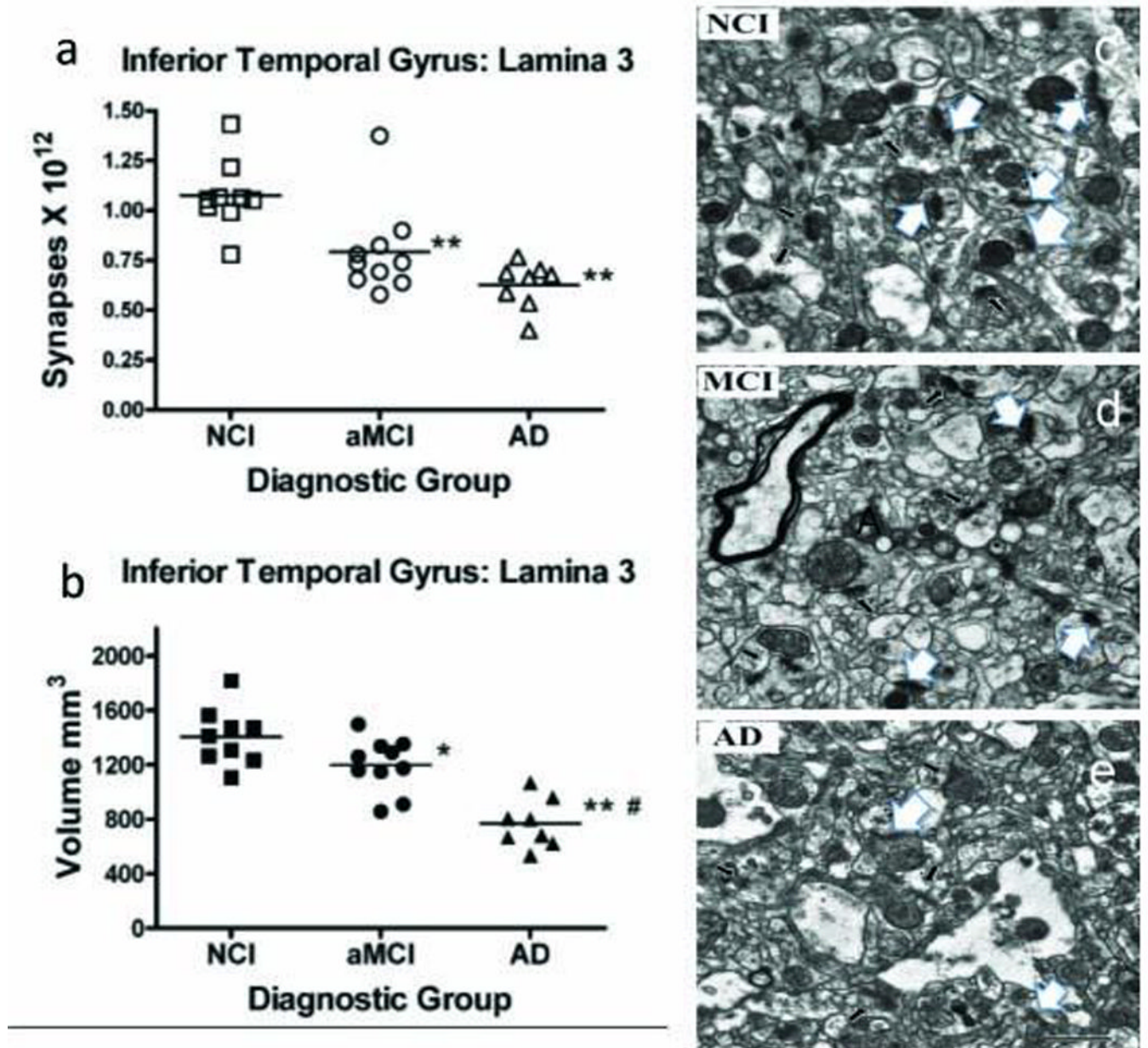
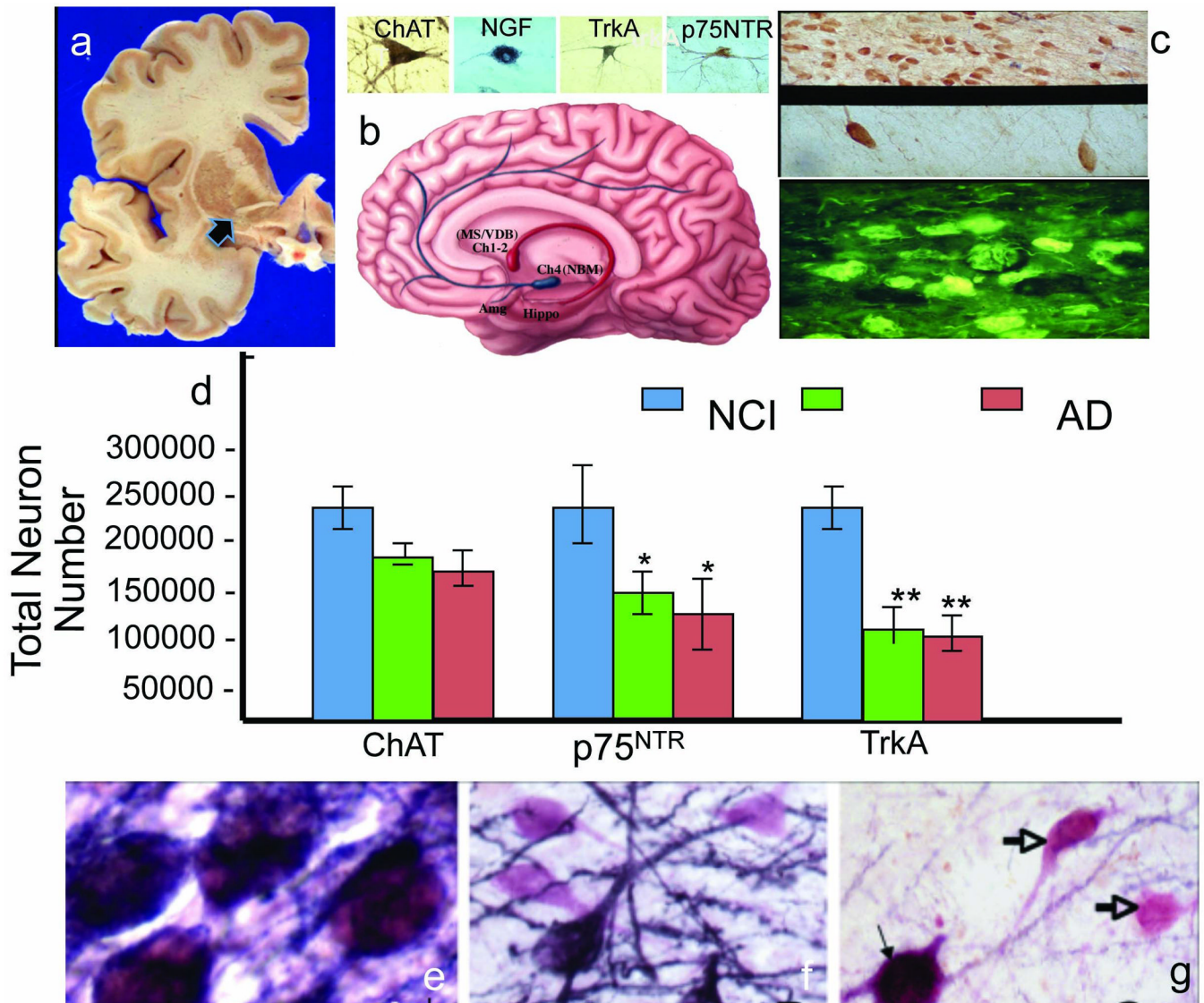


Fig. 3.

Graphs showing the estimate of total synapse number in layer 3 of the inferior temporal gyrus (ITG) across clinical diagnostic group using unbiased stereology coupled with electron microscopy. (a) Total volume of lamina 3 of the inferior temporal gyrus was estimated with the Cavalieri method from tissue sections immediately adjacent to regions used for synaptic counts (b). Single points represent individual subjects. Horizontal lines indicates group median. * $p < 0.05$; ** $p < 0.005$ compared to NCI; # $p < 0.005$ compared to MCI. Electron micrographs of ITG layer 3 showing synaptic complexes (white arrows) in tissue from (c) no cognitive impairment (NCI), (d) amnesic mild cognitive impairment (aMCI) and (e) AD cases. In all tissue, the synaptic complexes appeared normal with synaptic vesicles observed in the presynaptic component and the synaptic density observed in the postsynaptic component. Scale bar = 1.0 mm Adapted with permission from [152]

**Fig. 4.**

(a) Coronal brain slab showing the location of cholinergic neurons within the anterior nucleus basalis subfield (arrow). (b) Artist's drawing showing the cholinergic innervation from the medial septal/vertical limb of the diagonal band (MS/VDB; Ch1-2) to the hippocampus (red) and the nucleus basalis of Meynert (NBM) to the cortex and amygdala (blue). Above the schematic are images showing nucleus basalis neurons containing choline acetyltransferase (ChAT), nerve growth factor (NGF), trkA and p75^{NTR}. (c) Low power image showing the extensive loss of cholinergic neurons in the nucleus basalis in AD, higher magnification of the cholinergic cell loss (d) and NFT containing cholinergic neurons revealed by thioflavin-S histochemistry (yellow) (e). (f) Composite histogram showing phenotypic differences in the number of ChAT, vesicular acetylcholine transporter (VAChT), trkA and p75^{NTR}-immunopositive neurons in MCI and AD individuals. Note a significant reduction in NGF receptor and not ChAT containing neurons within the NB of people with MCI and AD. This difference is not exacerbated during the transition from MCI to early AD. (g) All NB neurons were dual immunostained for ChAT (pink) and p75^{NTR} (blue) in aged non-demented controls, which appeared as purple. (h) In contrast, in MCI (i, pink

neurons) and AD (J; open arrows) many more ChAT-only immune reactive neurons compared to dual stained neurons (thin arrow in j)

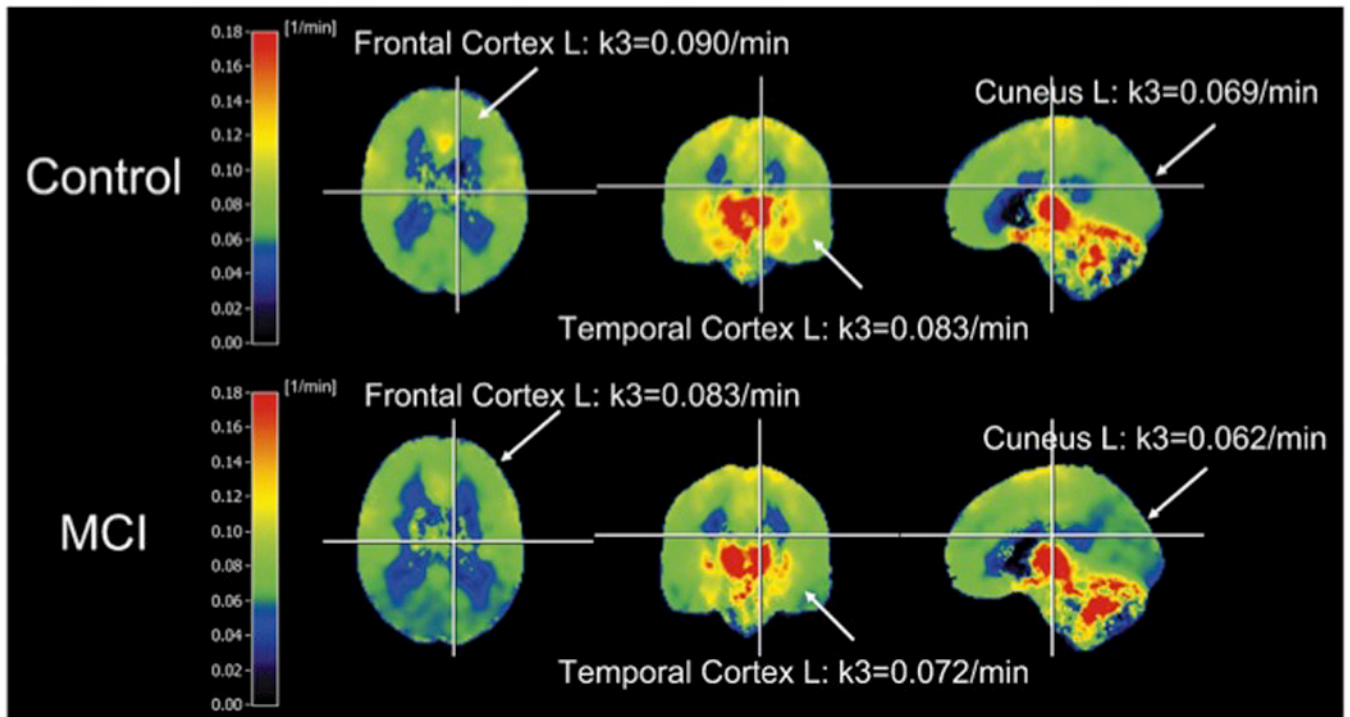


Fig. 5. Mean parametric images of $[^{11}\text{C}]$ MP4A hydrolysis rates [1/minute] used to measure acetylcholinesterase activity in controls and mild cognitive impairment (MCI) patients (orthogonal slices, L= left hemisphere, R = right hemisphere). Reproduced with permission from [59]

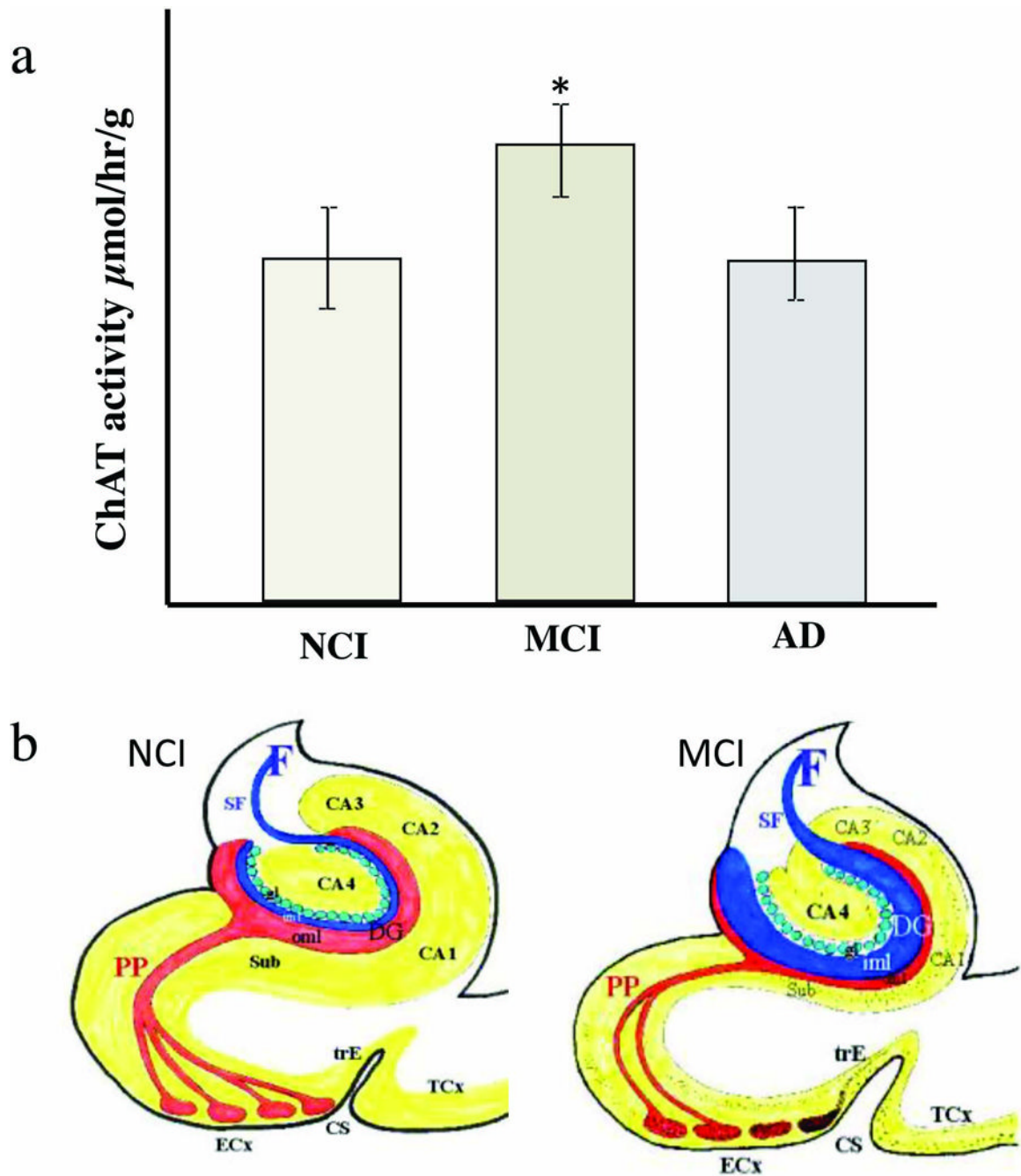
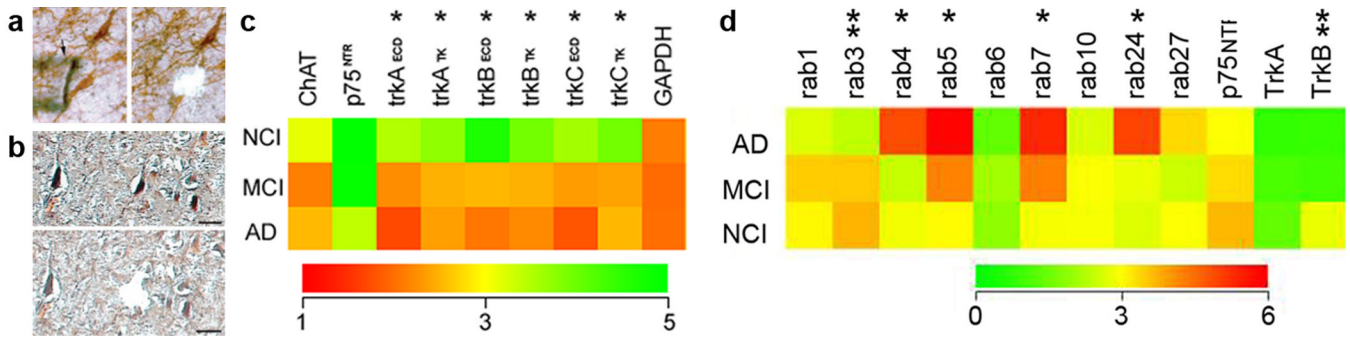


Fig. 6.
 (a) ChAT activity increased in the hippocampus in MCI and returned to control levels in mild AD. (b) Schematic drawing illustration the loss of innervation to the hippocampus arising from the glutamatergic layer II entorhinal cortex neurons (red) triggering a cholinergic plasticity response (blue) which likely originates from the septal cholinergic projection neurons into the denervated glutamatergic sites in the hippocampus in MCI

**Fig. 7.**

(a) Pre and post microaspirated cholinergic nucleus basalis (NB) and (b) neurofilament (NF)-immunoreactive hippocampal CA1 neurons employed for microarray analysis following terminal continuation (TC) RNA amplification. Scale bar: 50 μ m. (c) Color-coded heatmap illustrating no significant differences in ChAT, p75^{NTR} and GAPDH gene expression compared to a significant down regulation (asterisk) of trkA, trkB, and trkC in MCI and AD. Expressed sequence-tagged cDNAs (ESTs) identifying extracellular domains (ECD) and tyrosine kinase (TK) domains display down regulation in MCI and AD. The decrement of trk gene expression in MCI is intermediate relative to AD. (d) Color-coded heatmap illustrating a significant up regulation (asterisk) for endosomal and trafficking compartments rab GTPases rab4, rab5, rab7, and rab24 in MCI and AD compared to a down regulation (double asterisk) for the synaptic marker rab3 and the BDNF receptor TrkB within NF-immunoreactive CA1 neurons

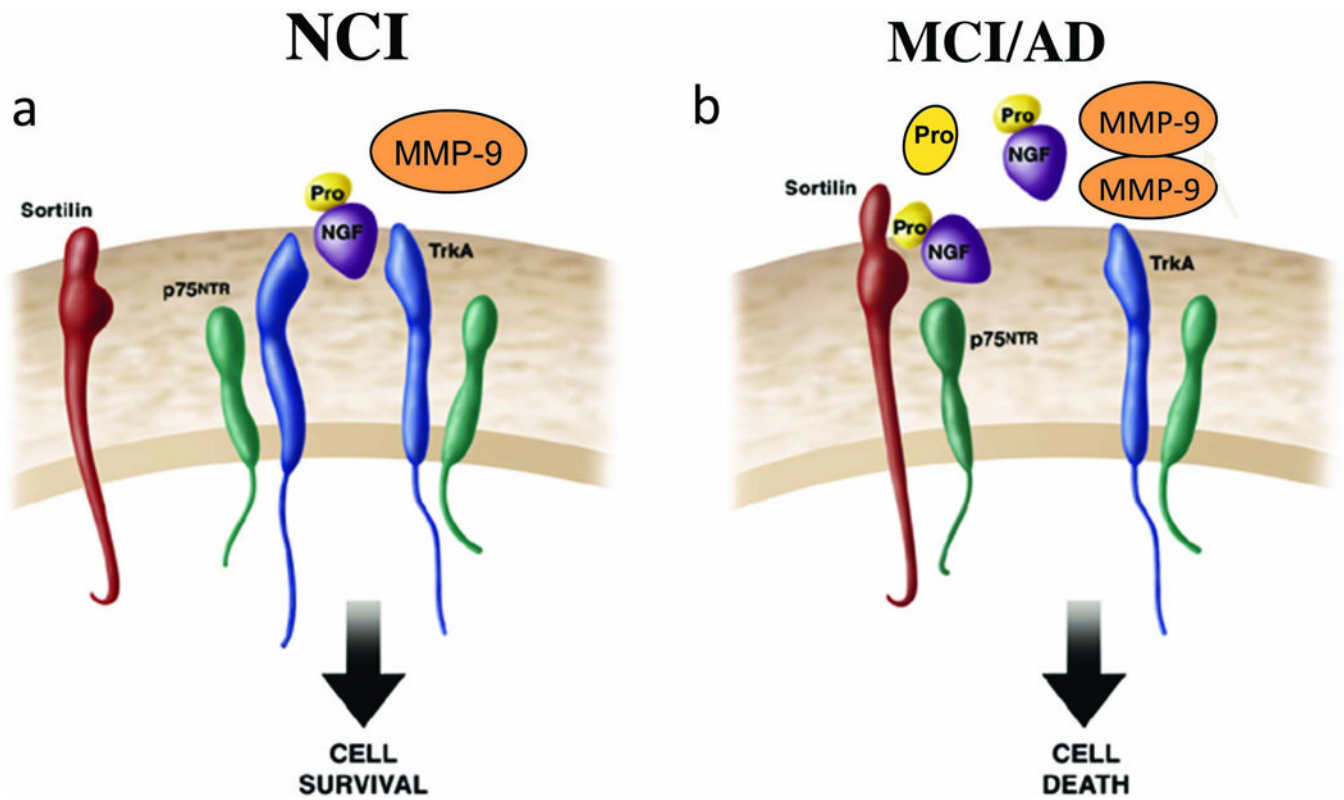


Fig. 8.

Illustration of changes related to the nerve growth factor system during the progression from non-demented (a) to MCI (b) within the cortex. During the progression of AD, elevated cortical proNGF in the face of reduced TrkA may enhance binding of proNGF to p75^{NTR}/sortilin complex. Since sortilin acts as molecular switch governing a p75^{NTR} mediated pro-apoptotic signal, increased proNGF may trigger cell death in the face of decreased TrkA. In addition, increased MMP-9 may also facilitate the degradation of mature NGF further impairing cholinergic trophic support